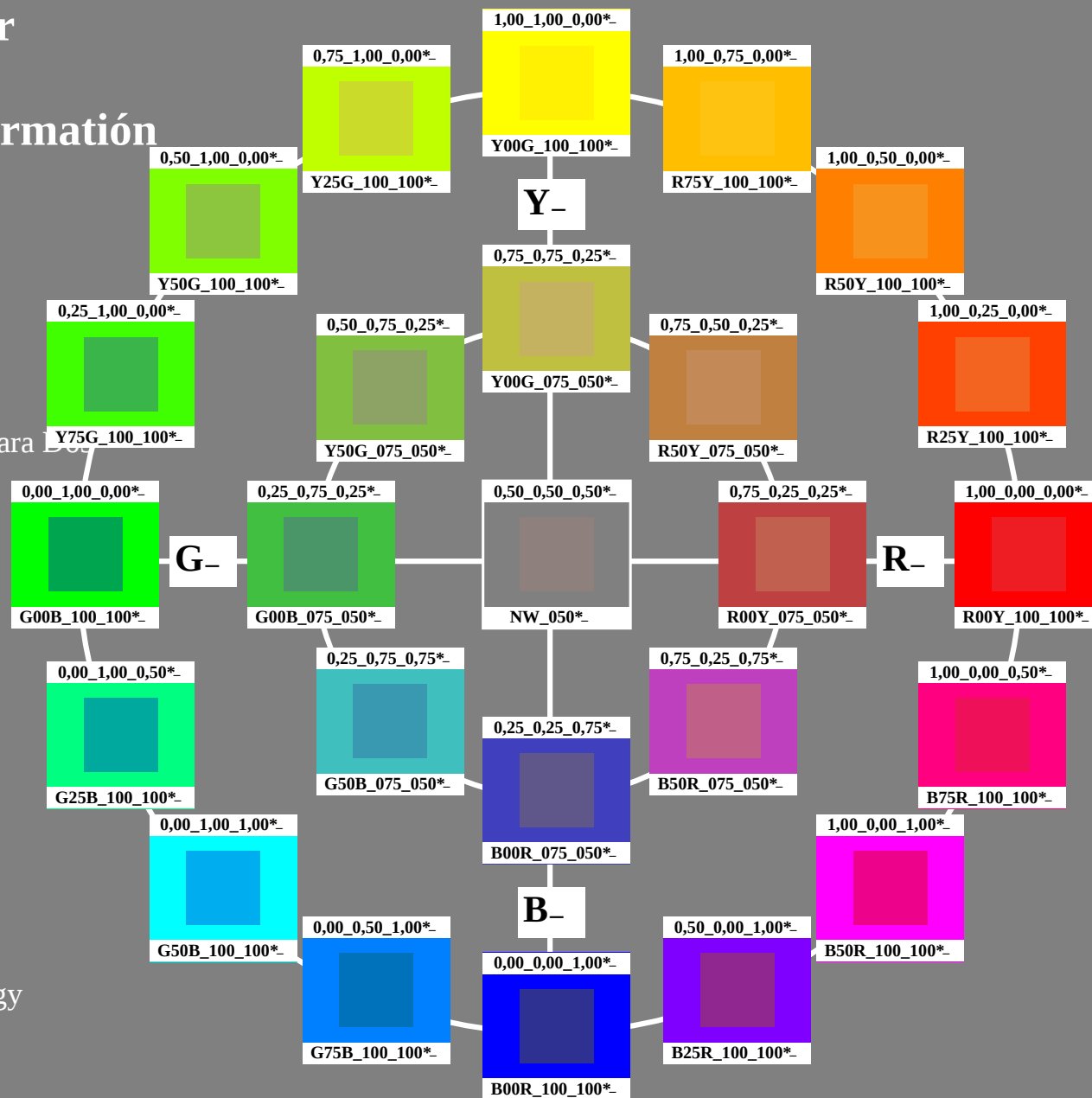


Color e Visión del Color Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar sRGB
rgb data: rgb^*_e (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : HIC^*_e (derecha)

Impresión especial par la exposición
El Color y la Visión del Color
Section Lighting Technology
of the Berlin University of Technology
Einsteinufer 19, D-10587 Berlin
ver: <http://www.li.tu-berlin.de>
y <http://130.149.60.45/~farbmetrik>



2-003030-L0

PS790-7N

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65

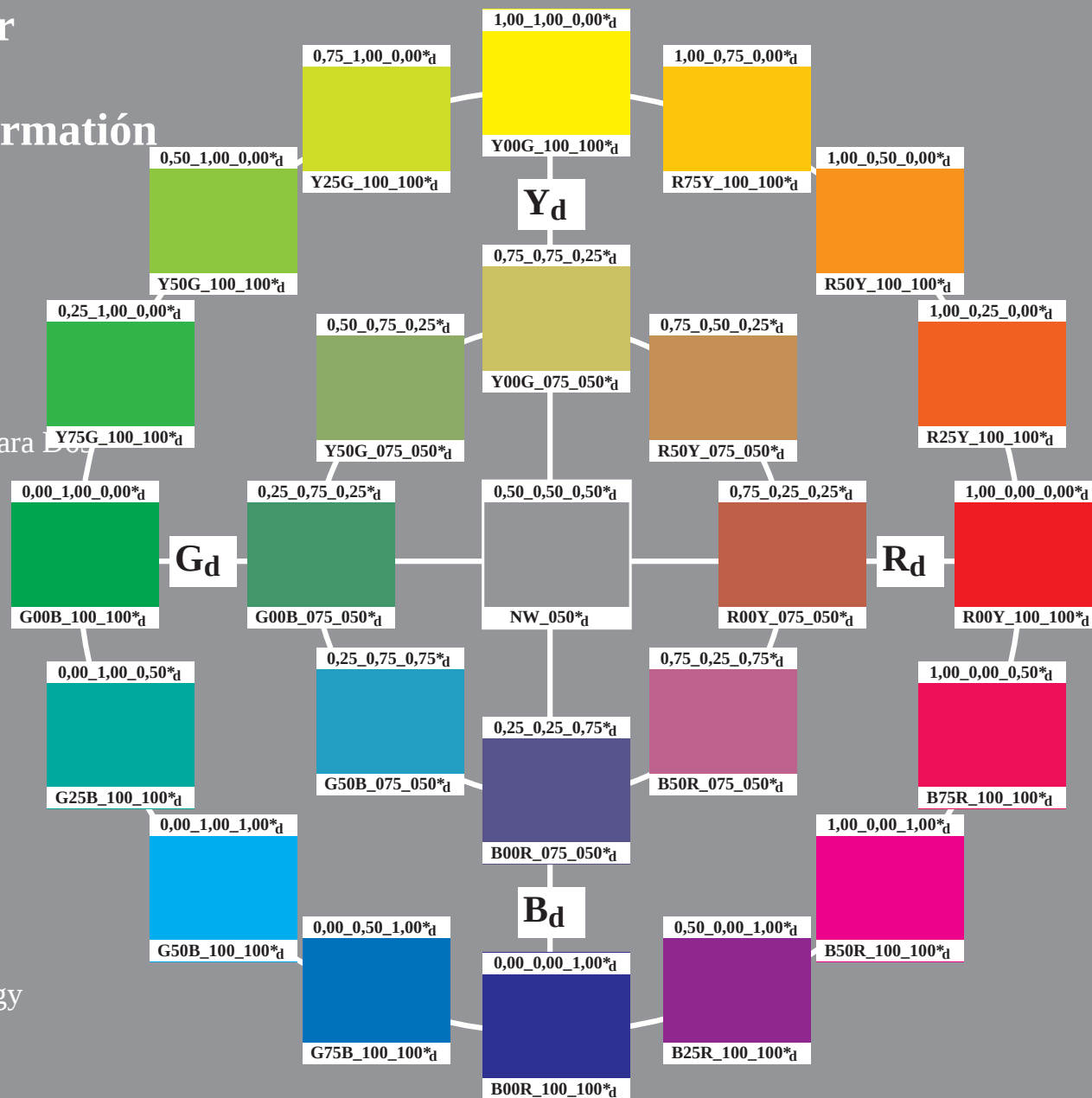
entrada: $rgb/cmyk \rightarrow rgb/cmyk$
salida: ningún cambio

Color e Visión del Color Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar sRGB
rgb data: rgb^*_e (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : HIC^*_e (derecha)

Impresión especial par la exposición
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y <http://130.149.60.45/~farbmetrik>



2-003130-L0

PS790-70

PE4300P_120901.TXT, 1080 colors, Separation cmykn6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=0, cmyk

entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmyk_d$

2-003130-F0

C

M

Y

O

L

V

C

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS TUB material: code=rha4ta
aplicación para la medida salida de impresora láser, separación cmyñ6 (CMYK)

<http://www.li.tu-berlin.de>
<http://130.149.60.45/~farbmetrik>

2-003230-L0

PS790-70

PE4300P_120901.TXT, 1080 colors, Separation cmyñ6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=0, cmyk

entrada: *rgb/cmyk* -> *rgb_d*
salida: transfiera a *cmyk_d*

2-003230-E0

C

M

Y

O

L

V

C

<http://www.li.tu-berlin.de>
<http://130.149.60.45/~farbmetrik>

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=0, cmyk

PE4300P_120901.TXT, 1080 colors, Separation cmyñ6*
entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmyk_d$

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS TUB material: code=rha4ta
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=0, cmyk

entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmyk_d$

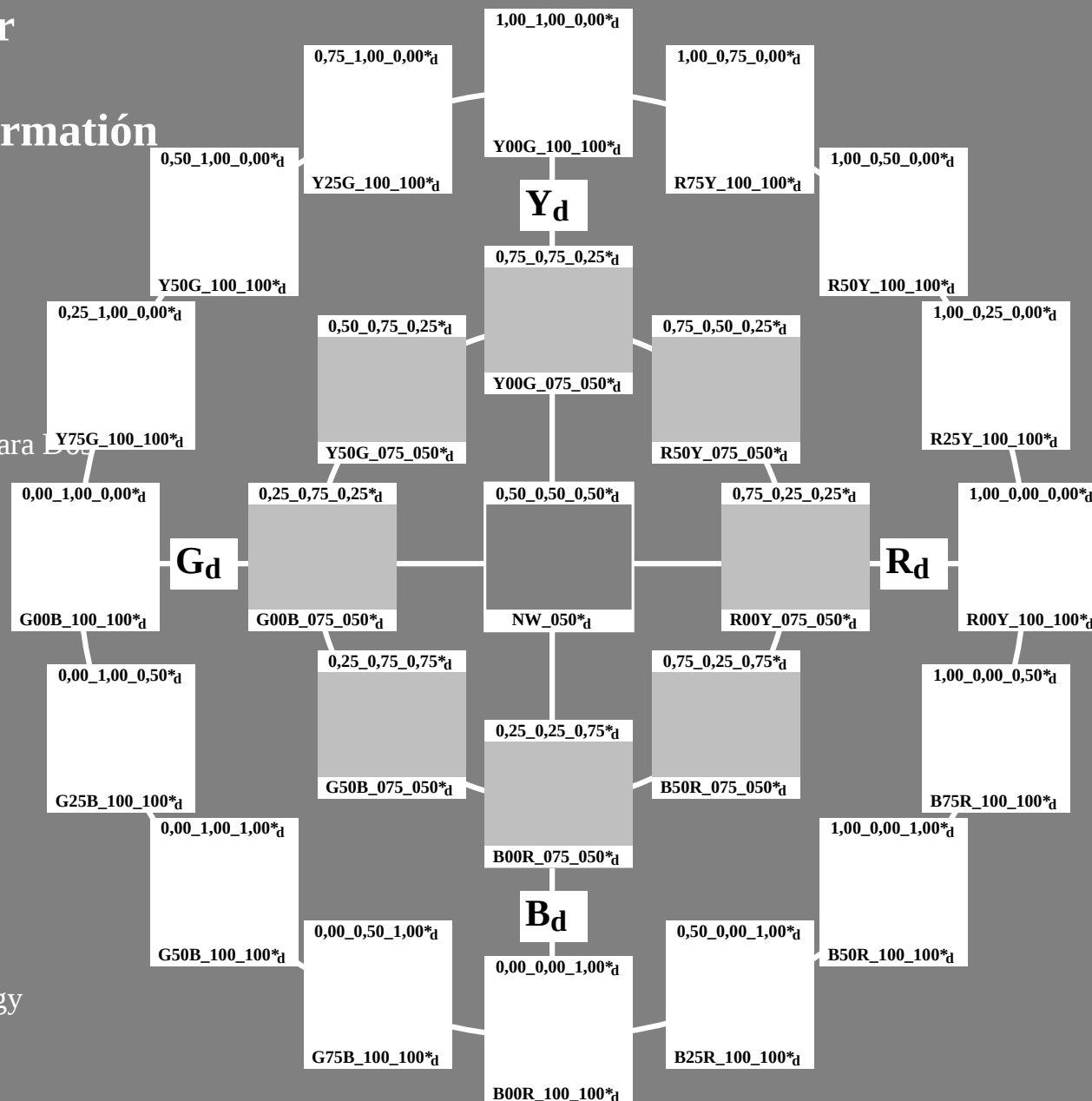
PS790-70 PE4300P_120901.TXT, 1080 colors, Separation cmykn6*

Color e Visión del Color Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar *sRGB*
rgb data: rgb^*_e (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : HIC^*_e (derecha)

Impresión especial par la exposición
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Section Lighting Technology
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y <http://130.149.60.45/~farbmetrik>



2-003530-L0

PS790-70

PE4300P_120901.TXT, 1080 colors, Separation cmykn6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=0, *cmyk*

entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmyk_d$

2-003530-F0

Color e Visión del Color

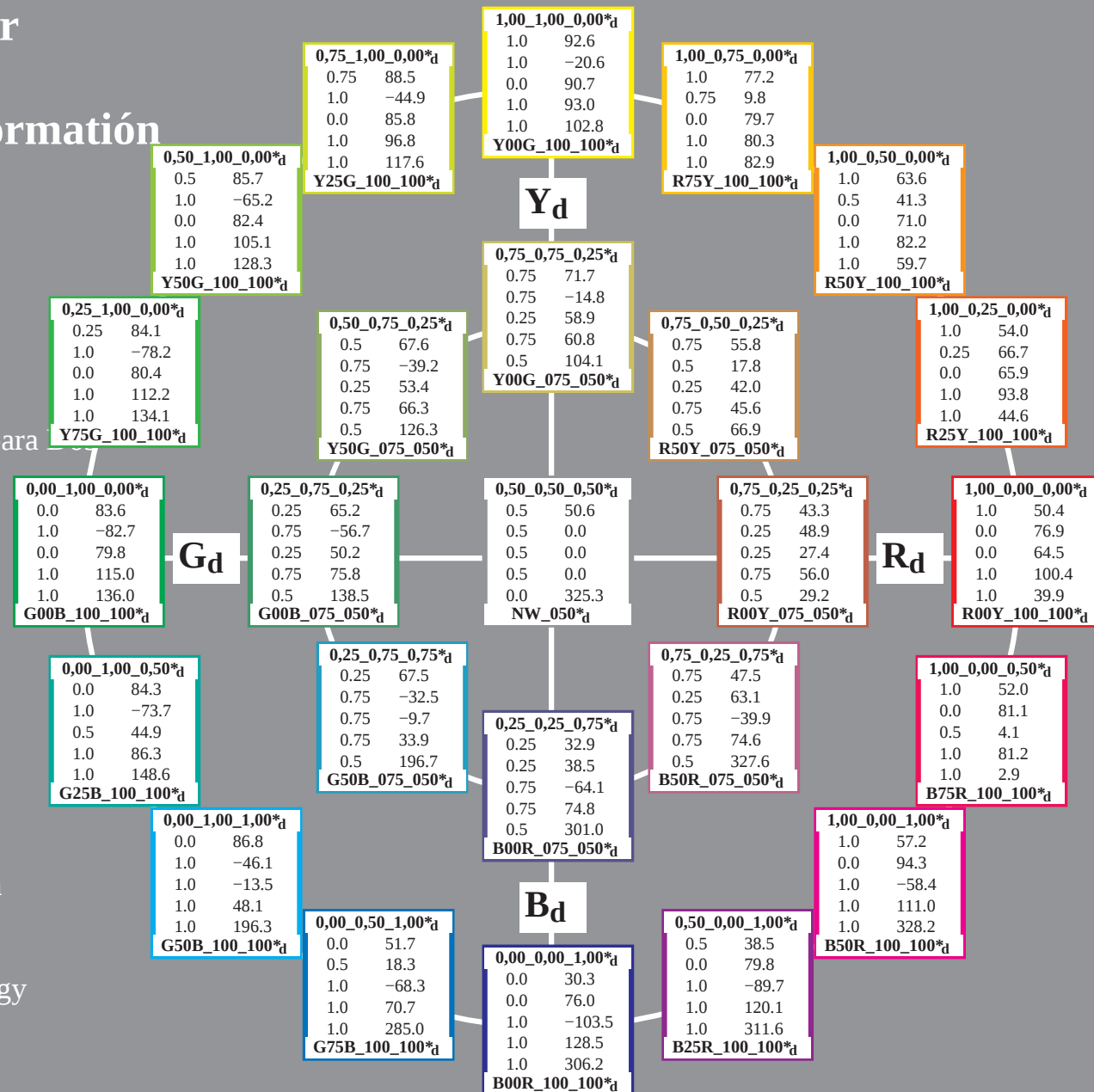
Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para círculo di tono con 16 y 8 pasos pantalla estándar sRGB

rgb data: rgb^*_e (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : HIC^*_e (derecha)
código color:
 $rgbicd$; $LabCh^*_d$

Impresión especial par la exposición
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2-003630-L0

PS790-70

PE4300P_120901.TXT, 1080 colors, Separation cmyn6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=0, cmyk

entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmyk_d$

2-003630-F0

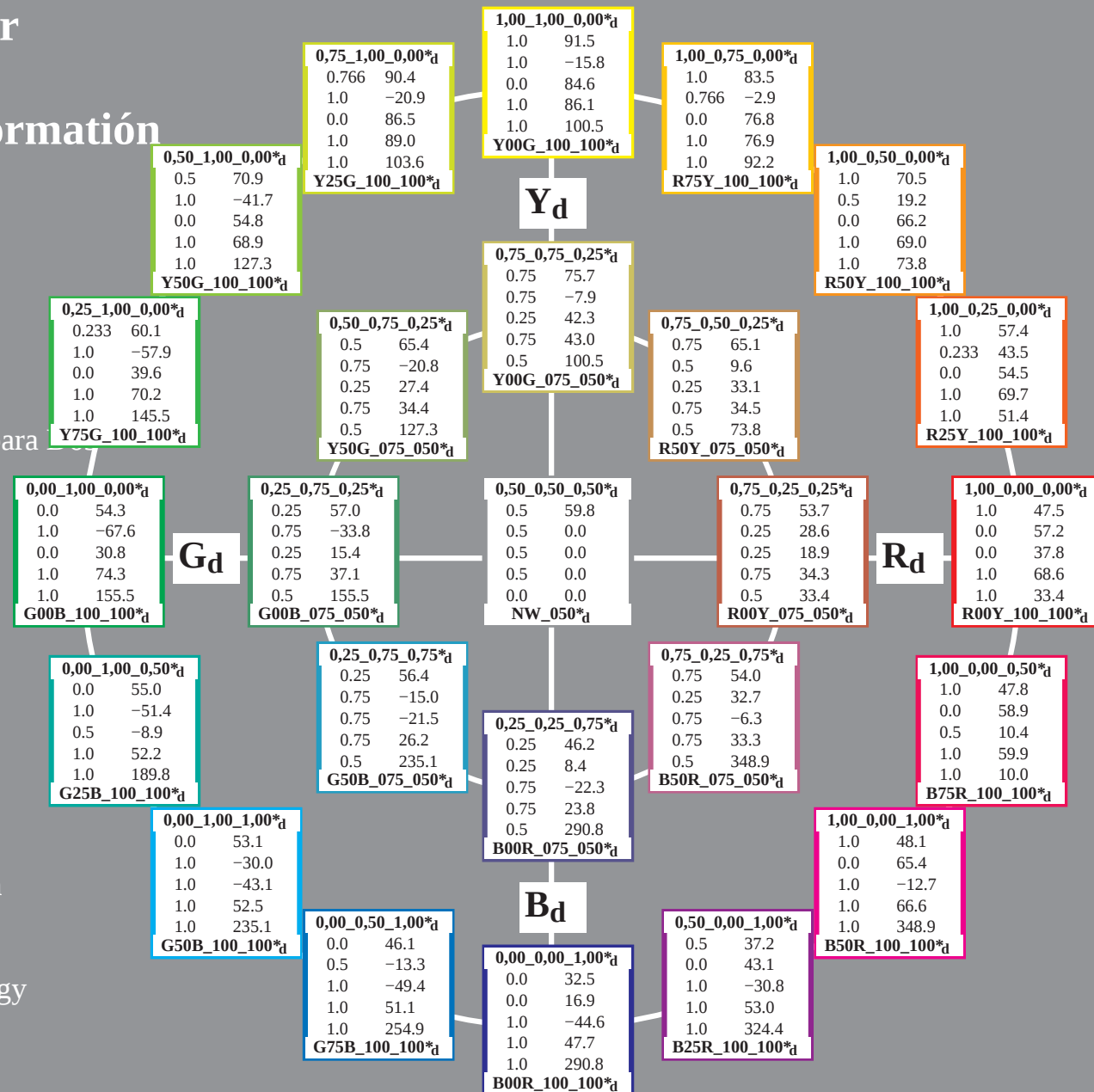
Color e Visión del Color Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar sRGB

rgb data: $rgb * e$ (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : $HIC * e$ (derecha)
código color:
 $rgbic * d$; $LabCh * d$

Impresión especial par la exposición
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Color e Visión del Color

Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar sRGB

rgb data: *rgb**_e (izquierda)
tonos elementales *H**, billantez *I**,
croma *C**: *HIC**_e (derecha)
código color:
*rgbic**_d; *rgbic*'*_d

Impresión especial par la exposición
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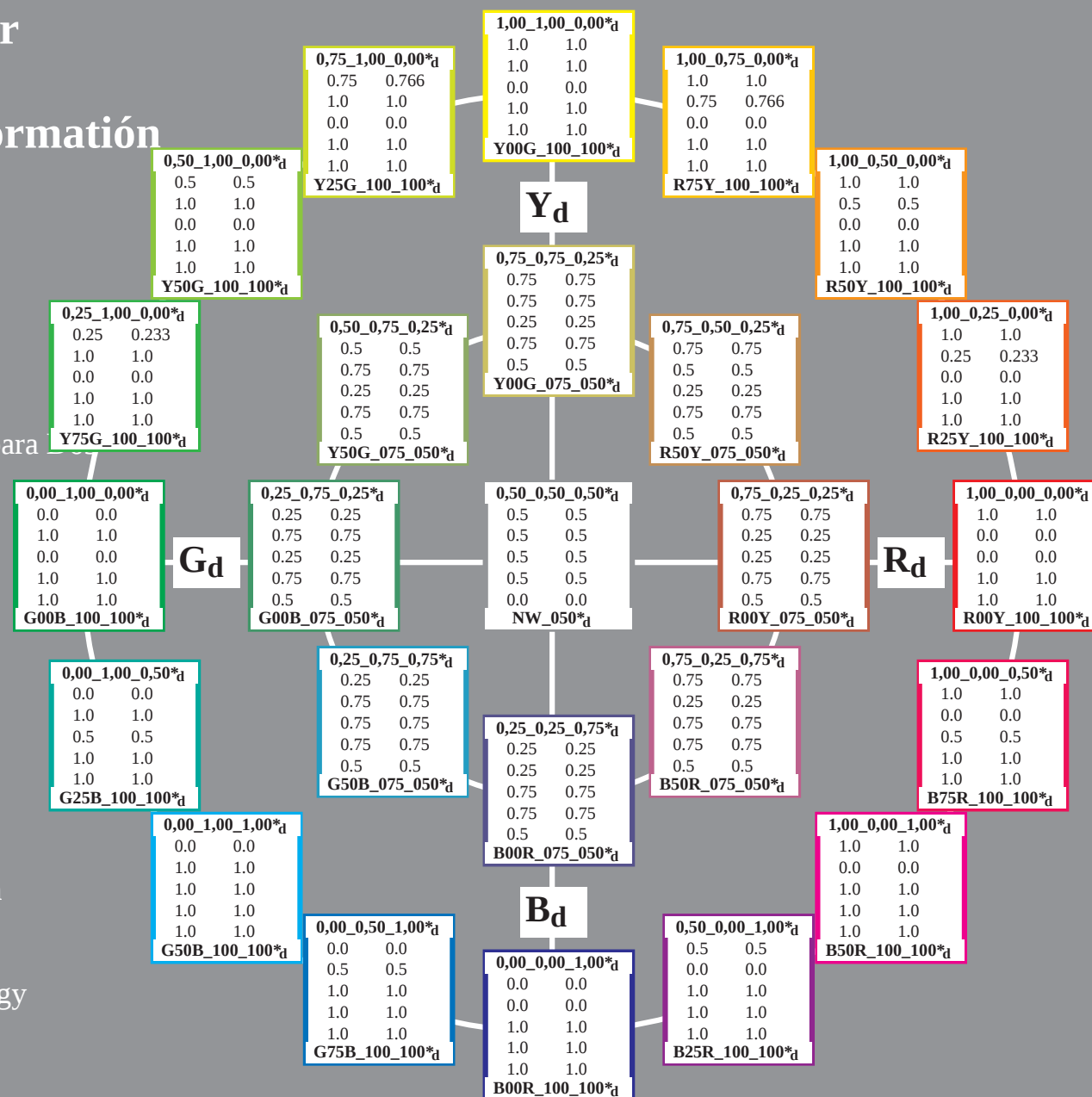


gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=0, *cm* *yk*

entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmyk_d$

Color e Visión del Color

Colores Elementales en la Tecnología de la Información

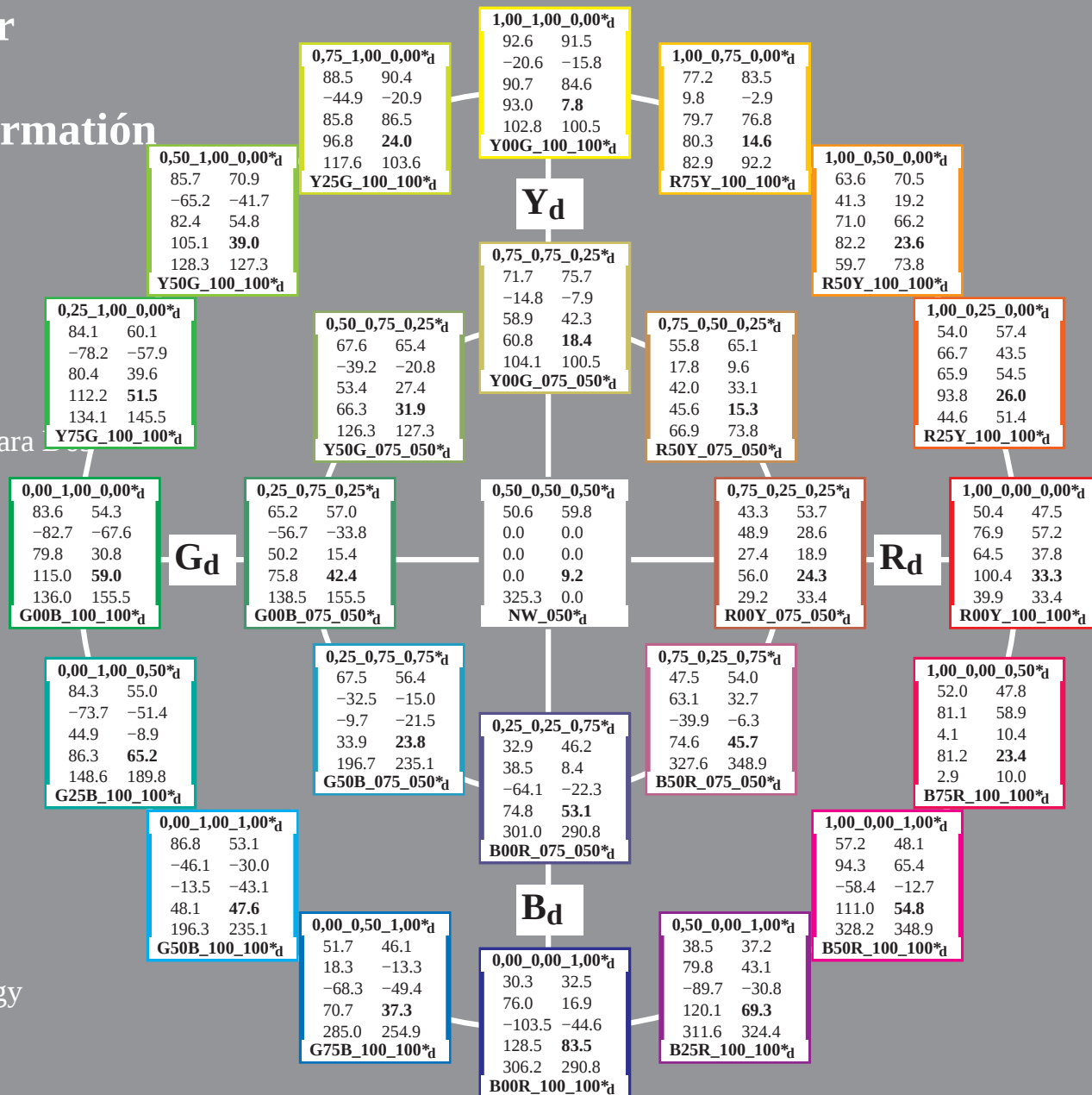
Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar sRGB

rgb data: $rgb * e$ (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : $HIC * e$ (derecha)
código color:

$LabCh * d$; $Lab * / DE * / h * d$

Impresión especial par la exposición
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vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS TUB material: code=rh4ta
aplicación para la medida salida de impresora láser, separación cmyñ6 (CMYK)

n/f	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
0/648	R00Y_100_100d	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4 0.0	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
1/657	R13Y_100_100d	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	51.6 54.5 48.4	72.9 41.6	1.0 0.125 0.0	51.9 54.3 49.2	73.2 42.1 0.8	36	1.0 0.116 0.0	51.6 54.5 48.4	72.9 41.6
2/666	R25Y_100_100d	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	57.4 43.5 54.5	69.7 51.4	1.0 0.25 0.0	58.2 41.8 55.1	69.2 52.8 1.9	42	1.0 0.233 0.0	57.4 43.5 54.5	69.7 51.4
3/675	R38Y_100_100d	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	64.2 30.6 60.1	67.5 63.0	1.0 0.375 0.0	64.6 29.8 60.4	67.3 63.7 0.9	51	1.0 0.366 0.0	64.2 30.6 60.1	67.5 63.0
4/684	R50Y_100_100d	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	70.5 19.2 66.2	69.0 73.8	1.0 0.5 0.0	70.5 19.2 66.2	69.0 73.8 0.0	59	1.0 0.5 0.0	70.5 19.2 66.2	69.0 73.8
5/693	R63Y_100_100d	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	75.4 10.6 71.2	72.0 81.5	1.0 0.625 0.0	74.9 11.4 70.7	71.6 80.7 1.1	68	1.0 0.633 0.0	75.4 10.6 71.2	72.0 81.5
6/702	R75Y_100_100d	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	83.5 -2.9 76.8	76.9 92.2	1.0 0.75 0.0	82.9 -2.0 76.9	77.0 91.5 1.1	77	1.0 0.766 0.0	83.5 -2.9 76.8	76.9 92.2
7/711	R88Y_100_100d	1.0 0.875 0.0	1.0 1.0 0.5	83	1.0 0.883 0.0	87.8 -9.4 76.3	76.9 97.0	1.0 0.875 0.0	87.6 -9.0 75.7	76.3 96.8 0.7	83	1.0 0.883 0.0	87.8 -9.4 76.3	76.9 97.0
8/720	Y00G_100_100d	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5 0.0	89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
9/639	Y13G_100_100d	0.875 1.0 0.0	1.0 1.0 0.5	97	0.883 1.0 0.0	92.7 -18.0 89.1	90.9 101.4	0.875 1.0 0.0	92.8 -18.1 89.4	91.2 101.4 0.3	96	0.883 1.0 0.0	92.7 -18.0 89.1	90.9 101.4
10/558	Y25G_100_100d	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	90.4 -20.9 86.5	89.0 103.6	0.75 1.0 0.0	90.1 -21.3 86.0	88.6 103.9 0.7	102	0.766 1.0 0.0	90.4 -20.9 86.5	89.0 103.6
11/477	Y38G_100_100d	0.625 1.0 0.0	1.0 1.0 0.5	112	0.633 1.0 0.0	80.5 -31.2 69.2	75.9 114.2	0.625 1.0 0.0	79.9 -31.7 67.9	75.0 115.0 1.4	111	0.633 1.0 0.0	80.5 -31.2 69.2	75.9 114.2
12/396	Y50G_100_100d	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	70.9 -41.7 54.8	68.9 127.3	0.5 1.0 0.0	70.9 -41.7 54.8	68.9 127.3 0.0	119	0.5 1.0 0.0	70.9 -41.7 54.8	68.9 127.3
13/315	Y63G_100_100d	0.375 1.0 0.0	1.0 1.0 0.5	128	0.366 1.0 0.0	66.1 -48.2 47.5	67.7 135.3	0.375 1.0 0.0	66.5 -47.5 48.0	67.6 134.7 0.8	128	0.366 1.0 0.0	66.1 -48.2 47.5	67.7 135.3
14/234	Y75G_100_100d	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	60.1 -57.9 39.6	70.2 145.5	0.25 1.0 0.0	60.6 -57.2 40.4	70.1 144.7 1.1	137	0.233 1.0 0.0	60.1 -57.9 39.6	70.2 145.5
15/153	Y88G_100_100d	0.125 1.0 0.0	1.0 1.0 0.5	143	0.116 1.0 0.0	56.8 -62.5 34.1	71.3 151.3	0.125 1.0 0.0	57.0 -62.2 34.4	71.1 151.0 0.4	143	0.116 1.0 0.0	56.8 -62.5 34.1	71.3 151.3
16/72	G00C_100_100d	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	54.3 -67.6 30.8	74.3 155.5	0.0 1.0 0.0	54.3 -67.6 30.8	74.3 155.5 0.0	149	0.0 1.0 0.0	54.3 -67.6 30.8	74.3 155.5
17/73	G13C_100_100d	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.116	53.8 -66.5 23.5	70.5 160.5	0.0 1.0 0.125	53.8 -66.4 23.0	70.2 160.8 0.5	156	0.0 1.0 0.116	53.8 -66.5 23.5	70.5 160.5
18/74	G25C_100_100d	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.233	53.7 -63.6 14.1	65.2 167.4	0.0 1.0 0.25	53.7 -63.1 12.8	64.4 168.5 1.3	162	0.0 1.0 0.233	53.7 -63.6 14.1	65.2 167.4
19/75	G38C_100_100d	0.0 1.0 0.375	1.0 1.0 0.5	172	0.0 1.0 0.366	54.7 -57.3 0.8	57.3 179.1	0.0 1.0 0.375	54.7 -56.8 0.0	56.8 179.9 0.9	171	0.0 1.0 0.366	54.7 -57.3 0.8	57.3 179.1
20/76	G50C_100_100d	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	55.0 -51.4 -8.9	52.2 189.8	0.0 1.0 0.5	55.0 -51.4 -8.9	52.2 189.8 0.0	180	0.0 1.0 0.5	55.0 -51.4 -8.9	52.2 189.8
21/77	G63C_100_100d	0.0 1.0 0.625	1.0 1.0 0.5	188	0.0 1.0 0.633	55.3 -43.8 -20.5	48.4 205.1	0.0 1.0 0.625	55.3 -44.1 -20.0	48.5 204.4 0.5	188	0.0 1.0 0.633	55.3 -43.8 -20.5	48.4 205.1
22/78	G75C_100_100d	0.0 1.0 0.75	1.0 1.0 0.5	196	0.0 1.0 0.766	55.1 -39.2 -27.9	48.1 215.4	0.0 1.0 0.75	55.2 -39.5 -27.1	47.9 214.4 0.8	197	0.0 1.0 0.766	55.1 -39.2 -27.9	48.1 215.4
23/79	G88C_100_100d	0.0 1.0 0.875	1.0 1.0 0.5	203	0.0 1.0 0.883	54.3 -36.4 -33.7	49.6 222.8	0.0 1.0 0.875	54.4 -36.7 -33.0	49.4 221.9 0.7	203	0.0 1.0 0.883	54.3 -36.4 -33.7	49.6 222.8
24/80	C00B_100_100d	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1	0.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1 0.0	210	0.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
25/71	C13B_100_100d	0.0 0.875 1.0	1.0 1.0 0.5	217	0.0 0.883 1.0	53.1 -28.1 -44.6	52.7 237.7	0.0 0.875 1.0	53.1 -27.9 -44.7	52.7 237.9 0.1	216	0.0 0.883 1.0	53.1 -28.1 -44.6	52.7 237.7
26/62	C25B_100_100d	0.0 0.75 1.0	1.0 1.0 0.5	224	0.0 0.766 1.0	52.9 -26.2 -47.2	53.9 240.9	0.0 0.75 1.0	52.9 -25.9 -47.5	54.1 241.3 0.4	222	0.0 0.766 1.0	52.9 -26.2 -47.2	53.9 240.9
27/53	C38B_100_100d	0.0 0.625 1.0	1.0 1.0 0.5	232	0.0 0.633 1.0	50.7 -21.1 -49.4	53.7 246.8	0.0 0.625 1.0	50.5 -20.8 -49.5	53.7 247.2 0.3	231	0.0 0.633 1.0	50.7 -21.1 -49.4	53.7 246.8
28/44	C50B_100_100d	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	46.1 -13.3 -49.4	51.1 254.9	0.0 0.5 1.0	46.1 -13.3 -49.4	51.1 254.9 0.0	240	0.0 0.5 1.0	46.1 -13.3 -49.4	51.1 254.9
29/35	C63B_100_100d	0.0 0.375 1.0	1.0 1.0 0.5	248	0.0 0.366 1.0	41.1 -5.7 -49.2	49.6 263.3	0.0 0.375 1.0	41.4 -6.3 -49.2	49.6 262.6 0.6	248	0.0 0.366 1.0	41.1 -5.7 -49.2	49.6 263.3
30/26	C75B_100_100d	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.233 1.0	36.6 3.2 -48.3	48.4 273.8	0.0 0.25 1.0	36.8 2.2 -48.5	48.6 272.6 1.0	257	0.0 0.233 1.0	36.6 3.2 -48.3	48.4 273.8
31/17	C88B_100_100d	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.116 1.0	34.9 9.9 -46.3	47.3 282.0	0.0 0.125 1.0	35.0 9.4 -46.3	47.3 281.4 0.5	263	0.0 0.116 1.0	34.9 9.9 -46.3	47.3 282.0
32/8	B00M_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8 0.0	270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
33/89	B13M_100_100d	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	31.6 23.1 -42.4	48.3 298.6	0.125 0.0 1.0	31.6 23.6 -42.2	48.4 299.2 0.4	276	0.116 0.0 1.0	31.6 23.1 -42.4	48.3 298.6
34/170	B25M_100_100d	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	31.1 29.6 -39.8	49.6 306.6	0.25 0.0 1.0	31.0 30.5 -39.3	49.8 307.8 1.0	282	0.233 0.0 1.0	31.1 29.6 -39.8	49.6 306.6
35/251	B38M_100_100d	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	34.0 37.7 -35.3	51.7 316.8	0.375 0.0 1.0	34.2 38.2 -35.0	51.8 317.5 0.6	291	0.366 0.0 1.0	34.0 37.7 -35.3	51.7 316.8
36/332	B50M_100_100d	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	37.2 43.1 -30.8	53.0 324.4	0.5 0.0 1.0	37.2 43.1 -30.8	53.0 324.4 0.0	300	0.5 0.0 1.0	37.2 43.1 -30.8	53.0 324.4
37/413	B63M_100_100d	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	39.2 48.9 -26.9	55.8 331.1	0.625 0.0 1.0	39.1 48.4 -27.2	55.6 330.6 0.6	308	0.633 0.0 1.0	39.2 48.9 -26.9	55.8 331.1
38/494	B75M_100_100d	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	42.4 55.8 -20.9	59.6 339.4	0.75 0.0 1.0	41.8 55.1 -21.4	59.1 338.7 0.9	317	0.766 0.0 1.0	42.4 55.8 -20.9	59.6 339.4
39/575	B88M_100_100d	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	45.8 60.5 -17.0	62.8 344.2	0.875 0.0 1.0	45.6 60.1 -17.3	62.6 343.9 0.4	323	0.883 0.0 1.0	45.8 60.5 -17.0	62.8 344.2
40/656	M00R_100_100d	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9 0.0	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
41/655	M13R_100_100d	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	49.4 66.1 -10.9	67.0 350.6	1.0 0.0 0.875	49.5 66.1 -10.7	67.0 350.7 0.1	336	1.0 0.0 0.883	49.4 66.1 -10.9	67.0 350.6
42/654	M25R_100_100d	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	49.3 64.7 -7.1	65.1 353.7	1.0 0.0 0.75	49.3 64.5 -6.5	64.8 354.2 0.5	342	1.0 0.0 0.766	49.3 64.7 -7.1	65.1 353.7
43/653	M38R_100_100d	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	48.0 62.0 1.5	62.0 1.4	1.0 0.0 0.625	48.0 61.8 2.1	61.8 361.9 0.6	351	1.0 0.0 0.633	48.0 62.0 1.5	62.0 1.4
44/652	M50R_100_100d	1.0 0.0 0.5	1.0 1											

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)

TUB material: code=rha4ta

n/f	HIC*Fd	rgb_Fd		ict_Fd		hsi_Fd	rgb*Fd		LabCh*Fd		rgb*Fd		LabCh*Fd		DE**Fd		hsiMd	rgb*Md		LabCh*Md						
0/648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	47.5	57.2	37.8	68.6	33.4	0.0	389	1.0	0.0	0.0	47.5	57.2	37.8	68.6	33.4
1/666	R25Y_100_100a	1.0	0.25	0.0	1.0	1.0	0.5	44	1.0	0.233	0.0	57.4	43.5	54.5	69.7	51.4	0.0	42	1.0	0.233	0.0	57.4	43.5	54.5	69.7	51.4
2/684	R50Y_100_100a	1.0	0.5	0.0	1.0	1.0	0.5	60	1.0	0.5	0.0	70.5	19.2	66.2	69.0	73.8	0.0	59	1.0	0.5	0.0	70.5	19.2	66.2	69.0	73.8
3/702	R75Y_100_100a	1.0	0.75	0.0	1.0	1.0	0.5	76	1.0	0.766	0.0	83.5	-2.9	76.8	76.9	92.2	1.0	77	1.0	0.766	0.0	83.5	-2.9	76.8	76.9	92.2
4/720	Y00G_100_100a	1.0	1.0	0.0	1.0	1.0	0.5	90	1.0	1.0	0.0	91.5	-15.8	84.6	86.1	100.5	0.0	89	1.0	1.0	0.0	91.5	-15.8	84.6	86.1	100.5
5/558	Y25G_100_100a	0.75	1.0	0.0	1.0	1.0	0.5	104	0.766	1.0	0.0	90.4	-20.9	86.5	89.0	103.6	0.7	102	0.766	1.0	0.0	90.4	-20.9	86.5	89.0	103.6
6/396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	120	0.5	1.0	0.0	70.9	-41.7	54.8	68.9	127.3	0.0	119	0.5	1.0	0.0	70.9	-41.7	54.8	68.9	127.3
7/234	Y75G_100_100a	0.25	1.0	0.0	1.0	1.0	0.5	136	0.233	1.0	0.0	60.1	-57.9	39.6	70.2	145.5	1.1	137	0.233	1.0	0.0	60.1	-57.9	39.6	70.2	145.5
8/72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	54.3	-67.6	30.8	74.3	155.5	0.0	149	0.0	1.0	0.0	54.3	-67.6	30.8	74.3	155.5
9/72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	54.3	-67.6	30.8	74.3	155.5	0.0	149	0.0	1.0	0.0	54.3	-67.6	30.8	74.3	155.5
10/76	G25B_100_100a	0.0	1.0	0.5	1.0	1.0	0.5	180	0.0	1.0	0.5	55.0	-51.4	-8.9	52.2	189.8	0.0	180	0.0	1.0	0.5	55.0	-51.4	-8.9	52.2	189.8
11/80	G50B_100_100a	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	1.0	53.1	-30.0	-43.1	52.5	235.1	0.0	210	0.0	1.0	1.0	53.1	-30.0	-43.1	52.5	235.1
12/44	G75B_100_100a	0.0	0.5	1.0	1.0	1.0	0.5	240	0.0	0.5	1.0	46.1	-13.3	-49.4	51.1	254.9	0.0	240	0.0	0.5	1.0	46.1	-13.3	-49.4	51.1	254.9
13/8	B00R_100_100a	0.0	0.0	1.0	1.0	1.0	0.5	270	0.0	0.0	1.0	32.5	16.9	-44.6	47.7	290.8	0.0	270	0.0	0.0	1.0	32.5	16.9	-44.6	47.7	290.8
14/332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	300	0.5	0.0	1.0	37.2	43.1	-30.8	53.0	324.4	0.0	300	0.5	0.0	1.0	37.2	43.1	-30.8	53.0	324.4
15/656	B50R_100_100a	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	48.1	65.4	-12.7	66.6	348.9	0.0	330	1.0	0.0	1.0	48.1	65.4	-12.7	66.6	348.9
16/652	B75R_100_100a	1.0	0.0	0.5	1.0	1.0	0.5	360	1.0	0.0	0.5	47.8	58.9	10.4	59.9	10.0	0.0	360	1.0	0.0	0.5	47.8	58.9	10.4	59.9	10.0
17/648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	47.5	57.2	37.8	68.6	33.4	0.0	389	1.0	0.0	0.0	47.5	57.2	37.8	68.6	33.4
18/688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	390	1.0	0.5	0.5	71.7	28.6	18.9	34.3	33.4	1.0	0.5	0.5	71.4	24.0	27.4	36.4	48.8	9.6	389
19/706	R50Y_100_050a	1.0	0.75	0.5	1.0	0.5	0.75	60	1.0	0.75	0.5	83.1	9.6	33.1	34.5	73.8	1.0	0.75	0.5	83.6	2.8	39.3	39.4	85.7	9.1	59
20/724	Y00G_100_050a	1.0	1.0	0.5	1.0	0.5	0.75	90	1.0	1.0	0.5	93.7	-7.9	42.3	43.0	100.5	1.0	1.0	0.5	93.1	-11.8	45.5	47.0	104.6	5.0	89
21/562	Y50G_100_050a	0.75	1.0	0.5	1.0	0.5	0.75	120	0.75	1.0	0.5	83.4	-20.8	27.4	34.4	127.3	0.75	1.0	0.5	86.2	-21.9	37.8	43.7	120.1	10.8	119
22/490	G00B_100_050a	0.5	1.0	0.5	1.0	0.5	0.75	150	0.5	1.0	0.5	75.0	-33.8	15.4	37.1	155.5	0.5	1.0	0.5	74.1	-30.5	11.7	32.6	158.9	5.0	149
23/404	G50B_100_050a	0.5	1.0	1.0	1.0	0.5	0.75	210	0.5	1.0	1.0	74.4	-15.0	-21.5	26.2	235.1	0.5	1.0	1.0	73.7	-17.1	-27.4	32.3	238.0	6.3	210
24/368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	270	0.5	0.5	1.0	64.2	8.4	-22.3	23.8	290.8	0.5	0.5	1.0	54.8	11.5	-32.2	34.2	289.7	14.0	270
25/692	B50R_100_050a	1.0	0.5	1.0	1.0	0.5	0.75	330	1.0	0.5	1.0	72.0	32.7	-6.3	33.3	348.9	1.0	0.5	1.0	71.6	36.1	-8.9	37.2	346.1	4.3	330
26/688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	390	1.0	0.5	0.5	71.7	28.6	18.9	34.3	33.4	1.0	0.5	0.5	71.4	24.0	27.4	36.4	48.8	9.6	389
27/506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	390	0.75	0.25	0.25	53.7	28.6	18.9	34.3	33.4	0.75	0.25	0.25	52.4	27.1	25.3	37.1	43.0	6.7	389
28/524	R50Y_075_050a	0.75	0.5	0.25	0.75	0.5	0.5	60	0.75	0.5	0.25	65.1	9.6	33.1	34.5	73.8	0.75	0.5	0.25	66.1	6.5	36.1	36.6	79.7	4.3	59
29/542	Y00G_075_050a	0.75	0.75	0.25	0.75	0.5	0.5	90	0.75	0.75	0.25	75.7	-7.9	42.3	43.0	100.5	0.75	0.75	0.25	81.7	-11.5	50.7	52.0	102.8	10.9	89
30/380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	120	0.5	0.75	0.25	65.4	-20.8	27.4	34.4	127.3	0.5	0.75	0.25	70.5	-23.0	31.5	39.0	126.1	6.9	119
31/218	G00B_075_050a	0.25	0.75	0.25	0.75	0.5	0.5	150	0.25	0.75	0.25	57.0	-33.8	15.4	37.1	155.5	0.25	0.75	0.25	57.2	-36.4	13.9	39.0	159.0	2.9	149
32/222	G50B_075_050a	0.25	0.75	0.75	0.75	0.5	0.5	210	0.25	0.75	0.75	56.4	-15.0	-21.5	26.2	235.1	0.25	0.75	0.75	60.1	-19.6	-29.7	35.6	236.5	10.1	210
33/186	B00R_075_050a	0.25	0.25	0.75	0.75	0.5	0.5	270	0.25	0.25	0.75	46.2	8.4	-22.3	23.8	290.8	0.25	0.25	0.75	43.1	11.1	-34.5	36.3	287.8	12.9	270
34/510	B50R_075_050a	0.75	0.25	0.75	0.75	0.5	0.5	330	0.75	0.25	0.75	54.0	32.7	-6.3	33.3	348.9	0.75	0.25	0.75	53.9	38.1	-12.4	40.1	341.9	8.1	330
35/506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	390	0.75	0.25	0.25	53.7	28.6	18.9	34.3	33.4	0.75	0.25	0.25	52.4	27.1	25.3	37.1	43.0	6.7	389
36/324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.25	390	0.5	0.0	0.0	35.7	28.6	18.9	34.3	33.4	0.5	0.0	0.0	33.0	34.7	23.4	41.8	34.0	8.0	389
37/342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	60	0.5	0.25	0.0	47.1	9.6	33.1	34.5	73.8	0.5	0.25	0.0	42.9	9.5	37.9	39.1	75.8	6.3	59
38/360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	90	0.5	0.5	0.0	57.7	-7.9	42.3	43.0	100.5	0.5	0.5	0.0	58.4	-9.8	54.3	55.2	100.3	12.1	89
39/198	Y50G_050_050a	0.25	0.5	0.0	0.5	0.5	0.25	120	0.25	0.5	0.0	47.4	-20.8	27.4	34.4	127.3	0.25	0.5	0.0	43.9	-28.1	32.6	43.1	130.7	9.6	119
40/36	G00B_050_050a	0.0	0.5	0.0	0.5	0.5	0.25	150	0.0	0.5	0.0	39.0	-33.8	15.4	37.1	155.5	0.0	0.5	0.0	42.9	-59.6	21.5	63.4	160.1	26.8	149
41/40	G50B_050_050a	0.0	0.5	0.5	0.5	0.5	0.25	210	0.0	0.5	0.5	38.4	-15.0	-21.5	26.2	235.1	0.0	0.5	0.5	44.1	-23.4	-34.5	41.7	235.7	16.4	210
42/4	B00R_050_050a	0.0	0.0	0.5	0.5	0.5	0.25	270	0.0	0.0	0.5	28.2	8.4	-22.3	23.8	290.8	0.0	0.0	0.5	30.3	13.1	-38.9	41.0	288.6	17.3	270
43/328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.25																			

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n=j	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md
0	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	1.0
1	B00R_012_012a	0.0	0.0	0.125	0.125	0.125	0.062	0.062	0.0	0.0	1.0	1.0
2	B00R_025_025a	0.0	0.0	0.25	0.25	0.25	0.125	0.125	0.0	0.0	1.0	1.0
3	B00R_037_037a	0.0	0.0	0.375	0.375	0.375	0.187	0.187	0.0	0.0	1.0	1.0
4	B00R_050_050a	0.0	0.0	0.5	0.5	0.5	0.25	0.25	0.0	0.0	1.0	1.0
5	B00R_062_062a	0.0	0.0	0.625	0.625	0.625	0.312	0.312	0.0	0.0	1.0	1.0
6	B00R_075_075a	0.0	0.0	0.75	0.75	0.75	0.375	0.375	0.0	0.0	1.0	1.0
7	B00R_087_087a	0.0	0.0	0.875	0.875	0.875	0.437	0.437	0.0	0.0	1.0	1.0
8	B00R_100_100a	0.0	0.0	1.0	1.0	1.0	0.5	0.5	0.0	0.0	1.0	1.0
9	G00B_012_012a	0.0	0.125	0.0	0.125	0.125	0.062	0.062	0.0	0.125	0.0	0.125
10	G50B_012_012a	0.0	0.125	0.125	0.125	0.125	0.062	0.062	0.0	0.125	0.125	0.125
11	G75B_025_025a	0.0	0.125	0.25	0.25	0.25	0.125	0.125	0.0	0.125	0.25	0.25
12	G84B_037_037a	0.0	0.125	0.375	0.375	0.375	0.187	0.187	0.0	0.125	0.375	0.375
13	G88B_050_050a	0.0	0.125	0.5	0.5	0.5	0.25	0.25	0.0	0.125	0.5	0.5
14	G90B_062_062a	0.0	0.125	0.625	0.625	0.625	0.312	0.312	0.0	0.125	0.625	0.625
15	G92B_075_075a	0.0	0.125	0.75	0.75	0.75	0.375	0.375	0.0	0.125	0.75	0.75
16	G93B_087_087a	0.0	0.125	0.875	0.875	0.875	0.437	0.437	0.0	0.125	0.875	0.875
17	G94B_100_100a	0.0	0.125	1.0	1.0	1.0	0.5	0.5	0.0	0.125	1.0	1.0
18	G00B_025_025a	0.0	0.25	0.0	0.25	0.25	0.125	0.125	0.0	0.25	0.0	0.25
19	G25B_025_025a	0.0	0.25	0.125	0.25	0.25	0.125	0.125	0.0	0.25	0.125	0.125
20	G50B_025_025a	0.0	0.25	0.25	0.25	0.25	0.125	0.125	0.0	0.25	0.25	0.25
21	G65B_037_037a	0.0	0.25	0.375	0.375	0.375	0.187	0.187	0.0	0.25	0.375	0.375
22	G75B_050_050a	0.0	0.25	0.5	0.5	0.5	0.25	0.25	0.0	0.25	0.5	0.5
23	G80B_062_062a	0.0	0.25	0.625	0.625	0.625	0.312	0.312	0.0	0.25	0.625	0.625
24	G84B_075_075a	0.0	0.25	0.75	0.75	0.75	0.375	0.375	0.0	0.25	0.75	0.75
25	G86B_087_087a	0.0	0.25	0.875	0.875	0.875	0.437	0.437	0.0	0.25	0.875	0.875
26	G88B_100_100a	0.0	0.25	1.0	1.0	1.0	0.5	0.5	0.0	0.25	1.0	1.0
27	G00B_037_037a	0.0	0.375	0.0	0.375	0.375	0.187	0.187	0.0	0.375	0.0	0.375
28	G15B_037_037a	0.0	0.375	0.125	0.375	0.375	0.187	0.187	0.0	0.375	0.125	0.125
29	G34B_037_037a	0.0	0.375	0.25	0.375	0.375	0.187	0.187	0.0	0.375	0.25	0.25
30	G50B_037_037a	0.0	0.375	0.375	0.375	0.375	0.187	0.187	0.0	0.375	0.375	0.375
31	G61B_050_050a	0.0	0.375	0.5	0.5	0.5	0.25	0.25	0.0	0.375	0.5	0.5
32	G69B_062_062a	0.0	0.375	0.625	0.625	0.625	0.312	0.312	0.0	0.375	0.625	0.625
33	G75B_075_075a	0.0	0.375	0.75	0.75	0.75	0.375	0.375	0.0	0.375	0.75	0.75
34	G79B_087_087a	0.0	0.375	0.875	0.875	0.875	0.437	0.437	0.0	0.375	0.875	0.875
35	G81B_100_100a	0.0	0.375	1.0	1.0	1.0	0.5	0.5	0.0	0.375	1.0	1.0
36	G00B_050_050a	0.0	0.5	0.0	0.5	0.5	0.25	0.25	0.0	0.5	0.0	0.5
37	G11B_050_050a	0.0	0.5	0.125	0.5	0.5	0.25	0.25	0.0	0.5	0.125	0.125
38	G25B_050_050a	0.0	0.5	0.25	0.5	0.5	0.25	0.25	0.0	0.5	0.25	0.25
39	G38B_050_050a	0.0	0.5	0.375	0.5	0.5	0.25	0.25	0.0	0.5	0.375	0.375
40	G50B_050_050a	0.0	0.5	0.5	0.5	0.5	0.25	0.25	0.0	0.5	0.5	0.5
41	G59B_062_062a	0.0	0.5	0.625	0.625	0.625	0.312	0.312	0.0	0.5	0.625	0.625
42	G65B_075_075a	0.0	0.5	0.75	0.75	0.75	0.375	0.375	0.0	0.5	0.75	0.75
43	G70B_087_087a	0.0	0.5	0.875	0.875	0.875	0.437	0.437	0.0	0.5	0.875	0.875
44	G75B_100_100a	0.0	0.5	1.0	1.0	1.0	0.5	0.5	0.0	0.5	1.0	1.0
45	G00B_062_062a	0.0	0.625	0.0	0.625	0.625	0.312	0.312	0.0	0.625	0.0	0.625
46	G09B_062_062a	0.0	0.625	0.125	0.625	0.625	0.312	0.312	0.0	0.625	0.125	0.125
47	G19B_062_062a	0.0	0.625	0.25	0.625	0.625	0.312	0.312	0.0	0.625	0.25	0.25
48	G30B_062_062a	0.0	0.625	0.375	0.625	0.625	0.312	0.312	0.0	0.625	0.375	0.375
49	G40B_062_062a	0.0	0.625	0.5	0.625	0.625	0.312	0.312	0.0	0.625	0.5	0.5
50	G50B_062_062a	0.0	0.625	0.625	0.625	0.625	0.312	0.312	0.0	0.625	0.625	0.625
51	G57B_075_075a	0.0	0.625	0.75	0.75	0.75	0.375	0.375	0.0	0.625	0.75	0.75
52	G63B_087_087a	0.0	0.625	0.875	0.875	0.875	0.437	0.437	0.0	0.625	0.875	0.875
53	G68B_100_100a	0.0	0.625	1.0	1.0	1.0	0.5	0.5	0.0	0.625	1.0	1.0
54	G00B_075_075a	0.0	0.75	0.0	0.75	0.75	0.375	0.375	0.0	0.75	0.0	0.75
55	G07B_075_075a	0.0	0.75	0.125	0.75	0.75	0.375	0.375	0.0	0.75	0.125	0.125
56	G15B_075_075a	0.0	0.75	0.25	0.75	0.75	0.375	0.375	0.0	0.75	0.25	0.25
57	G25B_075_075a	0.0	0.75	0.375	0.75	0.75	0.375	0.375	0.0	0.75	0.375	0.375
58	G34B_075_075a	0.0	0.75	0.5	0.75	0.75	0.375	0.375	0.0	0.75	0.5	0.5
59	G42B_075_075a	0.0	0.75	0.625	0.75	0.75	0.375	0.375	0.0	0.75	0.625	0.625
60	G50B_075_075a	0.0	0.75	0.75	0.75	0.75	0.375	0.375	0.0	0.75	0.75	0.75
61	G56B_087_087a	0.0	0.75	0.875	0.875	0.875	0.437	0.437	0.0	0.75	0.875	0.875
62	G61B_100_100a	0.0	0.75	1.0	1.0	1.0	0.5	0.5	0.0	0.75	1.0	1.0
63	G00B_087_087a	0.0	0.875	0.0	0.875	0.875	0.437	0.437	0.0	0.875	0.0	0.875
64	G06B_087_087a	0.0	0.875	0.125	0.875	0.875	0.437	0.437	0.0	0.875	0.125	0.125
65	G13B_087_087a	0.0	0.875	0.25	0.875	0.875	0.437	0.437	0.0	0.875	0.25	0.25
66	G20B_087_087a	0.0	0.875	0.375	0.875	0.875	0.437	0.437	0.0	0.875	0.375	0.375
67	G29B_087_087a	0.0	0.875	0.5	0.875	0.875	0.437	0.437	0.0	0.875	0.5	0.5
68	G36B_087_087a	0.0	0.875	0.625	0.875	0.875	0.437	0.437	0.0	0.875	0.625	0.625
69	G43B_087_087a	0.0	0.875	0.75	0.875	0.875	0.437	0.437	0.0	0.875	0.75	0.75
70	G50B_087_087a	0.0	0.875	0.875	0.875	0.875	0.437	0.437	0.0	0.875	0.875	0.875
71	G55B_100_100a	0.0	0.875	1.0	1.0	1.0	0.5	0.5	0.0	0.875	1.0	1.0
72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	0.5	0.0	1.0	0.0	1.0
73	G05B_100_100a	0.0	1.0	0.125	1.0	1.0	0.5	0.5	0.0	1.0	0.125	0.125
74	G11B_100_100a	0.0	1.0	0.25	1.0	1.0	0.5	0.5	0.0	1.0	0.25	0.25
75	G18B_100_100a	0.0	1.0	0.375	1.0	1.0	0.5	0.5	0.0	1.0	0.375	0.375
76	G25B_100_100a	0.0	1.0	0.5	1.0	1.0	0.5	0.5	0.0	1.0	0.5	0.5
77	G31B_100_100a	0.0	1.0	0.625	1.0	1.0	0.5	0.5	0.0	1.0	0.625	0.625
78	G38B_100_100a	0.0	1.0	0.75	1.0	1.0	0.5	0.5	0.0	1.0	0.75	0.75
79	G44B_100_100a	0.0	1.0	0.875	1.0	1.0	0.5	0.5	0.0	1.0	0.875	0.875
80	G50B_100_100a	0.0	1.0	1.0	1.0	1.0	0.5	0.5	0.0	1.0	1.0	1.0

2-0031230-F0

PS790-7N, 13/26-F

PE4300P_120901.TXT, 1080 colors, Separation cmykn6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
colores y diferencia en color, ΔE^* , 3D=0, de=0, cmyk

entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmyk_d$

2-0031230-F0

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)
TUB material: code=rh4ta

delta E* = 10.8

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd		
81	R00Y_012_012a	0.125 0.0 0.0	0.125 0.125 0.062	390	0.125 0.0 0.0	26.8 7.1 4.7	8.5 33.4	0.125 0.0 0.0	24.7 7.7	8.0 11.1	45.9 3.9	389
82	B50R_012_012a	0.125 0.0 0.125	0.125 0.125 0.062	330	0.125 0.0 0.125	26.8 8.1 -1.5	8.3 348.9	0.125 0.0 0.125	26.6 12.3	-6.3 13.8	332.7 6.3	330
83	B25R_025_025a	0.125 0.0 0.25	0.25 0.25 0.125	300	0.125 0.0 0.25	27.1 10.7 -7.7	13.2 324.4	0.125 0.0 0.25	25.7 13.6	-17.5 22.2	307.9 10.3	300
84	B15R_037_037a	0.125 0.0 0.375	0.375 0.375 0.187	289	0.118 0.0 0.375	27.1 13.0 -13.9	19.1 312.9	0.125 0.0 0.375	27.6 14.3	-24.2 28.1	300.5 10.3	288
85	B11R_050_050a	0.125 0.0 0.5	0.5 0.5 0.25	284	0.116 0.0 0.5	27.4 14.8 -19.9	24.8 306.6	0.125 0.0 0.5	29.0 21.5	-32.8 39.2	303.2 14.6	282
86	B09R_062_062a	0.125 0.0 0.625	0.625 0.625 0.312	281	0.114 0.0 0.625	28.5 16.8 -25.6	30.6 303.2	0.125 0.0 0.625	29.5 25.2	-36.3 44.2	304.7 13.6	279
87	B07R_075_075a	0.125 0.0 0.75	0.75 0.75 0.375	279	0.112 0.0 0.75	29.5 18.7 -31.3	36.5 300.9	0.125 0.0 0.75	30.6 25.4	-39.9 47.4	302.5 11.0	278
88	B06R_087_087a	0.125 0.0 0.875	0.875 0.875 0.437	278	0.110 0.0 0.875	30.6 21.0 -36.8	42.4 299.8	0.125 0.0 0.875	30.8 25.7	-41.3 48.7	301.9 6.5	277
89	B05R_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	31.6 23.1 -42.4	48.3 298.6	0.125 0.0 1.0	31.6 23.6	-42.2 48.4	299.2 0.4	276
90	Y00G_012_012a	0.125 0.125 0.0	0.125 0.125 0.062	90	0.125 0.125 0.0	32.3 -1.9 10.5	10.7 100.5	0.125 0.125 0.0	32.9 -5.2	16.0 16.9	108.2 6.4	89
91	NW_012a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	32.8 0.0 0.0	0.0 0.0	0.0 0.125 0.125 0.125	27.8 0.0	-0.5 0.5	273.6 5.0	360
92	B00R_025_012a	0.125 0.125 0.25	0.25 0.125 0.187	270	0.124 0.124 0.25	33.9 2.1 -5.5	5.9 290.8	0.125 0.125 0.25	27.8 2.1	-17.8 17.9	276.9 13.6	270
93	B00R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.124 0.375	35.0 4.2 -11.1	11.9 290.8	0.125 0.125 0.375	30.4 4.1	-25.6 25.9	279.1 15.1	270
94	B00R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.124 0.5	36.1 6.3 -16.7	17.8 290.8	0.125 0.125 0.5	33.6 7.2	-30.1 31.0	283.5 13.6	270
95	B00R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	37.2 8.4 -22.3	23.8 290.8	0.125 0.125 0.625	36.0 11.5	-35.5 37.4	287.9 13.6	270
96	B00R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	38.2 10.5 -27.8	29.8 290.8	0.125 0.125 0.75	37.8 12.8	-40.0 42.0	287.7 12.4	270
97	B00R_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	39.3 12.7 -33.4	35.7 290.8	0.125 0.125 0.875	35.7 15.8	-43.8 46.5	289.8 11.4	270
98	B00R_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	40.4 14.8 -39.0	41.7 290.8	0.125 0.125 1.0	34.1 17.5	-43.8 47.2	291.8 8.4	270
99	Y50G_025_025a	0.125 0.25 0.0	0.25 0.25 0.125	120	0.125 0.25 0.0	35.6 -10.4 13.7	17.2 127.3	0.125 0.25 0.0	36.4 -13.3 17.0	21.6 127.9	4.4 119	0.5
100	G00B_025_012a	0.125 0.25 0.125	0.25 0.125 0.187	150	0.124 0.25 0.124	36.6 -8.4 3.8	9.2 155.5	0.125 0.25 0.125	37.3 -12.7 1.8	12.8 121.8	4.7 149	0.5
101	G50B_025_012a	0.125 0.25 0.25	0.25 0.125 0.187	210	0.124 0.25 0.25	36.5 -3.7 -5.3	6.5 235.1	0.125 0.25 0.25	34.1 -10.5 -16.7	19.7 237.8	13.4 210	0.0
102	G75B_037_025a	0.125 0.25 0.375	0.375 0.25 0.25	240	0.124 0.25 0.375	38.4 -3.3 -12.3	12.7 254.9	0.125 0.25 0.375	36.2 -6.8 -24.4	25.4 254.3	12.8 240	0.0
103	G84B_050_037a	0.125 0.25 0.5	0.5 0.375 0.312	251	0.124 0.243 0.5	38.6 -0.8 -18.4	18.4 267.3	0.125 0.25 0.5	37.9 -3.1 -29.4	29.6 263.8	11.2 251	0.0
104	G88B_062_050a	0.125 0.25 0.625	0.625 0.5 0.375	256	0.125 0.241 0.625	39.2 1.6 -24.1	24.2 273.8	0.125 0.25 0.625	40.5 -0.2 -32.8	32.8 269.6	8.9 257	0.0
105	G90B_075_062a	0.125 0.25 0.75	0.75 0.625 0.437	259	0.125 0.239 0.75	40.3 3.8 -29.7	29.9 277.3	0.125 0.25 0.75	41.5 2.5 -38.3	38.4 273.7	8.8 260	0.0
106	G92B_087_075a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.237 0.875	41.5 6.0 -35.1	35.6 279.6	0.125 0.25 0.875	37.9 7.9	-45.5 46.2	279.9 11.1	262
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.241 1.0	42.7 7.8 -40.7	41.4 280.8	0.125 0.25 1.0	36.4 10.6	-45.2 46.4	283.2 8.2	262
108	Y68G_037_037a	0.125 0.375 0.0	0.375 0.375 0.187	131	0.118 0.375 0.0	38.8 -19.5 16.7	25.7 139.4	0.125 0.375 0.0	39.5 -30.9 19.8	36.7 147.2	11.7 131	0.316
109	G00B_037_025a	0.125 0.375 0.125	0.375 0.25 0.25	150	0.124 0.375 0.124	40.4 -16.9 7.7	18.5 155.5	0.125 0.375 0.125	39.1 -24.1 4.4	24.5 169.4	8.0 149	0.0
110	G25B_037_025a	0.125 0.375 0.25	0.375 0.25 0.25	180	0.124 0.375 0.25	40.6 -12.8 -2.2	13.0 189.8	0.125 0.375 0.25	37.7 -18.6 -9.6	20.9 207.3	9.8 180	0.0
111	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	40.1 -7.5 -10.7	13.1 235.1	0.125 0.375 0.375	41.9 -14.5 -23.1	27.3 237.7	14.3 210	0.0
112	G65B_050_037a	0.125 0.375 0.5	0.5 0.375 0.312	229	0.124 0.381 0.5	43.2 -8.7 -18.2	20.2 244.5	0.125 0.375 0.5	42.5 -11.5 -28.8	31.0 248.1	10.9 228	0.0
113	G75B_062_050a	0.125 0.375 0.625	0.625 0.5 0.375	240	0.125 0.375 0.625	44.0 -6.6 -24.7	25.5 254.9	0.125 0.375 0.625	43.5 -8.6 -31.7	32.8 254.7	7.3 240	0.0
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.364 0.75	44.0 -4.2 -30.8	31.1 262.1	0.125 0.375 0.75	44.9 -6.0 -36.4	36.9 260.5	5.9 247	0.0
115	G84B_087_075a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.362 0.875	44.4 -1.7 -36.8	36.8 267.3	0.125 0.375 0.875	43.8 -4.2 -45.9	46.1 264.7	9.4 251	0.0
116	G86B_100_087a	0.125 0.375 1.0	1.0 0.875 0.562	254	0.125 0.358 1.0	44.7 0.9 -42.6	42.6 271.5	0.125 0.375 1.0	39.2 1.2	-47.2 47.3	271.4 7.2	255
117	Y76G_050_050a	0.125 0.5 0.0	0.5 0.5 0.25	136	0.116 0.5 0.0	42.0 -28.9 19.8	35.1 145.5	0.125 0.5 0.0	41.4 -44.7 24.0	50.7 151.6	16.3 137	0.233
118	G00B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.124	44.2 -25.3 11.5	27.8 155.5	0.125 0.5 0.125	42.7 -31.6 8.9	32.8 164.2	6.9 149	0.0
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.243	44.2 -22.5 2.1	22.6 174.6	0.125 0.5 0.25	43.1 -29.5 -2.1	29.5 184.0	8.2 168	0.0
120	G34B_050_037a	0.125 0.5 0.375	0.5 0.375 0.312	191	0.124 0.5 0.381	44.6 -15.8 -8.8	18.0 209.1	0.125 0.5 0.375	44.4 -22.2 -17.9	28.6 218.8	11.2 191	0.0
121	G50B_050_037a	0.125 0.5 0.5	0.5 0.375 0.312	210	0.124 0.5 0.5	43.8 -11.2 -16.1	19.6 235.1	0.125 0.5 0.5	45.7 -17.6 -27.9	33.0 237.6	13.5 210	0.0
122	G61B_062_050a	0.125 0.5 0.625	0.625 0.5 0.375	224	0.125 0.508 0.625	47.3 -13.1 -23.6	26.9 240.9	0.125 0.5 0.625	47.3 -14.4 -32.2	35.2 245.8	8.7 222	0.0
123	G69B_075_062a	0.125 0.5 0.75	0.75 0.625 0.437	233	0.125 0.51 0.75	49.3 -12.6 -30.9	33.4 247.7	0.125 0.5 0.75	47.8 -13.4 -35.8	38.3 249.4	5.1 232	0.0
124	G75B_087_075a	0.125 0.5 0.875	0.875 0.75 0.5	240	0.125 0.5 0.875	49.5 -9.9 -37.0	38.3 254.9	0.125 0.5 0.875	48.0 -12.0 -43.4	45.0 254.4	6.8 240	0.0
125	G79B_100_087a	0.125 0.5 1.0	1.0 0.875 0.562	245	0.125 0.489 1.0	49.6 -7.5 -43.2	43.9 260.0	0.125 0.5 1.0	44.3 -8.4 -48.1	48.9 260.0	7.2 245	0.0
126	Y81G_062_062a	0.125 0.625 0.0	0.625 0.625 0.312	139	0.114 0.625 0.0	45.6 -37.4 23.3	44.1 148.1	0.125 0.625 0.0	47.0 -55.6 28.1	62.3 153.1	18.8 140	0.183
127	G00B_062_050a	0.125 0.625 0.125	0.625 0.5 0.375	150	0.125 0.625 0.125	48.0 -33.8 15.4	37.1 155.5	0.125 0.625 0.125	47.4 -40.2 14.8	42.9 157.6	8.4 149	0.0
128	G11B_062_050a	0.125 0.625 0.25	0.625 0.5 0.375	164	0.125 0.625 0.241	47.8 -31.8 7.0	32.6 167.4	0.125 0.625 0.25	47.8 -37.6 3.8	37.8 174.1	6.6 162	0.0
129	G25B_062_050a	0.125 0.625 0.375	0.625 0.5 0.375	180	0.125 0.625 0.375	48.4 -25.7 -4.4	26.1 189.8	0.125 0.625 0.375	49.3 -31.0 -11.2	33.0 199.9	8.6 180	0.0
130	G38B_062_050a	0.125 0.625 0.5	0.625 0.5 0.375	196	0.125 0.625 0.508	48.4 -19.6 -13.9	24.0 215.4	0.125 0.625 0.5	50.1 -24.7 -22.9	33.7 222.8	10.4 197	0.0
131	G50B_062_050a	0.125 0.625 0.625	0.625 0.5 0.375	210	0.125 0.625 0.625	47.4 -15.0 -21.5	26.2 235.1	0.125 0.625 0.625	50.9 -2			

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	
162	R00Y_025_025a	0.25 0.0 0.0	0.25 0.25 0.125	390	0.25 0.0 0.0	29.7 14.3 9.4	17.1 33.4	0.25 0.0 0.0	26.6 14.4 12.1	18.9 40.1 4.1	389
163	R00Y_025_025a	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.125	29.8 14.7 2.6	14.9 10.0	0.25 0.0 0.125	26.4 17.6 2.2	17.7 7.1 4.4	360
164	B50R_025_025a	0.25 0.0 0.25	0.25 0.25 0.125	330	0.25 0.0 0.25	29.9 16.3 -3.1	16.6 348.9	0.25 0.0 0.25	27.9 22.7 -10.0	24.8 336.0 9.6	330
165	B34R_037_037a	0.25 0.0 0.375	0.375 0.375 0.187	311	0.256 0.0 0.375	30.0 19.3 -9.2	21.4 334.4	0.25 0.0 0.375	28.8 24.9 -17.6	30.5 324.6 10.1	311
166	B25R_050_050a	0.25 0.0 0.5	0.5 0.5 0.25	300	0.25 0.0 0.5	30.5 21.5 -15.4	26.5 324.4	0.25 0.0 0.5	28.9 25.7 -25.1	35.9 315.6 10.6	300
167	B19R_062_062a	0.25 0.0 0.625	0.625 0.625 0.312	293	0.239 0.0 0.625	30.4 24.1 -21.7	32.4 317.9	0.25 0.0 0.625	29.8 28.8 -31.0	42.3 312.9 10.4	292
168	B15R_075_075a	0.25 0.0 0.75	0.75 0.75 0.375	289	0.237 0.0 0.75	30.5 26.0 -27.9	38.2 312.9	0.25 0.0 0.75	31.0 30.1 -35.2	46.3 310.5 8.3	288
169	B13R_087_087a	0.25 0.0 0.875	0.875 0.875 0.437	286	0.233 0.0 0.875	30.5 27.6 -34.0	43.8 309.1	0.25 0.0 0.875	31.5 30.7 -38.1	48.9 308.8 5.2	284
170	B11R_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	31.1 29.6 -39.8	49.6 306.6	0.25 0.0 1.0	31.0 30.5 -39.3	49.8 307.8 1.0	282
171	R50Y_025_025a	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.125 0.0	35.5 4.8 16.5	17.2 73.8	0.25 0.125 0.0	32.1 3.1 19.8	20.0 81.0 5.0	59
172	R00Y_025_012a	0.25 0.125 0.125	0.25 0.125 0.187	390	0.25 0.124 0.124	35.8 7.1 4.7	8.5 33.4	0.25 0.125 0.125	30.7 8.8 7.3	11.5 39.8 5.9	389
173	B50R_025_012a	0.25 0.125 0.25	0.25 0.125 0.187	330	0.25 0.124 0.25	35.8 8.1 -1.5	8.3 348.9	0.25 0.125 0.25	32.9 13.4 -8.4	15.9 327.8 9.1	330
174	B25R_037_025a	0.25 0.125 0.375	0.375 0.25 0.25	300	0.25 0.124 0.375	36.1 10.7 -7.7	13.2 324.4	0.25 0.125 0.375	33.3 13.6 -17.6	22.3 307.5 10.7	300
175	B15R_050_037a	0.25 0.125 0.5	0.5 0.375 0.312	289	0.243 0.124 0.5	36.1 13.0 -13.9	19.1 312.9	0.25 0.125 0.5	34.1 15.5 -25.6	30.0 301.2 12.1	288
176	B11R_062_050a	0.25 0.125 0.625	0.625 0.5 0.375	284	0.241 0.125 0.625	36.4 14.8 -19.9	24.8 306.6	0.25 0.125 0.625	37.0 18.0 -31.1	36.0 300.1 11.7	282
177	B09R_075_062a	0.25 0.125 0.75	0.75 0.625 0.437	281	0.239 0.125 0.75	37.5 16.8 -25.6	30.6 303.2	0.25 0.125 0.75	38.2 19.8 -36.3	41.4 298.5 11.1	279
178	B07R_087_075a	0.25 0.125 0.875	0.875 0.75 0.5	279	0.237 0.125 0.875	38.5 18.7 -31.3	36.5 300.9	0.25 0.125 0.875	35.5 24.3 -40.4	47.1 301.0 11.1	278
179	B06R_100_087a	0.25 0.125 1.0	1.0 0.875 0.562	278	0.241 0.125 1.0	39.6 21.0 -36.8	42.4 299.8	0.25 0.125 1.0	34.9 24.2 -40.6	47.3 300.7 6.8	277
180	Y00G_025_025a	0.25 0.25 0.0	0.25 0.25 0.125	90	0.25 0.25 0.0	40.7 -3.9 21.1	21.5 100.5	0.25 0.25 0.0	39.4 -7.8 29.7	30.8 104.8 9.5	89
181	Y00G_025_012a	0.25 0.25 0.125	0.25 0.125 0.187	90	0.25 0.25 0.124	41.3 -1.9 10.5	10.7 100.5	0.25 0.25 0.125	39.8 -5.3 15.2	16.1 109.2 5.8	89
182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	41.8 0.0 0.0	0.0 0.0	0.25 0.25 0.25	39.8 -0.1 -1.2	1.2 265.1 2.3	360
183	B00R_037_012a	0.25 0.25 0.375	0.375 0.125 0.312	270	0.249 0.249 0.375	42.9 2.1 -5.5	5.9 290.8	0.25 0.25 0.375	37.2 2.3 -18.7	18.9 276.9 14.3	270
184	B00R_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.249 0.5	44.0 4.2 -11.1	11.9 290.8	0.25 0.25 0.5	39.7 5.0 -25.0	25.4 281.3 14.5	270
185	B00R_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	45.1 6.3 -16.7	17.8 290.8	0.25 0.25 0.625	41.8 8.2 -29.3	30.4 285.7 13.1	270
186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	46.2 8.4 -22.3	23.8 290.8	0.25 0.25 0.75	43.1 11.1 -34.5	36.3 287.8 12.9	270
187	B00R_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	47.2 10.5 -27.8	29.8 290.8	0.25 0.25 0.875	44.2 15.3 -41.4	44.1 290.2 15.9	270
188	B00R_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	48.3 12.7 -33.4	35.7 290.8	0.25 0.25 1.0	38.0 16.5 -42.0	45.1 291.5 14.0	270
189	Y31G_037_037a	0.25 0.375 0.0	0.375 0.375 0.187	109	0.256 0.375 0.0	46.6 -10.3 28.7	30.5 109.8	0.25 0.375 0.0	41.6 -14.3 30.8	34.0 114.9 6.7	108
190	Y50G_037_025a	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	44.6 -10.4 13.7	17.2 127.3	0.25 0.375 0.125	44.8 -13.9 15.6	20.9 131.6 4.0	119
191	G00B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	45.6 -8.4 3.8	9.2 155.5	0.25 0.375 0.25	43.9 -11.7 -0.1	11.7 180.5 5.4	149
192	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	45.5 -3.7 -5.3	6.5 235.1	0.25 0.375 0.375	44.7 -10.3 -17.0	19.9 238.7 13.3	210
193	G75B_050_025a	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.375 0.5	47.4 -3.3 -12.3	12.7 254.9	0.25 0.375 0.5	46.4 -5.5 -23.0	23.6 256.5 10.9	240
194	G84B_062_037a	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.368 0.625	47.6 -0.8 -18.4	18.4 267.3	0.25 0.375 0.625	46.8 -2.5 -27.6	27.7 264.7 9.4	251
195	G88B_075_050a	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.366 0.75	48.2 1.6 -24.1	24.2 273.8	0.25 0.375 0.75	46.6 0.6 -33.1	33.1 271.0 9.1	257
196	G90B_087_062a	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.364 0.875	49.3 3.8 -29.7	29.9 277.3	0.25 0.375 0.875	45.7 4.1 -41.8	42.1 275.7 12.7	260
197	G92B_100_075a	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.362 1.0	50.5 6.0 -35.1	35.6 279.6	0.25 0.375 1.0	42.5 7.0 -42.9	43.5 279.3 11.2	262
198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	47.4 -20.8 27.4	34.4 127.3	0.25 0.5 0.0	43.9 -28.1 32.6	43.1 130.7 9.6	119
199	Y68G_050_037a	0.25 0.5 0.125	0.5 0.375 0.312	131	0.243 0.5 0.124	47.8 -19.5 16.7	25.7 139.4	0.25 0.5 0.125	47.6 -24.8 16.6	29.9 146.1 5.2	131
200	G00B_050_025a	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.249	49.4 -16.9 7.7	18.5 155.5	0.25 0.5 0.25	46.4 -22.2 3.8	22.6 170.1 7.2	149
201	G25B_050_025a	0.25 0.5 0.375	0.5 0.25 0.375	180	0.249 0.5 0.375	49.6 -12.8 -2.2	13.0 189.8	0.25 0.5 0.375	48.2 -16.8 -9.9	19.5 210.3 8.7	180
202	G50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	49.1 -7.5 -10.7	13.1 235.1	0.25 0.5 0.5	50.5 -14.1 -21.4	25.7 236.7 12.6	210
203	G65B_062_037a	0.25 0.5 0.625	0.625 0.375 0.437	229	0.25 0.506 0.625	52.2 -8.7 -18.2	20.2 244.5	0.25 0.5 0.625	51.2 -10.1 -27.6	29.4 249.8 9.5	228
204	G75B_075_050a	0.25 0.5 0.75	0.75 0.5 0.5	240	0.25 0.5 0.75	53.0 -6.6 -24.7	25.5 254.9	0.25 0.5 0.75	51.3 -8.5 -31.4	32.6 254.7 7.2	240
205	G80B_087_062a	0.25 0.5 0.875	0.875 0.625 0.562	247	0.25 0.489 0.875	53.0 -4.2 -30.8	31.1 262.1	0.25 0.5 0.875	51.1 -6.6 -39.5	40.0 260.4 9.2	247
206	G84B_100_075a	0.25 0.5 1.0	1.0 0.75 0.625	251	0.25 0.487 1.0	53.4 -1.7 -36.8	36.8 267.3	0.25 0.5 1.0	45.0 -0.5 -44.4	44.4 269.3 11.3	251
207	Y61G_062_062a	0.25 0.625 0.0	0.625 0.625 0.312	127	0.239 0.625 0.0	50.7 -29.9 30.3	42.3 134.2	0.25 0.625 0.0	49.0 -39.1 34.5	52.1 138.5 10.6	127
208	Y76G_062_050a	0.25 0.625 0.125	0.625 0.5 0.375	136	0.241 0.625 0.125	51.0 -28.9 19.8	35.1 145.5	0.25 0.625 0.125	51.3 -33.6 22.4	40.4 146.3 5.3	137
209	G00B_062_037a	0.25 0.625 0.25	0.625 0.375 0.437	150	0.25 0.625 0.25	53.2 -25.3 11.5	27.8 155.5	0.25 0.625 0.25	51.5 -29.9 9.9	31.5 161.5 5.1	149
210	G15B_062_037a	0.25 0.625 0.375	0.625 0.375 0.437	169	0.25 0.625 0.368	53.2 -22.5 2.1	22.6 174.6	0.25 0.625 0.375	53.7 -24.9 -3.1	25.1 187.5 5.7	168
211	G34B_062_037a	0.25 0.625 0.5	0.625 0.375 0.437	191	0.25 0.625 0.506	53.6 -11.8 -8.8	18.0 209.1	0.25 0.625 0.5	54.8 -20.1 -15.5	25.4 217.5 8.0	191
212	G50B_062_037a	0.25 0.625 0.625	0.625 0.375 0.437	210	0.25 0.625 0.625	52.8 -11.2 -16.1	19.6 235.1	0.25 0.625 0.625	54.2 -16.5 -26.0	30.8 237.5 11.2	210
213	G61B_075_050a	0.25 0.625 0.75	0.75 0.5 0.5	224	0.25 0.633 0.75	56.3 -13.1 -23.6	26.9 240.9	0.25 0.625 0.75	55.1 -13.9 -31.3	34.2 245.9 7.8	222
214	G69B_087_062a	0.25 0.625 0.875	0.875 0.625 0.562	233	0.25 0.635 0.875	58.3 -12.6 -30.9	33.4 247.7	0.25 0.625 0.875	55.5 -12.8 -36.5	38.7 250.6 6.3	232
215	G75B_100_075a	0.25 0.625 1.0	1.0 0.75 0.625	240	0.25 0.625 1.0	58.5 -9.9 -37.0	38.3 254.9	0.25 0.625 1.0	52.2 -11.8 -43.9	45.5 254.8 9.5	240
216											

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información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md	
243	R00Y_037_037a	0.375 0.0 0.0	0.375 0.375 0.187	390	0.375 0.0 0.0	32.7 21.4 14.1	25.7 33.4	0.375 0.0 0.0	31.0 26.1 19.8	32.8 37.1 7.5	389	1.0 0.0 0.0	47.5 57.2 37.8 68.6 33.4
244	R18Y_037_037a	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.118	32.7 21.2 8.7	22.9 22.3	0.375 0.0 0.125	33.0 25.3 9.6	27.1 20.9 4.2	371	1.0 0.0 0.316	47.4 56.5 23.2 61.1 22.3
245	B65R_037_037a	0.375 0.0 0.25	0.375 0.375 0.187	349	0.375 0.0 0.256	33.1 23.7 -0.6	23.7 358.3	0.375 0.0 0.25	32.8 30.2 -2.5	30.3 355.1 6.8	348	1.0 0.0 0.683	48.6 63.2 -1.8 63.2 358.3
246	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	32.9 24.5 -4.7	24.9 348.9	0.375 0.0 0.375	33.5 35.7 -11.0	37.4 342.8 12.8	330	1.0 0.0 1.0	48.1 65.4 -12.7 66.6 348.9
247	B38R_050_050a	0.375 0.0 0.5	0.5 0.5 0.25	316	0.383 0.0 0.5	33.1 27.9 -10.4	29.8 339.4	0.375 0.0 0.5	32.8 37.7 -16.7	41.2 336.0 11.6	317	0.766 0.0 1.0	42.4 55.8 -20.9 59.6 339.4
248	B30R_062_062a	0.375 0.0 0.625	0.625 0.625 0.312	307	0.385 0.0 0.625	33.3 30.0 -17.2	34.6 330.2	0.375 0.0 0.625	32.2 37.2 -24.4	44.5 326.7 10.2	307	0.616 0.0 1.0	38.9 48.1 -27.5 55.4 330.2
249	B25R_075_075a	0.375 0.0 0.75	0.75 0.75 0.375	300	0.375 0.0 0.75	33.8 32.3 -23.1	39.8 324.4	0.375 0.0 0.75	33.6 38.8 -29.6	48.6 322.4 9.0	300	0.5 0.0 1.0	37.2 43.1 -30.8 53.0 324.4
250	B20R_087_087a	0.375 0.0 0.875	0.875 0.875 0.437	295	0.364 0.0 0.875	33.8 34.9 -29.5	45.7 319.8	0.375 0.0 0.875	34.5 38.5 -33.1	51.0 319.5 5.3	294	0.416 0.0 1.0	35.2 39.9 -33.7 52.2 319.8
251	B18R_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	34.0 37.7 -35.3	51.7 316.8	0.375 0.0 1.0	34.2 38.2 -35.0	51.8 317.5 0.6	291	0.366 0.0 1.0	34.0 37.7 -35.3 51.7 316.8
252	R31Y_037_037a	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.118 0.0	38.0 13.3 21.8	25.5 58.6	0.375 0.125 0.0	36.3 11.9 25.7	28.3 65.0 4.4	48	1.0 0.316 0.0	61.6 35.5 58.2 68.2 58.6
253	R00Y_037_025a	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.124	38.7 14.3 9.4	17.1 33.4	0.375 0.125 0.125	38.6 13.7 16.1	21.2 49.5 6.7	389	1.0 0.0 0.0	47.5 57.2 37.8 68.6 33.4
254	R00Y_037_025a	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.25	38.8 14.7 2.6	14.9 10.0	0.375 0.125 0.25	37.5 19.2 1.8	19.3 5.6 4.7	360	1.0 0.0 0.5	47.8 58.9 10.4 59.9 10.0
255	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.124 0.375	38.9 16.3 -3.1	16.6 348.9	0.375 0.125 0.375	38.9 24.5 -10.7	26.7 336.2 11.1	330	1.0 0.0 1.0	48.1 65.4 -12.7 66.6 348.9
256	B34R_050_037a	0.375 0.125 0.5	0.5 0.375 0.312	311	0.381 0.124 0.5	39.0 19.3 -9.2	21.4 334.4	0.375 0.125 0.5	38.4 26.0 -16.9	31.0 326.8 10.2	311	0.683 0.0 1.0	40.4 51.6 -24.7 57.2 334.4
257	B25R_062_050a	0.375 0.125 0.625	0.625 0.5 0.375	300	0.375 0.125 0.625	39.5 21.5 -15.4	26.5 324.4	0.375 0.125 0.625	38.1 26.4 -24.3	35.9 317.3 10.2	300	0.5 0.0 1.0	37.2 43.1 -30.8 53.0 324.4
258	B19R_075_062a	0.375 0.125 0.75	0.75 0.625 0.437	293	0.364 0.125 0.75	39.4 24.1 -21.7	32.4 317.9	0.375 0.125 0.75	39.9 26.8 -29.2	39.7 312.5 8.0	292	0.383 0.0 1.0	34.4 38.5 -34.7 51.9 317.9
259	B15R_087_075a	0.375 0.125 0.875	0.875 0.75 0.5	289	0.362 0.125 0.875	39.5 26.0 -27.9	38.2 312.9	0.375 0.125 0.875	36.6 33.1 -35.8	48.7 312.7 10.9	288	0.316 0.0 1.0	32.7 34.7 -37.2 50.9 312.9
260	B13R_100_087a	0.375 0.125 1.0	1.0 0.875 0.562	286	0.358 0.125 1.0	39.5 27.6 -34.0	43.8 309.1	0.375 0.125 1.0	36.6 31.2 -36.5	48.1 310.4 5.2	284	0.266 0.0 1.0	31.4 31.6 -38.8 50.1 309.1
261	R68Y_037_037a	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.256 0.0	44.4 2.0 27.7	27.8 85.7	0.375 0.25 0.0	41.9 0.3 36.3	36.3 89.4 9.1	71	1.0 0.683 0.0	78.6 5.4 73.9 74.1 85.7
262	R50Y_037_025a	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.25 0.124	44.5 4.8 16.5	17.2 73.8	0.375 0.25 0.125	46.0 19.9 22.4	22.4 89.9 6.6	59	1.0 0.5 0.0	70.5 19.2 66.2 69.0 73.8
263	R00Y_037_012a	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.249	44.8 7.1 4.7	8.5 33.4	0.375 0.25 0.25	44.2 8.2 6.2	10.3 37.2 2.0	389	1.0 0.0 0.0	47.5 57.2 37.8 68.6 33.4
264	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.249 0.375	44.8 8.1 -1.5	8.3 348.9	0.375 0.25 0.375	44.6 12.6 -8.8	15.4 324.8 8.5	330	1.0 0.0 1.0	48.1 65.4 -12.7 66.6 348.9
265	B25R_050_025a	0.375 0.25 0.5	0.5 0.25 0.375	300	0.375 0.249 0.5	45.1 10.7 -7.7	13.2 324.4	0.375 0.25 0.5	43.9 14.9 -17.7	23.1 310.0 10.9	300	0.5 0.0 1.0	37.2 43.1 -30.8 53.0 324.4
266	B15R_062_037a	0.375 0.25 0.625	0.625 0.375 0.437	289	0.368 0.25 0.625	45.1 13.0 -13.9	19.1 312.9	0.375 0.25 0.625	43.8 16.7 -24.9	30.0 303.9 11.6	288	0.316 0.0 1.0	32.7 34.7 -37.2 50.9 312.9
267	B11R_075_050a	0.375 0.25 0.75	0.75 0.5 0.5	284	0.366 0.25 0.75	45.4 14.8 -19.9	24.8 306.6	0.375 0.25 0.75	45.3 17.2 -29.4	34.1 300.3 9.8	282	0.233 0.0 1.0	31.1 29.6 -39.8 49.6 306.6
268	B09R_087_062a	0.375 0.25 0.875	0.875 0.625 0.562	281	0.364 0.25 0.875	46.5 16.8 -25.6	30.6 303.2	0.375 0.25 0.875	41.4 23.7 -37.0	43.9 302.7 14.2	279	0.183 0.0 1.0	31.3 26.8 -41.0 49.0 303.2
269	B07R_100_075a	0.375 0.25 1.0	1.0 0.75 0.625	279	0.362 0.25 1.0	47.5 18.7 -31.3	36.5 300.9	0.375 0.25 1.0	40.0 23.6 -38.0	44.7 301.8 11.1	278	0.15 0.0 1.0	31.4 25.0 -41.7 48.6 300.9
270	Y00G_037_037a	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.375 0.0	49.2 -5.9 31.7	32.3 100.5	0.375 0.375 0.0	51.3 -10.5 44.1	45.3 103.3 13.3	89	1.0 1.0 0.0	91.5 -15.8 84.6 86.1 100.5
271	Y00G_037_025a	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.375 0.124	49.7 -3.9 21.1	21.5 100.5	0.375 0.375 0.125	54.5 -9.1 32.2	33.5 105.8 13.1	89	1.0 1.0 0.0	91.5 -15.8 84.6 86.1 100.5
272	Y00G_037_012a	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.375 0.249	50.3 -1.9 10.5	10.7 100.5	0.375 0.375 0.25	52.7 -5.7 13.0	14.2 113.7 5.0	89	1.0 1.0 0.0	91.5 -15.8 84.6 86.1 100.5
273	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	50.8 0.0 0.0	0.0 0.0	0.375 0.375 0.375	51.9 0.0 -1.3	1.3 267.6 1.7	360	1.0 1.0 1.0	95.8 0.0 0.0 0.0 0.0
274	B00R_050_012a	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.375 0.5	51.9 2.1 -5.5	5.9 290.8	0.375 0.375 0.5	49.1 2.6 -17.1	17.3 278.8 11.8	270	0.0 0.0 1.0	32.5 16.9 -44.6 47.7 290.8
275	B00R_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	53.0 4.2 -11.1	11.9 290.8	0.375 0.375 0.625	49.0 5.4 -23.7	24.3 282.9 13.2	270	0.0 0.0 1.0	32.5 16.9 -44.6 47.7 290.8
276	B00R_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	54.1 6.3 -16.7	17.8 290.8	0.375 0.375 0.75	50.1 8.2 -28.1	29.3 286.2 12.2	270	0.0 0.0 1.0	32.5 16.9 -44.6 47.7 290.8
277	B00R_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	55.2 8.4 -22.3	23.8 290.8	0.375 0.375 0.875	48.1 13.7 -36.3	38.8 286.0 16.5	270	0.0 0.0 1.0	32.5 16.9 -44.6 47.7 290.8
278	B00R_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	56.2 10.5 -27.8	29.8 290.8	0.375 0.375 1.0	45.6 15.1 -37.4	40.3 291.9 14.9	270	0.0 0.0 1.0	32.5 16.9 -44.6 47.7 290.8
279	Y23G_050_050a	0.375 0.5 0.0	0.5 0.5 0.25	104	0.383 0.5 0.0	57.1 -10.4 43.2	44.5 103.6	0.375 0.5 0.0	52.5 -14.9 46.5	48.9 107.8 7.2	102	0.766 1.0 0.0	90.4 -20.9 86.5 89.0 103.6
280	Y31G_050_037a	0.375 0.5 0.125	0.5 0.375 0.312	109	0.381 0.5 0.124	55.6 -10.3 28.7	30.5 109.8	0.375 0.5 0.125	58.4 -15.5 32.6	36.1 115.5 7.0	108	0.683 1.0 0.0	84.6 -27.6 76.5 81.3 109.8
281	Y50G_050_025a	0.375 0.5 0.25	0.5 0.25 0.375	120	0.375 0.5 0.249	53.6 -10.4 13.7	17.2 127.3	0.375 0.5 0.25	57.6 -13.8 15.4	20.6 118.8 5.5	119	0.5 1.0 0.0	70.9 -41.7 54.8 68.9 127.3
282	G00B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	54.6 -8.4 3.8	9.2 155.5	0.375 0.5 0.375	57.3 -10.5 1.3	10.6 172.6 4.2	149	0.0 1.0 0.0	54.3 -67.6 30.8 74.3 155.5
283	G50B_050_012a	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.5	54.5 -3.7 -5.3	6.5 235.1	0.375 0.5 0.5	57.8 -8.1 -14.6	16.7 240.7 10.7	210	0.0 1.0 1.0	

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)

TUB material: code=rh4ta

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md
324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.0	0.0	33.0	34.7	23.4	41.8
325	R26Y_050_050a	0.5	0.0	0.125	0.5	0.5	0.116	35.7	28.0	18.9	34.3	33.4
326	R00Y_050_050a	0.5	0.0	0.25	0.5	0.5	0.25	35.8	29.4	5.2	29.9	10.0
327	B61R_050_050a	0.5	0.0	0.375	0.5	0.5	0.383	36.6	32.3	-3.5	32.5	353.7
328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.5	36.0	32.7	-6.3	33.3	348.9
329	B40R_062_062a	0.5	0.0	0.625	0.625	0.625	0.312	31.9	0.51	0.0	0.625	36.3
330	B34R_075_075a	0.5	0.0	0.75	0.75	0.75	0.375	31.1	0.512	0.0	0.75	36.2
331	B29R_087_087a	0.5	0.0	0.875	0.875	0.875	0.437	30.5	0.51	0.0	0.875	36.6
332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	30.0	0.5	0.0	1.0	37.2
333	R23Y_050_050a	0.5	0.125	0.0	0.5	0.5	0.25	44	0.5	0.116	0.0	40.6
334	R00Y_050_037a	0.5	0.125	0.125	0.5	0.375	0.312	390	0.5	0.124	0.124	41.7
335	R18Y_050_037a	0.5	0.125	0.25	0.5	0.375	0.312	371	0.5	0.124	0.243	41.7
336	B65R_050_037a	0.5	0.125	0.375	0.5	0.375	0.312	349	0.5	0.124	0.381	42.1
337	B50R_050_037a	0.5	0.125	0.5	0.5	0.375	0.312	330	0.5	0.124	0.5	41.9
338	B38R_062_050a	0.5	0.125	0.625	0.625	0.5	0.375	316	0.508	0.125	0.625	42.1
339	B30R_075_062a	0.5	0.125	0.75	0.75	0.625	0.437	307	0.51	0.125	0.75	42.3
340	B25R_087_075a	0.5	0.125	0.875	0.875	0.75	0.5	300	0.5	0.125	0.875	42.8
341	B20R_100_087a	0.5	0.125	1.0	1.0	0.875	0.562	295	0.489	0.125	1.0	42.8
342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	60	0.5	0.25	0.0	47.1
343	R31Y_050_037a	0.5	0.25	0.125	0.5	0.375	0.312	49	0.5	0.243	0.124	47.0
344	R00Y_050_025a	0.5	0.25	0.25	0.5	0.25	0.375	390	0.5	0.249	0.249	47.7
345	R00Y_050_025a	0.5	0.25	0.375	0.5	0.25	0.375	360	0.5	0.249	0.375	47.8
346	B50R_050_025a	0.5	0.25	0.5	0.5	0.25	0.375	330	0.5	0.249	0.5	47.9
347	B34R_062_037a	0.5	0.25	0.625	0.625	0.375	0.437	311	0.506	0.25	0.625	48.0
348	B25R_075_050a	0.5	0.25	0.75	0.75	0.5	0.5	300	0.5	0.25	0.75	48.5
349	B19R_087_062a	0.5	0.25	0.875	0.875	0.625	0.562	293	0.489	0.25	0.875	48.4
350	B15R_100_075a	0.5	0.25	1.0	1.0	0.75	0.625	289	0.487	0.25	1.0	48.5
351	R76Y_050_050a	0.5	0.375	0.0	0.5	0.5	0.25	76	0.5	0.383	0.0	53.7
352	R68Y_050_037a	0.5	0.375	0.125	0.5	0.375	0.312	71	0.5	0.381	0.124	53.4
353	R50Y_050_025a	0.5	0.375	0.25	0.5	0.25	0.375	60	0.5	0.375	0.249	53.5
354	R00Y_050_012a	0.5	0.375	0.375	0.5	0.125	0.437	390	0.5	0.375	0.375	53.8
355	B50R_050_012a	0.5	0.375	0.5	0.5	0.125	0.437	330	0.5	0.375	0.5	53.8
356	B25R_062_025a	0.5	0.375	0.625	0.625	0.25	0.5	300	0.5	0.375	0.625	54.1
357	B15R_075_037a	0.5	0.375	0.75	0.75	0.375	0.562	289	0.493	0.375	0.75	54.1
358	B11R_087_050a	0.5	0.375	0.875	0.875	0.5	0.625	284	0.491	0.375	0.875	54.4
359	B09R_100_062a	0.5	0.375	1.0	1.0	0.625	0.687	281	0.489	0.375	1.0	55.5
360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	90	0.5	0.5	0.0	57.7
361	Y00G_050_037a	0.5	0.5	0.125	0.5	0.375	0.312	90	0.5	0.5	0.124	58.2
362	Y00G_050_025a	0.5	0.5	0.25	0.5	0.25	0.375	90	0.5	0.5	0.249	58.7
363	Y00G_050_012a	0.5	0.5	0.375	0.5	0.125	0.437	90	0.5	0.5	0.375	59.3
364	NW_050a	0.5	0.5	0.5	0.5	0.0	0.5	360	0.5	0.5	0.5	59.8
365	B00R_062_012a	0.5	0.625	0.625	0.625	0.125	0.562	270	0.5	0.5	0.625	60.9
366	B00R_075_025a	0.5	0.5	0.75	0.75	0.25	0.625	270	0.5	0.5	0.75	62.0
367	B00R_087_037a	0.5	0.5	0.875	0.875	0.375	0.687	270	0.5	0.5	0.875	63.1
368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	270	0.5	0.5	1.0	64.2
369	Y18G_062_062a	0.5	0.625	0.0	0.625	0.625	0.312	101	0.51	0.625	0.0	66.1
370	Y23G_062_050a	0.5	0.625	0.125	0.625	0.5	0.375	104	0.508	0.625	0.125	66.1
371	Y31G_062_037a	0.5	0.625	0.25	0.625	0.375	0.437	109	0.506	0.625	0.25	66.6
372	Y50G_062_025a	0.5	0.625	0.375	0.625	0.25	0.5	120	0.5	0.625	0.375	62.6
373	G00B_062_012a	0.5	0.625	0.5	0.625	0.125	0.562	150	0.5	0.625	0.5	63.6
374	G50B_062_012a	0.5	0.625	0.625	0.625	0.125	0.562	210	0.5	0.625	0.625	63.5
375	G75B_075_025a	0.5	0.625	0.75	0.75	0.25	0.625	240	0.5	0.625	0.75	65.4
376	G84B_087_037a	0.5	0.625	0.875	0.875	0.375	0.687	251	0.5	0.618	0.875	65.6
377	G88B_100_050a	0.5	0.625	1.0	1.0	0.5	0.75	256	0.5	0.616	1.0	66.2
378	Y31G_075_075a	0.5	0.75	0.0	0.75	0.75	0.375	109	0.512	0.75	0.0	69.4
379	Y38G_075_062a	0.5	0.75	0.125	0.75	0.625	0.437	113	0.51	0.75	0.125	67.5
380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	120	0.5	0.75	0.25	65.4
381	Y68G_075_037a	0.5	0.75	0.375	0.75	0.375	0.562	131	0.493	0.75	0.375	65.8
382	G00B_075_025a	0.5	0.75	0.5	0.75	0.25	0.625	150	0.5	0.75	0.5	65.4
383	G25B_075_025a	0.5	0.75	0.625	0.75	0.25	0.625	180	0.5	0.75	0.625	67.6
384	G50B_075_025a	0.5	0.75	0.75	0.75	0.25	0.625	210	0.5	0.75	0.75	67.1
385	G65B_087_037a	0.5	0.75	0.875	0.875	0.375	0.687	229	0.5	0.756	0.875	70.2
386	G75B_100_050a	0.5	0.75	1.0	1.0	0.5	0.75	240	0.5	0.75	1.0	70.9
387	Y41G_087_087a	0.5	0.875	0.0	0.875	0.75	0.437	115	0.51	0.875	0.0	70.3
388	Y50G_087_075a	0.5	0.875	0.125	0.875	0.75	0.5	120	0.5	0.875	0.125	68.2
389	Y61G_087_062a	0.5	0.875	0.25	0.875	0.625	0.562	127	0.489	0.875	0.25	68.7
390	Y76G_087_050a	0.5	0.875	0.375	0.875	0.5	0.625	136	0.491	0.875	0.375	68.9
391	G00B_087_037a	0.5	0.875	0.5	0.875	0.375	0.687	150	0.5	0.875	0.5	71.2
392	G15B_087_037a	0.5	0.875	0.625	0.875	0.375	0.687	169	0.5	0.875	0.618	71.2
393	G34B_087_037a	0.5	0.875	0.75	0.875	0.375	0.687	191	0.5	0.875	0.75	71.6
394	G50B_087_037a	0.5	0.875	0.875	0.875	0.375	0.687	210	0.5	0.875	0.875	70.8
395	G61B_100_050a	0.5	0.875	1.0	1.0	0.5	0.75	224	0.5	0.883	1.0	74.3
396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	120	0.5	1.0	0.0	70.9
397	Y58G_100_087a	0.5	1.0	0.125	1.0	0.875	0.562	125	0.489	1.0	0.125	71.5
398	Y68G_100_075a	0.5	1.0	0.25	1.0	0.75	0.625	131	0.487	1.0	0.25	71.8
399	Y81G_100_062a	0.5	1.0	0.375	1.0	0.625	0.687	139	0.489	1.0	0.375	72.6
400	G00B_100_050a	0.5	1.0	0.5	1.0	0.5	0.75	150	0.5	1.0	0.5	75.0
401	G11B_100_050a	0.5	1.0	0.625	1.0	0.5	0.75	164	0.5	1.0	0.616	74.8
402	G25B_100_050a	0.5	1.0	0.75	1.0	0.5	0.75	186	0.5	1.0	0.75	75.4
403	G38B_100_050a	0.5	1.0	0.875	1.0	0.5	0.75	190	0.5	1.0	0.883	75.4
404	G50B_100_050a	0.5	1.0	1.0	1.0	0.5	0.75	210	0.5	1.0	1.0	74.4

delta E* = 7.3

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
colores y diferencia en color, ΔE^* , 3D=0, de=0, cmyk

entrada: *rgbcmyk* -> *rgb*_d
salida: transfiera a *cmyk*_d

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)
TUB material: code=rha4ta

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md		
405	R00Y_062_062a	0.625 0.0 0.0	0.625 0.625 0.312	390	0.625 0.0 0.0	38.6 35.7 23.6	42.8 33.4	0.625 0.0 0.0	36.3 40.2 26.2	48.0 33.1 5.6	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
406	R31Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.114	38.7 35.1 19.4	40.1 28.9	0.625 0.0 0.125	37.1 40.1 21.9	45.7 28.6 5.8	380	1.0 0.0 0.183	47.6 56.2 31.1	64.2 28.9
407	R11Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.239	38.5 35.6 11.8	37.5 18.3	0.625 0.0 0.25	36.4 41.6 12.6	43.4 16.8 6.3	367	1.0 0.0 0.383	47.4 57.0 18.9	60.0 18.3
408	B69R_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.385	38.9 38.5 1.6	38.5 2.5	0.625 0.0 0.375	36.9 45.2 0.9	45.2 1.1 6.9	352	1.0 0.0 0.616	47.9 61.6 2.7	67.1 2.5
409	B59R_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.625 0.0 0.51	39.8 40.9 -5.4	41.2 352.3	0.625 0.0 0.5	37.2 49.0 -7.9	49.6 350.7 8.8	339	1.0 0.0 0.816	49.4 65.4 -8.7	66.0 352.3
410	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	39.0 40.8 -7.9	41.6 348.9	0.625 0.0 0.625	37.9 49.8 -14.9	52.0 343.3 11.4	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
411	B42R_075_075a	0.625 0.0 0.75	0.75 0.75 0.375	321	0.637 0.0 0.75	39.6 44.3 -13.6	46.4 342.9	0.625 0.0 0.75	38.9 50.3 -20.0	54.1 338.3 8.7	322	0.85 0.0 1.0	44.9 59.1 -18.2	61.9 342.9
412	B36R_087_087a	0.625 0.0 0.875	0.875 0.875 0.437	314	0.641 0.0 0.875	39.3 47.5 -19.5	51.3 337.6	0.625 0.0 0.875	39.3 49.4 -24.6	55.2 333.4 5.4	315	0.733 0.0 1.0	41.5 54.3 -22.3	58.7 337.6
413	B31R_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	39.2 48.9 -26.9	55.8 331.1	0.625 0.0 1.0	39.1 48.4 -27.2	55.6 330.6 0.6	308	0.633 0.0 1.0	39.2 48.9 -26.9	55.8 331.1
414	R18Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	41	0.625 0.114 0.0	43.2 30.3 32.7	44.6 47.1	0.625 0.125 0.0	40.3 31.8 34.0	46.6 46.8 3.5	39	1.0 0.183 0.0	54.9 48.5 52.3	71.4 47.1
415	R00Y_062_050a	0.625 0.125 0.125	0.625 0.5 0.375	390	0.625 0.125 0.125	44.7 28.6 18.9	34.3 33.4	0.625 0.125 0.125	44.2 28.5 28.7	40.5 45.1 9.8	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
416	R26Y_062_050a	0.625 0.125 0.25	0.625 0.5 0.375	376	0.625 0.125 0.241	44.7 28.0 14.9	31.4 26.9	0.625 0.125 0.25	43.5 29.7 15.8	33.7 28.0 2.6	377	1.0 0.0 0.233	47.5 56.0 28.4	62.8 26.9
417	R00Y_062_050a	0.625 0.125 0.375	0.625 0.5 0.375	360	0.625 0.125 0.375	44.8 29.4 5.2	29.9 10.0	0.625 0.125 0.375	43.7 33.1 4.7	33.5 8.0 3.8	360	1.0 0.0 0.5	47.8 58.9 10.4	59.9 10.0
418	B61R_062_050a	0.625 0.125 0.5	0.625 0.5 0.375	344	0.625 0.125 0.508	45.6 32.3 -3.5	32.5 353.7	0.625 0.125 0.5	43.8 38.9 -5.6	39.3 351.7 7.1	342	1.0 0.0 0.766	49.3 64.7 -7.1	65.1 353.7
419	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	45.0 32.7 -6.3	33.3 348.9	0.625 0.125 0.625	44.4 41.4 -12.9	43.4 342.7 10.9	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
420	B40R_075_062a	0.625 0.125 0.75	0.75 0.625 0.437	319	0.635 0.125 0.75	45.3 36.1 -12.0	38.1 341.5	0.625 0.125 0.75	44.8 43.5 -17.8	47.0 337.7 9.3	320	0.816 0.0 1.0	43.9 57.8 -19.3	61.0 341.5
421	B34R_087_075a	0.625 0.125 0.875	0.875 0.75 0.5	311	0.637 0.125 0.875	45.2 38.7 -18.5	42.9 334.4	0.625 0.125 0.875	44.1 46.8 -24.2	52.7 332.6 9.9	311	0.683 0.0 1.0	40.4 51.6 -24.7	57.2 334.4
422	B29R_100_087a	0.625 0.125 1.0	1.0 0.875 0.562	305	0.635 0.125 1.0	45.6 40.8 -24.9	47.9 328.5	0.625 0.125 1.0	41.0 43.2 -29.0	52.0 326.1 6.5	305	0.583 0.0 1.0	38.4 46.7 -28.5	54.7 328.5
423	R38Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	53	0.625 0.239 0.0	49.5 18.2 38.0	42.1 64.4	0.625 0.25 0.0	45.9 19.0 40.0	44.3 64.5 4.2	52	1.0 0.383 0.0	65.0 29.1 60.8	67.4 64.4
424	R23Y_062_050a	0.625 0.25 0.125	0.625 0.5 0.375	44	0.625 0.241 0.125	49.6 21.7 27.2	34.8 51.4	0.625 0.25 0.125	50.5 16.9 33.1	37.2 62.8 7.5	42	1.0 0.233 0.0	57.4 43.5 54.5	69.7 51.4
425	R00Y_062_037a	0.625 0.25 0.25	0.625 0.375 0.437	390	0.625 0.25 0.25	50.7 21.4 14.1	25.7 33.4	0.625 0.25 0.25	50.5 18.8 20.3	27.7 47.2 6.7	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
426	R18Y_062_037a	0.625 0.25 0.375	0.625 0.375 0.437	371	0.625 0.25 0.368	50.7 21.2 8.7	22.9 22.3	0.625 0.25 0.375	50.6 22.5 8.2	24.0 19.9 1.4	371	1.0 0.0 0.316	47.4 56.5 23.2	61.1 22.3
427	B65R_062_037a	0.625 0.25 0.5	0.625 0.375 0.437	349	0.625 0.25 0.506	51.1 23.7 -0.6	23.7 358.3	0.625 0.25 0.5	51.8 27.5 -4.1	27.8 351.4 5.2	348	1.0 0.0 0.683	48.6 63.2 -1.8	63.2 358.3
428	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	50.9 24.5 -4.7	24.9 348.9	0.625 0.25 0.625	52.0 30.3 -11.7	32.5 338.7 9.1	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
429	B38R_075_050a	0.625 0.25 0.75	0.75 0.5 0.5	316	0.633 0.25 0.75	51.1 27.9 -10.4	29.8 339.4	0.625 0.25 0.75	52.3 31.0 -15.4	34.7 333.4 6.0	317	0.766 0.0 1.0	42.4 55.8 -20.9	59.6 339.4
430	B30R_087_062a	0.625 0.25 0.875	0.875 0.625 0.562	307	0.635 0.25 0.875	51.3 30.0 -17.2	34.6 330.2	0.625 0.25 0.875	50.3 37.1 -24.3	44.3 326.7 10.0	307	0.616 0.0 1.0	38.9 48.1 -27.5	55.4 330.2
431	B25R_100_075a	0.625 0.25 1.0	1.0 0.75 0.625	300	0.625 0.25 1.0	51.8 32.3 -23.1	39.8 324.4	0.625 0.25 1.0	45.0 36.4 -29.0	46.6 321.4 9.9	300	0.5 0.0 1.0	37.2 43.1 -30.8	53.0 324.4
432	R61Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	67	0.625 0.385 0.0	55.5 7.5 44.0	44.6 80.3	0.625 0.375 0.0	51.0 6.3 45.7	46.2 82.0 4.9	67	1.0 0.616 0.0	74.6 12.0 70.4	71.4 80.3
433	R50Y_062_050a	0.625 0.375 0.125	0.625 0.5 0.375	60	0.625 0.375 0.125	56.1 9.6 33.1	34.5 73.8	0.625 0.375 0.125	57.4 6.1 39.1	39.6 81.0 7.0	59	1.0 0.5 0.0	70.5 19.2 66.2	69.0 73.8
434	R31Y_062_037a	0.625 0.375 0.25	0.625 0.375 0.437	49	0.625 0.368 0.25	56.0 13.3 21.8	25.5 58.6	0.625 0.375 0.25	57.6 8.9 25.0	26.5 70.2 5.6	48	1.0 0.316 0.0	61.6 35.5 58.2	68.2 58.6
435	R00Y_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	56.7 14.3 9.4	17.1 33.4	0.625 0.375 0.375	58.1 11.8 11.2	16.3 43.4 3.3	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
436	R00Y_062_025a	0.625 0.375 0.5	0.625 0.25 0.5	360	0.625 0.375 0.5	56.8 14.7 2.6	14.9 10.0	0.625 0.375 0.5	59.0 15.5 -0.3	15.5 358.8 3.7	360	1.0 0.0 0.5	47.8 58.9 10.4	59.9 10.0
437	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	56.9 16.3 -3.1	16.6 348.9	0.625 0.375 0.625	59.5 19.7 -8.9	21.7 335.6 7.2	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
438	B34R_075_037a	0.625 0.375 0.75	0.75 0.375 0.562	311	0.631 0.375 0.75	57.0 19.3 -9.2	21.4 334.4	0.625 0.375 0.75	58.0 21.5 -14.7	26.1 325.7 5.9	311	0.683 0.0 1.0	40.4 51.6 -24.7	57.2 334.4
439	B25R_087_050a	0.625 0.375 0.875	0.875 0.5 0.625	300	0.625 0.375 0.875	57.5 21.5 -15.4	26.5 324.4	0.625 0.375 0.875	56.8 27.0 -23.8	36.0 318.5 10.0	300	0.5 0.0 1.0	37.2 43.1 -30.8	53.0 324.4
440	B19R_100_062a	0.625 0.375 1.0	1.0 0.625 0.687	293	0.614 0.375 1.0	57.4 24.1 -21.7	32.4 317.9	0.625 0.375 1.0	51.8 26.1 -28.3	38.6 312.6 8.9	292	0.383 0.0 1.0	34.4 38.5 -34.7	51.9 317.9
441	R81Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	79	0.625 0.51 0.0	62.3 -3.6 47.7	47.9 94.3	0.625 0.5 0.0	60.5 -3.4 56.6	56.7 93.4 9.0	80	1.0 0.816 0.0	85.4 -5.8 76.4	76.6 94.3
442	R76Y_062_050a	0.625 0.5 0.125	0.625 0.5 0.375	76	0.625 0.508 0.125	62.7 -1.4 38.4	38.4 92.2	0.625 0.5 0.125	64.8 -4.4 49.8	50.0 95.0 11.9	77	1.0 0.766 0.0	83.5 -2.9 76.8	76.9 92.2
443	R68Y_062_037a	0.625 0.5 0.25	0.625 0.375 0.437	71	0.625 0.506 0.25	62.4 2.0 27.7	27.8 85.7	0.625 0.5 0.25	66.0 -2.3 33.6	33.7 94.0 8.2	71	1.0 0.683 0.0	78.6 5.4 73.9	74.1 85.7
444	R50Y_062_025a	0.625 0.5 0.375	0.625 0.25 0.5	60	0.625 0.5 0.375	62.5 4.8 16.5	17.2 73.8	0.625 0.5 0.375	65.6 1.4 17.9	17.9 85.5 4.8	59	1.0 0.5 0.0	70.5 19.2 66.2	69.0 73.8
445	R00Y_062_012a	0.625 0.5 0.5	0.625 0.125 0.562	390	0.625 0.5 0.5	62.8 7.1 4.7	8.5 33.4	0.625 0.5 0.5	65.7 5					

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
486	R00Y_075_075a	0.75 0.0 0.0	0.75 0.75 0.375	390	0.75 0.0 0.0	41.6 42.9 28.3	51.4 33.4	0.75 0.0 0.0	39.7 47.0 29.4	55.5 32.0 4.6	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
487	R35Y_075_075a	0.75 0.0 0.125	0.75 0.75 0.375	381	0.75 0.0 0.112	41.7 42.2 24.6	48.9 30.3	0.75 0.0 0.125	39.0 46.5 23.4	52.0 26.7 5.1	382	1.0 0.0 0.15	47.6 56.3 32.9	65.2 30.3
488	R18Y_075_075a	0.75 0.0 0.25	0.75 0.75 0.375	371	0.75 0.0 0.237	41.5 42.4 17.4	45.8 22.3	0.75 0.0 0.25	39.4 47.0 16.3	49.8 19.1 5.2	371	1.0 0.0 0.316	47.4 56.5 23.2	61.1 22.3
489	R00Y_075_075a	0.75 0.0 0.375	0.75 0.75 0.375	360	0.75 0.0 0.375	41.8 44.2 7.8	44.9 10.0	0.75 0.0 0.375	39.3 48.8 6.0	49.2 7.1 5.5	360	1.0 0.0 0.5	47.8 58.9 10.4	59.9 10.0
490	B65R_075_075a	0.75 0.0 0.5	0.75 0.75 0.375	349	0.75 0.0 0.512	42.4 47.4 -1.3	47.4 358.3	0.75 0.0 0.5	39.5 52.9 -4.1	53.1 355.4 6.8	348	1.0 0.0 0.683	48.6 63.2 -1.8	63.2 358.3
491	B57R_075_075a	0.75 0.0 0.625	0.75 0.75 0.375	339	0.75 0.0 0.637	43.0 49.4 -7.4	49.9 351.4	0.75 0.0 0.625	39.9 55.3 -11.1	56.4 348.6 7.7	337	1.0 0.0 0.85	49.4 65.8 -9.9	66.6 351.4
492	B50R_075_075a	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	42.1 49.0 -9.5	49.9 348.9	0.75 0.0 0.75	41.1 54.2 -16.4	56.6 343.1 8.6	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
493	B43R_087_087a	0.75 0.0 0.875	0.875 0.875 0.437	322	0.758 0.0 0.875	42.7 52.3 -15.4	54.5 343.5	0.75 0.0 0.875	42.0 54.2 -21.0	58.1 338.8 5.9	322	0.866 0.0 1.0	45.4 59.8 -17.6	62.3 343.5
494	B38R_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	42.4 55.8 -20.9	59.6 339.4	0.75 0.0 1.0	41.8 55.1 -21.4	59.1 338.7 0.9	317	0.766 0.0 1.0	42.4 55.8 -20.9	59.6 339.4
495	R15Y_075_075a	0.75 0.125 0.0	0.75 0.75 0.375	39	0.75 0.112 0.0	45.8 38.8 37.9	54.3 44.3	0.75 0.125 0.0	44.9 42.3 40.8	58.8 43.9 4.6	37	1.0 0.15 0.0	53.2 51.8 50.6	72.4 44.3
496	R00Y_075_062a	0.75 0.125 0.125	0.75 0.625 0.437	390	0.75 0.125 0.125	47.6 35.7 23.6	42.8 33.4	0.75 0.125 0.125	45.5 36.3 31.1	47.8 40.6 7.8	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
497	R31Y_075_062a	0.75 0.125 0.25	0.75 0.625 0.437	379	0.75 0.125 0.239	47.7 35.1 19.4	40.1 28.9	0.75 0.125 0.25	45.6 37.0 22.6	43.3 41.4 4.2	380	1.0 0.0 0.183	47.6 56.2 31.1	64.2 28.9
498	R11Y_075_062a	0.75 0.125 0.375	0.75 0.625 0.437	367	0.75 0.125 0.364	47.5 35.6 11.8	37.5 18.3	0.75 0.125 0.375	45.0 38.4 14.5	41.1 20.7 4.7	367	1.0 0.0 0.383	47.4 57.0 18.9	60.0 18.3
499	B69R_075_062a	0.75 0.125 0.5	0.75 0.625 0.437	353	0.75 0.125 0.51	47.9 38.5 1.6	38.5 2.5	0.75 0.125 0.5	45.2 42.6 3.5	42.7 4.7 5.2	352	1.0 0.0 0.616	47.9 61.6 2.7	61.7 2.5
500	B59R_075_062a	0.75 0.125 0.625	0.75 0.625 0.437	341	0.75 0.125 0.635	48.8 40.9 -5.4	41.2 352.3	0.75 0.125 0.625	45.8 46.9 -5.5	47.2 353.2 6.7	339	1.0 0.0 0.816	49.4 65.4 -8.7	66.0 352.3
501	B50R_075_062a	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	48.0 40.8 -7.9	41.6 348.9	0.75 0.125 0.75	46.6 48.7 -13.3	50.5 344.6 9.6	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
502	B42R_087_075a	0.75 0.125 0.875	0.875 0.75 0.5	321	0.762 0.125 0.875	48.6 44.3 -13.6	46.4 342.9	0.75 0.125 0.875	47.1 50.2 -17.9	53.3 340.2 7.4	322	0.85 0.0 1.0	44.9 59.1 -18.2	61.9 342.9
503	B36R_100_087a	0.75 0.125 1.0	1.0 0.875 0.562	314	0.766 0.125 1.0	48.3 47.5 -19.5	51.3 337.6	0.75 0.125 1.0	45.0 50.7 -22.1	55.3 336.3 5.2	315	0.733 0.0 1.0	41.5 54.3 -22.3	58.7 337.6
504	R31Y_075_075a	0.75 0.25 0.0	0.75 0.75 0.375	49	0.75 0.237 0.0	52.2 26.6 43.7	51.1 58.6	0.75 0.25 0.0	52.2 26.4 47.8	54.6 61.1 4.1	48	1.0 0.316 0.0	61.6 35.5 58.2	68.2 58.6
505	R18Y_075_062a	0.75 0.25 0.125	0.75 0.625 0.437	41	0.75 0.239 0.125	52.2 30.3 32.7	44.6 47.1	0.75 0.25 0.125	52.1 26.6 36.8	45.4 54.0 5.4	39	1.0 0.183 0.0	54.9 48.5 52.3	71.4 47.1
506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	53.7 28.6 18.9	34.3 33.4	0.75 0.25 0.25	52.4 27.1 25.3	37.1 43.0 6.7	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
507	R26Y_075_050a	0.75 0.25 0.375	0.75 0.5 0.5	376	0.75 0.25 0.366	53.7 28.0 14.2	31.4 26.9	0.75 0.25 0.375	52.3 29.3 16.1	33.5 28.8 2.7	377	1.0 0.0 0.233	47.5 56.0 28.4	62.8 26.9
508	R00Y_075_050a	0.75 0.25 0.5	0.75 0.5 0.5	360	0.75 0.25 0.5	53.8 29.4 5.2	29.9 10.0	0.75 0.25 0.5	53.2 30.6 5.9	31.2 10.9 1.4	360	1.0 0.0 0.5	47.8 58.9 10.4	59.9 10.0
509	B61R_075_050a	0.75 0.25 0.625	0.75 0.5 0.5	344	0.75 0.25 0.633	54.5 32.3 -3.5	32.5 353.7	0.75 0.25 0.625	53.9 34.9 -4.4	35.2 352.6 2.7	342	1.0 0.0 0.766	49.3 64.7 -7.1	65.1 353.7
510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	54.0 32.7 -6.3	33.3 348.9	0.75 0.25 0.75	53.9 38.1 -12.4	40.1 341.9 8.1	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
511	B40R_087_062a	0.75 0.25 0.875	0.875 0.625 0.562	319	0.76 0.25 0.875	54.3 36.1 -12.0	38.1 341.5	0.75 0.25 0.875	54.3 40.1 -17.3	43.7 336.6 6.5	320	0.816 0.0 1.0	43.9 57.8 -19.3	61.0 341.5
512	B34R_100_075a	0.75 0.25 1.0	1.0 0.75 0.625	311	0.762 0.25 1.0	54.2 38.7 -18.5	42.9 334.4	0.75 0.25 1.0	50.3 42.9 -22.8	48.6 332.0 7.1	311	0.683 0.0 1.0	40.4 51.6 -24.7	57.2 334.4
513	R50Y_075_075a	0.75 0.375 0.0	0.75 0.75 0.375	60	0.75 0.375 0.0	58.8 14.4 49.7	51.7 73.8	0.75 0.375 0.0	57.7 15.3 53.6	55.7 74.0 4.1	59	1.0 0.5 0.0	70.5 19.2 66.2	69.0 73.8
514	R38Y_075_062a	0.75 0.375 0.125	0.75 0.625 0.437	53	0.75 0.364 0.125	58.5 18.2 38.0	42.1 64.4	0.75 0.375 0.125	57.9 16.7 44.5	47.5 69.3 6.6	52	1.0 0.383 0.0	65.0 29.1 60.8	67.4 64.4
515	R23Y_075_050a	0.75 0.375 0.25	0.75 0.5 0.5	44	0.75 0.366 0.25	58.6 21.7 27.2	34.8 51.4	0.75 0.375 0.25	58.5 17.2 29.7	34.3 59.9 5.1	42	1.0 0.233 0.0	57.4 43.5 54.5	69.7 51.4
516	R00Y_075_037a	0.75 0.375 0.375	0.75 0.375 0.562	390	0.75 0.375 0.375	59.7 21.4 14.1	25.7 33.4	0.75 0.375 0.375	59.1 18.5 19.5	26.9 46.3 6.0	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
517	R18Y_075_037a	0.75 0.375 0.5	0.75 0.375 0.562	371	0.75 0.375 0.493	59.7 21.2 8.7	22.9 22.3	0.75 0.375 0.5	60.1 19.9 9.2	21.9 24.9 1.4	371	1.0 0.0 0.316	47.4 56.5 23.2	61.1 22.3
518	B65R_075_037a	0.75 0.375 0.625	0.75 0.375 0.562	349	0.75 0.375 0.631	60.1 23.7 -0.6	23.7 358.3	0.75 0.375 0.625	60.2 24.2 -2.3	24.3 354.5 1.7	348	1.0 0.0 0.683	48.6 63.2 -1.8	63.2 358.3
519	B50R_075_037a	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	59.9 24.5 -4.7	24.9 348.9	0.75 0.375 0.75	60.1 28.2 -10.2	30.0 339.9 6.6	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
520	B38R_087_050a	0.75 0.375 0.875	0.875 0.5 0.625	316	0.758 0.375 0.875	60.1 27.9 -10.4	29.8 339.4	0.75 0.375 0.875	60.4 31.7 -15.9	35.5 333.4 6.6	317	0.766 0.0 1.0	42.4 55.8 -20.9	59.6 339.4
521	B30R_100_062a	0.75 0.375 1.0	1.0 0.625 0.687	307	0.76 0.375 1.0	60.3 30.0 -17.2	34.6 330.2	0.75 0.375 1.0	55.2 36.2 -21.7	42.2 329.0 9.1	307	0.616 0.0 1.0	38.9 48.1 -27.5	55.4 330.2
522	R68Y_075_075a	0.75 0.5 0.0	0.75 0.75 0.375	71	0.75 0.512 0.0	64.9 4.0 55.4	55.6 85.7	0.75 0.5 0.0	64.3 4.5 59.9	60.1 85.6 4.5	71	1.0 0.683 0.0	78.6 54 73.9	74.1 85.7
523	R61Y_075_062a	0.75 0.5 0.125	0.75 0.625 0.437	67	0.75 0.51 0.125	64.5 7.5 44.0	44.6 80.3	0.75 0.5 0.125	65.5 5.4 54.3	54.6 84.2 10.5	67	1.0 0.616 0.0	74.6 12.0 70.4	71.4 80.3
524	R50Y_075_050a	0.75 0.5 0.25	0.75 0.5 0.5	60	0.75 0.5 0.25	65.1 9.6 33.1	34.5 73.8	0.75 0.5 0.25	66.1 6.5 36.1	36.6 79.7 4.3	59	1.0 0.5 0.0	70.5 19.2 66.2	69.0 73.8
525	R31Y_075_037a	0.75 0.5 0.375	0.75 0.375 0.562	49	0.75 0.493 0.375	65.0 13.3 21.8	25.5 58.6	0.75 0.5 0.375	65.9 8.9 24.5	26.0 70.0 5.2	48	1.0 0.316 0.0	61.6 35.5 58.2	68.2 58.6
526	R00Y_075_025a	0.75 0.5 0.5	0.75 0.25 0.625	390	0.75 0.5 0.5	65.7 14.3 9.4	17.1 33.4	0.75 0.5 0.5	67.2 10.6 12.7	16.5 50.1 5.1	389	1.0 0.0 0.0	47.5 57.	

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md		
567	R00Y_087_087a	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.0	44.6 50.0 33.1	60.0 33.4	0.875 0.0 0.0	43.6 51.3 31.6	60.2 31.6 2.1	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
568	R36Y_087_087a	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.116	44.6 49.3 29.6	57.5 30.9	0.875 0.0 0.125	43.2 51.0 26.3	57.5 27.3 3.9	382	1.0 0.0 0.133	47.6 56.3 33.8	65.7 30.9
569	R23Y_087_087a	0.875 0.0 0.25	0.875 0.875 0.437	374	0.875 0.0 0.233	44.5 49.0 23.1	54.3 25.2	0.875 0.0 0.25	43.0 51.3 20.1	55.1 21.4 4.0	375	1.0 0.0 0.266	47.5 56.1 26.5	62.0 25.2
570	R08Y_087_087a	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.364	44.5 50.5 14.4	52.5 15.9	0.875 0.0 0.375	43.3 53.1 10.6	54.2 11.3 4.7	365	1.0 0.0 0.416	47.5 57.7 16.5	60.0 15.9
571	B70R_087_087a	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.51	44.9 53.3 4.3	53.5 4.6	0.875 0.0 0.5	43.4 55.6 1.4	55.6 1.4 4.0	354	1.0 0.0 0.583	47.9 60.9 4.9	61.1 4.6
572	B63R_087_087a	0.875 0.0 0.625	0.875 0.875 0.437	346	0.875 0.0 0.641	45.9 56.2	-4.6 56.4 355.2	0.875 0.0 0.625	43.8 59.4	-6.7 59.8 353.5	4.3	1.0 0.0 0.733	49.1 64.2	-5.3 64.4 355.2
573	B56R_087_087a	0.875 0.0 0.75	0.875 0.875 0.437	338	0.875 0.0 0.758	46.3 57.8	-9.1 58.5 350.9	0.875 0.0 0.75	45.4 60.7	-12.7 62.1 348.1	4.7	1.0 0.0 0.866	49.5 66.0	-10.4 66.9 350.9
574	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	45.1 57.2	-11.1 58.3 348.9	0.875 0.0 0.875	45.5 59.0	-16.8 61.4 344.0	5.9	1.0 0.0 1.0	48.1 65.4	-12.7 66.6 348.9
575	B44R_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	45.8 60.5	-17.0 62.8 344.2	0.875 0.0 1.0	45.6 60.1	-17.3 62.6 343.9	0.4	0.883 0.0	1.0 45.8 60.5	-17.0 62.8 344.2
576	R13Y_087_087a	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.116 0.0	48.8 46.8	43.4 63.8 42.9	0.875 0.125 0.0	49.7 50.4	46.7 68.7 42.8 4.9	37	1.0 0.133 0.0	52.3 53.4	49.7 73.0 42.9
577	R00Y_087_075a	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.125	50.6 42.9	28.3 51.4 33.4	0.875 0.125 0.125	49.6 47.7	41.6 63.3 41.1 14.1	389	1.0 0.0 0.0	47.5 57.2	37.8 68.6 33.4
578	R35Y_087_075a	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.237	50.7 42.2	24.6 48.9 30.3	0.875 0.125 0.25	49.6 47.9	30.7 56.9 32.6 8.3	382	1.0 0.0 0.15	47.6 56.3	32.9 65.2 30.3
579	R18Y_087_075a	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.362	50.5 42.4	17.4 45.8 22.3	0.875 0.125 0.375	49.3 49.0	22.0 53.8 24.1 8.1	371	1.0 0.0 0.316	47.4 56.5	23.2 61.1 22.3
580	R00Y_087_075a	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.5	50.8 44.2	7.8 44.9 30.3	1.0 0.875 0.125 0.5	49.5 51.8	11.4 53.0 12.4 8.5	370	1.0 0.0 0.5	47.8 58.9	10.4 59.9 10.0
581	B65R_087_075a	0.875 0.125 0.625	0.875 0.75 0.5	349	0.875 0.125 0.637	51.4 47.4	-1.3 47.4 358.3	0.875 0.125 0.625	49.1 55.8	0.5 55.8 0.6 8.9	348	1.0 0.0 0.683	48.6 63.2	-1.8 63.2 358.3
582	B57R_087_075a	0.875 0.125 0.75	0.875 0.75 0.5	339	0.875 0.125 0.762	52.0 49.4	-7.4 49.9 351.4	0.875 0.125 0.75	50.0 58.8	-10.1 59.7 350.2 10.0	337	1.0 0.0 0.85	49.4 65.8	-9.9 66.6 351.4
583	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	51.1 49.0	-9.5 49.9 348.9	0.875 0.125 0.875	49.2 57.7	-16.2 60.0 344.2 11.1	330	1.0 0.0 1.0	48.1 65.4	-12.7 66.6 348.9
584	B43R_100_087a	0.875 0.125 1.0	1.0 1.0 0.875	322	0.883 0.125 1.0	51.7 52.3	-15.4 54.5 343.5	0.875 0.125 1.0	48.0 56.9	-17.9 59.6 342.4 6.3	322	0.866 0.0	1.0 45.4 59.8	-17.6 62.3 343.5
585	R26Y_087_087a	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.233 0.0	54.7 35.2	49.0 60.3 54.2	0.875 0.25 0.0	56.8 35.4	52.6 63.5 56.0 4.2	44	1.0 0.266 0.0	59.1 40.2	56.0 69.0 54.2
586	R15Y_087_075a	0.875 0.25 0.125	0.875 0.75 0.5	39	0.875 0.237 0.125	54.8 38.8	37.9 54.3 44.3	0.875 0.25 0.125	55.4 36.3	46.0 58.6 51.6 8.4	37	1.0 0.15 0.0	53.2 51.8	50.6 72.4 44.3
587	R00Y_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	56.6 35.7	23.6 42.8 33.4	0.875 0.25 0.25	56.9 34.5	32.7 47.5 43.4 9.1	389	1.0 0.0 0.0	47.5 57.2	37.8 68.6 33.4
588	R31Y_087_062a	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.364	56.7 35.1	19.4 40.1 28.9	0.875 0.25 0.375	56.3 36.2	24.8 43.9 34.4 5.5	380	1.0 0.0 0.183	47.6 56.2	31.1 64.2 28.9
589	R11Y_087_062a	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.489	56.5 35.6	11.8 37.5 18.3	0.875 0.25 0.5	56.0 38.3	15.0 41.1 21.4 4.2	367	1.0 0.0 0.383	47.4 57.0	18.9 60.0 18.3
590	B69R_087_062a	0.875 0.25 0.625	0.875 0.625 0.562	353	0.875 0.25 0.635	56.9 38.5	1.6 38.5 2.5	0.875 0.25 0.625	56.4 42.0	3.3 42.1 4.6 3.9	352	1.0 0.0 0.616	47.9 61.6	2.7 61.7 2.5
591	B59R_087_062a	0.875 0.25 0.75	0.875 0.625 0.562	341	0.875 0.25 0.76	57.8 40.9	-5.4 41.2 352.3	0.875 0.25 0.75	57.2 45.5	-6.6 46.0 351.7 4.8	339	1.0 0.0 0.816	49.4 65.4	-8.7 66.0 352.3
592	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	57.0 40.8	-7.9 41.6 348.9	0.875 0.25 0.875	56.8 46.5	-14.0 48.6 343.1 8.2	330	1.0 0.0 1.0	48.1 65.4	-12.7 66.6 348.9
593	B42R_100_075a	0.875 0.25 1.0	1.0 1.0 0.75	321	0.887 0.25 1.0	57.6 44.3	-13.6 46.4 342.9	0.875 0.25 1.0	54.7 48.9	-16.3 51.6 341.5 6.0	322	0.85 0.0	1.0 44.9 59.1	-18.2 61.9 342.9
594	R41Y_087_087a	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.364 0.0	61.2 23.1	54.7 59.4 67.0	0.875 0.375 0.0	63.1 24.3	58.4 63.2 67.4 4.2	54	1.0 0.416 0.0	66.6 26.4	62.5 67.9 67.0
595	R31Y_087_075a	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.362 0.125	61.2 26.6	43.7 51.1 58.6	0.875 0.375 0.125	61.5 24.8	49.6 55.5 63.3 6.1	48	1.0 0.316 0.0	61.6 35.5	58.2 68.2 58.6
596	R18Y_087_062a	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.364 0.25	61.2 30.3	32.7 44.6 47.1	0.875 0.375 0.25	62.3 24.9	36.0 43.8 55.3 6.4	39	1.0 0.183 0.0	54.9 48.5	52.3 71.4 47.1
597	R00Y_087_050a	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	62.7 28.6	18.9 34.3 33.4	0.875 0.375 0.375	62.9 25.1	26.0 36.2 45.9 7.9	389	1.0 0.0 0.0	47.5 57.2	37.8 68.6 33.4
598	R26Y_087_050a	0.875 0.375 0.5	0.875 0.5 0.625	376	0.875 0.375 0.491	62.7 28.0	14.2 31.4 26.9	0.875 0.375 0.5	63.1 27.5	16.1 31.9 30.3 1.9	377	1.0 0.0 0.233	47.5 56.0	28.4 62.8 26.9
599	R00Y_087_050a	0.875 0.375 0.625	0.875 0.5 0.625	360	0.875 0.375 0.625	62.8 29.4	5.2 29.9 10.0	0.875 0.375 0.625	63.0 30.1	6.9 30.9 12.9 1.9	360	1.0 0.0 0.5	47.8 58.9	10.4 59.9 10.0
600	B61R_087_050a	0.875 0.375 0.75	0.875 0.5 0.625	344	0.875 0.375 0.758	63.5 32.3	-3.5 32.5 353.7	0.875 0.375 0.75	63.3 35.2	-4.9 35.6 352.0 3.2	342	1.0 0.0 0.766	49.3 64.7	-7.1 65.1 353.7
601	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	63.0 32.7	-6.3 33.3 348.9	0.875 0.375 0.875	63.7 36.7	-11.8 38.6 342.1 6.8	330	1.0 0.0 1.0	48.1 65.4	-12.7 66.6 348.9
602	B40R_100_062a	0.875 0.375 1.0	1.0 1.0 0.625	319	0.885 0.375 1.0	63.3 36.1	-12.0 38.1 341.5	0.875 0.375 1.0	61.4 37.8	-15.8 41.0 337.2 4.6	320	0.816 0.0	1.0 43.9 57.8	-19.3 61.0 341.5
603	R58Y_087_087a	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.51 0.0	67.2 12.3	60.6 61.9 78.4	0.875 0.5 0.0	69.2 12.9	63.8 65.1 78.5 3.7	65	1.0 0.583 0.0	73.4 14.1	69.3 70.7 78.4
604	R50Y_087_075a	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.5 0.125	67.8 14.4	49.7 51.7 73.8	0.875 0.5 0.125	68.1 14.1	57.3 59.0 76.1 7.6	59	1.0 0.5 0.0	70.5 19.2	66.2 69.0 73.8
605	R38Y_087_062a	0.875 0.5 0.25	0.875 0.625 0.562	53	0.875 0.489 0.25	67.5 18.2	38.0 42.1 64.4	0.875 0.5 0.25	68.9 13.2	41.0 43.0 72.1 5.9	52	1.0 0.383 0.0	65.0 29.1	60.8 67.4 64.4
606	R23Y_087_050a	0.875 0.5 0.375	0.875 0.5 0.625	44	0.875 0.491 0.375	67.6 21.7	27.2 34.8 51.4	0.875 0.5 0.375	68.6 15.1	30.6 34.1 63.7 7.5	42	1.0 0.233 0.0	57.4 43.5	54.5 69.7 51.4
607	R00Y_087_037a	0.875 0.5 0.5	0.875 0.375 0.687	390										

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md
648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4	1.0 0.0 0.0	47.5 57.2 37.8
649	R38Y_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	47.6 56.4 34.5	66.1 31.4	1.0 0.0 0.125	47.6 56.4 34.5	66.1 31.4	1.0 0.0 0.116	47.6 56.4 34.5
650	R26Y_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	47.5 56.0 28.4	62.8 26.9	1.0 0.0 0.25	47.5 56.0 28.4	62.8 26.9	1.0 0.0 0.233	47.5 56.0 28.4
651	R13Y_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	47.4 56.8 20.0	60.2 19.4	1.0 0.0 0.375	47.4 56.8 20.0	60.2 19.4	1.0 0.0 0.366	47.4 56.8 20.0
652	R00Y_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	47.8 58.9 10.4	59.9 10.0	1.0 0.0 0.5	47.8 58.9 10.4	59.9 10.0	1.0 0.0 0.5	47.8 58.9 10.4
653	B68R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	48.0 62.0 1.5	62.0 1.4	1.0 0.0 0.625	48.0 61.8 2.1	61.8 1.9	1.0 0.0 0.633	48.0 62.0 1.5
654	B61R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	49.3 64.7 -7.1	65.1 353.7	1.0 0.0 0.75	49.3 64.5 -6.5	64.8 354.2	1.0 0.0 0.766	49.3 64.7 -7.1
655	B55R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	49.4 66.1 -10.9	67.0 350.6	1.0 0.0 0.875	49.5 66.1 -10.7	67.0 350.7	1.0 0.0 0.883	49.4 66.1 -10.9
656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9	1.0 0.0 1.0	48.1 65.4 -12.7
657	R11Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	51.6 54.5 48.4	72.9 41.6	1.0 0.125 0.0	51.9 54.3 49.2	73.2 42.1	1.0 0.116 0.0	51.6 54.5 48.4
658	R00Y_100_087a	1.0 0.125 0.125	1.0 0.875 0.562	390	1.0 0.125 0.125	53.6 50.0 33.1	60.0 33.4	1.0 0.125 0.125	50.9 56.4 46.0	72.8 39.1	1.0 0.0 0.0	47.5 57.2 37.8
659	R36Y_100_087a	1.0 0.125 0.25	1.0 0.875 0.562	382	1.0 0.125 0.241	53.6 49.3 29.6	57.5 30.9	1.0 0.125 0.25	51.1 57.3 36.9	68.2 32.7	1.0 0.0 0.0	47.5 57.2 37.8
660	R23Y_100_087a	1.0 0.125 0.375	1.0 0.875 0.562	374	1.0 0.125 0.358	53.5 49.0 23.1	54.3 25.2	1.0 0.125 0.375	50.9 59.2 26.9	65.0 24.4	1.0 0.0 0.0	47.5 57.2 37.8
661	R08Y_100_087a	1.0 0.125 0.5	1.0 0.875 0.562	365	1.0 0.125 0.489	53.5 50.5 14.4	52.5 15.9	1.0 0.125 0.5	51.5 60.5 17.6	63.0 16.2	1.0 0.0 0.0	47.5 57.2 37.8
662	B70R_100_087a	1.0 0.125 0.625	1.0 0.875 0.562	355	1.0 0.125 0.635	53.9 53.3 4.3	53.5 4.6	1.0 0.125 0.625	51.1 64.7 5.0	64.9 4.4	1.0 0.0 0.0	47.5 57.2 37.8
663	B63R_100_087a	1.0 0.125 0.75	1.0 0.875 0.562	346	1.0 0.125 0.766	54.9 56.2 -4.6	56.4 355.2	1.0 0.125 0.75	51.5 67.6 -5.3	67.8 355.4	1.0 0.0 0.0	47.5 57.2 37.8
664	B56R_100_087a	1.0 0.125 0.875	1.0 0.875 0.562	338	1.0 0.125 0.883	55.3 57.8 -9.1	58.5 350.9	1.0 0.125 0.875	52.1 68.4 -9.9	69.1 351.7	1.0 0.0 0.0	47.5 57.2 37.8
665	B50R_100_087a	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	54.1 57.2 -11.1	58.3 348.9	1.0 0.125 1.0	51.6 65.2 -11.3	66.1 350.0	1.0 0.0 0.0	47.5 57.2 37.8
666	R23Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	57.4 43.5 54.5	69.7 51.9	1.0 0.25 0.0	58.2 41.8 55.1	69.2 52.8	1.0 0.233 0.0	57.4 43.5 54.5
667	R13Y_100_087a	1.0 0.25 0.125	1.0 0.875 0.562	38	1.0 0.241 0.125	57.8 46.8 43.4	63.8 42.9	1.0 0.25 0.125	57.8 42.8 53.7	68.7 54.4	1.0 0.133 0.0	52.3 53.4 49.7
668	R00Y_100_075a	1.0 0.25 0.25	1.0 0.75 0.625	390	1.0 0.25 0.25	59.6 42.9 28.3	51.4 33.4	1.0 0.25 0.25	58.2 43.1 44.9	62.3 46.1	1.0 0.0 0.0	47.5 57.2 37.8
669	R35Y_100_075a	1.0 0.25 0.375	1.0 0.75 0.625	381	1.0 0.25 0.362	59.7 42.2 24.6	48.9 30.3	1.0 0.25 0.375	57.6 45.9 33.2	56.7 35.9	1.0 0.0 0.0	47.5 57.2 37.8
670	R18Y_100_075a	1.0 0.25 0.5	1.0 0.75 0.625	371	1.0 0.25 0.487	59.5 42.4 17.4	45.8 22.3	1.0 0.25 0.5	56.8 49.3 23.4	54.6 25.4	1.0 0.0 0.0	47.5 57.2 37.8
671	R00Y_100_075a	1.0 0.25 0.625	1.0 0.75 0.625	360	1.0 0.25 0.625	59.8 44.2 7.8	44.9 10.0	1.0 0.25 0.625	57.2 52.2 10.3	53.2 11.2	1.0 0.0 0.0	47.5 57.2 37.8
672	B65R_100_075a	1.0 0.25 0.75	1.0 0.75 0.625	349	1.0 0.25 0.762	60.4 47.4 -1.3	47.4 358.3	1.0 0.25 0.75	58.2 54.5 -1.8	54.5 358.0	1.0 0.0 0.0	47.5 57.2 37.8
673	B57R_100_075a	1.0 0.25 0.875	1.0 0.75 0.625	339	1.0 0.25 0.887	61.0 49.4 -7.4	49.9 351.4	1.0 0.25 0.875	58.8 56.4 -9.7	57.2 350.2	1.0 0.0 0.0	47.5 57.2 37.8
674	B50R_100_075a	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	60.1 49.0 -9.5	49.9 348.9	1.0 0.25 1.0	59.2 53.0 -11.1	54.2 348.1	1.0 0.0 0.0	47.5 57.2 37.8
675	R36Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	64.2 30.6 60.1	67.5 63.0	1.0 0.375 0.0	64.6 29.8 60.4	67.3 63.7	1.0 0.366 0.0	64.2 30.6 60.1
676	R26Y_100_087a	1.0 0.375 0.125	1.0 0.875 0.562	46	1.0 0.358 0.125	63.7 35.2 49.0	60.3 54.2	1.0 0.375 0.125	63.2 32.5 57.2	65.9 60.3	1.0 0.266 0.0	59.1 40.2 56.0
677	R15Y_100_075a	1.0 0.375 0.25	1.0 0.75 0.625	39	1.0 0.362 0.25	63.8 38.8 37.9	54.3 44.3	1.0 0.375 0.25	63.3 33.3 49.0	59.3 55.8	1.0 0.15 0.0	53.2 51.8 50.6
678	R00Y_100_062a	1.0 0.375 0.375	1.0 0.625 0.687	390	1.0 0.375 0.375	65.6 35.7 23.6	42.8 33.4	1.0 0.375 0.375	64.5 33.3 35.8	48.9 47.0	1.0 0.0 0.0	47.5 57.2 37.8
679	R31Y_100_062a	1.0 0.375 0.5	1.0 0.625 0.687	379	1.0 0.375 0.489	65.7 35.1 19.4	40.1 28.9	1.0 0.375 0.5	64.4 35.7 25.6	44.0 35.7	1.0 0.0 0.0	47.5 57.2 37.8
680	R11Y_100_062a	1.0 0.375 0.625	1.0 0.625 0.687	367	1.0 0.375 0.614	65.5 35.6 11.8	37.5 18.3	1.0 0.375 0.625	64.1 39.5 13.1	41.6 18.3	1.0 0.0 0.0	47.5 57.2 37.8
681	B69R_100_062a	1.0 0.375 0.75	1.0 0.625 0.687	353	1.0 0.375 0.76	65.9 38.5 1.6	38.5 2.5	1.0 0.375 0.75	64.9 42.1 1.3	42.1 1.8	1.0 0.0 0.0	47.5 57.2 37.8
682	B59R_100_062a	1.0 0.375 0.875	1.0 0.625 0.687	341	1.0 0.375 0.885	66.8 40.9 -5.4	41.2 352.3	1.0 0.375 0.875	65.6 44.6 -7.7	45.2 350.4	1.0 0.0 0.0	47.5 57.2 37.8
683	B50R_100_062a	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	66.0 40.8 -7.9	41.6 348.9	1.0 0.375 1.0	66.3 43.9 -9.3	44.9 348.0	1.0 0.0 0.0	47.5 57.2 37.8
684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	70.5 19.2 66.2	69.0 73.8	1.0 0.5 0.0	70.5 19.2 66.2	69.0 73.8	1.0 0.5 0.0	70.5 19.2 66.2
685	R41Y_100_087a	1.0 0.5 0.125	1.0 0.875 0.562	55	1.0 0.489 0.125	70.2 23.1 54.7	59.4 67.0	1.0 0.5 0.125	68.8 22.5 62.1	66.1 70.0	1.0 0.416 0.0	66.6 26.4 62.5
686	R31Y_100_075a	1.0 0.5 0.25	1.0 0.75 0.625	49	1.0 0.487 0.25	70.2 26.6 43.7	51.1 58.6	1.0 0.5 0.25	69.3 22.8 50.9	55.8 65.8	1.0 0.316 0.0	61.6 25.5 58.2
687	R18Y_100_062a	1.0 0.5 0.375	1.0 0.625 0.687	41	1.0 0.489 0.375	70.2 30.3 32.7	44.6 47.1	1.0 0.5 0.375	69.7 24.5 38.1	45.3 57.2	1.0 0.183 0.0	54.9 48.5 52.3
688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	71.7 28.6 18.9	39.3 34.4	1.0 0.5 0.5	71.4 24.0 27.4	36.4 48.8	1.0 0.0 0.0	47.5 57.2 37.8
689	R26Y_100_050a	1.0 0.5 0.625	1.0 0.5 0.75	376	1.0 0.5 0.616	71.7 28.0 14.2	31.4 26.9	1.0 0.5 0.625	71.6 27.0 15.7	31.2 30.2	1.0 0.233 0.0	47.5 57.2 37.8
690	R00Y_100_050a	1.0 0.5 0.75	1.0 0.5 0.75	360	1.0 0.5 0.75	71.8 29.4 5.2	29.9 10.0	1.0 0.5 0.75	71.9 30.4 4.7	30.7 8.8	1.0 0.0 0.0	47.5 57.2 37.8
691	B61R_100_050a	1.0 0.5 0.875	1.0 0.5 0.75	344	1.0 0.5 0.883	72.5 32.3 -3.5	32.5 353.7	1.0 0.5 0.875	71.8 34.7 -5.9	35.2 350.3	1.0 0.0 0.0	47.5 57.2 37.8
692	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	72.0 32.7 -6.3	33.3 348.9	1.0 0.5 1.0	71.6 36.1 -8.9	37.2 346.1	1.0 0.0 0.0	47.5 57.2 37.8
693	R63Y_100_100a	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	75.4 10.6 71.2	72.0 81.5	1.0 0.625 0.0	74.9 11.4 70.7	71.6 80.7	1.0 0.633 0.0	75.4 10.6 71.2
694	R58Y_100_087a	1.0 0.625 0.125	1.0 0.875 0.562	65	1.0 0.635 0.125	76.2 12.3 60.6	61.9 78.4	1.0 0.625 0.125	75.7 10.3 67.8	68.6 81.3	1.0 0.583 0.0	73.4 14.1 69.3
695	R50Y_100_075a	1.0 0.625 0.25	1.0 0.75 0.625	60	1.0 0.625 0.25	76.8 14.4 49.7	51.7 73.8	1.0 0.625 0.25	75.9 11.1 56.5	57.6 78.8	1.0 0.5 0.0	70.5 19.2 66.2
696	R38Y_100_062a	1.0 0.625 0.375	1.0 0.625 0.687	53	1.0 0.614 0.375	76.5 18.2 38.0	42.1 64.4	1.0 0.625 0.375	76.0 13.2 42.5	44.6 72.7	1.0 0.383 0.0	65.0 29.1 60.8
697	R23Y_100_050a	1.0 0.625 0.5	1.0 0.5 0.75	44	1.0 0.616 0.5	76.6 21.7 27.2	34.8 51.4	1.0 0.625 0.5	77.1 14.1 31.2	34.3 65.5	1.0 0.233 0.0	57.4 43.5 54.5
698	R00Y_100_037a	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	77.7 21.4 14.1	25.7 33.4	1.0 0.625 0.625	78.4 15.6 19.6	25.1 51.4	1.0 0.0 0.0	47.5 57.2 37.8
699	R18Y_100_037a	1.0 0.625 0.75	1.0 0.375 0.812	371	1.0 0.625 0.743	77.7 21.2 8.7	22.9 22.3	1.0 0.625 0.75	78.6 19.2 8.4	21.0 23.7	1.0 0.0 0.0	47.5 57.2 37.8
700	B65R_100_037a	1.0 0.625 0.875	1.0 0.375 0.812	349	1.0 0.625 0.881	78.1 23.7 -0.6	23.7 358.3	1.0 0.625 0.875	79.4 23.4 -4.3	23.8 349.5	1.0 0.0 0.0	47.5 57.2 37.8
701	B50R_100_037a	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	77.9 24.5 -4.7	24.9 348.9	1.0 0.625 1.0	78.5 25.6 -7.2	26.6 344.3	1.0 0.0 0.0	47.5 57.2 37.8
702	R76Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	83.5 -2.9 76.8	76.9 92.2	1.0 0.75 0.0	82.9 -2.0 76.9	77.0 91.5	1.0 0.766 0.0	83.5 -2.9 76.8
703	R73Y_100_087a	1.0 0.75 0.125	1.0 0.875 0.562	74	1.0 0.766 0.125	83.6 4.0 66.7	90.0 1.0	1.0 0.75 0.125	82.6 -0.5 66.7	90.5 1.0	1.0 0.733 0.0	81.8 -0.1 76.3
704	R68Y_100_075a	1.0 0.75 0.25	1.0 0.75 0.625	71	1.0 0.762 0.25	82.9 4.0 55.4	55.6 85.7	1.0 0.75 0.25	82.4 1.1 57.8	88.5 4.3	1.0 0.683 0.0	78.6 5.4 73.9
705	R61Y_100_062a	1.0 0.75 0.375	1.0 0.625 0.687	67	1.0 0.76							

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md		
729	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	96.1 0.0 0.0	0.0 178.6 0.2	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
730	G50B_100_012d	0.875 1.0 1.0	1.0 0.125 0.937	210	0.875 1.0 1.0	90.4 -3.7 -5.3	6.5 235.1	0.875 1.0 1.0	90.5 -5.7 -7.6	9.6 232.8 3.0	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
731	G50B_100_025d	0.75 1.0 1.0	1.0 0.25 0.875	210	0.75 1.0 1.0	85.1 -7.5 -10.7	13.1 235.1	0.75 1.0 1.0	84.8 -9.9 -14.7	17.7 236.0 4.6	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
732	G50B_100_037d	0.625 1.0 1.0	1.0 0.375 0.812	210	0.625 1.0 1.0	79.8 -11.2 -16.1	19.6 235.1	0.625 1.0 1.0	78.6 -14.0 -21.7	25.8 237.2 6.3	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
733	G50B_100_050d	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	74.4 -15.0 -21.5	26.2 235.1	0.5 1.0 1.0	73.1 -16.9 -27.8	32.5 238.7 6.7	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
734	G50B_100_062d	0.375 1.0 1.0	1.0 0.625 0.687	210	0.375 1.0 1.0	69.1 -18.7 -26.9	32.8 235.1	0.375 1.0 1.0	67.3 -20.0 -32.8	38.5 238.6 6.3	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
735	G50B_100_075d	0.25 1.0 1.0	1.0 0.75 0.625	210	0.25 1.0 1.0	63.8 -22.5 -32.3	39.3 235.1	0.25 1.0 1.0	59.2 -26.0 -38.7	46.6 236.0 8.6	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
736	G50B_100_087d	0.125 1.0 1.0	1.0 0.875 0.562	210	0.125 1.0 1.0	58.4 -26.2 -37.7	45.9 235.1	0.125 1.0 1.0	54.7 -28.9 -42.5	51.4 235.7 6.6	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
737	G50B_100_100d	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1	0.0 1.0 1.0	52.2 -29.2 -44.1	52.9 236.4 1.5	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
738	ROOY_100_012d	1.0 0.875 0.875	1.0 0.125 0.937	390	1.0 0.875 0.875	89.8 7.1 4.7	8.5 33.4	1.0 0.875 0.875	91.5 4.1 3.9	5.7 43.0 3.5	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
739	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.8 0.0 0.0	0.0 0.0	0.875 0.875 0.875	90.3 0.0 -0.2	0.2 277.5 3.5	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
740	G50B_087_012d	0.75 0.875 0.875	0.875 0.125 0.812	210	0.75 0.875 0.875	81.4 -3.7 -5.3	6.5 235.1	0.75 0.875 0.875	85.5 -7.5 -10.5	12.9 234.5 7.5	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
741	G50B_087_025d	0.625 0.875 0.875	0.875 0.25 0.75	210	0.625 0.875 0.875	76.1 -7.5 -10.7	13.1 235.1	0.625 0.875 0.875	80.6 -12.5 -18.7	22.5 236.2 10.4	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
742	G50B_087_037d	0.5 0.875 0.875	0.875 0.375 0.687	210	0.5 0.875 0.875	70.8 -11.2 -16.1	19.6 235.1	0.5 0.875 0.875	73.7 -17.3 -25.2	30.7 235.4 11.3	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
743	G50B_087_050d	0.375 0.875 0.875	0.875 0.5 0.625	210	0.375 0.875 0.875	65.4 -15.0 -21.5	26.2 235.1	0.375 0.875 0.875	68.3 -20.0 -31.6	37.4 237.7 11.6	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
744	G50B_087_062d	0.25 0.875 0.875	0.875 0.625 0.562	210	0.25 0.875 0.875	60.1 -18.7 -26.9	32.8 235.1	0.25 0.875 0.875	63.0 -23.7 -34.2	41.7 235.2 9.3	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
745	G50B_087_075d	0.125 0.875 0.875	0.875 0.75 0.5	210	0.125 0.875 0.875	54.8 -22.5 -32.3	39.3 235.1	0.125 0.875 0.875	57.0 -28.4 -39.3	48.5 234.1 9.4	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
746	G50B_087_087d	0.0 0.875 0.875	0.875 0.875 0.437	210	0.0 0.875 0.875	49.4 -26.2 -37.7	45.9 235.1	0.0 0.875 0.875	52.0 -31.5 -42.1	52.6 233.1 7.3	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
747	ROOY_100_025d	1.0 0.75 0.75	1.0 0.25 0.875	390	1.0 0.75 0.75	83.7 14.3 9.4	17.1 33.4	1.0 0.75 0.75	86.6 8.0 10.0	12.8 51.3 6.9	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
748	ROOY_087_012d	0.875 0.75 0.75	0.875 0.125 0.812	390	0.875 0.75 0.75	80.8 7.1 4.7	8.5 33.4	0.875 0.75 0.75	83.1 5.7 3.9	7.0 34.1 2.8	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
749	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0	0.75 0.75 0.75	80.9 0.0 -0.5	0.5 269.9 3.1	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
750	G50B_075_012d	0.625 0.75 0.75	0.75 0.125 0.687	210	0.625 0.75 0.75	72.5 -3.7 -5.3	6.5 235.1	0.625 0.75 0.75	77.5 -6.9 -10.7	12.8 237.0 8.0	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
751	G50B_075_025d	0.5 0.75 0.75	0.75 0.25 0.625	210	0.5 0.75 0.75	67.1 -7.5 -10.7	13.1 235.1	0.5 0.75 0.75	71.5 -14.0 -19.5	24.0 234.3 11.7	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
752	G50B_075_037d	0.375 0.75 0.75	0.75 0.375 0.562	210	0.375 0.75 0.75	61.8 -11.2 -16.1	19.6 235.1	0.375 0.75 0.75	66.6 -17.0 -24.9	30.2 235.6 11.6	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
753	G50B_075_050d	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.75	56.4 -15.0 -21.5	26.2 235.1	0.25 0.75 0.75	60.0 -20.0 -30.4	36.4 236.6 10.8	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
754	G50B_075_062d	0.125 0.75 0.75	0.75 0.625 0.437	210	0.125 0.75 0.75	51.1 -18.7 -26.9	32.8 235.1	0.125 0.75 0.75	55.7 -25.2 -35.3	43.4 234.5 11.5	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
755	G50B_075_075d	0.0 0.75 0.75	0.75 0.75 0.375	210	0.0 0.75 0.75	45.8 -22.5 -32.3	39.3 235.1	0.0 0.75 0.75	52.0 -33.9 -38.5	51.3 228.5 14.4	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
756	ROOY_100_037d	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	77.7 21.4 14.1	25.7 33.4	1.0 0.625 0.625	79.0 15.4 17.7	23.4 48.9 7.1	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
757	ROOY_087_025d	0.875 0.625 0.625	0.875 0.25 0.75	390	0.875 0.625 0.625	74.7 14.3 9.4	17.1 33.4	0.875 0.625 0.625	76.0 9.9 12.8	16.2 52.1 5.6	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
758	ROOY_075_012d	0.75 0.625 0.625	0.75 0.125 0.687	390	0.75 0.625 0.625	71.8 7.1 4.7	8.5 33.4	0.75 0.625 0.625	74.8 5.4 4.1	6.8 37.0 3.5	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
759	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.8 0.0 0.0	0.0 0.0	0.625 0.625 0.625	71.8 0.0 -1.0	1.0 270.7 3.1	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
760	G50B_062_012d	0.5 0.625 0.625	0.625 0.125 0.562	210	0.5 0.625 0.625	63.5 -3.7 -5.3	6.5 235.1	0.5 0.625 0.625	67.1 -7.7 -13.2	15.3 239.7 9.5	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
761	G50B_062_025d	0.375 0.625 0.625	0.625 0.25 0.5	210	0.375 0.625 0.625	58.1 -7.5 -10.7	13.1 235.1	0.375 0.625 0.625	61.5 -13.9 -20.3	24.7 235.6 12.0	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
762	G50B_062_037d	0.25 0.625 0.625	0.625 0.375 0.437	210	0.25 0.625 0.625	52.8 -11.2 -16.1	19.6 235.1	0.25 0.625 0.625	55.0 -17.0 -27.0	31.9 237.6 12.5	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
763	G50B_062_050d	0.125 0.625 0.625	0.625 0.5 0.375	210	0.125 0.625 0.625	47.4 -15.0 -21.5	26.2 235.1	0.125 0.625 0.625	51.6 -22.0 -31.5	38.5 234.9 12.9	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
764	G50B_062_062d	0.0 0.625 0.625	0.625 0.625 0.312	210	0.0 0.625 0.625	42.1 -18.7 -26.9	32.8 235.1	0.0 0.625 0.625	49.9 -30.2 -39.0	49.4 232.3 18.4	210	1.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1
765	ROOY_100_050d	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	71.7 28.6 18.9	34.3 33.4	1.0 0.5 0.5	72.0 23.9 24.2	34.0 45.3 7.0	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
766	ROOY_087_037d	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.5	68.7 21.4 14.1	25.7 33.4	0.875 0.5 0.5	70.4 16.5 16.7	23.5 45.4 5.8	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
767	ROOY_075_025d	0.75 0.5 0.5	0.75 0.25 0.625	390	0.75 0.5 0.5	65.7 14.3 9.4	17.1 33.4	0.75 0.5 0.5	66.8 10.9 11.1	15.6 45.6 3.9	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
768	ROOY_062_012d	0.625 0.5 0.5	0.625 0.125 0.562	390	0.625 0.5 0.5	62.8 7.1 4.7	8.5 33.4	0.625 0.5 0.5	64.1 6.0 2.5	6.5 22.6 2.7	389	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4
769	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	59.8 0.0 0.0	0.0 0.0	0.5 0.5 0.5	61.1 0.0 -1.1	1.1 268.7 1.7	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
770	G50B_050_012d	0.375 0.5 0												

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md		
810	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	96.1 -0.1 0.0	0.1 188.0	0.3 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
811	BOOR_100_012d	0.875 0.875 1.0	1.0 0.125 0.937	270	0.875 0.875 1.0	87.9 2.1 -5.5	5.9 290.8	0.875 0.875 1.0	88.8 0.5 -7.8	7.8 273.8	2.9 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
812	BOOR_100_025d	0.75 0.75 1.0	1.0 0.25 0.875	270	0.75 0.75 1.0	80.0 4.2 -11.1	11.9 290.8	0.75 0.75 1.0	79.2 2.5 -15.9	16.1 278.9	5.2 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
813	BOOR_100_037d	0.625 0.625 1.0	1.0 0.375 0.812	270	0.625 0.625 1.0	72.1 6.3 -16.7	17.8 290.8	0.625 0.625 1.0	65.8 5.5 -25.7	26.3 282.2	10.9 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
814	BOOR_100_050d	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	64.2 8.4 -22.3	23.8 290.8	0.5 0.5 1.0	54.6 10.9 -32.6	34.4 288.5	14.3 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
815	BOOR_100_062d	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	56.2 10.5 -27.8	29.8 290.8	0.375 0.375 1.0	45.7 14.0 -37.9	40.4 290.3	14.9 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
816	BOOR_100_075d	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	48.3 12.7 -33.4	35.7 290.8	0.25 0.25 1.0	38.1 18.6 -41.2	45.2 294.2	14.1 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
817	BOOR_100_087d	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	40.4 14.8 -39.0	41.7 290.8	0.125 0.125 1.0	34.5 19.2 -42.9	47.0 294.1	8.3 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
818	BOOR_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8	0.0 0.0 1.0	30.8 20.2 -44.2	48.6 294.6	3.7 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
819	Y00G_100_012d	1.0 1.0 0.875	1.0 0.125 0.937	90	1.0 1.0 0.875	95.3 -1.9 10.5	10.7 100.5	1.0 1.0 0.875	95.9 -3.8 9.4	10.2 111.9	2.2 89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
820	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.8 0.0 0.0	0.0 0.0	0.875 0.875 0.875	91.1 0.0 -0.1	0.1 242.5	4.3 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0
821	BOOR_087_012d	0.75 0.75 0.875	0.875 0.125 0.812	270	0.75 0.75 0.875	78.9 2.1 -5.5	5.9 290.8	0.75 0.75 0.875	80.2 1.7 -11.6	11.7 278.4	6.2 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
822	BOOR_087_025d	0.625 0.625 0.875	0.875 0.25 0.75	270	0.625 0.625 0.875	71.0 4.2 -11.1	11.9 290.8	0.625 0.625 0.875	69.8 4.2 -20.4	20.8 281.7	9.3 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
823	BOOR_087_037d	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.5 0.875	63.1 6.3 -16.7	17.8 290.8	0.5 0.5 0.875	58.7 8.3 -29.7	30.9 285.6	13.9 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
824	BOOR_087_050d	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	55.2 8.4 -22.3	23.8 290.8	0.375 0.375 0.875	47.8 12.6 -36.7	38.8 288.9	16.7 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
825	BOOR_087_062d	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	47.2 10.5 -27.8	29.8 290.8	0.25 0.25 0.875	39.6 16.7 -41.1	44.4 292.2	16.5 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
826	BOOR_087_075d	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	39.3 12.7 -33.4	35.7 290.8	0.125 0.125 0.875	35.1 17.5 -43.1	46.5 292.1	11.5 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
827	BOOR_087_087d	0.0 0.0 0.875	0.875 0.875 0.437	270	0.0 0.0 0.875	31.4 14.8 -39.0	41.7 290.8	0.0 0.0 0.875	29.1 21.9 -44.4	49.5 296.2	9.2 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
828	Y00G_100_025d	1.0 1.0 0.75	1.0 0.25 0.875	90	1.0 1.0 0.75	94.7 -3.9 21.1	21.5 100.5	1.0 1.0 0.75	95.2 -8.7 24.8	26.3 109.2	6.0 89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
829	Y00G_087_012d	0.875 0.875 0.75	0.875 0.125 0.812	90	0.875 0.875 0.75	86.3 -1.9 10.5	10.7 100.5	0.875 0.875 0.75	91.4 -3.2 9.1	9.6 109.3	5.5 89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
830	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0	0.75 0.75 0.75	82.2 0.0 -0.5	0.5 270.3	4.4 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0
831	BOOR_075_012d	0.625 0.625 0.75	0.75 0.125 0.687	270	0.625 0.625 0.75	69.9 2.1 -5.5	5.9 290.8	0.625 0.625 0.75	71.5 2.2 -12.7	12.9 280.2	7.3 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
832	BOOR_075_025d	0.5 0.5 0.75	0.75 0.25 0.625	270	0.5 0.5 0.75	62.0 4.2 -11.1	11.9 290.8	0.5 0.5 0.75	60.6 4.4 -20.2	20.7 282.3	9.2 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
833	BOOR_075_037d	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	54.1 6.3 -16.7	17.8 290.8	0.375 0.375 0.75	50.8 6.5 -27.4	28.2 283.3	11.2 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
834	BOOR_075_050d	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	46.2 8.4 -22.3	23.8 290.8	0.25 0.25 0.75	44.3 10.9 -33.2	35.0 288.2	11.3 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
835	BOOR_075_062d	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	38.2 10.5 -27.8	29.8 290.8	0.125 0.125 0.75	38.2 13.3 -39.1	41.3 288.7	11.6 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
836	BOOR_075_075d	0.0 0.0 0.75	0.75 0.75 0.375	270	0.0 0.0 0.75	30.3 12.7 -33.4	35.7 290.8	0.0 0.0 0.75	29.0 20.6 -44.7	49.2 294.7	13.8 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
837	Y00G_100_037d	1.0 1.0 0.625	1.0 0.375 0.812	90	1.0 1.0 0.625	94.2 -5.9 31.7	32.3 100.5	1.0 1.0 0.625	93.4 -9.0 33.9	35.0 104.9	3.8 89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
838	Y00G_087_025d	0.875 0.875 0.625	0.875 0.25 0.75	90	0.875 0.875 0.625	85.7 -3.9 21.1	21.5 100.5	0.875 0.875 0.625	90.1 -6.3 24.9	25.7 104.2	6.2 89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
839	Y00G_075_012d	0.75 0.75 0.625	0.75 0.125 0.687	90	0.75 0.75 0.625	77.3 -1.9 10.5	10.7 100.5	0.75 0.75 0.625	83.0 -2.2 7.7	8.1 106.0	6.4 89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
840	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.8 0.0 0.0	0.0 0.0	0.625 0.625 0.625	72.5 0.0 -1.0	1.0 270.5	3.8 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0
841	BOOR_062_012d	0.5 0.5 0.625	0.625 0.125 0.562	270	0.5 0.5 0.625	60.9 2.1 -5.5	5.9 290.8	0.5 0.5 0.625	61.7 2.6 -14.3	14.5 280.2	8.8 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
842	BOOR_062_025d	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	53.0 4.2 -11.1	11.9 290.8	0.375 0.375 0.625	50.5 4.0 -22.7	23.0 280.0	11.8 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
843	BOOR_062_037d	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	45.1 6.3 -16.7	17.8 290.8	0.25 0.25 0.625	44.0 8.2 -27.8	29.0 286.5	11.3 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
844	BOOR_062_050d	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	37.2 8.4 -22.3	23.8 290.8	0.125 0.125 0.625	37.5 11.3 -34.2	36.1 288.3	12.3 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
845	BOOR_062_062d	0.0 0.0 0.625	0.625 0.625 0.312	270	0.0 0.0 0.625	29.2 10.5 -27.8	29.8 290.8	0.0 0.0 0.625	28.0 21.5 -45.0	49.9 295.5	20.4 270	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8
846	Y00G_100_050d	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	93.7 -7.9 42.3	43.0 100.5	1.0 1.0 0.5	92.9 -11.3	44.8 46.2	10.4 89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
847	Y00G_087_037d	0.875 0.875 0.5	0.875 0.375 0.687	90	0.875 0.875 0.5	85.2 -5.9 31.7	32.3 100.5	0.875 0.875 0.5	90.0 -8.5	36.5 37.5	10.3 89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
848	Y00G_075_025d	0.75 0.75 0.5	0.75 0.25 0.625	90	0.75 0.75 0.5	76.7 -3.9 21.1	21.5 100.5	0.75 0.75 0.5	82.9 -6.7	24.9 25.8	10.2 89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
849	Y00G_062_012d	0.625 0.625 0.5	0.625 0.125 0.562	90	0.625 0.625 0.5	68.3 -1.9 10.5	10.7 100.5	0.625 0.625 0.5	73.9 -3.5	7.3 8.1	115.9 89	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5
850	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	59.8 0.0 0.0	0.0 0.0	0.5 0.5 0.5	61.9 0.0 -1.2	1.2 270.4	2.4 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0
851	BOOR_050_012d	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.375									

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)

TUB material: code=rh4ta

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
891	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	96.1 -0.1 0.0	0.1 168.6 0.3	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
892	B50R_100_012d	1.0 0.875 1.0	1.0 0.125 0.937	330	1.0 0.875 1.0	89.8 8.1 -1.5	8.3 348.9	1.0 0.875 1.0	91.4 7.6 -3.0	8.1 338.4 2.1	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
893	B50R_100_025d	1.0 0.75 1.0	1.0 0.25 0.875	330	1.0 0.75 1.0	83.9 16.3 -3.1	16.6 348.9	1.0 0.75 1.0	85.5 16.1 -5.2	16.9 341.9 2.6	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
894	B50R_100_037d	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	77.9 24.5 -4.7	24.9 348.9	1.0 0.625 1.0	78.4 26.0 -7.1	26.9 344.6 2.7	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
895	B50R_100_050d	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	72.0 32.7 -6.3	33.3 348.9	1.0 0.5 1.0	71.7 36.3 -8.9	37.4 346.1 4.4	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
896	B50R_100_062d	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	66.0 40.8 -7.9	41.6 348.9	1.0 0.375 1.0	66.1 44.6 -9.4	45.6 348.0 4.0	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
897	B50R_100_075d	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	60.1 49.0 -9.5	49.9 348.9	1.0 0.25 1.0	59.2 53.6 -11.4	54.8 347.9 4.9	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
898	B50R_100_087d	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	54.1 57.2 -11.1	58.3 348.9	1.0 0.125 1.0	52.3 65.2 -11.5	66.2 349.9 8.2	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
899	B50R_100_100d	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9	1.0 0.0 1.0	48.3 65.1 -12.7	66.3 348.9 0.3	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
900	GO0B_100_012d	0.875 1.0 0.875	1.0 0.125 0.937	150	0.875 1.0 0.875	90.6 -8.4 3.8	9.2 155.5	0.875 1.0 0.875	92.1 -6.9 1.8	7.1 165.1 2.9	149	0.0 1.0 0.0	54.3 -67.6	30.8 74.3 155.5
901	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.8 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	91.6 0.0 -0.2	0.2 263.1 4.8	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
902	B50R_087_012d	0.875 0.75 0.875	0.875 0.125 0.812	330	0.875 0.75 0.875	80.8 8.1 -1.5	8.3 348.9	0.875 0.75 0.875	84.3 8.6 -5.6	10.2 326.9 5.3	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
903	B50R_087_025d	0.875 0.625 0.875	0.875 0.25 0.75	330	0.875 0.625 0.875	74.9 16.3 -3.1	16.6 348.9	0.875 0.625 0.875	77.2 18.7 -8.0	20.4 336.6 5.9	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
904	B50R_087_037d	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	68.9 24.5 -4.7	24.9 348.9	0.875 0.5 0.875	70.8 27.7 -10.0	29.5 340.0 6.4	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
905	B50R_087_050d	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	63.0 32.7 -6.3	33.3 348.9	0.875 0.375 0.875	64.0 37.0 -11.8	38.8 342.2 7.0	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
906	B50R_087_062d	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	57.0 40.8 -7.9	41.6 348.9	0.875 0.25 0.875	57.7 45.0 -13.6	47.1 343.1 7.1	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
907	B50R_087_075d	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	51.1 49.0 -9.5	49.9 348.9	0.875 0.125 0.875	49.8 55.9 -16.4	58.3 343.6 9.8	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
908	B50R_087_087d	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	45.1 57.2 -11.1	58.3 348.9	0.875 0.0 0.875	44.7 59.0 -15.8	61.1 344.9 4.9	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
909	GO0B_100_025d	0.75 1.0 0.75	1.0 0.25 0.875	150	0.75 1.0 0.75	85.4 -16.9 7.7	18.5 155.5	0.75 1.0 0.75	87.7 -14.1 8.3	16.4 149.3 3.6	149	0.0 1.0 0.0	54.3 -67.6	30.8 74.3 155.5
910	GO0B_087_012d	0.75 0.875 0.75	0.875 0.125 0.812	150	0.75 0.875 0.75	81.6 -8.4 3.8	9.2 155.5	0.75 0.875 0.75	87.2 -9.9 2.7	10.3 164.7 5.9	149	0.0 1.0 0.0	54.3 -67.6	30.8 74.3 155.5
911	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	82.6 0.0 -0.4	0.4 270.1 4.8	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
912	B50R_075_012d	0.75 0.625 0.75	0.75 0.125 0.687	330	0.75 0.625 0.75	71.8 8.1 -1.5	8.3 348.9	0.75 0.625 0.75	75.8 9.0 -5.9	10.8 326.6 5.9	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
913	B50R_075_025d	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	65.9 16.3 -3.1	16.6 348.9	0.75 0.5 0.75	68.2 19.3 -8.4	21.0 336.3 6.4	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
914	B50R_075_037d	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	59.9 24.5 -4.7	24.9 348.9	0.75 0.375 0.75	60.9 27.8 -9.9	29.5 340.2 6.2	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
915	B50R_075_050d	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	54.0 32.7 -6.3	33.3 348.9	0.75 0.25 0.75	54.4 37.2 -12.4	39.2 341.5 7.5	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
916	B50R_075_062d	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	48.0 40.8 -7.9	41.6 348.9	0.75 0.125 0.75	47.9 47.7 -13.2	49.5 344.5 8.6	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
917	B50R_075_075d	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	42.1 49.0 -9.5	49.9 348.9	0.75 0.0 0.75	41.5 53.5 -15.6	55.7 343.6 7.5	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
918	GO0B_100_037d	0.625 1.0 0.625	1.0 0.375 0.812	150	0.625 1.0 0.625	80.2 -25.3 11.5	27.8 155.5	0.625 1.0 0.625	81.7 -21.9 10.5	24.3 154.3 3.9	149	0.0 1.0 0.0	54.3 -67.6	30.8 74.3 155.5
919	GO0B_087_025d	0.625 0.875 0.625	0.875 0.25 0.75	150	0.625 0.875 0.625	76.4 -16.9 7.7	18.5 155.5	0.625 0.875 0.625	82.1 -17.2 9.2	19.6 151.7 5.8	149	0.0 1.0 0.0	54.3 -67.6	30.8 74.3 155.5
920	GO0B_075_012d	0.625 0.75 0.625	0.75 0.125 0.687	150	0.625 0.75 0.625	72.6 -8.4 3.8	9.2 155.5	0.625 0.75 0.625	77.8 -10.0 1.7	10.1 169.8 5.7	149	0.0 1.0 0.0	54.3 -67.6	30.8 74.3 155.5
921	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.8 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	72.1 0.0 -0.9	0.9 268.7 3.4	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
922	B50R_062_012d	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	62.8 8.1 -1.5	8.3 348.9	0.625 0.5 0.625	65.6 9.6 -6.6	11.7 325.4 5.9	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
923	B50R_062_025d	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	56.9 16.3 -3.1	16.6 348.9	0.625 0.375 0.625	57.2 19.8 -8.9	21.7 335.8 6.7	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
924	B50R_062_037d	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	50.9 24.5 -4.7	24.9 348.9	0.625 0.25 0.625	50.0 29.9 -11.7	32.1 338.6 8.8	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
925	B50R_062_050d	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	45.0 32.7 -6.3	33.3 348.9	0.625 0.125 0.625	43.2 39.6 -12.4	41.5 342.5 9.3	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
926	B50R_062_062d	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	39.0 40.8 -7.9	41.6 348.9	0.625 0.0 0.625	37.3 48.6 -13.9	50.5 343.9 9.9	330	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9
927	GO0B_100_050d	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	75.0 -33.8 15.4	37.1 155.5	0.5 1.0 0.5	75.2 -29.8 11.7	32.0 158.4 5.4	149	0.0 1.0 0.0	54.3 -67.6	30.8 74.3 155.5
928	GO0B_087_037d	0.5 0.875 0.5	0.875 0.375 0.687	150	0.5 0.875 0.5	71.2 -25.3 11.5	27.8 155.5	0.5 0.875 0.5	74.3 -28.1 10.0	29.8 160.3 4.4	149	0.0 1.0 0.0	54.3 -67.6	30.8 74.3 155.5
929	GO0B_075_025d	0.5 0.75 0.5	0.75 0.25 0.625	150	0.5 0.75 0.5	67.4 -16.9 7.7	18.5 155.5	0.5 0.75 0.5	71.8 -17.6 5.6	18.5 162.3 4.8	149	0.0 1.0 0.0	54.3 -67.6	30.8 74.3 155.5
930	GO0B_062_012d	0.5 0.625 0.5	0.625 0.125 0.562	150	0.5 0.625 0.5	63.6 -8.4 3.8	9.2 155.5	0.5 0.625 0.5	68.1 -10.1 1.4	10.2 171.5 5.3	149	0.0 1.0 0.0	54.3 -67.6	30.8 74.3 155.5
931	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	59.8 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	61.3 0.0 -1.1	1.1 271.0 1.9	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
932	B50R_050_012d	0.5 0.375 0.5	0.5 0.125 0.437	330	0.5 0.375 0.5	53.8 8.1 -1.5	8.3 348.9	0.5 0.375 0.5	53.3 1					

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

		HIC*Fd				rgb_Fd				ict_Fd				hsi_Fd				rgb*Fd				LabCh*Fd				rgb*Fd				LabCh*Fd				DE**Fd				hsiMd	rgb*Md				LabCh*Md			
972	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	23.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	49.6	1.3	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
973	NW_012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	360	0.125	0.125	0.125	32.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	272.9	5.9	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
974	NW_025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	360	0.25	0.25	0.25	41.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	266.3	2.4	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0				
975	NW_037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	360	0.375	0.375	0.375	50.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	265.7	1.2	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0			
976	NW_050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	360	0.5	0.5	0.5	59.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	268.6	1.4	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0		
977	NW_062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	360	0.625	0.625	0.625	68.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	266.5	3.5	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0		
978	NW_075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	360	0.75	0.75	0.75	77.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	266.9	4.3	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0	
979	NW_087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	360	0.875	0.875	0.875	86.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	248.8	4.6	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0	
980	NW_100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0			
981	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	23.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0			
982	NW_012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	360	0.125	0.125	0.125	32.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
983	NW_025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	360	0.25	0.25	0.25	41.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
984	NW_037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	360	0.375	0.375	0.375	50.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
985	NW_050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	360	0.5	0.5	0.5	59.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
986	NW_062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	360	0.625	0.625	0.625	68.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
987	NW_075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	360	0.75	0.75	0.75	77.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
988	NW_087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	360	0.875	0.875	0.875	86.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
989	NW_100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
990	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	23.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
991	NW_012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	360	0.125	0.125	0.125	32.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
992	NW_025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	360	0.25	0.25	0.25	41.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
993	NW_037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	360	0.375	0.375	0.375	50.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
994	NW_050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	360	0.5	0.5	0.5	59.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
995	NW_062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	360	0.625	0.625	0.625	68.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
996	NW_075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	360	0.75	0.75	0.75	77.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
997	NW_087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	360	0.875	0.875	0.875	86.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0</										

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd		ict_Fd		hsi_Fd		rgb*Fd		LabCh*Fd		rgb**Fd		LabCh**Fd		DE**Fd		hsiM,d	rgb*Md		LabCh*Md												
1053	NW_086d	0.866	0.866	0.866	0.866	0.0	0.866	360	0.866	0.866	0.866	86.1	0.0	0.0	0.0	0.0	0.866	0.866	0.866	90.6	0.0	-0.1	0.1	266.5	4.4	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1054	NW_093d	0.933	0.933	0.933	0.933	0.0	0.933	360	0.933	0.933	0.933	91.0	0.0	0.0	0.0	0.0	0.933	0.933	0.933	94.4	0.0	-0.2	0.2	278.1	3.4	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1055	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.8	0.0	0.0	0.0	152.8	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1056	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	23.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	18.1	0.0	0.2	0.2	83.2	5.6	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1057	NW_006d	0.066	0.066	0.066	0.066	0.0	0.066	360	0.066	0.066	0.066	28.6	0.0	0.0	0.0	0.0	0.066	0.066	0.066	21.5	0.1	0.1	0.2	48.9	7.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1058	NW_013d	0.133	0.133	0.133	0.133	0.0	0.133	360	0.133	0.133	0.133	33.4	0.0	0.0	0.0	0.0	0.133	0.133	0.133	28.9	0.0	-0.7	0.7	268.2	4.4	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1059	NW_020d	0.2	0.2	0.2	0.2	0.0	0.2	360	0.2	0.2	0.2	38.2	0.0	0.0	0.0	0.0	0.2	0.2	0.2	37.3	0.0	-1.1	1.1	267.2	1.4	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1060	NW_026d	0.266	0.266	0.266	0.266	0.0	0.266	360	0.266	0.266	0.266	42.9	0.0	0.0	0.0	0.0	0.266	0.266	0.266	44.2	0.0	-1.1	1.1	269.1	1.7	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1061	NW_033d	0.333	0.333	0.333	0.333	0.0	0.333	360	0.333	0.333	0.333	47.8	0.0	0.0	0.0	0.0	0.333	0.333	0.333	49.9	0.0	-0.8	0.8	274.5	2.3	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1062	NW_040d	0.4	0.4	0.4	0.4	0.0	0.4	360	0.4	0.4	0.4	52.6	0.0	0.0	0.0	0.0	0.4	0.4	0.4	53.8	0.0	-0.9	0.9	273.2	1.4	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1063	NW_046d	0.466	0.466	0.466	0.466	0.0	0.466	360	0.466	0.466	0.466	57.3	0.0	0.0	0.0	0.0	0.466	0.466	0.466	59.7	0.0	-1.1	1.1	268.9	2.6	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1064	NW_053d	0.533	0.533	0.533	0.533	0.0	0.533	360	0.533	0.533	0.533	62.2	0.0	0.0	0.0	0.0	0.533	0.533	0.533	65.4	0.0	-0.9	0.9	273.1	3.3	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1065	NW_060d	0.6	0.6	0.6	0.6	0.0	0.6	360	0.6	0.6	0.6	67.0	0.0	0.0	0.0	0.0	0.6	0.6	0.6	70.2	0.0	-0.8	0.8	268.8	3.2	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1066	NW_066d	0.666	0.666	0.666	0.666	0.0	0.666	360	0.666	0.666	0.666	71.7	0.0	0.0	0.0	0.0	0.666	0.666	0.666	75.5	0.0	-0.7	0.7	271.9	3.8	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1067	NW_073d	0.734	0.734	0.734	0.734	0.0	0.734	360	0.734	0.734	0.734	76.6	0.0	0.0	0.0	0.0	0.734	0.734	0.734	80.8	0.0	-0.4	0.4	265.0	4.1	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1068	NW_080d	0.8	0.8	0.8	0.8	0.0	0.8	360	0.8	0.8	0.8	81.4	0.0	0.0	0.0	0.0	0.8	0.8	0.8	85.3	0.0	-0.3	0.3	279.5	3.9	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1069	NW_086d	0.866	0.866	0.866	0.866	0.0	0.866	360	0.866	0.866	0.866	86.1	0.0	0.0	0.0	0.0	0.866	0.866	0.866	90.2	0.0	0.0	0.0	252.2	4.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1070	NW_093d	0.933	0.933	0.933	0.933	0.0	0.933	360	0.933	0.933	0.933	91.0	0.0	0.0	0.0	0.0	0.933	0.933	0.933	94.2	0.0	-0.2	0.2	289.2	3.2	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1071	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.8	0.0	0.0	0.1	331.9	0.1	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1072	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	23.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	19.2	0.1	0.2	0.2	58.1	4.6	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1073	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.7	0.0	-0.2	0.2	284.6	0.2	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0
1074	R00Y_100_100d	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	47.5	57.2	37.8	68.6	33.4	1.0	0.0	0.0	47.0	56.3	40.2	69.2	35.5	2.6	389	1.0	0.0	0.0	47.5	57.2	37.8	68.6
1075	G50B_100_100d	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	1.0	53.1	-30.0	-43.1	52.5	235.1	0.0	1.0	1.0	54.9	-30.4	-42.0	51.8	234.0	2.1	210	0.0	1.0	1.0	53.1	-30.0	-43.1	52.5
1076	Y00G_100_100d	1.0	1.0	0.0	1.0	1.0	0.5	90	1.0	1.0	0.0	91.5	-15.8	84.6	86.1	100.5	1.0	1.0	0.0	91.5	-16.0	86.1	87.6	100.5	1.5	89	1.0	1.0	0.0	91.5	-15.8	84.6	86.1
1077	B00R_100_100d	0.0	0.0	1.0	1.0	1.0	0.5	270	0.0	0.0	1.0	32.5	16.9	-44.6	47.7	290.8	0.0	0.0	1.0	30.7	21.3	-44.1	49.0	295.7	4.7	270	0.0	0.0	1.0	32.5	16.9	-44.6	47.7
1078	G00B_100_100d	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	54.3	-67.6	30.8	74.3	155.5	0.0	1.0	0.0	54.6	-69.2	33.1	76.7	154.3	2.8	149	0.0	1.0	0.0	54.3	-67.6	30.8	74.3
1079	B50R_100_100d	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	48.1	65.4	-12.7	66.6	348.9	1.0	0.0	1.0	48.3	66.3	-13.8	67.7	348.1	1.4	330	1.0	0.0	1.0	48.1	65.4	-12.7	66.6

delta E* = 3.0

2-0032530-F0

PS790-7N, 26/26-F

PE4300P_120901.TXT, 1080 colors, Separation cmyrn6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
colores y diferencia en color, ΔE^* , 3D=0, de=0, *cmyk*

entrada: *rgb/cmyk* -> *rgb*
salida: transfiera a *cmyk*_d

2-0032530-F0

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmyrn6 (CMYK)

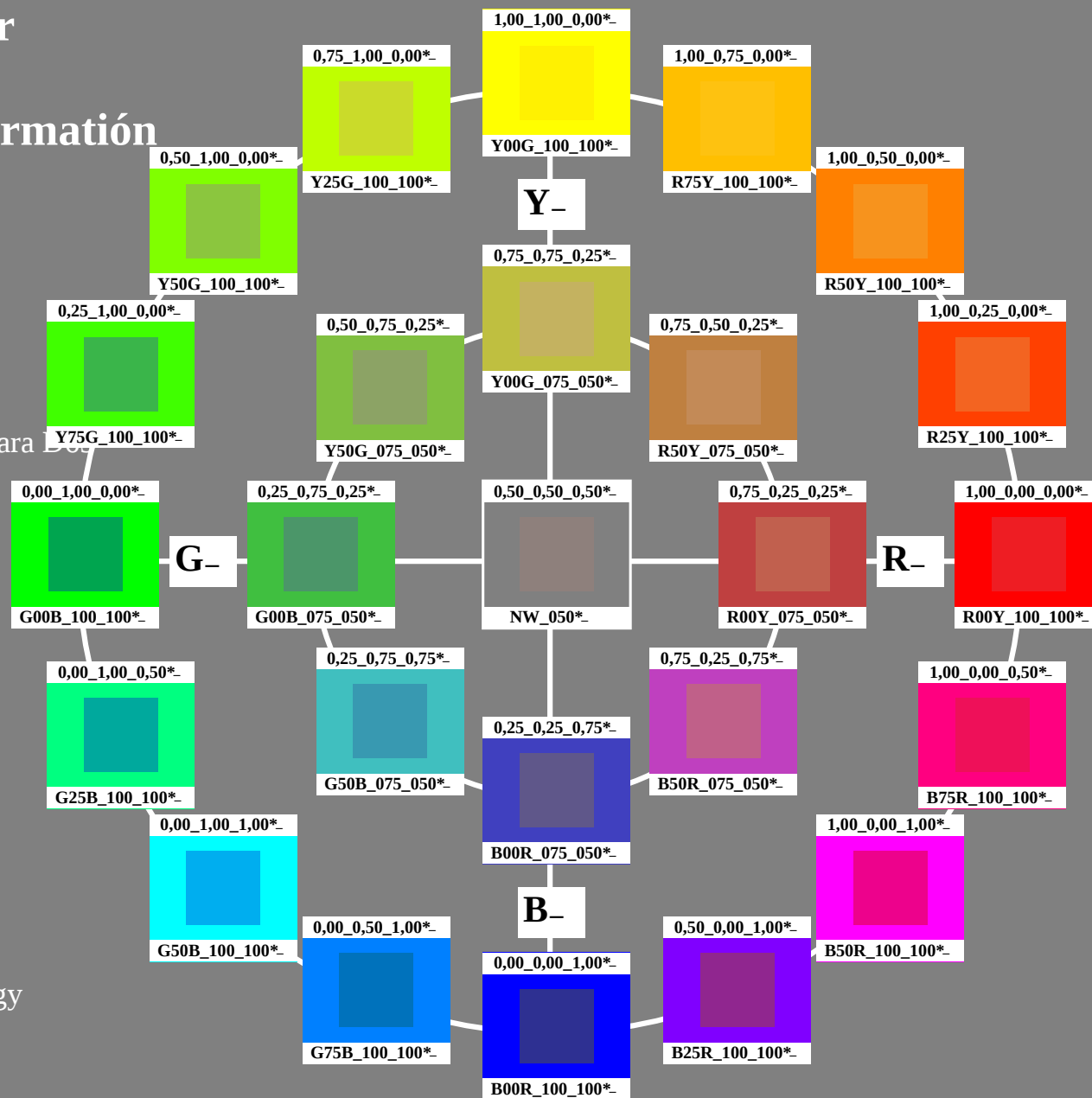
TUB material: code=rha4ta

Color e Visión del Color Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar sRGB
rgb data: rgb^*_e (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : HIC^*_e (derecha)

Impresión especial par la exposición
El Color y la Visión del Color
Section Lighting Technology
of the Berlin University of Technology
Einsteinufer 19, D-10587 Berlin
ver: <http://www.li.tu-berlin.de>
y <http://130.149.60.45/~farbmetrik>



2-013030-L0

PS790-7N

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65

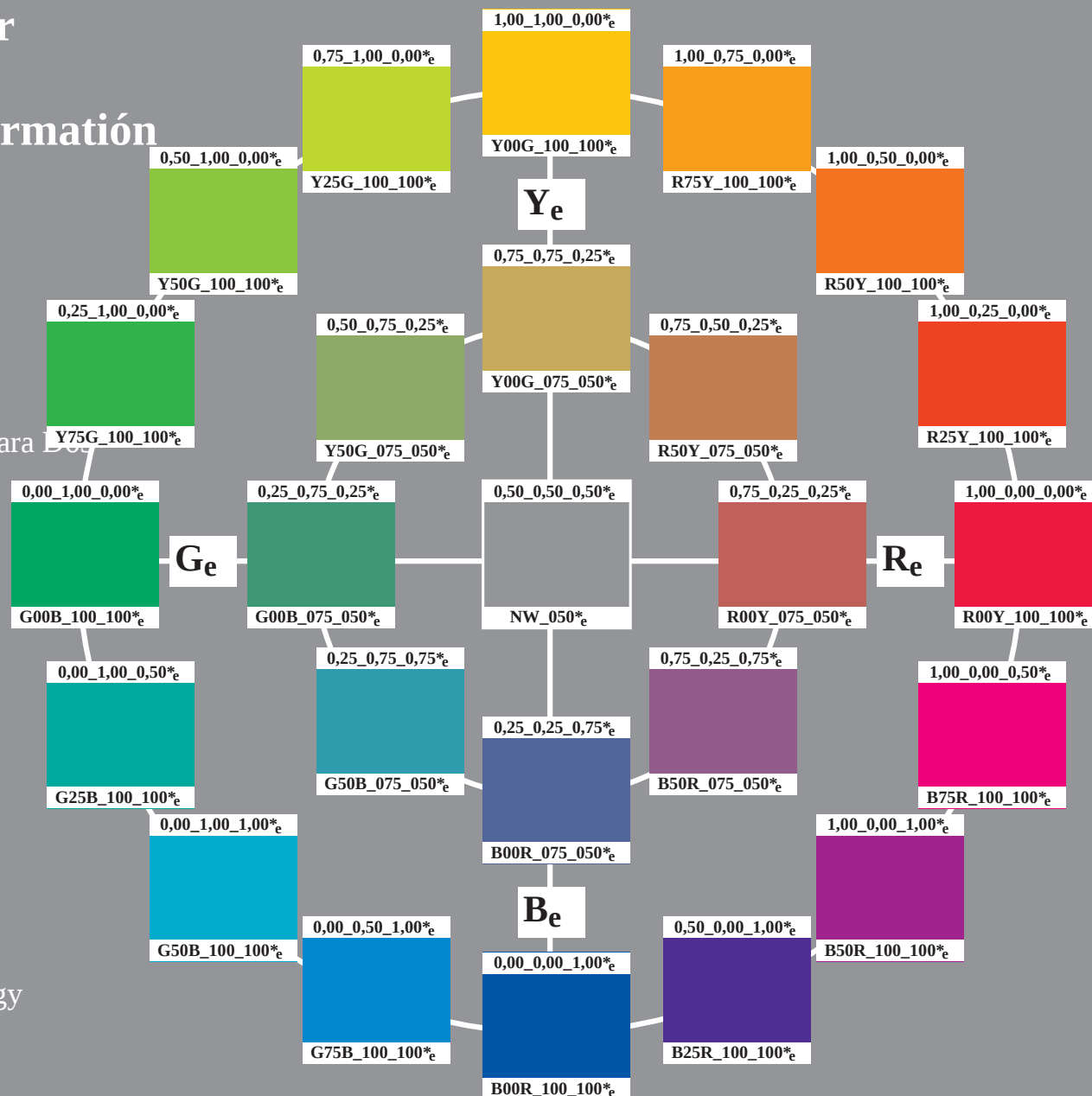
entrada: $rgb/cmyk \rightarrow rgb/cmyk$
salida: ningún cambio

Color e Visión del Color Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar sRGB
rgb data: rgb^*_e (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : HIC^*_e (derecha)

Impresión especial par la exposición
El Color y la Visión del Color
Section Lighting Technology
of the Berlin University of Technology
Einsteinufer 19, D-10587 Berlin
ver: <http://www.li.tu-berlin.de>
y <http://130.149.60.45/~farbmetrik>



2-013130-L0

PS790-71

PE4300P_120901.TXT, 1080 colors, Separation cmykn6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=1, cmyk

entrada: $rgb/cmyk \rightarrow rgb_e$
salida: transfiera a $cmyk_e$

2-013130-F0

C

M

Y

O

L

V

C

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS TUB material: code=rha4ta
aplicación para la medida salida de impresora láser, separación cmyk6 (CMYK)

<http://www.li.tu-berlin.de>
<http://130.149.60.45/~farbmetrik>

2-013230-L0

PS790-71

PE4300P_120901.TXT, 1080 colors, Separation cmyk6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=1, cmyk

entrada: *rgb/cmyk* -> *rgb_e*
salida: transfiera a *cmyk_e*

2-013230-E0

C

M

Y

O

L

V

C

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS TUB material: code=rha4ta
aplicación para la medida salida de impresora láser, separación cmyñ6 (CMYK)

<http://www.li.tu-berlin.de>
<http://130.149.60.45/~farbmetrik>

2-013330-L0

PS790-71

PE4300P_120901.TXT, 1080 colors, Separation cmyñ6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=1, cmyk

entrada: *rgb/cmyk* -> *rgb_e*
salida: transfiera a *cmyk_e*

2-013330-E0

C

M

Y

O

L

V

C

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS TUB material: code=rha4ta
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)

Color e Visión del Color

Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar sRGB
rgb data: *rgb**_e (izquierda)
tonos elementales *H**, billantez *I**,
croma *C**: *HIC**_e (derecha)

Impresión especial par la exposición
El Color y la Visión del Color
Section Lighting Technology
of the Berlin University of Technology
Einsteinufer 19, D-10587 Berlin
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y <http://130.149.60.45/~farbmetrik>

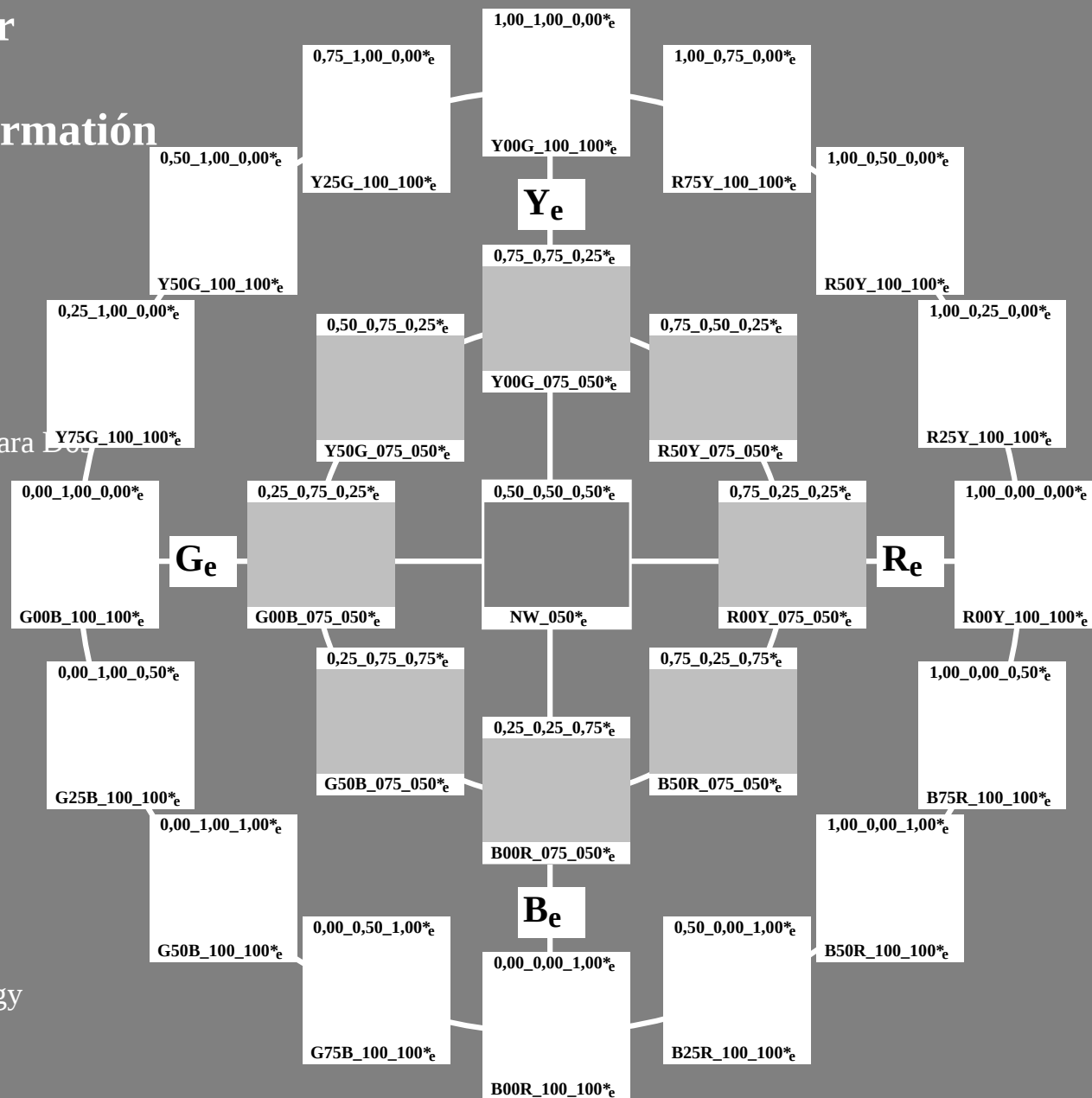


gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=1, *cm*yk

entrada: $rgb/cmyk \rightarrow rgb_e$
salida: transfiera a $cmyk_e$

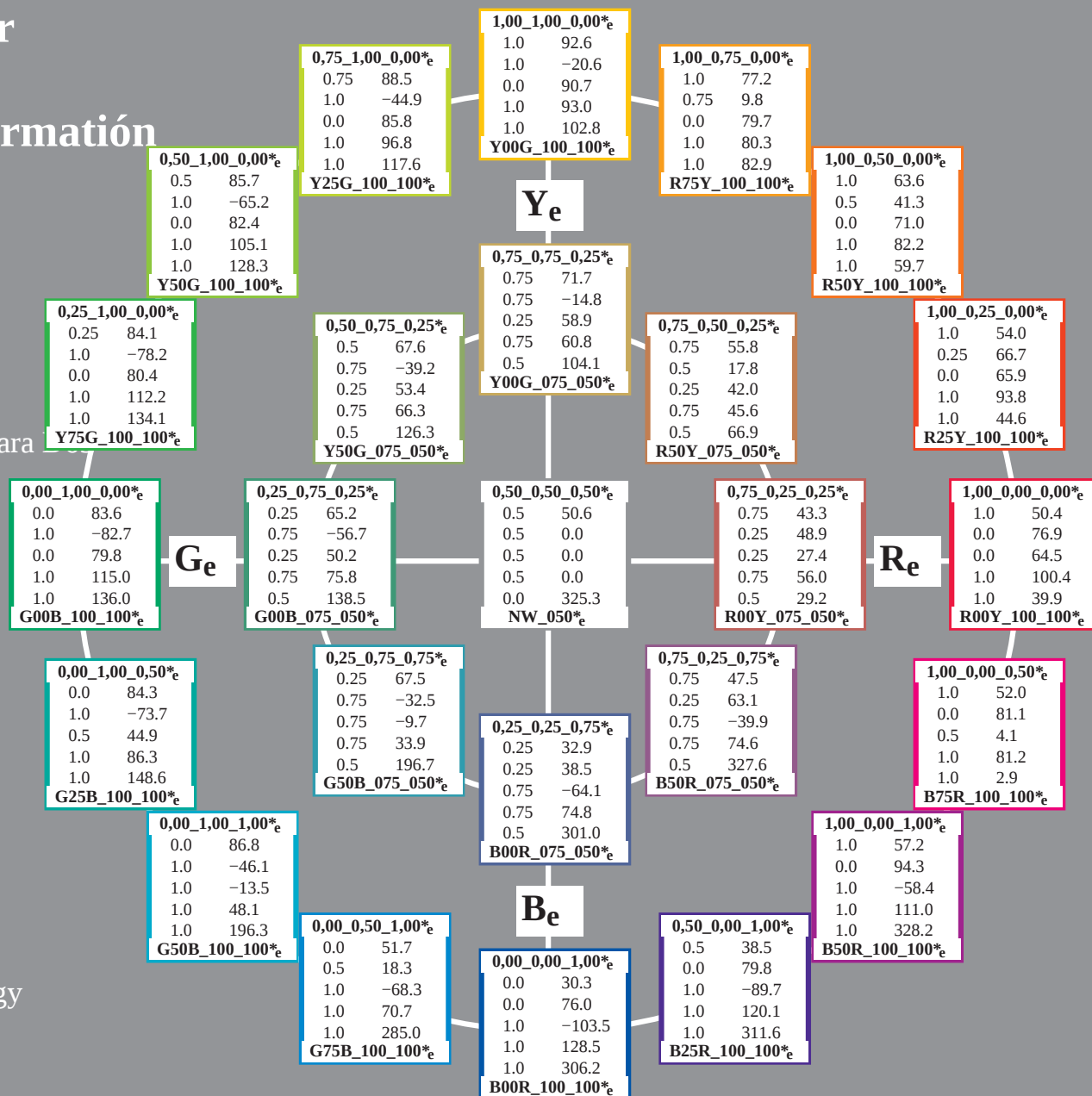
Color e Visión del Color Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar sRGB

rgb data: rgb^*_e (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : HIC^*_e (derecha)
código color:
 $rgbicd$; $LabCh^*_d$

Impresión especial par la exposición
El Color y la Visión del Color
Section Lighting Technology
of the Berlin University of Technology
Einsteinufer 19, D-10587 Berlin
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2-013630-L0

PS790-71

PE4300P_120901.TXT, 1080 colors, Separation cmyn6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=1, cmyk

entrada: $rgb/cmyk \rightarrow rgb_e$
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2-013630-F0

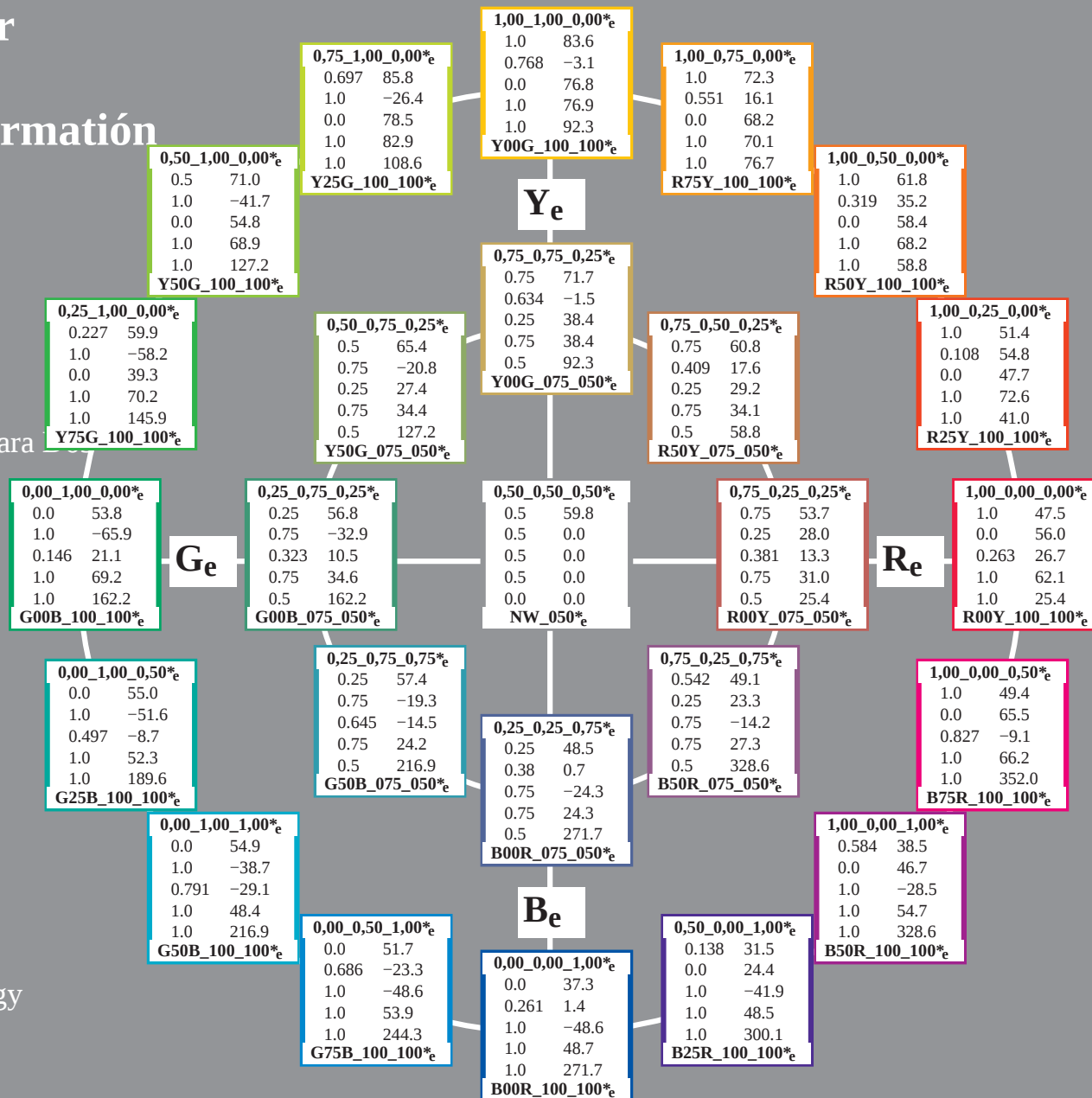
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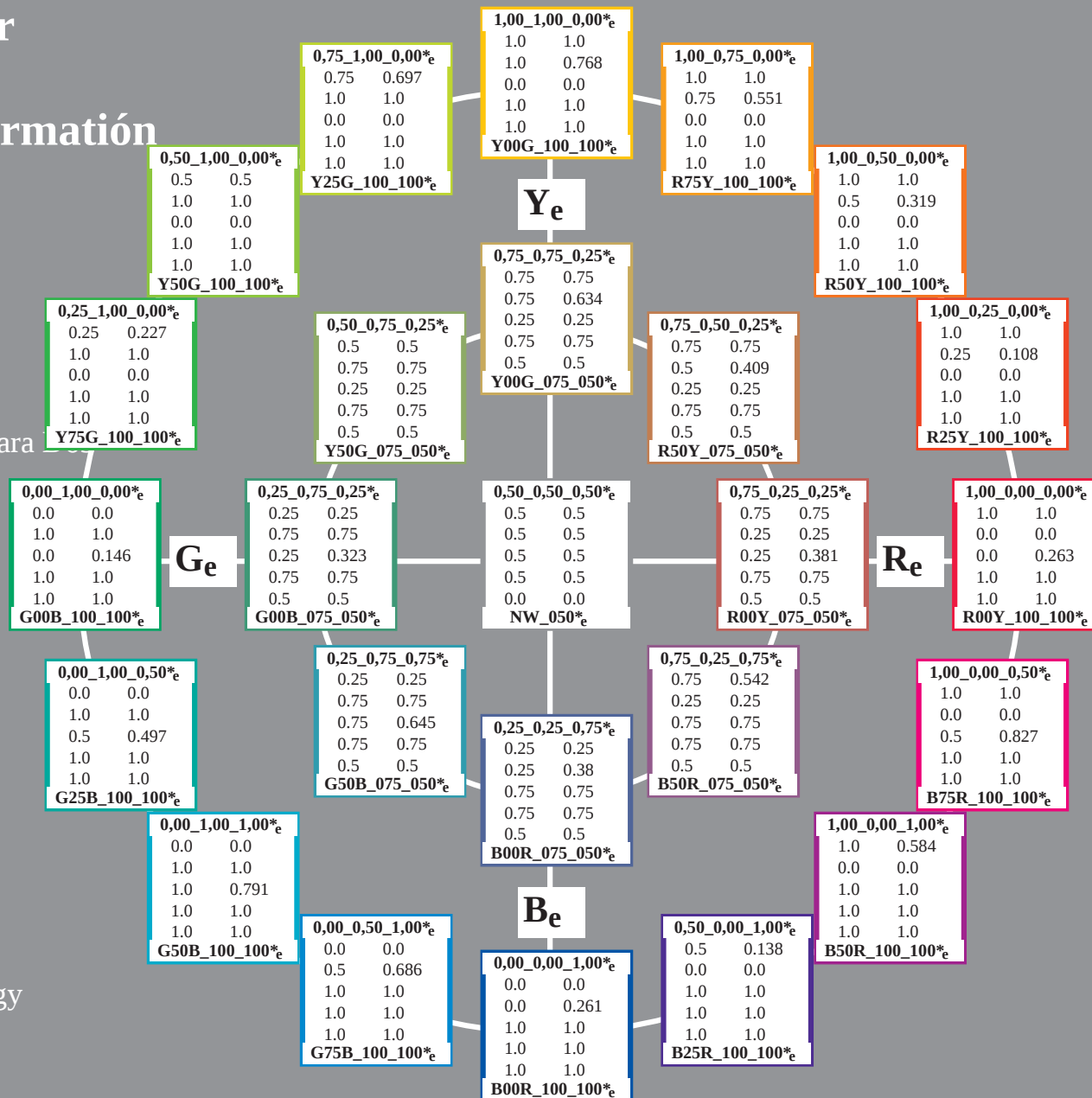
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croma C^* : HIC^*_e (derecha)
código color:
 $rgbic^*_d$; $rgbic^*_e$

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2-013830-L0

PS790-71

PE4300P_120901.TXT, 1080 colors, Separation cmyn6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=1, cmyk

entrada: $rgb/cmyk \rightarrow rgb_e$
salida: transfiera a $cmyk_e$

2-013830-F0

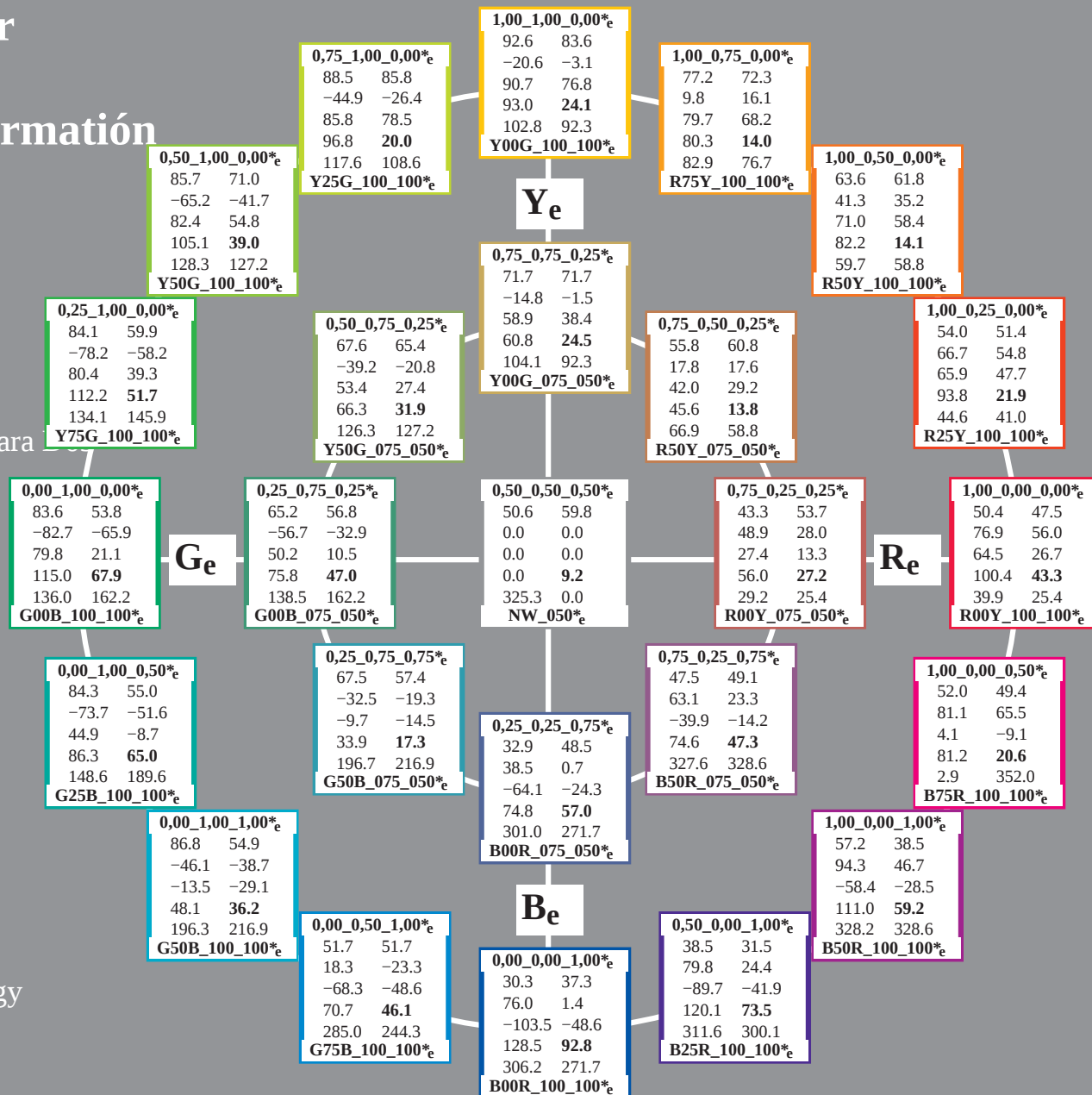
Color e Visión del Color Colores Elementales en la Tecnología de la Información

Author: Prof. Dr. Klaus Richter

25 colores para la medida del color para
círculo di tono con 16 y 8 pasos
pantalla estándar sRGB

rgb data: rgb^*_e (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : HIC^*_e (derecha)
código color:
 $LabCh^*_d$; $Lab^*/DE^*/h^*_e$

Impresión especial par la exposición
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2-013930-L0

PS790-71

PE4300P_120901.TXT, 1080 colors, Separation cmykn6*

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
25 colores del estándar para D65, 3D=0, de=1, cmyk

entrada: $rgb/cmyk \rightarrow rgb_e$
salida: transfiera a $cmyk_e$

2-013930-F0

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS TUB material: code=rha4ta
aplicación para la medida salida de impresora láser, separación cmyñ6 (CMYK)

n/j	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb**Fe	LabCh**Fe	DE**Fe	hsi_Me	rgb*Me	LabCh*Me
0/648	R00Y_100_100e	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.263	47.5 56.0 26.7 62.1 25.4	1.0 0.0 0.0	47.5 57.2 37.8 68.6 33.4 11.1 375	1.0 0.0 0.263	47.5 56.0 26.7 62.1 25.4		
1/657	R13Y_100_100e	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.0 0.012	47.5 57.1 37.5 68.3 33.2	1.0 0.125 0.0	51.9 54.3 49.2 73.2 42.1 12.8 389	1.0 0.0 0.012	47.5 57.1 37.5 68.3 33.2		
2/666	R25Y_100_100e	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.108 0.0	51.4 54.8 47.7 72.6 41.0	1.0 0.25 0.0	58.2 41.8 55.1 69.2 52.8 16.4 35	1.0 0.108 0.0	51.4 54.8 47.7 72.6 41.0		
3/675	R38Y_100_100e	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.216 0.0	56.5 45.2 53.8 70.3 49.9	1.0 0.375 0.0	64.6 29.8 60.4 67.3 63.7 18.5 41	1.0 0.216 0.0	56.5 45.2 53.8 70.3 49.9		
4/684	R50Y_100_100e	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.319 0.0	61.8 35.2 58.4 68.2 58.8	1.0 0.5 0.0	70.5 19.2 66.2 69.0 73.8 19.8 48	1.0 0.319 0.0	61.8 35.2 58.4 68.2 58.8		
5/693	R63Y_100_100e	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.425 0.0	67.0 25.7 63.0 68.0 67.8	1.0 0.625 0.0	74.9 11.4 70.7 71.6 80.7 18.0 55	1.0 0.425 0.0	67.0 25.7 63.0 68.0 67.8		
6/702	R75Y_100_100e	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.551 0.0	72.3 16.1 68.2 70.1 76.7	1.0 0.75 0.0	82.9 -2.0 76.9 77.0 91.5 22.7 63	1.0 0.551 0.0	72.3 16.1 68.2 70.1 76.7		
7/711	R88Y_100_100e	1.0 0.875 0.0	1.0 1.0 0.5	83	1.0 0.668 0.0	77.7 7.0 73.1 73.5 84.5	1.0 0.875 0.0	87.6 -9.0 75.7 76.3 96.8 19.0 70	1.0 0.668 0.0	77.7 7.0 73.1 73.5 84.5		
8/720	Y00G_100_100e	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 0.768 0.0	83.6 -3.1 76.8 76.9 92.3	1.0 1.0 0.0	91.5 -15.8 84.6 86.1 100.5 16.9 77	1.0 0.768 0.0	83.6 -3.1 76.8 76.9 92.3		
9/639	Y13G_100_100e	0.875 1.0 0.0	1.0 1.0 0.5	97	1.0 0.995 0.0	91.4 -15.5 84.4 85.8 100.4	0.875 1.0 0.0	92.8 -18.1 89.4 91.2 101.4 5.8 89	1.0 0.995 0.0	91.4 -15.5 84.4 85.8 100.4		
10/558	Y25G_100_100e	0.75 1.0 0.0	1.0 1.0 0.5	104	0.697 1.0 0.0	85.8 -26.4 78.5 82.9 108.6	0.75 1.0 0.0	90.1 -21.3 86.0 88.6 103.9 10.0 107	0.697 1.0 0.0	85.8 -26.4 78.5 82.9 108.6		
11/477	Y38G_100_100e	0.625 1.0 0.0	1.0 1.0 0.5	112	0.595 1.0 0.0	77.7 -34.4 64.9 73.5 117.9	0.625 1.0 0.0	79.9 -31.7 67.9 75.0 115.0 4.5 113	0.595 1.0 0.0	77.7 -34.4 64.9 73.5 117.9		
12/396	Y50G_100_100e	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	71.0 -41.7 54.8 68.9 127.2	0.5 1.0 0.0	70.9 -41.7 54.8 68.9 127.3 0.0 119	0.5 1.0 0.0	71.0 -41.7 54.8 68.9 127.2		
13/315	Y63G_100_100e	0.375 1.0 0.0	1.0 1.0 0.5	128	0.351 1.0 0.0	65.4 -49.4 46.7 68.0 136.5	0.375 1.0 0.0	66.5 -47.5 48.0 67.6 134.7 2.5 129	0.351 1.0 0.0	65.4 -49.4 46.7 68.0 136.5		
14/234	Y75G_100_100e	0.25 1.0 0.0	1.0 1.0 0.5	136	0.227 1.0 0.0	59.9 -58.2 39.3 70.2 145.9	0.25 1.0 0.0	60.6 -57.2 40.4 70.1 144.7 1.5 137	0.227 1.0 0.0	59.9 -58.2 39.3 70.2 145.9		
15/153	Y88G_100_100e	0.125 1.0 0.0	1.0 1.0 0.5	143	0.04 1.0 0.0	55.2 -65.9 32.0 73.3 154.0	0.125 1.0 0.0	57.0 -62.2 34.4 71.1 151.0 4.7 147	0.04 1.0 0.0	55.2 -65.9 32.0 73.3 154.0		
16/72	G00C_100_100e	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.146	53.8 -65.9 21.1 69.2 162.2	0.0 1.0 0.0	54.3 -67.6 30.8 74.3 155.5 9.8 157	0.0 1.0 0.146	53.8 -65.9 21.1 69.2 162.2		
17/73	G13C_100_100e	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.251	53.7 -63.1 12.7 64.3 168.6	0.0 1.0 0.125	53.8 -66.4 23.0 70.2 160.8 10.8 163	0.0 1.0 0.251	53.7 -63.1 12.7 64.3 168.6		
18/74	G25C_100_100e	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.32	54.3 -59.8 5.2 60.1 175.0	0.0 1.0 0.25	53.7 -63.1 12.8 64.4 168.5 8.2 168	0.0 1.0 0.32	54.3 -59.8 5.2 60.1 175.0		
19/75	G38C_100_100e	0.0 1.0 0.375	1.0 1.0 0.5	172	0.0 1.0 0.404	54.8 -55.6 -2.2 55.7 182.3	0.0 1.0 0.375	54.7 -56.8 0.0 56.8 179.9 2.5 173	0.0 1.0 0.404	54.8 -55.6 -2.2 55.7 182.3		
20/76	G50C_100_100e	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.497	55.0 -51.6 -8.7 52.3 189.6	0.0 1.0 0.5	55.0 -51.4 -8.9 52.2 189.8 0.2 179	0.0 1.0 0.497	55.0 -51.6 -8.7 52.3 189.6		
21/77	G63C_100_100e	0.0 1.0 0.625	1.0 1.0 0.5	188	0.0 1.0 0.56	55.1 -48.2 -14.6 50.4 196.9	0.0 1.0 0.625	55.3 -44.1 -20.0 48.5 204.4 6.7 183	0.0 1.0 0.56	55.1 -48.2 -14.6 50.4 196.9		
22/78	G75C_100_100e	0.0 1.0 0.75	1.0 1.0 0.5	196	0.0 1.0 0.622	55.3 -44.3 -19.9 48.5 204.2	0.0 1.0 0.75	55.2 -39.5 -27.1 47.9 214.4 8.6 188	0.0 1.0 0.622	55.3 -44.3 -19.9 48.5 204.2		
23/79	G88C_100_100e	0.0 1.0 0.875	1.0 1.0 0.5	203	0.0 1.0 0.701	55.2 -41.4 -24.5 48.1 210.5	0.0 1.0 0.875	54.4 -36.7 -33.0 49.4 221.9 9.7 193	0.0 1.0 0.701	55.2 -41.4 -24.5 48.1 210.5		
24/80	C00B_100_100e	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 0.791	54.9 -38.7 -29.1 48.4 216.9	0.0 1.0 1.0	53.1 -30.0 -43.1 52.5 235.1 16.5 198	0.0 1.0 0.791	54.9 -38.7 -29.1 48.4 216.9		
25/71	C13B_100_100e	0.0 0.875 1.0	1.0 1.0 0.5	217	0.0 1.0 0.888	54.3 -36.1 -34.1 49.7 223.3	0.0 0.875 1.0	53.1 -27.9 -44.7 52.7 237.9 13.4 204	0.0 1.0 0.888	54.3 -36.1 -34.1 49.7 223.3		
26/62	C25B_100_100e	0.0 0.75 1.0	1.0 1.0 0.5	224	0.0 1.0 0.948	53.6 -33.1 -39.1 51.2 229.7	0.0 0.75 1.0	52.9 -25.9 -47.5 54.1 241.3 11.1 207	0.0 1.0 0.948	53.6 -33.1 -39.1 51.2 229.7		
27/53	C38B_100_100e	0.0 0.625 1.0	1.0 1.0 0.5	232	0.0 0.915 1.0	53.1 -28.6 -44.2 52.6 237.0	0.0 0.625 1.0	50.5 -20.8 -49.5 53.7 247.2 9.8 214	0.0 0.915 1.0	53.1 -28.6 -44.2 52.6 237.0		
28/44	C50B_100_100e	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.686 1.0	51.7 -23.3 -48.6 53.9 244.3	0.0 0.5 1.0	46.1 -13.3 -49.4 51.1 254.9 11.4 227	0.0 0.686 1.0	51.7 -23.3 -48.6 53.9 244.3		
29/35	C63B_100_100e	0.0 0.375 1.0	1.0 1.0 0.5	248	0.0 0.552 1.0	48.0 -16.4 -49.6 52.2 251.6	0.0 0.375 1.0	41.4 -6.3 -49.2 49.6 262.6 12.0 236	0.0 0.552 1.0	48.0 -16.4 -49.6 52.2 251.6		
30/26	C75B_100_100e	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.434 1.0	43.6 -9.6 -49.4 50.3 258.9	0.0 0.25 1.0	36.8 2.2 -48.5 48.6 272.6 13.7 244	0.0 0.434 1.0	43.6 -9.6 -49.4 50.3 258.9		
31/17	C88B_100_100e	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.341 1.0	40.1 -4.0 -49.2 49.4 265.3	0.0 0.125 1.0	35.0 9.4 -46.3 47.3 281.4 14.6 250	0.0 0.341 1.0	40.1 -4.0 -49.2 49.4 265.3		
32/8	B00M_100_100e	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.261 1.0	37.3 1.4 -48.6 48.7 271.7	0.0 0.0 1.0	32.5 16.9 -44.6 47.7 290.8 16.6 255	0.0 0.261 1.0	37.3 1.4 -48.6 48.7 271.7		
33/89	B13M_100_100e	0.125 0.0 1.0	1.0 1.0 0.5	277	0.0 0.168 1.0	35.7 6.9 -47.2 47.7 278.3	0.125 0.0 1.0	31.6 23.6 -42.2 48.4 299.2 17.8 260	0.0 0.168 1.0	35.7 6.9 -47.2 47.7 278.3		
34/170	B25M_100_100e	0.25 0.0 1.0	1.0 1.0 0.5	284	0.0 0.077 1.0	34.1 12.2 -45.8 47.4 285.0	0.25 0.0 1.0	31.0 30.5 -39.3 49.8 307.8 19.6 266	0.0 0.077 1.0	34.1 12.2 -45.8 47.4 285.0		
35/251	B38M_100_100e	0.375 0.0 1.0	1.0 1.0 0.5	292	0.026 0.0 1.0	32.3 18.3 -44.1 47.8 292.5	0.375 0.0 1.0	34.2 38.2 -35.0 51.8 317.5 21.9 271	0.026 0.0 1.0	32.3 18.3 -44.1 47.8 292.5		
36/332	B50M_100_100e	0.5 0.0 1.0	1.0 1.0 0.5	300	0.138 0.0 1.0	31.5 24.4 -41.9 48.5 300.1	0.5 0.0 1.0	37.2 43.1 -30.8 53.0 324.4 22.5 277	0.138 0.0 1.0	31.5 24.4 -41.9 48.5 300.1		
37/413	B63M_100_100e	0.625 0.0 1.0	1.0 1.0 0.5	308	0.249 0.0 1.0	31.0 30.5 -39.4 49.8 307.7	0.625 0.0 1.0	39.1 48.4 -27.2 55.6 330.6 23.1 283	0.249 0.0 1.0	31.0 30.5 -39.4 49.8 307.7		
38/494	B75M_100_100e	0.75 0.0 1.0	1.0 1.0 0.5	316	0.347 0.0 1.0	33.5 36.5 -36.1 51.4 315.3	0.75 0.0 1.0	41.8 55.1 -21.4 59.1 338.7 25.1 289	0.347 0.0 1.0	33.5 36.5 -36.1 51.4 315.3		
39/575	B88M_100_100e	0.875 0.0 1.0	1.0 1.0 0.5	323	0.455 0.0 1.0	36.1 41.4 -32.4 52.6 321.9	0.875 0.0 1.0	45.6 60.1 -17.3 62.6 343.9 25.8 297	0.455 0.0 1.0	36.1 41.4 -32.4 52.6 321.9		
40/656	M00R_100_100e	1.0 0.0 1.0	1.0 1.0 0.5	330	0.584 0.0 1.0	38.5 46.7 -28.5 54.7 328.6	1.0 0.0 1.0	48.1 65.4 -12.7 66.6 348.9 26.2 305	0.584 0.0 1.0	38.5 46.7 -28.5 54.7 328.6		
41/655	M13R_100_100e	1.0 0.0 0.875	1.0 1.0 0.5	337	0.696 0.0 1.0	40.6 52.3 -24.1 57.6 335.2	1.0 0.0 0.875	49.5 66.1 -10.7 67.0 350.7 21.1 312	0.696 0.0 1.0	40.6 52.3 -24.1 57.6 335.2		
42/654	M25R_100_100e	1.0 0.0 0.75	1.0 1.0 0.5	344	0.825 0.0 1.0	44.1 58.2 -19.0 61.2 341.8	1.0 0.0 0.75	49.3 64.5 -6.5 64.8 354.2 14.9 320	0.825 0.0 1.0	44.1 58.2 -19.0 61.2 341.8		
43/653	M38R_100_100e	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.964	48.5 65.6 -12.2 66.7 349.4	1.0 0.0 0.625	48.0 61.8 2.1 61.8 361.9 14.8 331	1.0 0.0 0.964	48.5 65.6 -12.2 66.7 349.4		
44/652	M50R_100_100e	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.827	49.4 65.5 -9.1 66.2 352.0	1.0 0.0 0.5	47.8 58.9 10.4 59.9 370.0 20.7 339	1.0 0.0 0.827	49.4 65.5 -9.1 66.2 352.0		
45/651	M63R_100_100e	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0							

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)

TUB material: code=rha4ta

n/j	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb**Fe	LabCh**Fe	DE**Fe	hsi_Me	rgb*Me	LabCh*Me
0/648	R00Y_100_100e	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.263	47.5 56.0 26.7	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4 11.1	375	1.0 0.0 0.263	47.5 56.0 26.7
1/666	R25Y_100_100e	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.108 0.0	51.4 54.8 47.7	1.0 0.25 0.0	58.2 41.8 55.1	69.2 52.8 16.4	35	1.0 0.108 0.0	51.4 54.8 47.7
2/684	R50Y_100_100e	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.319 0.0	61.8 35.2 58.4	1.0 0.5 0.0	70.5 19.2 66.2	69.0 73.8 19.8	48	1.0 0.319 0.0	61.8 35.2 58.4
3/702	R75Y_100_100e	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.551 0.0	72.3 16.1 68.2	1.0 0.75 0.0	82.9 -2.0 76.9	77.0 91.5 22.7	63	1.0 0.551 0.0	72.3 16.1 68.2
4/720	Y00G_100_100e	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 0.768 0.0	83.6 -3.1 76.8	1.0 1.0 0.0	91.5 -15.8 84.6	86.1 100.5 16.9	77	1.0 0.768 0.0	83.6 -3.1 76.8
5/558	Y25G_100_100e	0.75 1.0 0.0	1.0 1.0 0.5	104	0.697 1.0 0.0	85.8 -26.4 78.5	0.75 1.0 0.0	90.1 -21.3 86.0	88.6 103.9 10.0	107	0.697 1.0 0.0	85.8 -26.4 78.5
6/396	Y50G_100_100e	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	71.0 -41.7 54.8	0.5 1.0 0.0	70.9 -41.7 54.8	68.9 127.3 0.0	119	0.5 1.0 0.0	71.0 -41.7 54.8
7/234	Y75G_100_100e	0.25 1.0 0.0	1.0 1.0 0.5	136	0.227 1.0 0.0	59.9 -58.2 39.3	0.25 1.0 0.0	60.6 -57.2 40.4	70.1 144.7 1.5	137	0.227 1.0 0.0	59.9 -58.2 39.3
8/72	G00B_100_100e	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.146	53.8 -65.9 21.1	0.0 1.0 0.0	54.3 -67.6 30.8	74.3 155.5 9.8	157	0.0 1.0 0.146	53.8 -65.9 21.1
9/72	G00B_100_100e	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.146	53.8 -65.9 21.1	0.0 1.0 0.0	54.3 -67.6 30.8	74.3 155.5 9.8	157	0.0 1.0 0.146	53.8 -65.9 21.1
10/76	G25B_100_100e	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.497	55.0 -51.6 -8.7	0.0 1.0 0.5	55.0 -51.4 -8.9	52.2 189.8 0.2	179	0.0 1.0 0.497	55.0 -51.6 -8.7
11/80	G50B_100_100e	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 0.791	54.9 -38.7 -29.1	0.0 1.0 1.0	53.1 -30.0 -43.1	52.5 235.1 16.5	198	0.0 1.0 0.791	54.9 -38.7 -29.1
12/44	G75B_100_100e	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.686 1.0	51.7 -23.3 -48.6	0.0 0.5 1.0	46.1 -13.3 -49.4	51.1 254.9 11.4	227	0.0 0.686 1.0	51.7 -23.3 -48.6
13/8	B00R_100_100e	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.261 1.0	37.3 1.4 -48.6	0.0 0.0 1.0	32.5 16.9 -44.6	47.7 290.8 16.6	255	0.0 0.261 1.0	37.3 1.4 -48.6
14/332	B25R_100_100e	0.5 0.0 1.0	1.0 1.0 0.5	300	0.138 0.0 1.0	31.5 24.4 -41.9	0.5 0.0 1.0	37.2 43.1 -30.8	53.0 324.4 22.5	277	0.138 0.0 1.0	31.5 24.4 -41.9
15/656	B50R_100_100e	1.0 0.0 1.0	1.0 1.0 0.5	330	0.584 0.0 1.0	38.5 46.7 -28.5	1.0 0.0 1.0	48.1 65.4 -12.7	66.6 348.9 26.2	305	0.584 0.0 1.0	38.5 46.7 -28.5
16/652	B75R_100_100e	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.827	49.4 65.5 -9.1	1.0 0.0 0.5	47.8 58.9 10.4	59.9 10.0 20.7	339	1.0 0.0 0.827	49.4 65.5 -9.1
17/648	R00Y_100_100e	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.263	47.5 56.0 26.7	1.0 0.0 0.0	47.5 57.2 37.8	68.6 33.4 11.1	375	1.0 0.0 0.263	47.5 56.0 26.7
18/688	R00Y_100_050e	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.631	71.6 28.0 13.3	1.0 0.5 0.5	71.4 24.0 27.4	36.4 48.8 14.6	375	1.0 0.5 0.631	71.6 28.0 13.3
19/706	R50Y_100_050e	1.0 0.75 0.5	1.0 0.5 0.75	60	1.0 0.659 0.5	78.8 17.6 29.2	1.0 0.75 0.5	83.6 2.8 39.3	39.4 85.7 18.5	48	1.0 0.319 0.0	61.8 35.2 58.4
20/724	Y00G_100_050e	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 0.884 0.5	89.7 -1.5 38.4	1.0 1.0 0.5	93.1 -11.8 45.5	47.0 104.6 12.9	77	1.0 0.768 0.0	83.6 -3.1 76.8
21/562	Y50G_100_050e	0.75 1.0 0.5	1.0 0.5 0.75	120	0.75 1.0 0.5	83.4 -20.8 27.4	0.75 1.0 0.5	86.2 -21.9 37.8	43.7 120.1 10.8	119	0.5 1.0 0.0	71.0 -41.7 54.8
22/400	G00B_100_050e	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.573	74.8 -32.9 10.5	0.5 1.0 0.5	74.1 -30.5 11.7	32.6 158.9 2.8	157	0.0 1.0 0.146	53.8 -65.9 21.1
23/404	G50B_100_050e	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 0.895	75.4 -19.3 -14.5	0.5 1.0 1.0	73.7 -17.1 -27.4	32.3 238.0 13.2	198	0.0 1.0 0.791	54.9 -38.7 -29.1
24/368	B00R_100_050e	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.63 1.0	66.5 0.7 -24.3	0.5 0.5 1.0	54.8 11.5 -32.2	34.2 289.7 17.8	255	0.0 0.261 1.0	37.3 1.4 -48.6
25/692	B50R_100_050e	1.0 0.5 1.0	1.0 0.5 0.75	330	0.792 0.5 1.0	67.1 23.3 -14.2	1.0 0.5 1.0	71.6 36.1 -8.9	37.2 346.1 14.5	305	0.584 0.0 1.0	38.5 46.7 -28.5
26/688	R00Y_100_050e	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.631	71.6 28.0 13.3	1.0 0.5 0.5	71.4 24.0 27.4	36.4 48.8 14.6	375	1.0 0.5 0.631	71.6 28.0 13.3
27/506	R00Y_075_050e	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.381	53.7 28.0 13.3	0.75 0.25 0.25	52.4 27.1 25.3	37.1 43.0 12.1	375	1.0 0.0 0.263	47.5 56.0 26.7
28/524	R50Y_075_050e	0.75 0.5 0.25	0.75 0.5 0.5	60	0.75 0.409 0.25	60.8 17.6 29.2	0.75 0.5 0.25	66.1 6.5 36.1	36.6 79.7 14.0	48	1.0 0.319 0.0	61.8 35.2 58.4
29/542	Y00G_075_050e	0.75 0.75 0.25	0.75 0.5 0.5	90	0.75 0.634 0.25	71.7 -1.5 38.4	0.75 0.75 0.25	81.7 -11.5 50.7	52.0 102.8 18.7	77	1.0 0.768 0.0	83.6 -3.1 76.8
30/380	Y50G_075_050e	0.5 0.75 0.25	0.75 0.5 0.5	120	0.5 0.75 0.25	65.4 -20.8 27.4	0.5 0.75 0.25	70.5 -23.0 31.5	39.0 126.1 6.8	119	0.5 1.0 0.0	71.0 -41.7 54.8
31/218	G00B_075_050e	0.25 0.75 0.25	0.75 0.5 0.5	150	0.25 0.75 0.323	56.8 -32.9 10.5	0.25 0.75 0.25	57.2 -36.4 13.9	39.0 159.0 4.8	157	0.0 1.0 0.146	53.8 -65.9 21.1
32/222	G50B_075_050e	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.645	57.4 -19.3 -14.5	0.25 0.75 0.75	60.1 -19.6 -29.7	35.6 236.5 15.4	198	0.0 1.0 0.791	54.9 -38.7 -29.1
33/186	B00R_075_050e	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.38 0.75	48.5 0.7 -24.3	0.25 0.25 0.75	43.1 11.1 -34.5	36.3 287.8 15.5	255	0.0 0.261 1.0	37.3 1.4 -48.6
34/510	B50R_075_050e	0.75 0.25 0.75	0.75 0.5 0.5	330	0.542 0.25 0.75	49.1 23.3 -14.2	0.75 0.25 0.75	53.9 38.1 -12.4	40.1 341.9 15.6	305	0.584 0.0 1.0	38.5 46.7 -28.5
35/506	R00Y_075_050e	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.381	53.7 28.0 13.3	0.75 0.25 0.25	52.4 27.1 25.3	37.1 43.0 12.1	375	1.0 0.0 0.263	47.5 56.0 26.7
36/324	R00Y_050_050e	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.131	35.7 28.0 13.3	0.5 0.0 0.0	33.0 34.7 23.4	41.8 34.0 12.3	375	1.0 0.0 0.263	47.5 56.0 26.7
37/342	R50Y_050_050e	0.5 0.25 0.0	0.5 0.5 0.25	60	0.5 0.159 0.0	42.8 17.6 29.2	0.5 0.25 0.0	42.9 9.5 37.9	39.1 75.8 11.8	48	1.0 0.319 0.0	61.8 35.2 58.4
38/360	Y00G_050_050e	0.5 0.5 0.0	0.5 0.5 0.25	90	0.5 0.384 0.0	53.7 -1.5 38.4	0.5 0.5 0.0	58.4 -9.8 54.3	55.2 100.3 18.5	77	1.0 0.768 0.0	83.6 -3.1 76.8
39/198	Y50G_050_050e	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	47.4 -20.8 27.4	0.25 0.5 0.0	43.9 -28.1 32.6	43.1 130.7 9.6	119	0.5 1.0 0.0	71.0 -41.7 54.8
40/36	G00B_050_050e	0.0 0.5 0.0	0.5 0.5 0.25	150	0.0 0.5 0.073	38.8 -32.9 10.5	0.0 0.5 0.0	42.9 -59.6 21.5	63.4 160.1 29.1	157	0.0 1.0 0.146	53.8 -65.9 21.1
41/40	G50B_050_050e	0.0 0.5 0.5	0.5 0.5 0.25	210	0.0 0.5 0.395	39.4 -19.3 -14.5	0.0 0.5 0.5	44.1 -23.4 -34.5	41.7 235.7 20.9	198	0.0 1.0 0.791	54.9 -38.7 -29.1
42/4	B00R_050_050e	0.0 0.0 0.5	0.5 0.5 0.25	270	0.0 0.13 0.5	30.5 0.7 -24.3	0.0 0.0 0.5	30.3 13.1 -38.9	41.0 288.6 19.1	255	0.0 0.261 1.0	37.3 1.4 -48.6
43/328	B50R_050_050e	0.5 0.0 0.5	0.5 0.5 0.25	330	0.292 0.0 0.5	31.1 23.3 -14.2	0.5 0.0 0.5	35.4 43.7 -12.1	45.3 344.4 20.8	305	0.584 0.0 1.0	38.5 46.7 -28.5
44/324	R00Y_050_050e	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.131	35.7 28.0 13.3	0.5 0.0 0.0	33.0 34.7 23.4	41.8 34.0 12.3	375	1.0 0.0 0.263	47.5 56.0 26.7
45/0	NW_000e	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	23.8 0.0 0.0	0.0 0.0 0.0	23.8 0.0 0.0	0.0 0.0 0.0	360	1.0 1.0 1.0	95.8 0.0 0.0
46/91	NW_013e	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	32.8 0.0 0.0	0.125 0.125 0.125	27.8 0.0 -0.5	0.5 273.6 5.0	360	1.0 1.0 1.0	95.8 0.0 0.0
47/182	NW_025e	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	41.8 0.0 0.0	0.25 0.25 0.25	39.8 -0.1 -1.2	1.2 265.1 2.3	360	1.0 1.0 1.0	95.8 0.0 0.0
48/273	NW_038e	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	50.8 0.0 0.0	0.375 0.375 0.375	51.9 0.0 -1.3	1.3 267.6 1.7	360	1.0 1.0 1.0	95.8 0.0 0.0
49/364	NW_050e	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	59.8 0.0 0.0	0.5 0.5 0.5	61.2 0.0 -1.2	1.2 268.4 1.9	360	1.0 1.0 1.0	95.8 0.0 0.0
50/455	NW_063e	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.8 0.0 0.0	0.0 0.625 0.625	72.6 0.0 -1.0	1.0 269.8 3.9	360	1.0 1.0 1.0	95.8 0.0 0.0
51/546	NW_075e	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.75 0.75 0.75	80.4 0.0 -0.3	0.3 272.2 2.6	360	1.0 1.0 1.0	95.8 0.0 0.0
52/637	NW_088e	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.8 0.0 0.0	0.0 0.875 0.875	90.8 0.0 -0.1	0.1 265.5 4.0	360	1.0 1.0 1.0	95.8 0.0 0.0
53/728	NW_100e	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.8 0.0 0.0	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0	360	1.0 1.0 1.0	95.8 0.0 0.0

delta E* = 12.1

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n=j	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb*Fe	LabCh*Fe	DE*Fe	hsi_Me	rgb*Me	LabCh*Me
0	NW_000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	1.0
1	B00R_012_012e	0.0	0.125	0.125	0.062	0.062	0.032	0.125	23.8	4.0	0.261	1.0
2	B00R_025_025e	0.0	0.25	0.25	0.125	0.125	0.065	0.25	27.2	0.3	0.261	1.0
3	B00R_037_037e	0.0	0.375	0.375	0.187	0.187	0.09	0.375	28.8	0.5	0.261	1.0
4	B00R_050_050e	0.0	0.5	0.5	0.25	0.25	0.13	0.5	30.5	0.7	0.261	1.0
5	B00R_062_062e	0.0	0.625	0.625	0.312	0.312	0.163	0.625	32.2	0.9	0.261	1.0
6	B00R_075_075e	0.0	0.75	0.75	0.375	0.375	0.195	0.75	33.9	1.1	0.261	1.0
7	B00R_087_087e	0.0	0.875	0.875	0.437	0.437	0.228	0.875	35.6	1.2	0.261	1.0
8	B00R_100_100e	0.0	1.0	1.0	0.5	0.5	0.261	1.0	37.3	1.4	0.261	1.0
9	G00B_012_012e	0.0	0.125	0.125	0.062	0.062	0.125	0.018	27.5	-8.2	0.1	0.146
10	G50B_012_012e	0.0	0.125	0.125	0.062	0.062	0.125	0.098	27.7	-4.8	0.1	0.791
11	G75B_025_025e	0.0	0.125	0.25	0.125	0.125	0.171	0.25	30.8	-5.8	0.1	0.686
12	G84B_037_037e	0.0	0.125	0.375	0.187	0.187	0.19	0.375	32.3	-5.1	0.1	0.508
13	G88B_050_050e	0.0	0.125	0.5	0.5	0.5	0.217	0.5	33.7	-4.8	0.1	0.434
14	G90B_062_062e	0.0	0.125	0.625	0.312	0.312	0.244	0.625	35.1	-4.5	0.1	0.39
15	G92B_075_075e	0.0	0.125	0.75	0.375	0.375	0.273	0.75	36.7	-4.1	0.1	0.364
16	G93B_087_087e	0.0	0.125	0.875	0.437	0.437	0.308	0.875	38.5	-4.1	0.1	0.352
17	G94B_100_100e	0.0	0.125	1.0	0.5	0.5	0.341	1.0	40.1	-4.9	0.1	0.341
18	G00B_025_025e	0.0	0.25	0.25	0.125	0.125	0.25	0.036	31.3	-16.4	0.1	0.146
19	G25B_025_025e	0.0	0.25	0.125	0.125	0.125	0.25	0.124	31.6	-12.9	0.1	0.497
20	G50B_025_025e	0.0	0.25	0.25	0.125	0.125	0.25	0.197	31.6	-9.6	0.1	0.791
21	G65B_037_037e	0.0	0.25	0.375	0.25	0.25	0.375	0.372	34.8	-11.4	0.1	0.992
22	G75B_050_050e	0.0	0.25	0.5	0.5	0.5	0.343	0.5	37.7	-11.6	0.1	0.686
23	G80B_062_062e	0.0	0.25	0.625	0.312	0.312	0.354	0.625	39.2	-10.8	0.1	0.567
24	G84B_075_075e	0.0	0.25	0.75	0.375	0.375	0.381	0.75	40.8	-10.3	0.1	0.508
25	G86B_087_087e	0.0	0.25	0.875	0.437	0.437	0.406	0.875	42.1	-9.8	0.1	0.464
26	G88B_100_100e	0.0	0.25	1.0	0.5	0.5	0.434	1.0	43.6	-9.6	0.1	0.434
27	G00B_037_037e	0.0	0.375	0.375	0.187	0.187	0.375	0.055	35.0	-24.7	0.1	0.146
28	G15B_037_037e	0.0	0.375	0.125	0.375	0.187	0.375	0.138	35.4	-21.4	0.1	0.37
29	G34B_037_037e	0.0	0.375	0.25	0.375	0.187	0.375	0.218	35.6	-17.5	0.1	0.583
30	G50B_037_037e	0.0	0.375	0.375	0.375	0.187	0.375	0.296	35.5	-14.5	0.1	0.791
31	G61B_050_050e	0.0	0.375	0.5	0.5	0.5	0.5	0.474	38.7	-16.5	0.1	0.948
32	G69B_062_062e	0.0	0.375	0.625	0.312	0.312	0.547	0.625	42.1	-17.4	0.1	0.875
33	G75B_075_075e	0.0	0.375	0.75	0.375	0.375	0.514	0.75	44.7	-17.5	0.1	0.686
34	G79B_087_087e	0.0	0.375	0.875	0.437	0.437	0.522	0.875	46.3	-16.7	0.1	0.597
35	G81B_100_100e	0.0	0.375	1.0	0.5	0.5	0.552	1.0	48.0	-16.4	0.1	0.552
36	G00B_050_050e	0.0	0.5	0.5	0.25	0.25	0.5	0.073	38.8	-32.9	0.1	0.146
37	G11B_050_050e	0.0	0.5	0.125	0.5	0.25	0.5	0.16	39.0	-29.9	0.1	0.32
38	G25B_050_050e	0.0	0.5	0.25	0.5	0.25	0.5	0.248	39.4	-25.8	0.1	0.497
39	G38B_050_050e	0.0	0.5	0.375	0.5	0.25	0.5	0.311	39.6	-22.1	0.1	0.622
40	G50B_050_050e	0.0	0.5	0.5	0.25	0.25	0.5	0.395	39.4	-19.3	0.1	0.791
41	G59B_062_062e	0.0	0.625	0.625	0.312	0.312	0.625	0.576	42.6	-21.5	0.1	0.922
42	G65B_075_075e	0.0	0.625	0.75	0.375	0.375	0.75	0.744	45.8	-22.8	0.1	0.992
43	G70B_087_087e	0.0	0.875	0.875	0.437	0.437	0.707	0.875	49.3	-23.5	0.1	0.808
44	G75B_100_100e	0.0	0.5	1.0	0.5	0.5	0.686	1.0	51.7	-23.3	0.1	0.686
45	G00B_062_062e	0.0	0.625	0.625	0.312	0.312	0.625	0.091	42.5	-41.2	0.1	0.146
46	G09B_062_062e	0.0	0.625	0.125	0.625	0.312	0.625	0.181	42.7	-38.3	0.1	0.29
47	G19B_062_062e	0.0	0.625	0.25	0.625	0.312	0.625	0.26	43.2	-34.5	0.1	0.416
48	G30B_062_062e	0.0	0.625	0.375	0.625	0.312	0.625	0.345	43.4	-30.4	0.1	0.552
49	G40B_062_062e	0.0	0.625	0.5	0.625	0.312	0.625	0.41	43.5	-26.9	0.1	0.656
50	G50B_062_062e	0.0	0.625	0.625	0.312	0.312	0.625	0.494	43.3	-24.1	0.1	0.791
51	G57B_075_075e	0.0	0.625	0.75	0.375	0.375	0.75	0.679	46.5	-26.5	0.1	0.905
52	G63B_087_087e	0.0	0.625	0.875	0.437	0.437	0.875	0.845	49.7	-28.1	0.1	0.966
53	G68B_100_100e	0.0	0.625	1.0	0.5	0.5	0.915	1.0	53.1	-28.6	0.1	0.915
54	G00B_075_075e	0.0	0.75	0.75	0.375	0.375	0.75	0.11	46.3	-49.4	0.1	0.146
55	G07B_075_075e	0.0	0.75	0.125	0.75	0.375	0.75	0.203	46.4	-46.7	0.1	0.271
56	G15B_075_075e	0.0	0.75	0.25	0.75	0.375	0.75	0.277	47.0	-42.8	0.1	0.37
57	G25B_075_075e	0.0	0.75	0.375	0.75	0.375	0.75	0.373	47.2	-39.7	0.1	0.497
58	G34B_075_075e	0.0	0.75	0.5	0.75	0.375	0.75	0.437	47.4	-35.1	0.1	0.583
59	G42B_075_075e	0.0	0.75	0.625	0.75	0.375	0.75	0.509	47.4	-31.7	0.1	0.678
60	G50B_075_075e	0.0	0.75	0.75	0.375	0.375	0.75	0.593	47.1	-29.0	0.1	0.791
61	G56B_087_087e	0.0	0.75	0.875	0.437	0.437	0.875	0.784	50.4	-31.3	0.1	0.896
62	G61B_100_100e	0.0	0.75	1.0	0.5	0.5	1.0	0.948	53.6	-33.1	0.1	0.948
63	G00B_087_087e	0.0	0.875	0.875	0.437	0.437	0.875	0.128	50.0	-57.7	0.1	0.146
64	G06B_087_087e	0.0	0.875	0.125	0.875	0.437	0.875	0.228	50.1	-54.8	0.1	0.261
65	G13B_087_087e	0.0	0.875	0.25	0.875	0.437	0.875	0.298	50.6	-51.4	0.1	0.34
66	G20B_087_087e	0.0	0.875	0.375	0.875	0.437	0.875	0.384	51.0	-47.4	0.1	0.439
67	G29B_087_087e	0.0	0.875	0.5	0.875	0.437	0.875	0.47	51.2	-43.3	0.1	0.537
68	G36B_087_087e	0.0	0.875	0.625	0.875	0.437	0.875	0.531	51.3	-39.6	0.1	0.607
69	G43B_087_087e	0.0	0.875	0.75	0.875	0.437	0.875	0.603	51.3	-36.6	0.1	0.69
70	G50B_087_087e	0.0	0.875	0.875	0.437	0.437	0.875	0.692	51.0	-33.8	0.1	0.791
71	G55B_100_100e	0.0	0.875	1.0	0.5	0.5	1.0	0.888	54.3	-36.1	0.1	0.888
72	G00B_100_100e	0.0	1.0	1.0	0.5	0.5	1.0	0.146	53.8	-65.9	0.1	0.146
73	G05B_100_100e	0.0	1.0	0.125	1.0	0.5	1.0	0.251	53.7	-63.1	0.1	0.251
74	G11B_100_100e	0.0	1.0	0.25	1.0	0.5	1.0	0.32	54.3	-59.8	0.1	0.32
75	G18B_100_100e	0.0	1.0	0.375	1.0	0.5	1.0	0.404	54.8	-55.6	0.1	0.404
76	G25B_100_100e	0.0	1.0	0.5	1.0	0.5	1.0	0.497	55.0	-51.6	0.1	0.497
77	G31B_100_100e	0.0	1.0	0.625	1.0	0.5	1.0	0.56	55.1	-48.2	0.1	0.56
78	G38B_100_100e	0.0	1.0	0.75	1.0	0.5	1.0	0.622	55.3	-44.3	0.1	0.622
79	G44B_100_100e	0.0	1.0	0.875	1.0	0.5	1.0	0.701	55.2	-41.4	0.1	0.701
80	G50B_100_100e	0.0	1.0	1.0	0.5	0.5	1.0	0.791	54.9	-38.7	0.1	0.791

delta E* = 15.2

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
colores y diferencia en color, ΔE^* , 3D=0, de=1, cmyk

entrada: $rgb/cmyk \rightarrow rgb_e$
salida: transfiera a $cmyk_e$

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)

TUB material: code=rh4ta

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb*Fe	LabCh*Fe	DE*Fe	hsiMe	rgb*Me	LabCh*Me
81	R00Y_012_012a	0.125 0.0 0.0	0.125 0.125 0.062	390	0.125 0.0 0.032	26.8 7.0 3.3	7.7 25.4	0.125 0.0 0.0	24.7 7.7 8.0	11.1 45.9 5.1	375	47.5 56.0 26.7 62.1 25.4
82	B50R_012_012a	0.125 0.0 0.125	0.125 0.125 0.062	330	0.073 0.0 0.125	25.6 5.8 -3.5	6.8 328.6	0.125 0.0 0.125	26.6 12.3 -6.3	13.8 332.7 7.1	305	38.5 46.7 -28.5 54.7 328.6
83	B25R_025_025a	0.125 0.0 0.25	0.25 0.25 0.125	300	0.034 0.0 0.25	25.7 6.1 -10.4	12.1 300.1	0.125 0.0 0.25	25.7 13.6 -17.5	22.2 307.9 10.3	277	0.138 0.0 1.0 31.5 24.4 -41.9 48.5 300.1
84	B15R_037_037a	0.125 0.0 0.375	0.375 0.375 0.187	289	0.0 0.005 0.375	27.2 6.0 -16.8	17.8 289.7	0.125 0.0 0.375	27.6 14.3 -24.2	28.1 300.5 11.0	269	0.0 0.014 1.0 32.8 16.1 -44.8 47.6 289.7
85	B11R_050_050a	0.125 0.0 0.5	0.5 0.5 0.25	284	0.0 0.038 0.5	28.9 6.1 -22.9	23.7 285.0	0.125 0.0 0.5	29.0 21.5 -32.8	39.2 303.2 18.3	266	0.0 0.077 1.0 34.1 12.2 -45.8 47.4 285.0
86	B09R_062_062a	0.125 0.0 0.625	0.625 0.625 0.312	281	0.0 0.072 0.625	30.7 6.2 -28.9	29.5 282.1	0.125 0.0 0.625	29.5 25.2 -36.3	44.2 304.7 20.4	263	0.0 0.115 1.0 34.8 9.9 -46.2 47.3 282.1
87	B07R_075_075a	0.125 0.0 0.75	0.75 0.75 0.375	279	0.0 0.106 0.75	32.4 6.3 -35.0	35.6 280.2	0.125 0.0 0.75	30.6 25.4 -39.9	47.4 302.5 19.8	262	0.0 0.141 1.0 35.3 8.4 -46.7 47.5 280.2
88	B06R_087_087a	0.125 0.0 0.875	0.875 0.875 0.437	278	0.0 0.135 0.875	34.0 6.7 -41.1	41.6 279.3	0.125 0.0 0.875	30.8 25.7 -41.3	48.7 301.9 19.3	261	0.0 0.155 1.0 35.5 7.7 -47.0 47.6 279.3
89	B05R_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.0 0.168 1.0	35.7 6.9 -47.2	47.7 278.3	0.125 0.0 1.0	31.6 23.6 -42.2	48.4 299.2 17.8	260	0.0 0.168 1.0 35.7 6.9 -47.2 47.7 278.3
90	Y00G_012_012a	0.125 0.125 0.0	0.125 0.125 0.062	90	0.125 0.096 0.0	31.3 -0.3 9.6	9.6 92.3	0.125 0.125 0.0	32.9 -5.2 16.0	16.9 108.2 8.2	77	1.0 0.768 0.0 83.6 -3.1 76.8 76.9 92.3
91	NW_012a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	32.8 0.0 0.0	0.0 0.0 0.0	0.125 0.125 0.125	27.8 0.0 -0.5	0.5 273.6 5.0	360	1.0 1.0 1.0 95.8 0.0 0.0 0.0 0.0
92	B00R_025_012a	0.125 0.125 0.25	0.25 0.125 0.187	270	0.124 0.157 0.25	34.5 0.1 -6.0	6.0 271.7	0.125 0.125 0.25	27.8 2.1 -17.8	17.9 276.9 13.6	255	0.0 0.261 1.0 37.3 1.4 -48.6 48.7 271.7
93	B00R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.19 0.375	36.2 0.3 -12.1	12.1 271.7	0.125 0.125 0.375	30.4 4.1 -25.6	25.9 279.1 15.0	255	0.0 0.261 1.0 37.3 1.4 -48.6 48.7 271.7
94	B00R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.222 0.5	37.8 0.5 -18.2	18.2 271.7	0.125 0.125 0.5	33.6 7.2 -30.1	31.0 283.5 14.2	255	0.0 0.261 1.0 37.3 1.4 -48.6 48.7 271.7
95	B00R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.255 0.625	39.5 0.7 -24.3	24.3 271.7	0.125 0.125 0.625	36.0 11.5 -35.5	37.4 287.9 15.9	255	0.0 0.261 1.0 37.3 1.4 -48.6 48.7 271.7
96	B00R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.288 0.75	41.2 0.9 -30.4	30.4 271.7	0.125 0.125 0.75	37.8 12.8 -40.0	42.0 287.7 15.7	255	0.0 0.261 1.0 37.3 1.4 -48.6 48.7 271.7
97	B00R_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.32 0.875	42.9 1.1 -36.5	36.5 271.7	0.125 0.125 0.875	35.7 15.8 -43.8	46.5 289.8 17.9	255	0.0 0.261 1.0 37.3 1.4 -48.6 48.7 271.7
98	B00R_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.353 1.0	44.6 1.2 -42.6	42.6 271.7	0.125 0.125 1.0	34.1 17.5 -43.8	47.2 281.8 19.3	255	0.0 0.261 1.0 37.3 1.4 -48.6 48.7 271.7
99	Y50G_025_025a	0.125 0.25 0.0	0.25 0.25 0.125	120	0.125 0.25 0.0	35.6 -10.4 13.7	17.2 127.2	0.125 0.25 0.0	36.4 -13.3 17.0	21.6 127.9 4.4	119	0.5 1.0 0.0 71.0 -41.7 54.8 68.9 127.2
100	G00B_025_012a	0.125 0.25 0.125	0.25 0.125 0.187	150	0.124 0.25 0.143	36.5 -8.2 2.6	8.6 162.2	0.125 0.25 0.125	37.3 -12.7 1.8	12.8 171.8 4.6	157	0.0 1.0 0.146 53.8 -65.9 21.1 69.2 162.2
101	G50B_025_012a	0.125 0.25 0.25	0.25 0.125 0.187	210	0.124 0.25 0.223	36.7 -4.8 -3.6	6.0 216.9	0.125 0.25 0.25	34.1 -10.5 -16.7	19.7 237.8 14.5	198	0.0 1.0 0.791 54.9 -38.7 -29.1 48.4 216.9
102	G75B_037_025a	0.125 0.25 0.375	0.375 0.25 0.25	240	0.124 0.296 0.375	39.8 -5.8 -12.1	13.4 244.3	0.125 0.25 0.375	36.2 -6.8 -24.4	25.4 254.3 12.8	227	0.0 0.686 1.0 51.7 -23.3 -48.6 53.9 244.3
103	G84B_050_037a	0.125 0.25 0.5	0.5 0.375 0.312	251	0.124 0.315 0.5	41.3 -5.1 -18.5	19.2 254.3	0.125 0.25 0.5	37.9 -3.1 -29.4	29.6 263.8 11.5	239	0.0 0.508 1.0 46.4 -13.8 -49.4 51.3 254.3
104	G88B_062_050a	0.125 0.25 0.625	0.625 0.5 0.375	256	0.125 0.342 0.625	42.7 -4.8 -24.7	25.1 259.9	0.125 0.25 0.625	40.5 -0.2 -32.8	32.8 269.6 9.5	244	0.0 0.434 1.0 43.6 -9.6 -49.4 50.3 259.9
105	G90B_075_062a	0.125 0.25 0.75	0.75 0.625 0.437	259	0.125 0.369 0.75	44.1 -4.5 -30.8	31.1 261.6	0.125 0.25 0.75	41.5 2.5 -38.3	38.4 273.7 10.6	247	0.0 0.39 1.0 42.0 -7.2 -49.3 49.8 261.6
106	G92B_087_075a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.398 0.875	45.7 -4.1 -36.9	37.1 263.5	0.125 0.25 0.875	37.9 7.9 -45.5	46.2 279.9 16.8	248	0.0 0.364 1.0 41.0 -9.5 -49.2 49.5 263.5
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.433 1.0	47.5 -4.1 -43.3	43.3 264.4	0.125 0.25 1.0	36.4 10.6 -45.2	46.4 283.2 18.6	249	0.0 0.352 1.0 40.6 -4.7 -49.2 49.4 264.4
108	Y68G_037_037a	0.125 0.375 0.0	0.375 0.375 0.187	131	0.115 0.375 0.0	38.6 -19.8 16.5	25.8 140.0	0.125 0.375 0.0	39.5 -30.9 19.8	36.7 147.2 11.5	132	0.308 1.0 0.0 63.4 -52.8 44.2 68.9 140.0
109	G00B_037_025a	0.125 0.375 0.125	0.375 0.25 0.25	150	0.124 0.375 0.161	40.3 -16.4 5.2	17.3 162.2	0.125 0.375 0.125	39.1 -24.1 4.4	24.5 169.4 7.8	157	0.0 1.0 0.146 53.8 -65.9 21.1 69.2 162.2
110	G25B_037_025a	0.125 0.375 0.25	0.375 0.25 0.25	180	0.124 0.375 0.249	40.6 -12.9 -2.1	13.0 189.6	0.125 0.375 0.25	37.7 -18.6 -9.6	20.9 207.3 9.8	179	0.0 1.0 0.497 55.0 -51.6 -8.7 52.3 189.6
111	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.322	40.6 -9.6 -7.2	12.1 216.9	0.125 0.375 0.375	41.9 -14.5 -23.1	27.3 237.7 16.6	198	0.0 1.0 0.791 54.9 -38.7 -29.1 48.4 216.9
112	G65B_050_037a	0.125 0.375 0.5	0.5 0.375 0.312	229	0.124 0.5 0.497	43.8 -11.4 -15.9	19.6 234.3	0.125 0.375 0.5	42.5 -11.5 -28.8	31.0 248.1 12.9	209	0.0 1.0 0.992 53.2 -30.5 -42.5 52.3 234.3
113	G75B_062_050a	0.125 0.375 0.625	0.625 0.5 0.375	240	0.125 0.468 0.625	46.7 -11.6 -24.3	26.9 244.3	0.125 0.375 0.625	43.5 -8.6 -31.7	32.8 254.7 8.6	227	0.0 0.686 1.0 51.7 -23.3 -48.6 53.9 244.3
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.479 0.75	48.2 -10.8 -31.0	32.8 250.7	0.125 0.375 0.75	44.9 -6.0 -36.4	36.9 260.5 7.9	235	0.0 0.567 1.0 48.5 -17.3 -49.6 52.5 250.7
115	G84B_087_075a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.506 0.875	49.7 -10.3 -37.0	38.5 254.3	0.125 0.375 0.875	43.8 -4.2 -45.9	46.1 264.7 12.3	239	0.0 0.508 1.0 46.4 -13.8 -49.4 51.3 254.3
116	G86B_100_087a	0.125 0.375 1.0	1.0 0.875 0.562	254	0.125 0.531 1.0	51.1 -9.8 -43.2	44.4 257.1	0.125 0.375 1.0	39.2 1.2 -47.2	47.3 271.4 16.7	242	0.0 0.464 1.0 44.7 -11.2 -49.4 50.7 257.1
117	Y76G_050_050a	0.125 0.5 0.0	0.5 0.5 0.25	136	0.113 0.5 0.0	41.9 -29.1 19.6	35.1 145.9	0.125 0.5 0.0	41.4 -44.7 24.0	50.7 151.6 16.2	137	0.227 1.0 0.0 59.9 -58.2 39.3 70.2 145.9
118	G00B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.18	44.0 -24.7 7.9	25.9 162.2	0.125 0.5 0.125	42.7 -31.6 8.9	32.8 164.2 7.0	157	0.0 1.0 0.146 53.8 -65.9 21.1 69.2 162.2
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.263	44.4 -21.4 0.1	21.4 179.5	0.125 0.5 0.25	43.1 -29.5 -2.1	29.5 184.0 8.5	171	0.0 1.0 0.37 54.7 -57.0 0.4 57.0 179.5
120	G34B_050_037a	0.125 0.5 0.375	0.5 0.375 0.312	191	0.124 0.5 0.343	44.6 -17.5 -6.2	18.6 199.6	0.125 0.5 0.375	44.4 -22.2 -17.9	28.6 218.8 12.6	185	0.0 1.0 0.583 55.2 -46.8 -16.7 49.7 199.6
121	G50B_050_037a	0.125 0.5 0.5	0.5 0.375 0.312	210	0.124 0.5 0.421	44.5 -14.5 -10.9	18.1 216.9	0.125 0.5 0.5	45.7 -17.6 -27.9	33.0 237.6 17.3	198	0.0 1.0 0.791 54.9 -38.7 -29.1 48.4 216.9
122	G61B_062_050a	0.125 0.5 0.625	0.625 0.5 0.375	224	0.125 0.625 0.599	47.7 -16.5 -19.5	25.6 229.7	0.125 0.5 0.625	47.3 -14.4 -32.2	35.2 245.8 12.8	207	0.0 1.0 0.948 53.6 -33.1 -39.1 51.2 229.7
123	G69B_075_062a	0.125 0.5 0.75	0.75 0.625 0.437	233	0.125 0.672 0.75	51.1 -17.4 -27.9	32.9 237.9	0.125 0.5 0.75	47.8 -13.4 -35.8	38.3 249.4 9.4	216	0.0 0.875 1.0 53.1 -27.9 -44.7 52.7 237.9
124	G75B_087_075a	0.125 0.5 0.875	0.875 0.75 0.5	240	0.125 0.639 0.875	53.7 -17.5 -36.4	40.4 244.3	0.125 0.5 0.875	48.0 -12.0 -43.4	45.0 254.4 10.5	227	0.0 0.686 1.0 51.7 -23.3 -48.6 53.9 244.3
125	G79B_100_087a	0.125 0.5 1.0	1.0 0.875 0.562	245	0.125 0.647 1.0	55.3 -16.7 -43.4	46.5 248.9	0.125 0.5 1.0	44.3 -8.4 -48.1	48.9 260.0 14.5	233	0.0 0.597 1.0 49.6 -19.1 -49.6 53.1 248.9
126	Y81G_062_062a	0.125 0.625 0.0	0.625 0.625 0.312	139	0.098 0.625 0.0	45.1 -38.1 22.5	44.2 149.4	0.125 0.625 0.0	47.0 -55.6 28.1	62.3 153.1 18.4	141	0.157 1.0 0.0 57.9 -60.9 36.0 70.8 149.4
127	G00B_062_050a	0.125 0.625 0.125	0.625 0.5 0.375	150	0.125 0.625 0.198	47.8 -32.9 10.5	34.6 162.2	0.125 0.625 0.125	47.4 -40.2 14.8	42.9 159.7 8.4	157	0.0 1.0 0.146 53.8 -65.9 21.1 69.2 162.2
128	G11B_062_050a	0.125 0.625 0.25	0.625 0.5 0.375	164	0.125 0.625 0.285	48.0 -29.9 2.6	30.0 175.0	0.125 0.625 0.25	47.8 -37.6 3.8	37.8 174.1 7.7	178	0.0 1.0 0.32 54.3 -59.8 5.2 60.1 175.0
129	G25B_062_050a	0.125 0.625 0.375	0.625 0.5 0.375	180	0.125 0.625 0.373	48.4 -25.8 -4.3	26.1 189.6	0.125 0.625 0.375	49.3 -31.0 -11.2	33.0 199.9 8.6	179	0.0 1.0 0.497 55.0 -51.6 -8.7 52.3 189.6
130	G38B_062_050a	0.125 0.625 0.5	0.625 0.5 0.375	196	0.125 0.625 0.436	48.5 -22.1 -9.9	24.2 204.2	0.125 0.625 0.5	50.1 -24.7 -22.9	33.7 222.8 13.2	188	0.0 1.0 0.622 55.3 -44.3 -19.9

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información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb**Fe	LabCh**Fe	DE**Fe	hsiMe	rgb*Me	LabCh*Me
162	R00Y_025_025e	0.25 0.0 0.0	0.25 0.25 0.125	390	0.25 0.0 0.065	29.7 14.0 6.6	15.5 25.4	0.25 0.0 0.0	26.6 14.4 12.1	18.9 40.1 6.3	375 1.0 0.0 0.263	47.5 56.0 26.7 62.1 25.4
163	R00Y_025_025e	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.206	30.2 16.3 -2.2	16.5 352.0	0.25 0.0 0.125	26.4 17.6 2.2	17.7 7.1 6.0	339 1.0 0.0 0.827	49.4 65.5 -9.1 66.2 352.0
164	B50R_025_025e	0.25 0.0 0.25	0.25 0.25 0.125	330	0.146 0.0 0.25	27.5 11.6 -7.1	13.6 328.6	0.25 0.0 0.25	27.9 22.7 -10.0	24.8 336.0 11.4	305 0.584 0.0 1.0	38.5 46.7 -28.5 54.7 328.6
165	B34R_037_037e	0.25 0.0 0.375	0.375 0.375 0.187	311	0.107 0.0 0.375	26.8 12.3 -14.3	18.9 310.5	0.25 0.0 0.375	28.8 24.9 -17.6	30.5 324.6 13.1	286 0.285 0.0 1.0	31.9 32.8 -38.2 50.4 310.5
166	B25R_050_050e	0.25 0.0 0.5	0.5 0.5 0.25	300	0.069 0.0 0.5	27.6 12.2 -20.9	24.2 300.1	0.25 0.0 0.5	28.9 25.7 -25.1	35.9 315.6 14.2	277 0.138 0.0 1.0	31.5 24.4 -41.9 48.5 300.1
167	B19R_062_062e	0.25 0.0 0.625	0.625 0.625 0.312	293	0.025 0.0 0.625	29.1 11.9 -27.4	29.9 293.5	0.25 0.0 0.625	29.8 28.8 -31.0	42.3 312.9 17.2	272 0.04 0.0 1.0	32.2 19.1 -43.9 47.9 293.5
168	B15R_075_075e	0.25 0.0 0.75	0.75 0.75 0.375	289	0.0 0.01 0.75	30.6 12.0 -33.6	35.7 289.7	0.25 0.0 0.75	31.0 30.1 -35.2	46.3 310.5 18.0	269 0.0 0.014 1.0	32.8 16.1 -44.8 47.6 289.7
169	B13R_087_087e	0.25 0.0 0.875	0.875 0.875 0.437	286	0.0 0.045 0.875	32.3 12.0 -39.8	41.6 286.9	0.25 0.0 0.875	31.5 30.7 -38.1	48.9 308.8 18.7	267 0.0 0.052 1.0	33.6 13.8 -45.5 47.5 286.9
170	B11R_100_100e	0.25 0.0 1.0	1.0 1.0 0.5	284	0.0 0.077 1.0	34.1 12.2 -45.8	47.4 285.0	0.25 0.0 1.0	31.0 30.5 -39.3	49.8 307.8 19.6	266 0.0 0.077 1.0	34.1 12.2 -45.8 47.4 285.0
171	R50Y_025_025e	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.079 0.0	33.3 8.8 14.6	17.0 58.8	0.25 0.125 0.0	32.1 3.1 19.8	20.0 81.0 7.8	48 1.0 0.319 0.0	61.8 35.2 58.4 68.2 58.8
172	R00Y_025_012e	0.25 0.125 0.125	0.25 0.125 0.187	390	0.25 0.124 0.157	35.8 7.0 3.3	7.7 25.4	0.25 0.125 0.125	30.7 8.8 7.3	11.5 39.8 6.7	375 1.0 0.0 0.263	47.5 56.0 26.7 62.1 25.4
173	B50R_025_012e	0.25 0.125 0.25	0.25 0.125 0.187	330	0.198 0.124 0.25	34.6 5.8 -3.5	6.8 328.6	0.25 0.125 0.25	32.9 13.4 -8.4	15.9 327.8 9.2	305 0.584 0.0 1.0	38.5 46.7 -28.5 54.7 328.6
174	B25R_037_025e	0.25 0.125 0.375	0.375 0.25 0.25	300	0.159 0.124 0.375	34.7 6.1 -10.4	12.1 300.1	0.25 0.125 0.375	33.3 13.6 -17.6	22.3 307.5 10.5	277 0.138 0.0 1.0	31.5 24.4 -41.9 48.5 300.1
175	B15R_050_037e	0.25 0.125 0.5	0.5 0.375 0.312	289	0.124 0.13 0.5	36.2 6.0 -16.8	17.8 289.7	0.25 0.125 0.5	34.1 15.5 -25.6	30.0 301.2 13.1	269 0.0 0.014 1.0	32.8 16.1 -44.8 47.6 289.7
176	B11R_062_050e	0.25 0.125 0.625	0.625 0.5 0.375	284	0.125 0.163 0.625	37.9 6.1 -22.9	23.7 285.0	0.25 0.125 0.625	37.0 18.0 -31.1	36.0 300.1 14.5	266 0.0 0.077 1.0	34.1 12.2 -45.8 47.4 285.0
177	B09R_075_062e	0.25 0.125 0.75	0.75 0.625 0.437	281	0.125 0.197 0.75	39.7 6.2 -28.9	29.5 282.1	0.25 0.125 0.75	38.2 19.8 -36.3	41.4 298.5 15.5	263 0.0 0.115 1.0	34.8 9.9 -46.2 47.3 282.1
178	B07R_087_075e	0.25 0.125 0.875	0.875 0.75 0.5	279	0.125 0.231 0.875	41.4 6.3 -35.0	35.6 280.2	0.25 0.125 0.875	35.5 24.3 -40.4	47.1 301.0 19.7	262 0.0 0.141 1.0	35.3 8.4 -46.7 47.5 280.2
179	B06R_100_087e	0.25 0.125 1.0	1.0 0.875 0.562	278	0.125 0.26 1.0	43.0 6.7 -41.1	41.6 279.3	0.25 0.125 1.0	34.9 24.2 -40.6	47.3 300.7 19.2	261 0.0 0.155 1.0	35.5 7.7 -47.0 47.6 279.3
180	Y00G_025_025e	0.25 0.25 0.0	0.25 0.25 0.125	90	0.25 0.192 0.0	38.8 -0.7 19.2	19.2 92.3	0.25 0.25 0.0	39.4 -7.8 29.7	30.8 104.8 12.7	77 1.0 0.768 0.0	83.6 -3.1 76.8 76.9 92.3
181	Y00G_025_012e	0.25 0.25 0.125	0.25 0.125 0.187	90	0.25 0.221 0.124	40.3 -0.3 9.6	9.6 92.3	0.25 0.25 0.125	39.8 -5.3 15.2	16.1 109.2 7.4	77 1.0 0.768 0.0	83.6 -3.1 76.8 76.9 92.3
182	NW_025e	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	41.8 0.0 0.0	0.0 0.0	0.25 0.25 0.25	39.8 -0.1 -1.2	1.2 265.1 2.3	360 1.0 1.0 1.0	95.8 0.0 0.0 0.0 0.0
183	B00R_037_012e	0.25 0.25 0.375	0.375 0.125 0.312	270	0.249 0.282 0.375	43.5 0.1 -6.0	6.0 271.7	0.25 0.25 0.375	37.2 2.3 -18.7	18.9 276.9 14.3	255 0.0 0.261 1.0	37.3 1.4 -48.6 48.7 271.7
184	B00R_050_025e	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.315 0.5	45.2 0.3 -12.1	12.1 271.7	0.25 0.25 0.5	39.7 5.0 -25.0	25.4 281.3 14.6	255 0.0 0.261 1.0	37.3 1.4 -48.6 48.7 271.7
185	B00R_062_037e	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.347 0.625	46.8 0.5 -18.2	18.2 271.7	0.25 0.25 0.625	41.8 8.2 -29.3	30.4 285.7 14.4	255 0.0 0.261 1.0	37.3 1.4 -48.6 48.7 271.7
186	B00R_075_050e	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.38 0.75	48.5 0.7 -24.3	24.3 271.7	0.25 0.25 0.75	43.1 11.1 -34.5	36.3 287.8 15.5	255 0.0 0.261 1.0	37.3 1.4 -48.6 48.7 271.7
187	B00R_087_062e	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.413 0.875	50.2 0.9 -30.4	30.4 271.7	0.25 0.25 0.875	40.2 15.3 -41.4	44.1 290.2 20.6	255 0.0 0.261 1.0	37.3 1.4 -48.6 48.7 271.7
188	B00R_100_075e	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.445 1.0	51.9 1.1 -36.5	36.5 271.7	0.25 0.25 1.0	38.0 16.5 -42.0	45.1 291.5 21.5	255 0.0 0.261 1.0	37.3 1.4 -48.6 48.7 271.7
189	Y31G_037_037e	0.25 0.375 0.0	0.375 0.375 0.187	109	0.236 0.375 0.0	45.0 -11.7 25.8	28.4 114.4	0.25 0.375 0.0	41.6 -14.3 30.8	34.0 114.9 6.5	111 0.631 1.0 0.0	80.4 -31.3 68.9 75.7 114.4
190	Y50G_037_025e	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	44.6 -10.4 13.7	17.2 127.2	0.25 0.375 0.125	44.8 -13.9 15.6	20.9 131.6 4.0	119 0.5 1.0 0.0	71.0 -41.7 54.8 68.9 127.2
191	G00B_037_012e	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.268	45.5 -8.2 2.6	8.6 162.2	0.25 0.375 0.25	43.9 -11.7 -0.1	11.7 180.5 4.7	157 0.0 1.0 0.146	53.8 -65.9 21.1 69.2 162.2
192	G50B_037_012e	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.348	45.7 -4.8 -3.6	6.0 216.9	0.25 0.375 0.375	44.7 -10.3 -17.0	19.9 238.7 14.5	198 0.0 1.0 0.791	54.9 -38.7 -29.1 48.4 216.9
193	G75B_050_025e	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.421 0.5	48.8 -5.8 -12.1	13.4 244.3	0.25 0.375 0.5	46.4 -5.5 -23.0	23.6 256.5 11.1	227 0.0 0.686 1.0	51.7 -23.3 -48.6 53.9 244.3
194	G84B_062_037e	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.44 0.625	50.3 -5.1 -18.5	19.2 254.3	0.25 0.375 0.625	46.8 -2.5 -27.6	27.7 264.7 10.1	239 0.0 0.508 1.0	46.4 -13.8 -49.4 51.3 254.3
195	G88B_075_050e	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.467 0.75	51.7 -4.8 -24.7	25.1 258.9	0.25 0.375 0.75	46.6 0.6 -33.1	33.1 271.0 11.2	244 0.0 0.434 1.0	43.6 -9.6 -49.4 50.3 258.9
196	G90B_087_062e	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.494 0.875	53.1 -4.5 -30.8	31.1 261.6	0.25 0.375 0.875	45.7 4.1 -41.8	42.1 275.7 15.8	247 0.0 0.39 1.0	42.0 -7.2 -49.3 49.8 261.6
197	G92B_100_075e	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.523 1.0	54.7 -4.1 -36.9	37.1 263.5	0.25 0.375 1.0	42.5 7.0 -42.9	43.5 279.3 17.6	248 0.0 0.364 1.0	41.0 -5.5 -49.2 49.5 263.5
198	Y50G_050_050e	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	47.4 -20.8 27.4	34.4 127.2	0.25 0.5 0.0	43.9 -28.1 32.6	43.1 130.7 9.6	119 0.5 1.0 0.0	71.0 -41.7 54.8 68.9 127.2
199	Y68G_050_037e	0.25 0.5 0.125	0.5 0.375 0.312	131	0.24 0.5 0.124	47.6 -19.8 16.5	25.8 140.0	0.25 0.5 0.125	47.6 -24.8 16.6	29.9 146.1 5.0	132 0.308 1.0 0.0	63.4 -52.8 44.2 68.9 140.0
200	G00B_050_025e	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.286	49.3 -16.4 5.2	17.3 162.2	0.25 0.5 0.25	46.4 -22.2 3.8	22.6 170.1 6.6	157 0.0 1.0 0.146	53.8 -65.9 21.1 69.2 162.2
201	G25B_050_025e	0.25 0.5 0.375	0.5 0.25 0.375	180	0.249 0.5 0.374	49.6 -12.9 -2.1	13.0 189.6	0.25 0.5 0.375	48.2 -16.8 -9.9	19.5 210.3 8.8	179 0.0 1.0 0.497	55.0 -51.6 -8.7 52.3 189.6
202	G50B_050_025e	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.447	49.6 -9.6 -7.2	12.1 216.9	0.25 0.5 0.5	50.5 -14.1 -21.4	25.7 236.7 14.9	198 0.0 1.0 0.791	54.9 -38.7 -29.1 48.4 216.9
203	G65B_062_037e	0.25 0.5 0.625	0.625 0.375 0.437	229	0.25 0.625 0.622	52.8 -11.4 -15.9	19.6 234.3	0.25 0.5 0.625	51.2 -10.1 -27.6	29.4 249.8 11.8	209 0.0 1.0 0.992	

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n	HIC* _{Fe}	rgb _{Fe}	ict _{Fe}	hsi _{Fe}	rgb* _{Fe}	LabCh* _{Fe}	rgb* _{Fe}	LabCh* _{Fe}	DE** _{Fe}	hsi _{Me}	rgb* _{Me}	LabCh* _{Me}
243	R00Y_037_037*	0.375 0.0 0.0	0.375 0.375 0.187	390	0.375 0.0 0.098	32.7 21.0 10.0	23.2 25.4	0.375 0.0 0.0	31.0 26.1 19.8	32.8 37.1 11.2	375 1.0 0.0	0.263 47.5 56.0
244	R18Y_037_037*	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.22	32.8 22.9 1.7	22.9 4.3	0.375 0.0 0.125	33.0 25.3 9.6	27.1 20.9 8.3	354 1.0 0.0	0.588 47.9 61.1
245	B65R_037_037*	0.375 0.0 0.25	0.375 0.375 0.187	349	0.353 0.0 0.375	32.5 23.6 15.6	24.2 346.6	0.375 0.0 0.25	32.8 30.2 2.5	30.3 355.1 7.3	327 0.941 0.0	1.0 47.0 63.0
246	B50R_037_037*	0.375 0.0 0.375	0.375 0.375 0.187	330	0.219 0.0 0.375	29.3 17.5 10.7	20.5 328.6	0.375 0.0 0.375	33.5 35.7 11.0	37.4 342.8 18.7	305 0.584 0.0	1.0 38.5 46.7
247	B38R_050_050*	0.375 0.0 0.5	0.5 0.5 0.25	316	0.173 0.0 0.5	28.6 18.2 10.8	25.7 315.3	0.375 0.0 0.5	32.8 37.7 16.7	41.2 336.0 19.9	289 0.347 0.0	1.0 33.5 36.5
248	B30R_062_062*	0.375 0.0 0.625	0.625 0.625 0.312	307	0.147 0.0 0.625	28.3 18.6 14.8	31.0 306.8	0.375 0.0 0.625	32.2 37.2 24.4	44.5 326.7 19.0	283 0.235 0.0	1.0 31.0 29.7
249	B25R_075_075*	0.375 0.0 0.75	0.75 0.75 0.375	300	0.104 0.0 0.75	29.6 18.3 13.4	36.4 300.1	0.375 0.0 0.75	33.6 38.5 29.6	48.6 322.4 20.7	277 0.138 0.0	1.0 31.5 24.4
250	B20R_087_087*	0.375 0.0 0.875	0.875 0.875 0.437	295	0.06 0.0 0.875	31.0 18.0 18.0	295.4 30.0	0.375 0.0 0.875	34.5 38.8 33.1	51.0 319.5 21.6	273 0.068 0.0	1.0 32.0 20.6
251	B18R_100_100*	0.375 0.0 1.0	1.0 1.0 0.5	292	0.026 0.0 1.0	32.3 18.3 14.4	292.5 30.0	0.375 0.0 1.0	34.2 38.2 35.0	51.8 317.5 21.9	271 0.026 0.0	1.0 32.3 18.3
252	R31Y_037_037*	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.066 0.0	35.3 18.4 19.5	26.8 46.6	0.375 0.125 0.0	36.3 11.9 25.7	28.3 65.0 9.0	39 1.0 0.177	0.0 54.6 49.1
253	R00Y_037_025*	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.19	38.7 14.0 6.6	15.5 25.4	0.375 0.125 0.125	38.6 13.7 16.1	21.2 49.5 9.4	375 1.0 0.0	0.263 47.5 56.0
254	R00Y_037_025*	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.331	39.2 16.3 14.0	22.2 35.2	0.375 0.125 0.25	37.5 19.2 1.8	19.3 5.6 5.3	339 1.0 0.0	0.827 49.4 65.5
255	B50R_037_025*	0.375 0.125 0.375	0.375 0.25 0.25	330	0.271 0.124 0.375	36.5 11.6 13.6	328.6 31.0	0.375 0.125 0.375	38.9 24.5 10.7	26.7 336.2 13.5	305 0.584 0.0	1.0 38.5 46.7
256	B34R_050_037*	0.375 0.125 0.5	0.5 0.375 0.312	311	0.232 0.124 0.5	35.8 12.3 14.3	18.9 310.5	0.375 0.125 0.5	38.4 26.0 16.9	31.0 326.8 14.2	286 0.285 0.0	1.0 31.9 32.8
257	B25R_062_050*	0.375 0.125 0.625	0.625 0.5 0.375	300	0.194 0.125 0.625	36.6 11.2 20.9	24.2 300.1	0.375 0.125 0.625	38.1 26.4 24.3	35.9 317.3 14.7	277 0.138 0.0	1.0 31.5 24.4
258	B19R_075_062*	0.375 0.125 0.75	0.75 0.625 0.437	293	0.15 0.125 0.75	38.1 11.9 27.4	29.3 30.0	0.375 0.125 0.75	39.9 26.8 29.7	31.2 315 15.1	272 0.04 0.0	1.0 32.2 19.1
259	B15R_087_075*	0.375 0.125 0.875	0.875 0.75 0.5	289	0.125 0.135 0.875	39.6 12.0 33.6	35.7 289.7	0.375 0.125 0.875	36.6 33.1 35.8	48.7 312.7 21.3	269 0.0 0.014	1.0 32.8 16.1
260	B13R_100_087*	0.375 0.125 1.0	1.0 0.875 0.562	286	0.125 0.17 1.0	41.3 12.0 39.8	41.6 286.9	0.375 0.125 1.0	36.6 31.2 36.5	48.1 310.4 19.9	267 0.0 0.052	1.0 33.6 13.8
261	R68Y_037_037*	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.175 0.0	40.7 8.3 24.3	25.7 71.1	0.375 0.25 0.0	41.9 0.3 36.3	36.3 89.4 14.4	57 1.0 0.466	0.0 68.9 22.1
262	R50Y_037_025*	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.204 0.124	42.3 8.8 14.6	17.0 58.8	0.375 0.25 0.125	44.6 0.9 22.4	22.4 84.9 10.9	48 1.0 0.319	0.0 61.8 35.2
263	R00Y_037_012*	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.282	44.8 7.0 3.3	7.7 25.4	0.375 0.25 0.25	44.2 8.2 6.2	10.3 37.2 3.2	375 1.0 0.0	0.263 47.5 56.0
264	B50R_037_012*	0.375 0.25 0.375	0.375 0.125 0.312	330	0.323 0.249 0.375	43.6 5.8 3.5	6.8 328.6	0.375 0.25 0.375	44.6 12.6 8.8	15.4 324.8 8.6	305 0.584 0.0	1.0 38.5 46.7
265	B25R_050_025*	0.375 0.25 0.5	0.5 0.25 0.375	300	0.284 0.249 0.5	43.7 6.1 10.4	12.1 300.1	0.375 0.25 0.5	43.9 14.9 17.7	23.1 310.0 11.4	277 0.138 0.0	1.0 31.5 24.4
266	B15R_062_037*	0.375 0.25 0.625	0.625 0.375 0.437	289	0.25 0.255 0.625	45.2 6.0 16.8	17.8 289.7	0.375 0.25 0.625	43.8 16.7 24.9	30.0 303.9 13.5	269 0.0 0.014	1.0 32.8 16.1
267	B11R_075_050*	0.375 0.25 0.75	0.75 0.5 0.5	284	0.25 0.288 0.75	46.9 6.1 22.9	23.7 285.0	0.375 0.25 0.75	45.3 17.2 29.4	34.1 300.3 12.9	266 0.0 0.077	1.0 34.1 12.2
268	B09R_087_062*	0.375 0.25 0.875	0.875 0.625 0.562	281	0.25 0.322 0.875	48.7 6.2 28.9	29.5 282.1	0.375 0.25 0.875	41.4 23.7 37.0	43.9 302.7 20.6	263 0.0 0.115	1.0 34.8 9.9
269	B07R_100_075*	0.375 0.25 1.0	1.0 0.75 0.625	279	0.25 0.356 1.0	50.4 6.3 35.0	35.6 280.2	0.375 0.25 1.0	40.0 23.6 38.0	44.7 301.8 20.3	262 0.0 0.141	1.0 35.3 8.4
270	Y00G_037_037*	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.288 0.0	46.2 1.1 28.8	28.8 92.3	0.375 0.375 0.0	51.3 10.5 44.1	45.3 103.3 18.6	77 1.0 0.768	0.0 83.6 3.1
271	Y00G_037_025*	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.317 0.124	47.8 0.7 19.2	19.2 92.3	0.375 0.375 0.125	54.5 9.1 32.2	33.5 105.8 16.9	77 1.0 0.768	0.0 83.6 3.1
272	Y00G_037_012*	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.346 0.249	49.3 0.3 9.6	9.6 92.3	0.375 0.375 0.25	52.7 5.7 13.0	14.2 113.7 7.1	77 1.0 0.768	0.0 83.6 3.1
273	NW_037*	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	50.8 0.0 0.0	0.0 0.0	0.375 0.375 0.375	51.9 0.0 1.3	267.6 1.7 360 1.0 1.0	1.0 95.8 0.0	0.0 0.0 0.0
274	B00R_050_012*	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.407 0.5	52.5 0.1 6.0	6.0 271.7	0.375 0.375 0.5	49.1 2.6 17.1	17.3 278.8 11.7	255 0.0 0.261	1.0 37.3 1.4
275	B00R_062_025*	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.44 0.625	54.2 0.3 12.1	12.1 271.7	0.375 0.375 0.625	49.0 5.4 23.7	24.3 282.9 13.6	255 0.0 0.261	1.0 37.3 1.4
276	B00R_075_037*	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.472 0.75	55.8 0.5 18.2	18.2 271.7	0.375 0.375 0.75	50.1 8.2 28.1	29.3 286.2 13.8	255 0.0 0.261	1.0 37.3 1.4
277	B00R_087_050*	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.505 0.875	57.5 0.7 24.3	24.3 271.7	0.375 0.375 0.875	48.1 13.7 36.3	38.8 290.6 20.0	255 0.0 0.261	1.0 37.3 1.4
278	B00R_100_062*	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.538 1.0	59.2 0.9 30.4	30.4 271.7	0.375 0.375 1.0	45.6 15.1 37.4	40.3 291.9 20.8	255 0.0 0.261	1.0 37.3 1.4
279	Y23G_050_050*	0.375 0.5 0.0	0.5 0.5 0.25	104	0.348 0.5 0.0	54.8 13.2 39.2	41.4 108.6	0.375 0.5 0.0	52.5 14.9 46.5	48.9 107.8 7.8	107 0.697	1.0 0.0 85.8
280	Y31G_050_037*	0.375 0.5 0.125	0.5 0.375 0.312	109	0.361 0.5 0.124	54.0 11.7 25.8	28.4 114.4	0.375 0.5 0.125	58.4 15.5 32.6	36.1 115.5 8.8	111 0.631	1.0 0.0 80.4
281	Y50G_050_025*	0.375 0.5 0.25	0.5 0.25 0.375	120	0.375 0.5 0.249	53.6 10.4 13.7	17.2 127.2	0.375 0.5 0.25	57.6 13.8 15.4	20.6 113.8 5.5	119 0.5 1.0	0.0 71.0
282	G00B_050_012*	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.393	54.5 8.2 2.6	8.6 162.2	0.375 0.5 0.375	57.3 10.5 1.3	10.6 172.6 3.8	157 0.0 1.0	0.146 53.8
283	G50B_050_012*	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.473	54.7 4.8 3.6	6.0 216.9	0.375 0.5 0.5	57.8 8.1 14.6	16.7 240.7 11.8	198 0.0 1.0	0.791 54.9
284	G75B_062_025*	0.375 0.5 0.625	0.625 0.25 0.5	240	0.375 0.546 0.625	57.8 5.8 12.1	13.4 244.3	0.375 0.5 0.625	55.5 5.3 22.4	23.0 256.4 10.4	227 0.0 0.686	1.0 51.7
285	G84B_075_037*	0.375 0.5 0.75	0.75 0.375 0.562	251	0.375 0.565 0.75	59.3 5.1 18.5	19.2 254.3	0.375 0.5 0.75	55.8 1.9 26.2	26.2 265.7 8.9	239 0.0 0.508	1.0 46.4
286	G88B_087_050*	0.375 0.5 0.875	0.875 0.5 0.625	256	0.375 0.592 0.875	60.7 4.8 24.7	25.1 258.9	0.375 0.5 0.875	55.6 1.9 34.3	34.3 273.1 12.7	244 0.0 0.434	1.0 43.6
287	G90B_100_062*	0.375 0.5 1.0	1.0 0.625 0.687	259	0.375 0.619 1.0							

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)

TUB material: code=rh4ta

n	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb**Fe	LabCh**Fe	DE**Fe	hsi_Me	rgb*Me	LabCh*Me	
324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.0	0.131	35.7	28.0	13.3	31.0	25.4
325	R26Y_050_050a	0.5	0.0	0.125	0.5	0.5	0.0	0.25	35.8	29.5	5.1	29.9	9.8
326	R00Y_050_050a	0.5	0.0	0.25	0.5	0.5	0.0	0.413	36.6	32.7	-4.5	33.1	352.0
327	B61R_050_050a	0.5	0.0	0.375	0.5	0.5	0.0	0.25	34.0	29.1	-9.5	30.6	341.8
328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.0	0.5	31.1	23.3	-14.2	27.3	328.6
329	B40R_062_062a	0.5	0.0	0.625	0.625	0.625	0.312	319	0.242	0.0	0.625	30.5	24.2
330	B34R_075_075a	0.5	0.0	0.75	0.75	0.75	0.375	311	0.214	0.0	0.75	29.9	24.6
331	B29R_087_087a	0.5	0.0	0.875	0.875	0.875	0.437	305	0.181	0.0	0.875	30.2	24.7
332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	300	0.138	0.0	1.0	31.5	24.4
333	R23Y_050_050a	0.5	0.125	0.0	0.5	0.5	0.25	44	0.5	0.054	0.0	37.6	27.4
334	R00Y_050_037a	0.5	0.125	0.125	0.5	0.375	0.312	390	0.5	0.124	0.223	41.7	21.0
335	R18Y_050_037a	0.5	0.125	0.25	0.5	0.375	0.312	371	0.5	0.124	0.345	41.8	22.9
336	B65R_050_037a	0.5	0.125	0.375	0.5	0.375	0.312	349	0.478	0.124	0.5	41.5	23.6
337	B50R_050_037a	0.5	0.125	0.5	0.5	0.375	0.312	330	0.344	0.124	0.5	38.3	17.5
338	B38R_062_050a	0.5	0.125	0.625	0.625	0.5	0.375	316	0.298	0.125	0.625	37.6	18.2
339	B30R_075_062a	0.5	0.125	0.75	0.75	0.625	0.437	307	0.272	0.125	0.75	37.3	18.6
340	B25R_087_075a	0.5	0.125	0.875	0.875	0.75	0.5	300	0.229	0.125	0.875	38.6	18.3
341	B20R_100_087a	0.5	0.125	1.0	1.0	0.875	0.562	295	0.185	0.125	1.0	40.0	18.0
342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	60	0.5	0.159	0.0	42.8	17.6
343	R31Y_050_037a	0.5	0.25	0.125	0.5	0.375	0.312	49	0.5	0.191	0.124	44.3	18.4
344	R00Y_050_025a	0.5	0.25	0.25	0.5	0.25	0.375	390	0.5	0.249	0.315	47.7	14.0
345	R00Y_050_025a	0.5	0.25	0.375	0.5	0.25	0.375	360	0.5	0.249	0.456	48.2	16.3
346	B50R_050_025a	0.5	0.25	0.5	0.5	0.25	0.375	330	0.396	0.249	0.5	45.5	11.6
347	B34R_062_037a	0.5	0.25	0.625	0.625	0.375	0.437	311	0.357	0.25	0.625	44.8	12.3
348	B25R_075_050a	0.5	0.25	0.75	0.75	0.5	0.5	300	0.319	0.25	0.75	45.6	12.2
349	B19R_087_062a	0.5	0.25	0.875	0.875	0.625	0.562	293	0.275	0.25	0.875	47.1	11.9
350	B15R_100_075a	0.5	0.25	1.0	1.0	0.75	0.625	289	0.25	0.26	1.0	48.5	12.0
351	R76Y_050_050a	0.5	0.375	0.0	0.5	0.5	0.25	76	0.5	0.275	0.0	48.0	8.0
352	R68Y_050_037a	0.5	0.375	0.125	0.5	0.375	0.312	71	0.5	0.3	0.124	49.7	8.3
353	R50Y_050_025a	0.5	0.375	0.25	0.5	0.25	0.375	60	0.5	0.329	0.249	51.3	8.8
354	R00Y_050_012a	0.5	0.375	0.375	0.5	0.125	0.437	390	0.5	0.375	0.407	53.8	7.0
355	B50R_050_012a	0.5	0.375	0.5	0.5	0.125	0.437	330	0.448	0.375	0.5	52.6	5.8
356	B25R_062_025a	0.5	0.375	0.625	0.625	0.25	0.5	300	0.409	0.375	0.625	52.7	6.1
357	B15R_075_037a	0.5	0.375	0.75	0.75	0.375	0.562	289	0.375	0.38	0.75	54.2	6.0
358	B11R_087_050a	0.5	0.375	0.875	0.875	0.5	0.625	284	0.375	0.413	0.875	55.9	6.1
359	B09R_100_062a	0.5	0.375	1.0	1.0	0.625	0.687	281	0.375	0.447	1.0	57.7	6.2
360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	90	0.5	0.384	0.0	53.7	-1.5
361	Y00G_050_037a	0.5	0.5	0.125	0.5	0.375	0.312	90	0.5	0.413	0.124	55.2	-1.1
362	Y00G_050_025a	0.5	0.5	0.25	0.5	0.25	0.375	90	0.5	0.442	0.249	56.8	-0.7
363	Y00G_050_012a	0.5	0.5	0.375	0.5	0.125	0.437	90	0.5	0.471	0.375	58.3	-0.3
364	NW_050a	0.5	0.5	0.5	0.5	0.0	0.5	360	0.5	0.5	0.5	59.8	0.0
365	B00R_062_012a	0.5	0.5	0.625	0.625	0.125	0.562	270	0.5	0.532	0.625	61.5	0.1
366	B00R_075_025a	0.5	0.5	0.75	0.75	0.25	0.625	270	0.5	0.565	0.75	63.2	0.3
367	B00R_087_037a	0.5	0.5	0.875	0.875	0.375	0.687	270	0.5	0.597	0.875	64.8	0.5
368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	270	0.5	0.63	1.0	66.5	0.7
369	Y18G_062_062a	0.5	0.625	0.0	0.625	0.625	0.312	101	0.46	0.625	0.0	64.5	-14.2
370	Y23G_062_050a	0.5	0.625	0.125	0.625	0.5	0.375	104	0.473	0.625	0.125	63.8	-13.2
371	Y31G_062_037a	0.5	0.625	0.25	0.625	0.375	0.437	109	0.486	0.625	0.25	63.0	-11.7
372	Y50G_062_025a	0.5	0.625	0.375	0.625	0.25	0.5	120	0.5	0.625	0.375	62.6	-10.4
373	G00B_062_012a	0.5	0.625	0.5	0.625	0.125	0.562	150	0.5	0.625	0.518	63.5	-8.2
374	G50B_062_012a	0.5	0.625	0.625	0.625	0.125	0.562	210	0.5	0.625	0.598	63.7	-4.8
375	G75B_075_025a	0.5	0.625	0.75	0.75	0.25	0.625	240	0.5	0.671	0.75	66.8	-5.8
376	G84B_087_037a	0.5	0.625	0.875	0.875	0.375	0.687	251	0.5	0.69	0.875	68.3	-5.1
377	G88B_100_050a	0.5	0.625	1.0	1.0	0.5	0.75	256	0.5	0.717	1.0	69.7	-4.8
378	Y31G_075_075a	0.5	0.75	0.0	0.75	0.75	0.375	109	0.473	0.75	0.0	66.2	-23.5
379	Y38G_075_062a	0.5	0.75	0.125	0.75	0.625	0.437	113	0.489	0.75	0.125	66.0	-22.1
380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	120	0.5	0.75	0.25	65.4	-20.8
381	Y68G_075_037a	0.5	0.75	0.375	0.75	0.375	0.562	131	0.49	0.75	0.375	65.6	-19.8
382	G00B_075_025a	0.5	0.75	0.5	0.75	0.25	0.625	150	0.5	0.75	0.536	65.3	-16.4
383	G25B_075_025a	0.5	0.75	0.625	0.75	0.25	0.625	180	0.5	0.75	0.624	67.6	-12.9
384	G50B_075_025a	0.5	0.75	0.75	0.75	0.25	0.625	210	0.5	0.75	0.697	67.6	-9.6
385	G65B_087_037a	0.5	0.75	0.875	0.875	0.375	0.687	229	0.5	0.875	0.872	70.8	-11.4
386	G75B_100_050a	0.5	0.75	1.0	1.0	0.5	0.75	240	0.5	0.843	1.0	73.7	-11.6
387	Y41G_087_087a	0.5	0.875	0.0	0.875	0.875	0.437	115	0.489	0.875	0.0	68.8	-32.7
388	Y50G_087_075a	0.5	0.875	0.125	0.875	0.75	0.5	120	0.5	0.875	0.125	68.2	-31.3
389	Y61G_087_062a	0.5	0.875	0.25	0.875	0.625	0.562	127	0.478	0.875	0.25	68.3	-30.1
390	Y76G_087_050a	0.5	0.875	0.375	0.875	0.5	0.625	136	0.488	0.875	0.375	68.9	-29.1
391	G00B_087_037a	0.5	0.875	0.5	0.875	0.375	0.687	150	0.5	0.875	0.555	71.0	-24.7
392	G15B_087_037a	0.5	0.875	0.625	0.875	0.375	0.687	169	0.5	0.875	0.638	71.4	-21.4
393	G34B_087_037a	0.5	0.875	0.75	0.875	0.375	0.687	191	0.5	0.875	0.718	71.6	-17.5
394	G50B_087_037a	0.5	0.875	0.875	0.875	0.375	0.687	210	0.5	0.875	0.796	71.5	-14.5
395	G61B_100_050a	0.5	0.875	1.0	1.0	0.5	0.75	224	0.5	1.0	0.974	74.7	-16.5
396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	120	0.5	1.0	0.0	71.0	-41.7
397	Y58G_100_087a	0.5	1.0	0.125	1.0	0.875	0.562	125	0.477	1.0	0.125	71.0	-40.5
398	Y68G_100_075a	0.5	1.0	0.25	1.0	0.75	0.625	131	0.481	1.0	0.25	71.5	-39.6
399	Y81G_100_062a	0.5	1.0	0.375	1.0	0.625	0.687	139	0.473	1.0	0.375	72.1	-38.1
400	G00B_100_050a	0.5	1.0	0.5	1.0	0.5	0.75	150	0.5	1.0	0.573	74.8	-32.9
401	G11B_100_050a	0.5	1.0	0.625	1.0	0.5	0.75	164	0.5	1.0	0.66	75.0	-29.9
402	G25B_100_050a	0.5	1.0	0.75	1.0	0.5	0.75	186	0.5	1.0	0.748	75.4	-25.8
403	G38B_100_050a	0.5	1.0	0.875	1.0	0.5	0.75	196	0.5	1.0	0.811	75.5	-22.1
404													

delta E* = 10.9

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
colores y diferencia en color, ΔE^* , 3D=0, de=1, cmyk

entrada: *rgbcmyk* -> *rgbe*
salida: transfiera a *cmyke*

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb*Fe	LabCh*Fe	DE*Fe	hsi_Me	rgb*Me	LabCh*Me
405	R00Y_062_062a	0.625 0.0 0.0	0.625 0.625 0.312	390	0.625 0.0 0.164	38.6 35.0 16.7	38.8 25.4	0.625 0.0 0.0	36.3 40.2 26.2	48.0 33.1 11.0	375	1.0 0.0 0.263 47.5 56.0 26.7 62.1 25.4
406	R30Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.284	38.7 36.4 8.5	37.4 13.2	0.625 0.0 0.125	37.1 40.1 21.9	45.7 28.6 13.9	362	1.0 0.0 0.454 47.6 58.3 13.7 59.9 13.2
407	R11Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.412	39.1 39.1 0.0	39.1 359.8	0.625 0.0 0.25	36.4 41.6 12.6	43.4 16.8 13.2	349	1.0 0.0 0.659 48.3 62.6 -0.1 62.6 359.8
408	B69R_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.562	39.7 41.2 -6.9	41.8 350.4	0.625 0.0 0.375	36.9 45.2 0.9	45.2 1.1 9.2	335	1.0 0.0 0.899 49.2 66.0 -11.1 66.9 350.4
409	B59R_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.473 0.0 0.625	35.2 34.6 -13.2	37.1 339.0	0.625 0.0 0.5	37.2 49.0 -7.9	49.6 350.7 15.4	316	0.756 0.0 1.0 42.1 55.4 -28.2 59.3 339.0
410	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.365 0.0 0.625	33.0 29.2 -17.8	34.2 328.6	0.625 0.0 0.625	37.9 49.8 -14.9	52.0 343.3 21.4	305	0.584 0.0 1.0 38.5 46.7 -21.5 54.7 328.6
411	B42R_075_075a	0.625 0.0 0.75	0.75 0.75 0.375	321	0.316 0.0 0.75	32.4 30.0 -25.1	39.2 320.0	0.625 0.0 0.75	38.9 50.3 -20.0	54.1 338.3 21.8	294	0.421 0.0 1.0 35.3 40.1 -33.5 52.3 320.0
412	B36R_087_087a	0.625 0.0 0.875	0.875 0.875 0.437	314	0.282 0.0 0.875	31.7 30.7 -32.4	44.6 313.4	0.625 0.0 0.875	39.3 49.4 -24.6	55.2 333.4 21.6	288	0.322 0.0 1.0 32.9 35.0 -37.0 51.0 313.4
413	B31R_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.249 0.0 1.0	31.0 30.5 -39.4	49.8 307.7	0.625 0.0 1.0	39.1 48.4 -27.2	55.6 330.6 23.1	283	0.249 0.0 1.0 31.0 30.5 -39.4 49.8 307.7
414	R18Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	41	0.625 0.038 0.0	40.0 35.0 27.1	44.3 37.7	0.625 0.125 0.0	40.3 31.8 34.0	46.6 46.8 7.5	33	1.0 0.006 0.0 49.7 56.0 43.3 70.8 37.7
415	R00Y_062_050a	0.625 0.125 0.125	0.625 0.5 0.375	390	0.625 0.125 0.256	44.7 28.0 13.3	31.0 25.4	0.625 0.125 0.125	44.2 28.5 28.7	40.5 45.1 15.3	375	1.0 0.0 0.263 47.5 56.0 26.7 62.1 25.4
416	R26Y_062_050a	0.625 0.125 0.25	0.625 0.5 0.375	376	0.625 0.125 0.375	44.8 29.5 5.1	29.9 9.8	0.625 0.125 0.25	43.5 29.7 15.8	33.7 28.0 10.8	359	1.0 0.0 0.501 47.8 59.0 10.2 59.9 9.8
417	R00Y_062_050a	0.625 0.125 0.375	0.625 0.5 0.375	360	0.625 0.125 0.538	45.6 32.7 -4.5	33.1 352.0	0.625 0.125 0.375	43.7 33.1 4.7	33.5 8.0 9.4	339	1.0 0.0 0.827 49.4 65.5 -9.1 66.2 352.0
418	B61R_062_050a	0.625 0.125 0.5	0.625 0.5 0.375	344	0.537 0.125 0.625	43.0 29.1 -9.5	30.6 341.8	0.625 0.125 0.5	43.8 38.9 -5.6	39.3 351.7 10.6	320	0.825 0.0 1.0 44.1 58.2 -19.0 61.2 341.8
419	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.417 0.125 0.625	40.1 23.3 -14.2	27.3 328.6	0.625 0.125 0.625	44.4 41.4 -12.9	43.4 342.7 18.6	305	0.584 0.0 1.0 38.5 46.7 -21.5 54.7 328.6
420	B40R_075_062a	0.625 0.125 0.75	0.75 0.625 0.437	319	0.367 0.125 0.75	39.5 24.2 -21.6	32.4 318.1	0.625 0.125 0.75	44.8 43.5 -17.8	47.0 337.7 20.4	292	0.387 0.0 1.0 34.5 38.7 -34.6 51.9 318.1
421	B34R_087_075a	0.625 0.125 0.875	0.875 0.75 0.5	311	0.339 0.125 0.875	38.9 24.6 -28.7	37.8 310.5	0.625 0.125 0.875	44.1 46.8 -24.2	52.7 332.6 23.2	286	0.285 0.0 1.0 31.9 32.8 -38.2 50.4 310.5
422	B29R_100_087a	0.625 0.125 1.0	1.0 0.875 0.562	305	0.306 0.125 1.0	39.2 24.7 -35.4	43.1 304.9	0.625 0.125 1.0	41.0 43.2 -29.0	52.0 326.1 19.6	281	0.207 0.0 1.0 31.2 28.2 -40.4 49.3 304.9
423	R38Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	53	0.625 0.143 0.0	44.7 27.4 34.0	43.6 51.0	0.625 0.25 0.0	45.9 19.0 40.0	44.3 64.5 10.4	42	1.0 0.229 0.0 57.2 43.9 54.4 69.9 51.0
424	R23Y_062_050a	0.625 0.25 0.125	0.625 0.5 0.375	44	0.625 0.179 0.125	46.6 27.4 23.8	36.3 41.0	0.625 0.25 0.125	50.5 16.9 33.1	37.2 62.8 14.4	35	1.0 0.108 0.0 51.4 54.8 47.7 72.6 41.0
425	R00Y_062_037a	0.625 0.25 0.25	0.625 0.375 0.437	390	0.625 0.25 0.348	50.7 21.0 10.0	23.2 25.4	0.625 0.25 0.25	50.5 18.8 20.3	27.7 47.2 10.5	375	1.0 0.0 0.263 47.5 56.0 26.7 62.1 25.4
426	R18Y_062_037a	0.625 0.25 0.375	0.625 0.375 0.437	371	0.625 0.25 0.47	50.8 22.9 1.7	22.9 4.3	0.625 0.25 0.375	50.6 22.5 8.2	24.0 19.9 6.4	354	1.0 0.0 0.588 47.9 61.1 4.6 61.2 4.3
427	B65R_062_037a	0.625 0.25 0.5	0.625 0.375 0.437	349	0.603 0.25 0.625	50.5 23.6 -5.6	24.2 346.6	0.625 0.25 0.5	51.8 27.5 -4.1	27.8 351.4 4.4	327	0.941 0.0 1.0 47.0 63.0 -14.9 64.7 346.6
428	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.469 0.25 0.625	47.3 17.5 -10.7	20.5 328.6	0.625 0.25 0.625	52.0 30.3 -11.7	32.5 338.7 13.6	305	0.584 0.0 1.0 38.5 46.7 -28.5 54.7 328.6
429	B38R_075_050a	0.625 0.25 0.75	0.75 0.5 0.5	316	0.423 0.25 0.75	46.6 18.2 -18.0	25.7 315.3	0.625 0.25 0.75	52.3 31.0 -15.4	34.7 333.4 14.2	289	0.347 0.0 1.0 33.5 36.5 -36.1 51.4 315.3
430	B30R_087_062a	0.625 0.25 0.875	0.875 0.625 0.562	307	0.397 0.25 0.875	46.3 18.6 -24.8	31.0 306.8	0.625 0.25 0.875	50.3 37.1 -24.3	44.3 326.7 18.9	283	0.235 0.0 1.0 31.0 29.7 -39.7 49.6 306.8
431	B25R_100_075a	0.625 0.25 1.0	1.0 0.75 0.625	300	0.354 0.25 1.0	47.6 18.3 -31.4	36.4 300.1	0.625 0.25 1.0	45.0 36.4 -29.0	46.6 321.4 18.5	277	0.138 0.0 1.0 31.5 24.4 -41.9 48.5 300.1
432	R61Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	67	0.625 0.257 0.0	50.4 16.7 38.9	42.4 66.6	0.625 0.375 0.0	51.0 6.3 45.7	46.2 82.0 12.4	54	1.0 0.411 0.0 66.3 26.8 62.3 67.8 66.6
433	R50Y_062_050a	0.625 0.375 0.125	0.625 0.5 0.375	60	0.625 0.284 0.125	51.8 17.6 29.2	34.1 58.8	0.625 0.375 0.125	57.4 6.1 39.1	39.6 81.0 16.1	48	1.0 0.319 0.0 61.8 35.2 58.4 68.2 58.8
434	R31Y_062_037a	0.625 0.375 0.25	0.625 0.375 0.437	49	0.625 0.316 0.25	53.3 18.4 19.5	26.8 46.6	0.625 0.375 0.25	57.6 8.9 25.0	26.5 70.2 11.7	39	1.0 0.177 0.0 54.6 49.1 52.0 71.6 46.6
435	R00Y_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.44	56.7 14.0 6.6	15.5 25.4	0.625 0.375 0.375	58.1 11.8 11.2	16.3 43.4 5.2	375	1.0 0.0 0.263 47.5 56.0 26.7 62.1 25.4
436	R00Y_062_025a	0.625 0.375 0.5	0.625 0.25 0.5	360	0.625 0.375 0.581	57.2 16.3 -2.2	16.5 352.0	0.625 0.375 0.5	59.0 15.5 -0.3	15.5 358.8 2.7	339	1.0 0.0 0.827 49.4 65.5 -9.1 66.2 352.0
437	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.521 0.375 0.625	54.5 11.6 -7.1	13.6 328.6	0.625 0.375 0.625	59.5 19.7 -8.9	21.7 335.6 9.7	305	0.584 0.0 1.0 38.5 46.7 -28.5 54.7 328.6
438	B34R_075_037a	0.625 0.375 0.75	0.75 0.375 0.562	311	0.482 0.375 0.75	53.8 12.3 -14.3	18.9 310.5	0.625 0.375 0.75	58.0 21.5 -14.7	26.1 325.7 10.1	286	0.285 0.0 1.0 31.9 32.8 -38.2 50.4 310.5
439	B25R_087_050a	0.625 0.375 0.875	0.875 0.5 0.625	300	0.444 0.375 0.875	54.6 12.2 -20.9	24.2 300.1	0.625 0.375 0.875	56.8 27.0 -23.8	36.0 318.5 15.2	277	0.138 0.0 1.0 31.5 24.4 -41.9 48.5 300.1
440	B19R_100_062a	0.625 0.375 1.0	1.0 0.625 0.687	293	0.4 0.375 1.0	56.0 11.9 -27.4	29.9 293.5	0.625 0.375 1.0	51.8 26.1 -28.3	38.6 312.6 14.8	272	0.04 0.0 1.0 32.2 19.1 -43.9 47.9 293.5
441	R18Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	79	0.625 0.382 0.0	55.4 7.7 43.9	44.6 80.0	0.625 0.5 0.0	60.5 -3.4	56.6 56.7 93.4	17.6 67	0.1 0.611 0.0 74.4 12.3 70.3 71.3 80.0
442	R76Y_062_050a	0.625 0.5 0.125	0.625 0.5 0.375	76	0.625 0.4 0.125	57.0 8.0 34.1	35.0 76.7	0.625 0.5 0.125	64.8 -4.4	49.8 50.0 95.0	21.5 63	1.0 0.551 0.0 72.3 16.1 68.2 70.1 76.7
443	R68Y_062_037a	0.625 0.5 0.25	0.625 0.375 0.437	71	0.625 0.425 0.25	58.7 8.3 24.3	25.7 71.1	0.625 0.5 0.25	66.0 -2.3	33.6 33.7 94.0	15.9 57	1.0 0.466 0.0 68.9 22.1 64.9 68.5 71.1
444	R50Y_062_025a	0.625 0.5 0.375	0.625 0.25 0.5	60	0.625 0.454 0.375	60.3 8.8 14.6	17.0 58.8	0.625 0.5 0.375	65.6 1.4 17.9	17.9 85.5 9.7 48	10	1.0 0.319 0.0 61.8 35.2 58.4 68.2 58.8
445	R00Y_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	390	0.625 0.5 0.532	62.8 7.0 3.3	7.7 25.4	0.625 0.5 0.5	65.7 5.9 3.3	6.8 29.2 3.0	375	1.0 0.0 0.263 47.5 56.0 26.7 62.1 25.4
446	B50R_062_012a	0.625 0.5 0.625	0.625 0.125 0.562									

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)
TUB material: code=rha4ta

n	HIC* _{Fe}	rgb_Fe	ict_Fe	hsi_Fe	rgb* _{Fe}	LabCh* _{Fe}	rgb** _{Fe}	LabCh** _{Fe}	DE** _{Fe}	hsi _{Me}	rgb* _{Me}	LabCh* _{Me}
486	R00Y_075_075a	0.75 0.0 0.0	0.75 0.75 0.375	390	0.75 0.0 0.197	41.6 42.0 20.0	46.5 25.4	0.75 0.0 0.0	39.7 47.0 29.4	55.5 32.0 10.8	375	1.0 0.0 0.263 47.5
487	R35Y_075_075a	0.75 0.0 0.125	0.75 0.75 0.375	381	0.75 0.0 0.317	41.6 43.3 11.9	45.0 15.4	0.75 0.0 0.125	39.0 46.5 23.4	52.0 26.7 12.1	365	1.0 0.0 0.423 47.5
488	R18Y_075_075a	0.75 0.0 0.25	0.75 0.75 0.375	371	0.75 0.0 0.441	41.9 45.8 3.4	45.9 4.3	0.75 0.0 0.25	39.4 47.0 16.3	49.8 19.1 13.1	354	1.0 0.0 0.588 47.9
489	R00Y_075_075a	0.75 0.0 0.375	0.75 0.75 0.375	360	0.75 0.0 0.62	43.0 49.1 -6.8	49.6 352.0	0.75 0.0 0.375	39.3 48.8 6.0	49.2 7.1 13.4	339	1.0 0.0 0.827 49.4
490	B65R_075_075a	0.75 0.0 0.5	0.75 0.75 0.375	349	0.706 0.0 0.75	41.2 47.2 -11.2	48.5 346.6	0.75 0.0 0.5	39.5 52.9 -4.1	53.1 35.4 9.2	327	0.941 0.0 1.0 47.0
491	B57R_075_075a	0.75 0.0 0.625	0.75 0.75 0.375	339	0.543 0.0 0.75	36.9 40.4 -17.0	43.8 337.1	0.75 0.0 0.625	39.9 55.3 -11.1	56.4 34.6 16.3	314	0.725 0.0 1.0 41.3
492	B50R_075_075a	0.75 0.0 0.75	0.75 0.75 0.375	330	0.438 0.0 0.75	34.8 35.0 -21.4	41.0 328.6	0.75 0.0 0.75	41.1 54.2 -16.4	56.6 34.3 20.7	305	0.584 0.0 1.0 38.5
493	B43R_087_087a	0.75 0.0 0.875	0.875 0.875 0.375	322	0.383 0.0 0.875	34.2 35.7 -28.8	45.9 321.0	0.75 0.0 0.875	42.0 54.2 -21.0	58.1 338.8 21.5	295	0.438 0.0 1.0 35.7
494	B38R_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.347 0.0 1.0	33.5 36.5 -36.1	51.4 315.3	0.75 0.0 1.0	41.8 55.1 -21.4	59.1 338.7 25.1	289	0.347 0.0 1.0 33.5
495	R15Y_075_075a	0.75 0.125 0.0	0.75 0.75 0.375	39	0.75 0.021 0.0	42.4 42.5 30.3	52.2 35.5	0.75 0.125 0.0	44.9 42.3 40.8	58.8 43.9 10.8	31	1.0 0.0028 0.0 48.6
496	R00Y_075_062a	0.75 0.125 0.125	0.75 0.625 0.437	390	0.75 0.125 0.289	47.6 35.0 16.7	38.8 25.4	0.75 0.125 0.125	45.5 36.3 31.1	47.8 40.6 14.6	375	1.0 0.0 0.263 47.5
497	R31Y_075_062a	0.75 0.125 0.25	0.75 0.625 0.437	379	0.75 0.125 0.409	47.7 36.4 8.5	37.4 13.2	0.75 0.125 0.25	45.6 37.0 22.6	43.3 31.4 14.2	362	1.0 0.0 0.454 47.6
498	R11Y_075_062a	0.75 0.125 0.375	0.75 0.625 0.437	367	0.75 0.125 0.537	48.1 39.1 0.0	39.1 359.8	0.75 0.125 0.375	45.0 38.4 14.5	41.1 20.7 15.0	349	1.0 0.0 0.659 48.3
499	B69R_075_062a	0.75 0.125 0.5	0.75 0.625 0.437	353	0.75 0.125 0.687	48.7 41.2 -6.9	41.8 350.4	0.75 0.125 0.5	45.2 42.6 3.5	42.7 4.7 11.1	335	1.0 0.0 0.899 49.2
500	B59R_075_062a	0.75 0.125 0.625	0.75 0.625 0.437	341	0.598 0.125 0.75	44.2 34.6 -13.2	37.1 339.0	0.75 0.125 0.625	45.8 46.9 -5.5	47.2 35.3 14.6	316	0.756 0.0 1.0 38.5
501	B50R_075_062a	0.75 0.125 0.75	0.75 0.625 0.437	330	0.49 0.125 0.75	42.0 29.2 -17.8	34.2 328.6	0.75 0.125 0.75	46.6 48.7 -13.3	50.5 34.6 20.5	305	0.584 0.0 1.0 38.5
502	B42R_087_075a	0.75 0.125 0.875	0.875 0.75 0.5	321	0.441 0.125 0.875	41.4 30.0 -25.1	39.2 320.0	0.75 0.125 0.875	47.1 50.2 -17.9	53.3 34.0 22.1	294	0.421 0.0 1.0 35.3
503	B36R_100_087a	0.75 0.125 1.0	1.0 0.875 0.562	314	0.407 0.125 1.0	40.7 30.7 -32.4	44.6 313.4	0.75 0.125 1.0	45.0 50.7 -22.1	55.3 33.6 22.8	288	0.322 0.0 1.0 32.9
504	R31Y_075_075a	0.75 0.25 0.0	0.75 0.75 0.375	49	0.75 0.132 0.0	46.9 36.8 39.0	53.7 46.6	0.75 0.25 0.0	52.2 26.4 47.8	54.6 61.1 14.6	39	1.0 0.177 0.0 54.6
505	R18Y_075_062a	0.75 0.25 0.125	0.75 0.625 0.437	41	0.75 0.163 0.125	49.0 35.0 27.1	44.3 37.7	0.75 0.25 0.125	52.1 26.6 36.8	45.4 54.0 13.1	33	1.0 0.06 0.0 49.7
506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.381	53.7 28.0 13.3	31.0 25.4	0.75 0.25 0.25	52.4 27.1 25.3	37.1 43.0 12.1	375	1.0 0.0 0.263 47.5
507	R26Y_075_050a	0.75 0.25 0.375	0.75 0.5 0.5	376	0.75 0.25 0.5	53.8 29.5 5.1	29.9 9.8	0.75 0.25 0.375	52.3 29.3 16.1	33.5 28.8 11.1	359	1.0 0.0 0.501 47.8
508	R00Y_075_050a	0.75 0.25 0.5	0.75 0.5 0.5	360	0.75 0.25 0.663	54.6 32.7 -4.5	33.1 352.0	0.75 0.25 0.5	53.2 30.6 5.9	31.2 10.9 10.8	339	1.0 0.0 0.827 49.4
509	B61R_075_050a	0.75 0.25 0.625	0.75 0.5 0.5	344	0.662 0.25 0.75	52.0 29.1 -9.5	30.6 341.8	0.75 0.25 0.625	53.9 34.9 -4.4	35.2 35.2 7.9	320	0.825 0.0 1.0 44.1
510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.542 0.25 0.75	49.1 23.3 -14.2	27.3 328.6	0.75 0.25 0.75	53.9 38.1 -12.4	40.1 341.9 15.6	305	0.584 0.0 1.0 38.5
511	B40R_087_062a	0.75 0.25 0.875	0.875 0.625 0.562	319	0.492 0.25 0.875	48.5 24.2 -21.6	32.4 318.1	0.75 0.25 0.875	54.3 40.1 -17.3	43.7 33.6 17.4	292	0.387 0.0 1.0 34.5
512	B34R_100_075a	0.75 0.25 1.0	1.0 0.75 0.625	311	0.464 0.25 1.0	47.9 24.6 -28.7	37.8 310.5	0.75 0.25 1.0	50.3 42.9 -22.8	48.6 33.0 19.4	286	0.285 0.0 1.0 31.9
513	R50Y_075_075a	0.75 0.375 0.0	0.75 0.75 0.375	60	0.75 0.239 0.0	52.3 26.4 43.8	51.1 58.8	0.75 0.375 0.0	57.7 15.3 53.6	55.7 74.0 15.7	48	1.0 0.319 0.0 61.8
514	R38Y_075_062a	0.75 0.375 0.125	0.75 0.625 0.437	53	0.75 0.268 0.125	53.7 27.4 34.0	43.6 51.0	0.75 0.375 0.125	57.9 16.7 44.5	47.5 69.3 15.5	42	1.0 0.229 0.0 57.2
515	R23Y_075_050a	0.75 0.375 0.25	0.75 0.5 0.5	44	0.75 0.304 0.25	55.6 27.4 23.8	36.3 41.0	0.75 0.375 0.25	58.5 17.2 29.7	34.3 59.9 12.1	35	1.0 0.108 0.0 51.4
516	R00Y_075_037a	0.75 0.375 0.375	0.75 0.375 0.562	390	0.75 0.375 0.473	59.7 21.0 10.0	23.2 25.4	0.75 0.375 0.375	59.1 18.5 19.5	26.9 46.3 9.8	375	1.0 0.0 0.263 47.5
517	R18Y_075_037a	0.75 0.375 0.5	0.75 0.375 0.562	371	0.75 0.375 0.595	59.8 22.9 1.7	22.9 4.3	0.75 0.375 0.5	60.1 19.9 9.2	21.9 24.9 8.1	354	1.0 0.0 0.588 47.9
518	B65R_075_037a	0.75 0.375 0.625	0.75 0.375 0.562	349	0.728 0.375 0.75	59.5 23.6 -5.6	24.2 346.6	0.75 0.375 0.625	60.2 24.2 -2.3	24.3 35.4 3.4	327	0.941 0.0 1.0 47.0
519	B50R_075_037a	0.75 0.375 0.75	0.75 0.375 0.562	330	0.594 0.375 0.75	56.3 17.5 -10.7	20.5 328.6	0.75 0.375 0.75	60.1 28.2 -10.2	30.0 339.9 11.3	305	0.584 0.0 1.0 38.5
520	B38R_087_050a	0.75 0.375 0.875	0.875 0.5 0.625	316	0.548 0.375 0.875	55.6 18.2 -18.0	25.7 315.3	0.75 0.375 0.875	60.4 31.7 -15.9	35.5 33.4 14.4	289	0.347 0.0 1.0 33.5
521	B30R_100_062a	0.75 0.375 1.0	1.0 0.625 0.687	307	0.522 0.375 1.0	55.3 18.6 -24.8	31.0 306.8	0.75 0.375 1.0	55.2 36.2 -21.7	42.2 329.0 17.9	283	0.235 0.0 1.0 31.0
522	R68Y_075_075a	0.75 0.5 0.0	0.75 0.75 0.375	71	0.75 0.35 0.0	57.6 16.6 48.6	51.4 71.1	0.75 0.5 0.0	64.3 4.5 59.9	60.1 85.6 17.8	57	1.0 0.466 0.0 68.9
523	R61Y_075_062a	0.75 0.5 0.125	0.75 0.625 0.437	67	0.75 0.382 0.125	59.4 16.7 38.9	42.4 66.6	0.75 0.5 0.125	65.5 5.4 54.3	54.6 84.2 20.0	54	1.0 0.411 0.0 66.3
524	R50Y_075_050a	0.75 0.5 0.25	0.75 0.5 0.5	60	0.75 0.409 0.25	60.8 17.6 29.2	34.1 58.8	0.75 0.5 0.25	66.1 6.5 36.1	36.6 79.7 14.0	48	1.0 0.319 0.0 61.8
525	R31Y_075_037a	0.75 0.5 0.375	0.75 0.375 0.562	49	0.75 0.441 0.375	62.3 18.4 19.5	26.8 46.6	0.75 0.5 0.375	65.9 8.9 24.5	26.0 70.0 11.3	39	1.0 0.177 0.0 54.6
526	R00Y_075_025a	0.75 0.5 0.5	0.75 0.25 0.625	390	0.75 0.5 0.565	65.7 14.0 6.6	15.5 25.4	0.75 0.5 0.5	67.2 10.6 12.7	16.5 50.1 7.0	375	1.0 0.0 0.263 47.5
527	R00Y_075_025a	0.75 0.625 0.5	0.75 0.25 0.625	360	0.75 0.5 0.706	66.2 16.3 -2.2	16.5 352.0	0.75 0.5 0.625	66.8 15.2 1.2	15.1 4.5 3.7	339	1.0 0.0 0.827 49.4
528	B50R_075_025a	0.75 0.5 0.75	0.75 0.25 0.625	330	0.646 0.5 0.75	63.5 11.6 -7.1	13.6 328.6	0.75 0.5 0.75	67.6 18.9 -8.5	20.7 335.7 8.4	305	0.584 0.0 1.0 38.5
529	B34R_087_037a	0.75 0.5 0.875	0.875 0.375 0.687	311	0.607 0.5 0.875	62.8 12.3 -14.3	18.9 310.5	0.75 0.5 0.875	67.3 22.2 -14.9	26.7 326.1 10.9	286	0.285 0.0 1.0 31.9
530	B25R_100_050a	0.75 0.5 1.0	1.0 0.5 0.75	300	0.569 0.5 1.0	63.6 12.2 -20.9	24.2 300.1	0.75 0.5 1.0	63.6 27.1 -18.9	33.0 325.2 15.0	277	0.138 0.0 1.0 31.5
531	R85Y_075_075a	0.75 0.625 0.0	0.75 0.75 0.375	81	0.75 0.481 0.0	62.9 7.2 53.8	54.3 82.2	0.75 0.625 0.0	74.6 -5.8 70.9	71.1 94.7 24.5	69	1.0 0.642 0.0 76.0

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb*Fe	LabCh*Fe	DE**Fe	hsiMe	rgb*Me	LabCh*Me		
567	R00Y_087_087_	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.23	44.5 49.0 23.3	54.3 25.4	0.875 0.0 0.0	43.6 51.3 31.6	60.2 31.6 8.6	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
568	R36Y_087_087_	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.357	44.5 50.3 14.9	52.5 16.5	0.875 0.0 0.125	43.2 51.0 26.3	57.5 27.3 11.5	366	1.0 0.0 0.408	47.5 57.5 17.1	60.0 16.5
569	R23Y_087_087_	0.875 0.0 0.25	0.875 0.875 0.437	374	0.875 0.0 0.469	44.8 52.4 7.0	52.9 7.6	0.875 0.0 0.25	43.0 51.3 20.1	55.1 21.4 13.2	357	1.0 0.0 0.536	47.8 59.9 8.0	60.4 7.6
570	R08Y_087_087_	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.608	45.6 55.5 -2.2	55.6 357.6	0.875 0.0 0.375	43.3 53.1 10.6	54.2 11.3 13.4	347	1.0 0.0 0.695	48.7 63.4 -2.6	63.5 357.6
571	B70R_087_087_	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.716	46.2 57.2 -7.7	57.8 352.3	0.875 0.0 0.5	43.4 55.6 1.4	55.6 1.4 9.6	339	1.0 0.0 0.818	49.4 65.4 -8.8	66.0 352.3
572	B63R_087_087_	0.875 0.0 0.625	0.875 0.875 0.437	346	0.762 0.0 0.875	42.8 52.5 -15.2	54.6 343.7	0.875 0.0 0.625	43.8 59.4 -6.7	59.8 353.5 11.0	323	0.87 0.0 1.0	45.5 60.0 -17.4	62.5 343.7
573	B56R_087_087_	0.875 0.0 0.75	0.875 0.875 0.437	338	0.621 0.0 0.875	38.8 46.4 -20.5	50.8 336.1	0.875 0.0 0.75	45.4 60.7 -12.7	62.1 348.1 17.5	313	0.71 0.0 1.0	41.0 53.1 -23.4	58.0 336.1
574	B50R_087_087_	0.875 0.0 0.875	0.875 0.875 0.437	330	0.511 0.0 0.875	36.6 40.9 -24.9	47.9 328.6	0.875 0.0 0.875	45.5 59.0 -16.8	61.4 344.0 21.8	305	0.584 0.0 1.0	38.5 46.7 -28.5	54.7 328.6
575	B44R_100_100_	0.875 0.0 1.0	1.0 1.0 0.5	323	0.455 0.0 1.0	36.1 41.4 -32.4	52.6 321.9	0.875 0.0 1.0	45.6 60.1 -17.3	62.6 343.9 25.8	297	0.455 0.0 1.0	36.1 41.4 -32.4	52.6 321.9
576	R13Y_087_087_	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.011 0.0	45.0 49.8 34.1	60.4 34.3	0.875 0.125 0.0	49.7 50.4 46.7	68.7 42.8 13.4	30	1.0 0.012 0.0	48.0 57.0 39.0	69.1 34.3
577	R00Y_087_075_	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.322	50.6 42.0 20.0	46.5 25.4	0.875 0.125 0.125	49.6 47.7 41.6	63.3 41.1 22.3	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
578	R35Y_087_075_	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.442	50.6 43.3 11.9	45.0 15.4	0.875 0.125 0.25	49.6 47.9 30.7	56.9 32.6 19.3	365	1.0 0.0 0.423	47.5 57.8 15.9	60.0 15.4
579	R18Y_087_075_	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.566	50.9 45.8 3.4	45.9 4.3	0.875 0.125 0.375	49.3 49.0 22.0	53.8 24.1 18.9	354	1.0 0.0 0.588	47.9 61.1 4.6	61.2 4.3
580	R00Y_087_075_	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.745	52.0 49.1 -6.8	49.6 352.0	0.875 0.125 0.5	49.5 51.8 11.4	53.0 12.4 18.6	339	1.0 0.0 0.827	49.4 65.5 -9.1	66.2 352.0
581	B65R_087_075_	0.875 0.125 0.625	0.875 0.75 0.5	349	0.831 0.125 0.875	50.2 47.2 -11.2	48.5 346.6	0.875 0.125 0.625	49.1 55.8 0.5	55.8 0.6 14.6	327	0.941 0.0 1.0	47.0 63.0 -14.9	64.7 346.6
582	B57R_087_075_	0.875 0.125 0.75	0.875 0.75 0.5	339	0.668 0.125 0.875	45.9 40.4 -17.0	43.8 337.1	0.875 0.125 0.75	50.0 58.8 -10.1	59.7 350.2 20.1	314	0.725 0.0 1.0	41.3 53.8 -22.7	58.4 337.1
583	B50R_087_075_	0.875 0.125 0.875	0.875 0.75 0.5	330	0.563 0.125 0.875	43.8 35.0 -21.4	41.0 328.6	0.875 0.125 0.875	49.2 57.7 -16.2	60.0 344.2 23.8	305	0.584 0.0 1.0	38.5 46.7 -28.5	54.7 328.6
584	B43R_100_087_	0.875 0.125 1.0	1.0 0.875 0.562	322	0.508 0.125 1.0	43.2 35.7 -28.8	45.9 321.0	0.875 0.125 1.0	48.0 56.9 -17.9	59.6 342.4 24.3	295	0.438 0.0 1.0	35.7 40.8 -33.0	52.4 321.0
585	R26Y_087_087_	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.12 0.0	49.0 46.4 43.7	63.7 43.3	0.875 0.25 0.0	56.8 35.4 52.6	63.5 56.0 16.1	37	1.0 0.138 0.0	52.6 53.0 49.9	72.8 43.3
586	R15Y_087_075_	0.875 0.25 0.125	0.875 0.75 0.5	39	0.875 0.146 0.125	51.4 42.5 30.3	52.2 35.5	0.875 0.25 0.125	55.4 36.3 46.0	58.6 51.6 17.3	31	1.0 0.028 0.0	48.6 56.7 40.4	69.6 35.5
587	R00Y_087_062_	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.414	56.6 35.0 16.7	38.8 25.4	0.875 0.25 0.25	56.9 34.5 32.7	47.5 43.4 16.0	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
588	R31Y_087_062_	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.534	56.7 36.4 8.5	37.4 13.2	0.875 0.25 0.375	56.3 36.2 24.8	43.9 43.4 16.3	362	1.0 0.0 0.454	47.6 58.3 13.7	59.9 13.2
589	R11Y_087_062_	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.662	57.1 39.1 0.0	39.1 359.8	0.875 0.25 0.5	56.0 38.3 15.0	41.1 21.4 15.2	349	1.0 0.0 0.659	48.3 62.6 -0.1	62.6 359.8
590	B69R_087_062_	0.875 0.25 0.625	0.875 0.625 0.562	353	0.875 0.25 0.812	57.7 41.2 -6.9	41.8 350.4	0.875 0.25 0.625	56.4 42.0 3.3	42.1 4.6 10.4	335	1.0 0.0 0.899	49.2 66.0 -11.1	66.9 350.4
591	B59R_087_062_	0.875 0.25 0.75	0.875 0.625 0.562	341	0.723 0.25 0.875	53.2 34.6 -13.2	37.1 339.0	0.875 0.25 0.75	57.2 45.5 -6.6	46.0 351.7 13.3	316	0.756 0.0 1.0	42.1 55.4 -21.2	59.3 339.0
592	B50R_087_062_	0.875 0.25 0.875	0.875 0.625 0.562	330	0.615 0.25 0.875	51.0 29.2 -17.8	34.2 328.6	0.875 0.25 0.875	56.8 46.5 -14.0	48.6 343.1 18.6	305	0.584 0.0 1.0	38.5 46.7 -28.5	54.7 328.6
593	B42R_100_075_	0.875 0.25 1.0	1.0 0.75 0.625	321	0.566 0.25 1.0	50.4 30.0 -25.1	39.2 320.0	0.875 0.25 1.0	54.7 48.9 -16.3	51.6 341.5 21.2	294	0.421 0.0 1.0	35.3 40.1 -33.5	52.3 320.0
594	R41Y_087_087_	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.223 0.0	54.2 36.1 48.5	60.5 53.3	0.875 0.375 0.0	63.1 24.3 58.4	63.2 67.4 17.8	44	1.0 0.255 0.0	58.5 41.3 55.4	69.1 53.3
595	R31Y_087_075_	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.257 0.125	55.9 36.8 39.0	53.7 46.6	0.875 0.375 0.125	61.5 24.8 49.6	55.5 63.3 16.9	39	1.0 0.177 0.0	54.6 49.1 52.0	71.6 46.6
596	R18Y_087_062_	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.288 0.25	58.0 35.0 27.1	44.3 37.7	0.875 0.375 0.25	62.3 24.9 36.0	43.8 55.3 14.1	33	1.0 0.06 0.0	49.7 56.0 43.3	70.8 37.7
597	R00Y_087_050_	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.506	62.7 28.0 13.3	31.0 25.4	0.875 0.375 0.375	62.9 25.1 26.0	36.2 45.9 12.9	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
598	R26Y_087_050_	0.875 0.375 0.5	0.875 0.5 0.625	376	0.875 0.375 0.625	62.8 29.5 5.1	29.9 9.8	0.875 0.375 0.5	63.1 27.5 16.1	31.9 30.3 11.1	359	1.0 0.0 0.501	47.8 59.0 10.2	59.9 9.8
599	R00Y_087_050_	0.875 0.375 0.625	0.875 0.5 0.625	360	0.875 0.375 0.788	63.6 32.7 -4.5	33.1 352.0	0.875 0.375 0.625	63.0 30.1 6.9	30.9 12.9 11.8	339	1.0 0.0 0.827	49.4 65.5 -9.1	66.2 352.0
600	B61R_087_050_	0.875 0.375 0.75	0.875 0.5 0.625	344	0.787 0.375 0.875	61.0 29.1 -9.5	30.6 341.8	0.875 0.375 0.75	63.3 35.2 -4.9	35.6 352.0 8.0	320	0.825 0.0 1.0	44.1 58.2 -19.0	61.2 341.8
601	B50R_087_050_	0.875 0.375 0.875	0.875 0.5 0.625	330	0.667 0.375 0.875	58.1 23.3 -14.2	27.3 328.6	0.875 0.375 0.875	63.7 36.7 -11.8	38.6 342.1 14.6	305	0.584 0.0 1.0	38.5 46.7 -28.5	54.7 328.6
602	B40R_100_062_	0.875 0.375 1.0	1.0 0.625 0.687	319	0.617 0.375 1.0	57.5 24.2 -21.6	32.4 318.1	0.875 0.375 1.0	61.4 37.8 -15.8	41.0 337.2 15.3	292	0.387 0.0 1.0	34.5 38.7 -34.6	51.9 318.1
603	R58Y_087_087_	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.336 0.0	59.9 25.4 53.2	59.0 64.4	0.875 0.5 0.0	69.2 12.9 63.8	65.1 78.5 18.8	52	1.0 0.384 0.0	65.0 29.0 60.9	67.5 64.4
604	R50Y_087_075_	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.364 0.125	61.3 26.4 43.8	51.1 58.8	0.875 0.5 0.125	68.1 14.1 57.3	59.0 76.1 19.5	48	1.0 0.319 0.0	61.8 35.2 58.4	68.2 58.8
605	R38Y_087_062_	0.875 0.5 0.25	0.875 0.625 0.562	53	0.875 0.393 0.25	62.7 27.4 34.0	43.6 51.0	0.875 0.5 0.25	68.9 13.2 41.0	43.0 72.1 17.0	42	1.0 0.229 0.0	57.2 43.9 54.4	69.9 51.0
606	R23Y_087_050_	0.875 0.5 0.375	0.875 0.5 0.625	44	0.875 0.429 0.375	64.6 27.4 23.8	36.3 41.0	0.875 0.5 0.375	68.6 15.1 30.6	34.1 63.7 14.6	35	1.0 0.108 0.0	51.4 54.8 47.	

n	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb*Fe	LabCh*Fe	DE**Fe	hsi_Me	rgb*Me	LabCh*Me		
648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4	47.5 57.2 37.8	68.6 33.4 11.1	375	1.0 0.0 0.263	47.5 56.0 26.7		
649	R38Y_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.392	47.4 57.2 18.2	60.0 17.6	47.6 56.3 34.2	65.9 31.3 16.0	367	1.0 0.0 0.392	47.4 57.2 18.2		
650	R26Y_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.501	47.8 59.0 10.2	59.9 9.8	47.5 55.9 27.5	62.3 26.2 17.5	359	1.0 0.0 0.501	47.8 59.0 10.2		
651	R13Y_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.641	48.1 62.2 1.0	62.2 0.9	47.4 56.8 19.5	60.0 18.9 19.2	350	1.0 0.0 0.641	48.1 62.2 1.0		
652	R00Y_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.827	49.4 65.5 -9.1	66.2 352.0	47.8 58.9 10.4	59.9 10.0 20.7	339	1.0 0.0 0.827	49.4 65.5 -9.1		
653	B68R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.964	48.5 65.6 -12.2	66.7 349.4	48.0 61.8 2.1	61.8 1.9 14.8	331	1.0 0.0 0.964	48.5 65.6 -12.2		
654	B61R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	0.825 0.0 1.0	44.1 58.2 -19.0	61.2 341.8	49.3 64.5 -6.5	64.8 354.2 14.9	320	0.825 0.0 1.0	44.1 58.2 -19.0		
655	B55R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	0.696 0.0 1.0	40.6 52.3 -24.1	57.6 335.2	49.5 66.1 -10.7	67.0 350.7 21.1	312	0.696 0.0 1.0	40.6 52.3 -24.1		
656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	0.584 0.0 1.0	38.5 46.7 -28.5	54.7 328.6	49.0 65.4 -12.7	66.6 348.9 26.2	305	0.584 0.0 1.0	38.5 46.7 -28.5		
657	R11Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.0 0.012	47.5 57.1 37.5	68.3 33.2	51.9 54.3 49.2	73.2 42.1 12.8	389	1.0 0.0 0.012	47.5 57.1 37.5		
658	R00Y_100_087e	1.0 0.125 0.125	1.0 0.875 0.562	390	1.0 0.125 0.355	53.5 49.0 23.3	54.3 25.4	50.9 56.4 46.0	72.8 39.1 23.9	375	1.0 0.0 0.263	47.5 56.0 26.7		
659	R36Y_100_087e	1.0 0.125 0.25	1.0 0.875 0.562	382	1.0 0.125 0.482	53.5 50.0 14.9	52.5 16.5	51.0 57.3 36.9	68.2 32.7 23.1	366	1.0 0.0 0.408	47.5 57.5 17.1		
660	R23Y_100_087e	1.0 0.125 0.375	1.0 0.875 0.562	374	1.0 0.125 0.594	53.8 52.4 7.0	52.9 7.6	50.9 59.2 26.9	65.0 24.4 21.1	357	1.0 0.0 0.536	47.8 59.9 8.0		
661	R08Y_100_087e	1.0 0.125 0.5	1.0 0.875 0.562	365	1.0 0.125 0.733	54.6 55.5 -2.2	55.6 357.6	51.0 57.5 17.6	63.0 16.2 20.7	347	1.0 0.0 0.695	48.7 63.4 -2.6		
662	B70R_100_087e	1.0 0.125 0.625	1.0 0.875 0.562	355	1.0 0.125 0.841	55.2 57.2 -7.7	57.8 352.3	51.1 64.7 5.0	64.9 4.4 15.3	339	1.0 0.0 0.818	49.4 65.4 -8.8		
663	B63R_100_087e	1.0 0.125 0.75	1.0 0.875 0.562	346	0.887 0.125 1.0	51.8 52.5 -15.2	54.6 343.7	51.5 67.6 -5.3	67.8 355.4 18.1	323	0.87 0.0 1.0	45.5 60.0 -17.4		
664	B56R_100_087e	1.0 0.125 0.875	1.0 0.875 0.562	338	0.746 0.125 1.0	47.8 46.4 -20.5	50.8 336.1	51.2 68.4 -9.9	69.1 351.7 24.7	313	0.71 0.0 1.0	41.0 53.1 -23.4		
665	B50R_100_087e	1.0 0.125 1.0	1.0 0.875 0.562	330	0.636 0.125 1.0	45.6 40.9 -24.9	47.9 328.6	51.0 65.2 -11.3	66.1 350.0 28.4	305	0.584 0.0 1.0	38.5 46.7 -28.5		
666	R23Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.108 0.0	51.4 54.8 47.7	72.6 41.0	58.2 41.8 55.1	69.2 52.8 16.4	35	1.0 0.108 0.0	51.4 54.8 47.7		
667	R13Y_100_087e	1.0 0.25 0.125	1.0 0.875 0.562	38	1.0 0.136 0.125	54.0 49.8 34.1	60.4 34.3	42.8 53.7 68.7	51.4 21.2 30	30	1.0 0.012 0.0	48.0 57.0 39.0		
668	R00Y_100_075e	1.0 0.25 0.25	1.0 0.75 0.625	390	1.0 0.25 0.447	59.6 42.0 20.0	46.5 25.4	58.2 43.1 44.9	62.3 46.1 24.9	375	1.0 0.0 0.263	47.5 56.0 26.7		
669	R35Y_100_075e	1.0 0.25 0.375	1.0 0.75 0.625	381	1.0 0.25 0.567	59.6 43.3 11.9	45.0 15.4	57.6 45.9 33.2	56.7 35.9 21.5	365	1.0 0.0 0.423	47.5 57.8 15.9		
670	R18Y_100_075e	1.0 0.25 0.5	1.0 0.75 0.625	371	1.0 0.25 0.691	59.9 45.8 3.4	45.9 4.3	56.8 49.3 23.4	54.6 25.4 20.5	354	1.0 0.0 0.588	47.9 61.1 4.6		
671	R00Y_100_075e	1.0 0.25 0.625	1.0 0.75 0.625	360	1.0 0.25 0.87	61.0 49.1 -6.8	49.6 352.0	57.2 52.2 10.3	53.2 11.2 17.9	339	1.0 0.0 0.827	49.4 65.5 -9.1		
672	B65R_100_075e	1.0 0.25 0.75	1.0 0.75 0.625	349	0.956 0.25 1.0	59.2 47.2 -11.2	48.5 346.6	54.5 -1.8	54.5 358.0 11.8	327	0.941 0.0 1.0	47.0 63.0 -14.9		
673	B57R_100_075e	1.0 0.25 0.875	1.0 0.75 0.625	339	0.793 0.25 1.0	54.9 40.4 -17.0	43.8 337.1	51.0 57.5 0.875	68.8 56.4 -9.7	314	0.725 0.0 1.0	41.3 53.8 -22.7		
674	B50R_100_075e	1.0 0.25 1.0	1.0 0.75 0.625	330	0.688 0.25 1.0	52.8 35.0 -21.4	41.0 32.6	59.2 53.0 -11.1	54.2 348.1 21.6	305	0.584 0.0 1.0	38.5 46.7 -28.5		
675	R36Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.216 0.0	56.5 45.2 53.8	70.3 49.9	0.375 0.0	64.6 29.8 60.4	67.3 63.7 18.5	41	1.0 0.216 0.0	56.5 45.2 53.8	
676	R26Y_100_087e	1.0 0.375 0.125	1.0 0.875 0.562	46	1.0 0.245 0.125	58.0 46.4 43.7	63.7 43.3	0.375 0.125	63.2 32.5 57.2	65.9 60.3 20.0	37	1.0 0.138 0.0	52.6 53.0 49.9	
677	R15Y_100_075e	1.0 0.375 0.25	1.0 0.75 0.625	39	1.0 0.271 0.25	60.4 42.5 30.3	52.2 35.5	0.375 0.25	63.3 33.3 49.0	59.3 55.8 21.0	31	1.0 0.028 0.0	48.6 56.7 40.4	
678	R00Y_100_062a	1.0 0.375 0.375	1.0 0.625 0.687	390	1.0 0.375 0.539	65.6 35.0 16.7	38.8 25.4	0.375 0.375	64.5 33.3 35.8	48.9 47.0 19.2	375	1.0 0.0 0.263	47.5 56.0 26.7	
679	R31Y_100_062a	1.0 0.375 0.5	1.0 0.625 0.687	379	1.0 0.375 0.659	65.7 36.4 8.5	37.4 13.2	0.375 0.5	64.4 35.7 25.6	44.0 35.7 17.1	362	1.0 0.0 0.454	47.6 58.3 13.7	
680	R11Y_100_062a	1.0 0.375 0.625	1.0 0.625 0.687	367	1.0 0.375 0.787	66.1 39.1 0.0	39.1 359.8	0.375 0.625	64.1 39.5 13.1	41.6 18.3 13.3	349	1.0 0.0 0.659	48.3 62.6 -0.1	
681	B69R_100_062a	1.0 0.375 0.75	1.0 0.625 0.687	353	1.0 0.375 0.937	66.7 41.2 -6.9	41.8 350.4	0.375 0.75	64.9 42.1 1.3	42.1 1.8 8.5	335	1.0 0.0 0.899	49.2 66.0 -11.1	
682	B59R_100_062a	1.0 0.375 0.875	1.0 0.625 0.687	341	0.848 0.375 1.0	62.2 34.6 -13.2	37.1 339.0	0.375 0.875	65.6 44.6 -7.7	45.2 350.1 11.8	316	0.756 0.0 1.0	42.1 55.4 -21.2	
683	B50R_100_062a	1.0 0.375 1.0	1.0 0.625 0.687	330	0.74 0.375 0.0	60.0 29.2 -17.8	34.2 328.6	0.375 1.0	66.3 43.9 -9.3	44.9 348.0 18.1	305	0.584 0.0 1.0	38.5 46.7 -28.5	
684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.319 0.0	61.8 35.2 58.4	68.2 58.8	0.5 0.0	70.5 19.2 66.2	69.0 73.8 19.8	48	1.0 0.319 0.0	61.8 35.2 58.4	
685	R41Y_100_087e	1.0 0.5 0.125	1.0 0.875 0.562	55	1.0 0.348 0.125	63.2 36.1 48.5	60.5 53.3	0.5 0.125	68.8 22.5 62.1	66.1 70.0 20.0	44	1.0 0.255 0.0	58.5 41.3 55.4	
686	R31Y_100_075e	1.0 0.5 0.25	1.0 0.75 0.625	49	1.0 0.382 0.25	64.9 36.8 39.0	53.7 46.6	0.5 0.25	69.3 22.8 50.9	55.8 65.8 18.8	39	1.0 0.177 0.0	54.6 49.1 52.0	
687	R18Y_100_062a	1.0 0.5 0.375	1.0 0.625 0.687	41	1.0 0.413 0.375	67.0 35.0 27.1	44.3 37.7	0.5 0.375	69.7 24.5 38.1	45.3 57.2 15.5	33	1.0 0.06 0.0	49.7 56.0 43.3	
688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.631	71.6 28.0 13.3	31.0 25.4	1.0 0.5	71.4 24.0 27.4	36.4 48.8 14.6	375	1.0 0.0 0.263	47.5 56.0 26.7	
689	R26Y_100_050a	1.0 0.5 0.625	1.0 0.5 0.75	376	1.0 0.5 0.75	71.6 29.5 5.1	29.9 9.8	1.0 0.5 0.625	71.6 27.0 15.7	31.2 30.2 10.8	359	1.0 0.0 0.501	47.8 59.0 10.2	
690	R00Y_100_050a	1.0 0.5 0.75	1.0 0.5 0.75	360	1.0 0.5 0.913	72.6 32.7 -4.5	33.1 352.0	1.0 0.5 0.75	71.9 30.4 47	30.7 8.8 9.6	339	1.0 0.0 0.827	49.4 65.5 -9.1	
691	B61R_100_050a	1.0 0.5 0.875	1.0 0.5 0.75	344	0.912 0.5 1.0	70.0 29.1 -9.5	30.6 341.8	1.0 0.5 0.875	71.8 34.7 -5.9	35.2 350.3 6.9	320	0.825 0.0 1.0	44.1 58.2 -19.0	
692	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	0.792 0.5 1.0	67.1 23.3 -14.2	27.3 328.6	1.0 0.5 1.0	71.6 36.1 -8.9	37.2 346.1 14.5	305	0.584 0.0 1.0	38.5 46.7 -28.5	
693	R63Y_100_100a	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.425 0.0	68.0 25.7 63.0	68.0 67.8	1.0 0.625	0.0 74.9 11.4	70.7 71.6 80.7	18.0 55	1.0 0.425 0.0	67.0 25.7 63.0	
694	R58Y_100_087e	1.0 0.625 0.125	1.0 0.875 0.562	65	1.0 0.461 0.125	68.9 25.4 53.2	59.0 64.4	1.0 0.625 0.125	75.7 10.3 67.8	68.6 81.3 22.0	52	1.0 0.384 0.0	65.0 29.0 60.9	
695	R50Y_100_075e	1.0 0.625 0.25	1.0 0.75 0.625	60	1.0 0.489 0.25	70.3 26.4 43.8	51.1 58.8	1.0 0.625 0.25	75.9 11.1 56.5	57.6 78.8 20.6	48	1.0 0.319 0.0	61.8 35.2 58.4	
696	R38Y_100_062a	1.0 0.625 0.375	1.0 0.625 0.687	53	1.0 0.518 0.375	71.7 27.4 34.0	43.6 51.0	1.0 0.625 0.375	76.0 13.2 42.5	44.6 72.7 17.1	42	1.0 0.229 0.0	57.2 43.9 54.4	
697	R23Y_100_050a	1.0 0.625 0.5	1.0 0.5 0.75	44	1.0 0.554 0.5	73.6 27.4 23.8	36.3 41.0	1.0 0.625 0.5	77.1 14.1 31.2	34.3 65.5 15.5	35	1.0 0.108 0.0	51.4 54.8 47.7	
698	R00Y_100_037e	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.723	77.7 21.0 10.0	23.2 25.4	1.0 0.625 0.625	78.4 15.6 19.6	25.1 51.4 11.0	375	1.0 0.0 0.263	47.5 56.0 26.7	
699	R18Y_100_037e	1.0 0.625 0.75	1.0 0.375 0.812	371	1.0 0.625 0.845	77.8 22.9 1.7	22.9 4.3	1.0 0.625 0.75	78.6 19.2 8.4	21.0 23.7 7.7	354	1.0 0.0 0.588	47.9 61.1 4.6	
700	B65R_100_037e	1.0 0.625 0.875	1.0 0.375 0.812	349	0.978 0.625 1.0	77.5 23.6 -5.6	24.2 346.6	1.0 0.625 0.875	79.4 23.4 -4.3	23.8 349.5 2.2	327	0.941 0.0 1.0	47.0 63.0 -14.9	
701	B50R_100_037e	1.0 0.625 1.0	1.0 0.375 0.812	330	0.844 0.625 1.0	74.3 17.5 -10.7	20.5 328.6	1.0 0.625 1.0	78.5 25.6 -7.2	26.6 344.3 9.8	305	0.584 0.0 1.0	38.5 46.7 -28.5	
702	R76Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.551 0.0	72.3 16.1 68.2	70.1 76.7	1.0 0.75 0.0	82.9 -2.0	76.9 77.0	91.5 22.7	63	1.0 0.551 0.0	72.3 16.1 68.2
703	R73Y_100_087e	1.0 0.75 0.125	1.0 0.875 0.562	74	1.0 0.572 0.125	74.0 16.2 58.4	60.6 74.4	1.0 0.75 0.125	82.6 -0.5	66.7 66.7	90.5 20.6	60	1.0 0.511 0.0	70.9 18.5 66.7
704	R68Y_100_075e	1.0 0.75 0												

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.L0NA.TXT> / .PS
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb*Fe	LabCh*Fe	DE**Fe	hsiMe	rgb*Me	LabCh*Me		
729	NW_100e	1.0 1.0 1.0	1.0 0.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	96.1 0.0 0.0	178.6 0.2 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
730	G50B_100_012e	0.875 1.0 1.0	1.0 0.125 0.937	210	0.875 1.0 0.973	90.7 -4.8 -3.6	6.0 216.9	0.875 1.0 1.0	90.5 -5.7 -7.6	9.6 232.8 4.1	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
731	G50B_100_025e	0.75 1.0 1.0	1.0 0.25 0.875	210	0.75 1.0 0.947	85.6 -9.6 -7.2	12.1 216.9	0.75 1.0 1.0	84.8 -9.9 -14.7	17.7 236.0 7.4	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
732	G50B_100_037e	0.625 1.0 1.0	1.0 0.375 0.812	210	0.625 1.0 0.921	80.5 -14.5 -10.9	18.1 216.9	0.625 1.0 1.0	78.6 -14.0 -21.7	25.8 237.2 11.0	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
733	G50B_100_050e	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 0.895	75.4 -19.3 -14.5	24.2 216.9	0.5 1.0 1.0	73.1 -16.9 -27.8	32.5 238.7 13.6	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
734	G50B_100_062e	0.375 1.0 1.0	1.0 0.625 0.687	210	0.375 1.0 0.869	70.3 -24.1 -18.2	30.2 216.9	0.375 1.0 1.0	67.3 -20.0 -32.8	38.5 238.6 15.5	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
735	G50B_100_075e	0.25 1.0 1.0	1.0 0.75 0.625	210	0.25 1.0 0.843	65.1 -29.0 -21.8	36.3 216.9	0.25 1.0 1.0	59.2 -26.0 -38.7	46.6 236.0 18.1	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
736	G50B_100_087e	0.125 1.0 1.0	1.0 0.875 0.562	210	0.125 1.0 0.817	60.0 -33.8 -25.5	42.3 216.9	0.125 1.0 1.0	54.7 -28.9 -42.5	51.4 235.7 18.5	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
737	G50B_100_100e	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9	0.0 1.0 1.0	52.2 -29.2 -44.1	52.9 236.4 17.9	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
738	RO0Y_100_012e	1.0 0.875 0.875	1.0 0.125 0.937	390	1.0 0.875 0.907	89.8 7.0 3.3	7.7 25.4	1.0 0.875 0.875	91.5 4.1 3.9	5.7 43.0 3.3	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
739	NW_087e	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.8 0.0 0.0	0.0 0.0	0.875 0.875 0.875	90.3 0.0 -0.2	0.2 277.5 3.5	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0
740	G50B_087_012e	0.75 0.875 0.875	0.875 0.125 0.812	210	0.75 0.875 0.848	81.7 -4.8 -3.6	6.0 216.9	0.75 0.875 0.875	85.5 -7.5 -10.5	12.9 234.5 8.3	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
741	G50B_087_025e	0.625 0.875 0.875	0.875 0.25 0.75	210	0.625 0.875 0.822	76.6 -9.6 -7.2	12.1 216.9	0.625 0.875 0.875	80.6 -12.5 -18.7	22.5 236.2 12.5	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
742	G50B_087_037e	0.5 0.875 0.875	0.875 0.375 0.687	210	0.5 0.875 0.796	71.5 -14.5 -10.9	18.1 216.9	0.5 0.875 0.875	73.7 -17.3 -25.2	30.7 235.4 14.8	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
743	G50B_087_050e	0.375 0.875 0.875	0.875 0.5 0.625	210	0.375 0.875 0.77	66.4 -19.3 -14.5	24.2 216.9	0.375 0.875 0.875	68.3 -20.0 -31.6	37.4 237.7 17.2	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
744	G50B_087_062e	0.25 0.875 0.875	0.875 0.625 0.562	210	0.25 0.875 0.744	61.3 -24.1 -18.2	30.2 216.9	0.25 0.875 0.875	63.0 -23.7 -34.2	41.7 235.2 16.1	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
745	G50B_087_075e	0.125 0.875 0.875	0.875 0.75 0.5	210	0.125 0.875 0.718	56.1 -29.0 -21.8	36.3 216.9	0.125 0.875 0.875	57.0 -28.4 -39.3	48.5 234.1 17.5	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
746	G50B_087_087e	0.0 0.875 0.875	0.875 0.875 0.437	210	0.0 0.875 0.692	51.0 -33.8 -25.5	42.3 216.9	0.0 0.875 0.875	52.0 -31.5 -42.1	52.6 233.1 16.8	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
747	RO0Y_100_025e	1.0 0.75 0.75	1.0 0.25 0.875	390	1.0 0.75 0.815	83.7 14.0 6.6	15.5 25.4	1.0 0.75 0.75	86.6 8.0 10.0	12.8 51.3 7.4	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
748	RO0Y_087_012e	0.875 0.75 0.75	0.75 0.875 0.125	812	0.875 0.75 0.782	80.8 7.0 3.3	7.7 25.4	0.875 0.75 0.75	83.1 5.7 3.9	7.0 34.1 2.7	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
749	NW_075e	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0	0.75 0.75 0.75	80.9 0.0 -0.5	0.5 269.9 3.1	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0
750	G50B_075_012e	0.625 0.75 0.75	0.75 0.125 0.687	210	0.625 0.75 0.723	72.7 -4.8 -3.6	6.0 216.9	0.625 0.75 0.75	77.5 -6.9 -10.7	12.8 237.0 8.8	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
751	G50B_075_025e	0.5 0.75 0.75	0.75 0.25 0.625	210	0.5 0.75 0.697	67.6 -9.6 -7.2	12.1 216.9	0.5 0.75 0.75	71.5 -14.0 -19.5	24.0 234.3 13.5	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
752	G50B_075_037e	0.375 0.75 0.75	0.75 0.375 0.562	210	0.375 0.75 0.671	62.5 -14.5 -10.9	18.1 216.9	0.375 0.75 0.75	66.6 -17.0 -24.9	30.2 235.6 14.8	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
753	G50B_075_050e	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.645	57.4 -19.3 -14.5	24.2 216.9	0.25 0.75 0.75	60.0 -20.0 -30.4	36.4 236.6 16.1	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
754	G50B_075_062e	0.125 0.75 0.75	0.75 0.625 0.437	210	0.125 0.75 0.619	52.3 -24.1 -18.2	30.2 216.9	0.125 0.75 0.75	55.7 -25.2 -35.3	43.4 234.5 17.5	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
755	G50B_075_075e	0.0 0.75 0.75	0.75 0.75 0.375	210	0.0 0.75 0.593	47.1 -29.0 -21.8	36.3 216.9	0.0 0.75 0.75	52.0 -33.9 -38.5	51.3 228.5 18.0	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
756	RO0Y_100_037e	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.723	77.7 21.0 10.0	23.2 25.4	1.0 0.625 0.625	79.0 15.4 17.7	23.4 48.9 9.6	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
757	RO0Y_087_025e	0.875 0.625 0.625	0.875 0.25 0.75	390	0.875 0.625 0.69	74.7 14.0 6.6	15.5 25.4	0.875 0.625 0.625	76.0 9.9 12.8	16.2 52.1 7.4	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
758	RO0Y_075_012e	0.75 0.625 0.625	0.75 0.125 0.687	390	0.75 0.625 0.657	71.8 7.0 3.3	7.7 25.4	0.75 0.625 0.625	74.8 5.4 4.1	6.8 37.0 3.4	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
759	NW_062e	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.8 0.0 0.0	0.0 0.0	0.625 0.625 0.625	71.8 0.0 -1.0	1.0 270.7 3.1	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0
760	G50B_062_012e	0.5 0.625 0.625	0.625 0.125 0.562	210	0.5 0.625 0.598	63.7 -4.8 -3.6	6.0 216.9	0.5 0.625 0.625	67.1 -7.7 -13.2	15.3 239.7 10.5	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
761	G50B_062_025e	0.375 0.625 0.625	0.625 0.25 0.5	210	0.375 0.625 0.572	58.6 -9.6 -7.2	12.1 216.9	0.375 0.625 0.625	61.5 -13.9 -20.3	24.7 235.6 14.0	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
762	G50B_062_037e	0.25 0.625 0.625	0.625 0.375 0.437	210	0.25 0.625 0.546	53.5 -14.5 -10.9	18.1 216.9	0.25 0.625 0.625	55.0 -17.0 -27.0	31.9 237.6 16.3	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
763	G50B_062_050e	0.125 0.625 0.625	0.625 0.5 0.375	210	0.125 0.625 0.52	48.4 -19.3 -14.5	24.2 216.9	0.125 0.625 0.625	51.6 -22.0 -31.5	38.5 234.9 17.4	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
764	G50B_062_062e	0.0 0.625 0.625	0.625 0.625 0.312	210	0.0 0.625 0.494	43.3 -24.1 -18.2	30.2 216.9	0.0 0.625 0.625	49.9 -30.2 -39.0	49.4 232.3 22.7	198	0.0 1.0 0.791	54.9 -38.7 -29.1	48.4 216.9
765	RO0Y_100_050e	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.631	71.6 28.0 13.3	31.0 25.4	1.0 0.5 0.5	72.0 23.9 24.2	34.0 45.3 11.6	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
766	RO0Y_087_037e	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.598	68.7 21.0 10.0	23.2 25.4	0.875 0.5 0.5	70.4 16.5 16.7	23.5 45.4 8.2	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
767	RO0Y_075_025e	0.75 0.5 0.5	0.75 0.25 0.625	390	0.75 0.5 0.565	65.7 14.0 6.6	15.5 25.4	0.75 0.5 0.5	66.8 10.9 11.1	15.6 45.6 5.5	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
768	RO0Y_062_012e	0.625 0.5 0.5	0.625 0.125 0.562	390	0.625 0.5 0.532	62.8 7.0 3.3	7.7 25.4	0.625 0.5 0.5	64.1 6.0 2.5	6.5 22.6 1.8	375	1.0 0.0 0.263	47.5 56.0 26.7	62.1 25.4
769	NW_050e	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	59.8 0.0 0.0	0.0 0.0	0.5 0.5 0.5	61.1 0.0 -1.1	1.1 268.7 1.7	360	1.0 1.0 1.0	95.8 0.0 0.0	0.

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)
TUB material: code=rh4ta

n	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb**Fe	LabCh**Fe	DE**Fe	hsi_Me	rgb*Me	LabCh*Me		
810	NW_100e	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	96.1 -0.1 0.0	0.1 188.0	0.3 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
811	B00R_100_012e	0.875 0.875 1.0	1.0 0.125 0.937	270	0.875 0.907 1.0	88.5 0.1 -6.0	0.0 0.0 0.0	0.875 0.875 1.0	88.8 0.5 -7.8	7.8 273.8	1.8 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
812	B00R_100_025e	0.75 0.75 1.0	1.0 0.25 0.875	270	0.75 0.815 1.0	81.2 0.3 -12.1	12.1 271.7	0.75 0.75 1.0	79.2 2.5 -15.9	16.1 278.9	4.8 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
813	B00R_100_037e	0.625 0.625 1.0	1.0 0.375 0.812	270	0.625 0.722 1.0	73.8 0.5 -18.2	18.2 271.7	0.625 0.625 1.0	65.8 5.5 -25.7	26.3 282.2	12.0 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
814	B00R_100_050e	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.63 1.0	66.5 0.7 -24.3	24.3 271.7	0.5 0.5 1.0	54.6 10.9 -32.6	34.4 288.5	17.8 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
815	B00R_100_062e	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.538 1.0	59.2 0.9 -30.4	30.4 271.7	0.375 0.375 1.0	45.7 14.0 -37.9	40.4 290.3	20.2 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
816	B00R_100_075e	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.445 1.0	51.9 1.1 -36.5	36.5 271.7	0.25 0.25 1.0	38.1 18.6 -41.2	45.2 294.2	22.7 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
817	B00R_100_087e	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.353 1.0	44.6 1.2 -42.6	42.6 271.7	0.125 0.125 1.0	34.5 19.2 -42.9	47.0 294.1	20.6 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
818	B00R_100_100e	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7	0.0 0.0 1.0	30.8 20.2 -44.2	48.6 294.6	20.3 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
819	Y00G_100_012e	1.0 1.0 0.875	1.0 0.125 0.937	90	1.0 0.971 0.875	94.3 -0.3 9.6	9.6 9.6	1.0 1.0 0.875	95.9 -3.8 9.4	10.2 111.9	3.8 77	1.0 0.768 0.0	83.6 -3.1 76.8	76.9 92.3
820	NW_087e	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.8 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	91.1 0.0 -0.1	0.1 242.5	4.3 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
821	B00R_087_012e	0.75 0.75 0.875	0.875 0.125 0.812	270	0.75 0.782 0.875	79.5 0.1 -6.0	6.0 0.0	0.75 0.75 0.875	80.2 1.7 -11.6	11.7 278.4	5.8 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
822	B00R_087_025e	0.625 0.625 0.875	0.875 0.25 0.75	270	0.625 0.69 0.875	72.2 0.3 -12.1	12.1 271.7	0.625 0.625 0.875	69.8 4.2 -20.4	20.8 281.7	9.3 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
823	B00R_087_037e	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.597 0.875	64.8 0.5 -18.2	18.2 271.7	0.5 0.5 0.875	58.7 8.3 -29.7	30.9 285.6	15.2 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
824	B00R_087_050e	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.505 0.875	57.5 0.7 -24.3	24.3 271.7	0.375 0.375 0.875	47.8 12.6 -36.7	38.8 288.9	19.7 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
825	B00R_087_062e	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.413 0.875	50.2 0.9 -30.4	30.4 271.7	0.25 0.25 0.875	39.6 16.7 -41.1	44.4 292.2	21.9 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
826	B00R_087_075e	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.32 0.875	42.9 1.1 -36.5	36.5 271.7	0.125 0.125 0.875	35.1 17.5 -43.1	46.5 292.1	19.3 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
827	B00R_087_087e	0.0 0.0 0.875	0.875 0.875 0.437	270	0.0 0.228 0.875	35.6 1.2 -42.6	42.6 271.7	0.0 0.0 0.875	29.1 21.9 -44.4	49.5 296.2	21.7 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
828	Y00G_100_025e	1.0 1.0 0.75	1.0 0.25 0.875	90	1.0 0.942 0.75	92.7 -0.7 19.2	19.2 92.3	1.0 1.0 0.75	95.2 -8.7 24.8	26.3 109.2	10.0 77	1.0 0.768 0.0	83.6 -3.1 76.8	76.9 92.3
829	Y00G_087_012e	0.875 0.875 0.75	0.875 0.125 0.812	90	0.875 0.846 0.75	85.3 -0.3 9.6	9.6 9.6	0.875 0.875 0.75	91.4 -3.2 9.1	9.6 109.3	6.7 77	1.0 0.768 0.0	83.6 -3.1 76.8	76.9 92.3
830	NW_075e	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	82.2 0.0 -0.5	0.5 270.3	4.4 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
831	B00R_075_012e	0.625 0.625 0.75	0.75 0.125 0.687	270	0.625 0.657 0.75	70.5 0.1 -6.0	6.0 0.0	0.625 0.625 0.75	71.5 2.2 -12.7	12.9 280.2	7.0 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
832	B00R_075_025e	0.5 0.5 0.75	0.75 0.25 0.625	270	0.5 0.565 0.75	63.2 0.3 -12.1	12.1 271.7	0.5 0.5 0.75	60.6 4.4 -20.2	20.7 282.3	9.4 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
833	B00R_075_037e	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.472 0.75	55.8 0.5 -18.2	18.2 271.7	0.375 0.375 0.75	50.8 6.5 -27.4	28.2 283.3	12.0 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
834	B00R_075_050e	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.38 0.75	48.5 0.7 -24.3	24.3 271.7	0.25 0.25 0.75	44.3 10.9 -33.2	35.0 288.2	14.2 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
835	B00R_075_062e	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.288 0.75	41.2 0.9 -30.4	30.4 271.7	0.125 0.125 0.75	38.2 13.3 -39.1	41.3 288.7	15.4 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
836	B00R_075_075e	0.0 0.0 0.75	0.75 0.75 0.375	270	0.0 0.195 0.75	33.9 1.1 -36.5	36.5 271.7	0.0 0.0 0.75	29.0 20.6 -44.7	49.2 294.7	21.7 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
837	Y00G_100_037e	1.0 1.0 0.625	1.0 0.375 0.812	90	1.0 0.913 0.625	91.2 -1.1 28.8	28.8 92.3	1.0 1.0 0.625	93.4 -9.0 33.9	35.0 104.9	9.6 77	1.0 0.768 0.0	83.6 -3.1 76.8	76.9 92.3
838	Y00G_087_025e	0.875 0.875 0.625	0.875 0.25 0.75	90	0.875 0.817 0.625	83.7 -0.7 19.2	19.2 92.3	0.875 0.875 0.625	90.1 -6.3 24.9	25.7 104.2	10.2 77	1.0 0.768 0.0	83.6 -3.1 76.8	76.9 92.3
839	Y00G_075_012e	0.75 0.75 0.625	0.75 0.125 0.687	90	0.75 0.721 0.625	76.3 -0.3 9.6	9.6 9.6	0.75 0.75 0.625	83.0 -2.2 7.7	8.1 106.0	7.2 77	1.0 0.768 0.0	83.6 -3.1 76.8	76.9 92.3
840	NW_062e	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.8 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	72.5 0.0 -1.0	1.0 270.5	3.8 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
841	B00R_062_012e	0.5 0.5 0.625	0.625 0.125 0.562	270	0.5 0.532 0.625	61.5 0.1 -6.0	6.0 0.0	0.5 0.5 0.625	61.7 2.6 -14.3	14.5 280.2	8.6 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
842	B00R_062_025e	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.44 0.625	54.2 0.3 -12.1	12.1 271.7	0.375 0.375 0.625	50.5 4.0 -22.7	23.0 280.0	11.7 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
843	B00R_062_037e	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.347 0.625	46.8 0.5 -18.2	18.2 271.7	0.25 0.25 0.625	44.0 8.2 -27.8	29.0 286.5	12.6 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
844	B00R_062_050e	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.255 0.625	39.5 0.7 -24.3	24.3 271.7	0.125 0.125 0.625	37.5 11.3 -34.2	36.1 288.3	14.7 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
845	B00R_062_062e	0.0 0.0 0.625	0.625 0.625 0.312	270	0.0 0.163 0.625	32.2 0.9 -30.4	30.4 271.7	0.0 0.0 0.625	28.0 21.5 -45.0	49.9 295.5	25.6 255	0.0 0.261 1.0	37.3 1.4 -48.6	48.7 271.7
846	Y00G_100_050e	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 0.884 0.5	89.7 -1.5 38.4	38.4 92.3	1.0 1.0 0.5	92.9 -11.3 44.8	46.2 104.1	12.1 77	1.0 0.768 0.0	83.6 -3.1 76.8	76.9 92.3
847	Y00G_087_037e	0.875 0.875 0.5	0.875 0.375 0.687	90	0.875 0.788 0.5	82.2 -1.1 28.8	28.8 92.3	0.875 0.875 0.5	90.0 -8.5 36.5	37.5 103.2	13.2 77	1.0 0.768 0.0	83.6 -3.1 76.8	76.9 92.3
848	Y00G_075_025e	0.75 0.75 0.5	0.75 0.25 0.625	90	0.75 0.692 0.5	74.7 -0.7 19.2	19.2 92.3	0.75 0.75 0.5	82.9 -6.7 24.9	25.8 105.2	11.6 77	1.0 0.768 0.0	83.6 -3.1 76.8	76.9 92.3
849	Y00G_062_012e	0.625 0.625 0.5	0.625 0.125 0.562	90	0.625 0.596 0.5	67.3 -0.3 9.6	9.6 9.6	0.625 0.625 0.5	73.9 -3.5 7.3	8.1 115.9	7.6 77	1.0 0.768 0.0	83.6 -3.1 76.8	76.9 92.3
850	NW_050e	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	59.8 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	61.9 0.0 -1.2	1.2 270.4	2.4 360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0
851	B00R_050_012e	0.375 0.375 0.5	0.5 0.125 0.437	270	0.0									

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS79/PS79L0NA.TXT /.PS
aplicación para la medida salida de impresora láser, separación cmykn6 (CMYK)
TUB material: code=rh4ta

n	HIC*Fe	rgb_Fe	ict_Fe	hsi_Fe	rgb*Fe	LabCh*Fe	rgb*Fe	LabCh*Fe	DE*Fe	hsi_Me	rgb*Me	LabCh*Me
891	NW_100e	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.8 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	96.1 -0.1 0.0	0.1 168.6 0.3	360	
892	B50R_100_012e	1.0 0.875 1.0	1.0 0.125 0.937	330	0.948 0.875 1.0	88.6 5.8 -3.5	6.8 328.6	1.0 0.875 1.0	91.4 7.6 -3.0	8.1 338.4 3.3	305	0.584 0.0 1.0
893	B50R_100_025e	1.0 0.75 1.0	1.0 0.25 0.875	330	0.896 0.75 1.0	81.5 11.6 -7.1	13.6 328.6	1.0 0.75 1.0	85.5 16.1 -5.2	16.9 341.9 6.2	305	0.584 0.0 1.0
894	B50R_100_037e	1.0 0.625 1.0	1.0 0.375 0.812	330	0.844 0.625 1.0	74.3 17.5 -10.7	20.5 328.6	1.0 0.625 1.0	78.4 26.0 -7.1	26.9 344.6 10.0	305	0.584 0.0 1.0
895	B50R_100_050e	1.0 0.5 1.0	1.0 0.5 0.75	330	0.792 0.5 1.0	67.1 23.3 -14.2	27.3 328.6	1.0 0.5 1.0	71.7 36.3 -8.9	37.4 346.1 14.7	305	0.584 0.0 1.0
896	B50R_100_062e	1.0 0.375 1.0	1.0 0.625 0.687	330	0.74 0.375 1.0	60.0 29.2 -17.8	34.2 328.6	1.0 0.375 1.0	66.1 44.6 -9.4	45.6 348.0 18.6	305	0.584 0.0 1.0
897	B50R_100_075e	1.0 0.25 1.0	1.0 0.75 0.625	330	0.688 0.25 1.0	52.8 35.0 -21.4	41.0 328.6	1.0 0.25 1.0	59.2 53.6 -11.4	54.8 347.9 22.0	305	0.584 0.0 1.0
898	B50R_100_087e	1.0 0.125 1.0	1.0 0.875 0.562	330	0.636 0.125 1.0	45.6 40.9 -24.9	47.9 328.6	1.0 0.125 1.0	52.3 65.2 -11.5	66.2 349.9 28.5	305	0.584 0.0 1.0
899	B50R_100_100e	1.0 0.0 1.0	1.0 1.0 0.5	330	0.584 0.0 1.0	38.5 46.7 -28.5	54.7 328.6	1.0 0.0 1.0	48.3 65.1 -12.7	66.3 348.9 26.1	305	0.584 0.0 1.0
900	GO0B_100_012e	0.875 1.0 0.875	1.0 0.125 0.937	150	0.875 1.0 0.893	90.5 -8.2 2.6	8.6 162.2	0.875 1.0 0.875	92.1 -6.9 1.8	7.1 165.1 2.1	157	0.0 1.0 1.0
901	NW_087e	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.8 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	91.6 0.0 -0.2	0.2 263.1 4.8	360	1.0 1.0 1.0
902	B50R_087_012e	0.875 0.75 0.875	0.875 0.125 0.812	330	0.823 0.75 0.875	79.6 5.8 -3.5	6.8 328.6	0.875 0.75 0.875	84.3 8.6 -5.6	10.2 326.9 5.8	305	0.584 0.0 1.0
903	B50R_087_025e	0.875 0.625 0.875	0.875 0.25 0.75	330	0.771 0.625 0.875	72.5 11.6 -7.1	13.6 328.6	0.875 0.625 0.875	77.2 18.7 -8.0	20.4 336.6 8.5	305	0.584 0.0 1.0
904	B50R_087_037e	0.875 0.5 0.875	0.875 0.375 0.687	330	0.719 0.5 0.875	65.3 17.5 -10.7	20.5 328.6	0.875 0.5 0.875	70.8 27.7 -10.0	29.5 340.0 11.6	305	0.584 0.0 1.0
905	B50R_087_050e	0.875 0.375 0.875	0.875 0.5 0.625	330	0.667 0.375 0.875	58.1 23.3 -14.2	27.3 328.6	0.875 0.375 0.875	64.0 37.0 -11.8	38.8 342.2 15.0	305	0.584 0.0 1.0
906	B50R_087_062e	0.875 0.25 0.875	0.875 0.625 0.562	330	0.615 0.25 0.875	51.0 29.2 -17.8	34.2 328.6	0.875 0.25 0.875	57.7 45.0 -13.6	47.1 341.3 17.7	305	0.584 0.0 1.0
907	B50R_087_075e	0.875 0.125 0.875	0.875 0.75 0.5	330	0.563 0.125 0.875	43.8 35.0 -21.4	41.0 328.6	0.875 0.125 0.875	49.8 55.9 -16.4	58.3 343.6 22.2	305	0.584 0.0 1.0
908	B50R_087_087e	0.875 0.0 0.875	0.875 0.875 0.437	330	0.511 0.0 0.875	36.6 40.9 -24.9	47.9 328.6	0.875 0.0 0.875	44.7 59.0 -15.8	61.1 344.9 21.8	305	0.584 0.0 1.0
909	GO0B_100_025e	0.75 1.0 0.75	1.0 0.25 0.875	150	0.75 1.0 0.786	85.3 -16.4 5.2	17.3 162.2	0.75 1.0 0.75	87.7 -14.1 8.3	16.4 149.3 4.6	157	0.0 1.0 1.0
910	GO0B_087_012e	0.75 0.875 0.75	0.875 0.125 0.812	150	0.75 0.875 0.768	81.5 -8.2 2.6	8.6 162.2	0.75 0.875 0.75	87.2 -9.9 2.7	10.3 164.7 5.9	157	0.0 1.0 1.0
911	NW_075e	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	82.6 0.0 -0.4	0.4 270.1 4.8	360	1.0 1.0 1.0
912	B50R_075_012e	0.75 0.625 0.75	0.75 0.125 0.687	330	0.698 0.625 0.75	70.6 5.8 -3.5	6.8 328.6	0.75 0.625 0.75	75.8 9.0 -5.9	10.8 326.6 6.5	305	0.584 0.0 1.0
913	B50R_075_025e	0.75 0.5 0.75	0.75 0.25 0.625	330	0.646 0.5 0.75	63.5 11.6 -7.1	13.6 328.6	0.75 0.5 0.75	68.2 19.3 -8.4	21.0 336.3 9.0	305	0.584 0.0 1.0
914	B50R_075_037e	0.75 0.375 0.75	0.75 0.375 0.562	330	0.594 0.375 0.75	56.3 17.5 -10.7	20.5 328.6	0.75 0.375 0.75	60.9 27.8 -9.9	29.5 340.2 11.3	305	0.584 0.0 1.0
915	B50R_075_050e	0.75 0.25 0.75	0.75 0.5 0.5	330	0.542 0.25 0.75	49.1 23.3 -14.2	27.3 328.6	0.75 0.25 0.75	54.4 37.2 -12.4	39.2 341.5 14.9	305	0.584 0.0 1.0
916	B50R_075_062e	0.75 0.125 0.75	0.75 0.625 0.437	330	0.49 0.125 0.75	42.0 29.2 -17.8	34.2 328.6	0.75 0.125 0.75	47.9 47.7 -13.2	49.5 344.5 19.9	305	0.584 0.0 1.0
917	B50R_075_075e	0.75 0.0 0.75	0.75 0.75 0.375	330	0.438 0.0 0.75	34.8 35.0 -21.4	41.0 328.6	0.75 0.0 0.75	41.5 53.5 -15.6	55.7 343.6 20.4	305	0.584 0.0 1.0
918	GO0B_100_037e	0.625 1.0 0.625	1.0 0.375 0.812	150	0.625 1.0 0.68	80.0 -24.7 7.9	25.9 162.2	0.625 1.0 0.625	81.7 -21.9 10.5	24.3 154.3 41.1	157	0.0 1.0 1.0
919	GO0B_087_025e	0.625 0.875 0.625	0.875 0.25 0.75	150	0.625 0.875 0.661	76.3 -16.4 5.2	17.3 162.2	0.625 0.875 0.625	82.1 -17.2 9.2	19.6 151.7 7.0	157	0.0 1.0 1.0
920	GO0B_075_012e	0.625 0.75 0.625	0.75 0.125 0.687	150	0.625 0.75 0.643	72.5 -8.2 2.6	8.6 162.2	0.625 0.75 0.625	77.8 -10.0 1.7	10.1 169.8 5.6	157	0.0 1.0 1.0
921	NW_062e	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.8 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	72.1 0.0 -0.9	0.9 268.7 3.4	360	1.0 1.0 1.0
922	B50R_062_012e	0.625 0.5 0.625	0.625 0.125 0.562	330	0.573 0.5 0.625	61.6 5.8 -3.5	6.8 328.6	0.625 0.5 0.625	65.6 9.6 -6.6	11.7 325.4 6.2	305	0.584 0.0 1.0
923	B50R_062_025e	0.625 0.375 0.625	0.625 0.25 0.5	330	0.521 0.375 0.625	54.5 11.6 -7.1	13.6 328.6	0.625 0.375 0.625	57.2 19.8 -8.9	21.7 335.8 8.8	305	0.584 0.0 1.0
924	B50R_062_037e	0.625 0.25 0.625	0.625 0.375 0.437	330	0.469 0.25 0.625	47.3 17.5 -10.7	20.5 328.6	0.625 0.25 0.625	50.0 29.9 -11.7	32.1 338.6 12.7	305	0.584 0.0 1.0
925	B50R_062_050e	0.625 0.125 0.625	0.625 0.5 0.375	330	0.417 0.125 0.625	40.1 23.3 -14.2	27.3 328.6	0.625 0.125 0.625	43.2 39.6 -12.4	41.5 342.5 16.6	305	0.584 0.0 1.0
926	B50R_062_062e	0.625 0.0 0.625	0.625 0.625 0.312	330	0.365 0.0 0.625	33.0 29.2 -17.8	34.2 328.6	0.625 0.0 0.625	37.3 48.6 -13.9	50.5 343.9 20.2	305	0.584 0.0 1.0
927	GO0B_100_050e	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.573	74.8 -32.9 10.5	34.6 162.2	0.5 1.0 0.5	75.2 -29.8 11.7	32.0 158.4 3.3	157	0.0 1.0 1.0
928	GO0B_087_037e	0.5 0.875 0.5	0.875 0.375 0.687	150	0.5 0.875 0.556	71.0 -24.7 7.9	25.9 162.2	0.5 0.875 0.5	74.3 -28.1 10.0	29.8 160.3 5.2	157	0.0 1.0 1.0
929	GO0B_075_025e	0.5 0.75 0.5	0.75 0.25 0.625	150	0.5 0.75 0.536	67.3 -16.4 5.2	17.3 162.2	0.5 0.75 0.5	71.8 -17.6 5.6	18.5 162.3 4.6	157	0.0 1.0 1.0
930	GO0B_062_012e	0.5 0.625 0.5	0.625 0.125 0.562	150	0.5 0.625 0.518	63.5 -8.2 2.6	8.6 162.2	0.5 0.625 0.5	68.1 -10.1 1.4	10.2 171.5 5.0	157	0.0 1.0 1.0
931	NW_050e	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	59.8 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	61.3 0.0 -1.1	1.1 271.0 1.9	360	1.0 1.0 1.0
932	B50R_050_012e	0.5 0.375 0.5	0.5 0.125 0.437	330	0.448 0.375 0.5	52.6 5.8 -3.5	6.8 328.6	0.5 0.375 0.5	53.3 10.5 -8.2	13.4 321.9 6.6	305	0.584 0.0 1.0
933	B50R_050_025e	0.5 0.25 0.5	0.5 0.25 0.375	330	0.396 0.249 0.5	45.5 11.6 -7.1	13.6 328.6	0.5 0.25 0.5	47.1 21.9 -10.7	24.4 333.8 10.9	305	0.584 0.0 1.0
934	B50R_050_037e	0.5 0.125 0.5	0.5 0.375 0.312	330	0.344 0.124 0.5	38.3 17.5 -10.7	20.5 328.6	0.5 0.125 0.5	40.5 32.1 -11.6	34.1 340.0 14.7	305	0.584 0.0 1.0
935	B50R_050_050e	0.5 0.0 0.5	0.5 0.5 0.25	330	0.292 0.0 0.5	31.1 23.3 -14.2	27.3 328.6	0.5 0.0 0.5	34.7 41.4 -11.9	43.1 343.9 18.5	305	0.584 0.0 1.0
936	GO0B_100_062e	0.375 1.0 0.375	1.0 0.625 0.687	150	0.375 1.0 0.466	69.5 -41.2 13.2	43.2 162.2	0.375 1.0 0.375	68.6 -39.6 17.6	43.4 156.0 4.7	157	0.0 1.0 1.0
937	GO0B_087_050e	0.375 0.875 0.375	0.875 0.5 0.625	150	0.375 0.875 0.448	65.8 -32.9 10.5	34.6 162.2	0.375 0.875 0.375	70.2 -36.7 16.1	40.2 156.2 8.1	157	0.0 1.0 1.0
938	GO0B_075_037e	0.375 0.75 0.375	0.75 0.375 0.562	150	0.375 0.75 0.43	62.0 -24.7 7.9	25.9 162.2	0.375 0.75 0.				

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información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

		HIC*Fe				rgb_Fe				ict_Fe				hsi_Fe				rgb*Fe				LabCh*Fe				rgb*Fe				LabCh*Fe				DE**Fe				hsiMe				rgb*Me				LabCh*Me																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																			
972	NW_000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS79/PS79.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fe	rgb_Fe		ict_Fe		hsi_F,e	rgb*Fe		LabCh*Fe						rgb**Fe		LabCh**Fe		DE**Fe		hsiM,e	rgb*Me		LabCh*Me										
1053	NW_086e	0.866	0.866	0.866	0.866	0.0	0.866	360	0.866	0.866	0.866	86.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1054	NW_093e	0.933	0.933	0.933	0.933	0.0	0.933	360	0.933	0.933	0.933	91.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1055	NW_100e	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1056	NW_000e	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	23.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1057	NW_006e	0.066	0.066	0.066	0.066	0.0	0.066	360	0.066	0.066	0.066	28.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1058	NW_013e	0.133	0.133	0.133	0.133	0.0	0.133	360	0.133	0.133	0.133	33.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1059	NW_020e	0.2	0.2	0.2	0.2	0.0	0.2	360	0.2	0.2	0.2	38.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1060	NW_026e	0.266	0.266	0.266	0.266	0.0	0.266	360	0.266	0.266	0.266	42.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1061	NW_033e	0.333	0.333	0.333	0.333	0.0	0.333	360	0.333	0.333	0.333	47.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1062	NW_040e	0.4	0.4	0.4	0.4	0.0	0.4	360	0.4	0.4	0.4	52.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1063	NW_046e	0.466	0.466	0.466	0.466	0.0	0.466	360	0.466	0.466	0.466	57.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1064	NW_053e	0.533	0.533	0.533	0.533	0.0	0.533	360	0.533	0.533	0.533	62.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1065	NW_060e	0.6	0.6	0.6	0.6	0.0	0.6	360	0.6	0.6	0.6	67.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1066	NW_066e	0.666	0.666	0.666	0.666	0.0	0.666	360	0.666	0.666	0.666	71.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1067	NW_073e	0.734	0.734	0.734	0.734	0.0	0.734	360	0.734	0.734	0.734	76.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1068	NW_080e	0.8	0.8	0.8	0.8	0.0	0.8	360	0.8	0.8	0.8	81.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1069	NW_086e	0.866	0.866	0.866	0.866	0.0	0.866	360	0.866	0.866	0.866	86.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1070	NW_093e	0.933	0.933	0.933	0.933	0.0	0.933	360	0.933	0.933	0.933	91.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1071	NW_100e	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1072	NW_000e	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	23.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1073	NW_100e	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.8	0.0	0.0	0.0	0.0					
1074	R00Y_100_100e	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.263	47.5	56.0	26.7	62.1	25.4	1.0	0.0	0.0	47.0	56.3	40.2	69.2	35.5	13.5	375	1.0	0.0	0.263	47.5	56.0	26.7	62.1	25.4
1075	G50B_100_100e	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	0.791	54.9	-38.7	-29.1	48.4	216.9	0.0	1.0	1.0	54.9	-30.4	-42.0	51.8	234.0	15.2	198	0.0	1.0	0.791	54.9	-38.7	-29.1	48.4	216.9
1076	Y00G_100_100e	1.0	1.0	0.0	1.0	1.0	0.5	90	1.0	0.768	0.0	83.6	-3.1	76.8	76.9	92.3	1.0	1.0	0.0	91.5	-16.0	86.1	87.6	100.5	17.8	77	1.0	0.768	0.0	83.6	-3.1	76.8	76.9	92.3
1077	B00R_100_100e	0.0	0.0	1.0	1.0	1.0	0.5	270	0.0	0.261	1.0	37.3	1.4	-48.6	48.7	271.7	0.0	0.0	1.0	30.7	21.3	-44.1	49.0	295.7	21.3	255	0.0	0.261	1.0	37.3	1.4	-48.6	48.7	271.7
1078	G00B_100_100e	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.146	53.8	-65.9	21.1	69.2	162.2	0.0	1.0	0.0	54.6	-69.2	33.1	76.7	154.3	12.4	157	0.0	1.0	0.146	53.8	-65.9	21.1	69.2	162.2
1079	B50R_100_100e	1.0	0.0	1.0	1.0	1.0	0.5	330	0.584	0.0	1.0	38.5	46.7	-28.5	54.7	328.6	1.0	0.0	1.0	48.3	66.3	-13.8	67.7	348.1	26.3	305	0.584	0.0	1.0	38.5	46.7	-28.5	54.7	328.6

delta E* = 6.3

gráfico TUB-PS79; círculo di tono; 16 y 8 pasos
colores y diferencia en color, ΔE^* , 3D=0, de=1, *cmYk*

entrada: *rgb/cmyk* -> *rgb_e*
salida: transfiera a *cmyk_e*