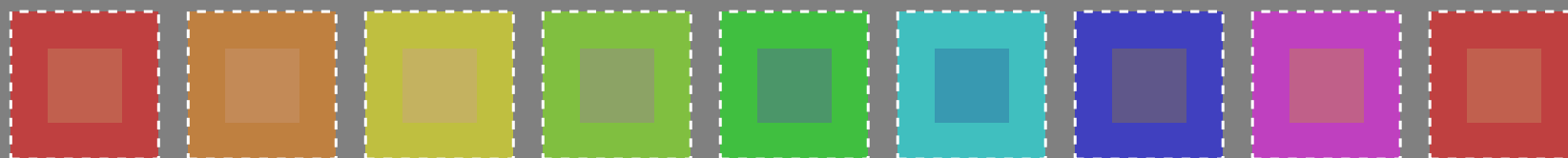
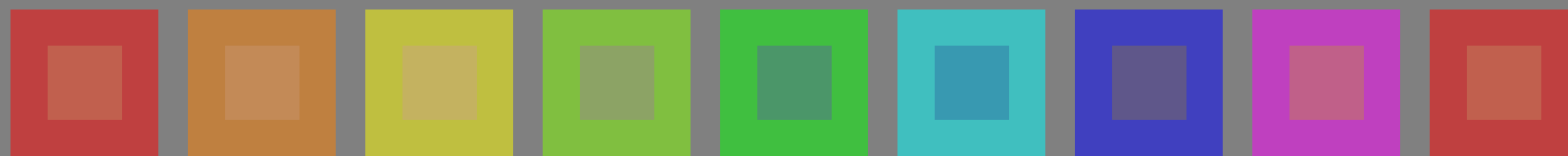


Test chart 3 for color rendering: metameric colours A and P4000



01: R00Y_075_050_ 02: R50Y_075_050_ 03: Y00G_075_050_ 04: Y50G_075_050_ 05: G00B_075_050_ 06: G50B_075_050_ 07: B00R_075_050_ 08: B50R_075_050_ 09: R00Y_075_050_



10: R00Y_075_050_ 11: R50Y_075_050_ 12: Y00G_075_050_ 13: Y50G_075_050_ 14: G00B_075_050_ 15: G50B_075_050_ 16: B00R_075_050_ 17: B50R_075_050_ 18: R00Y_075_050_

metameric colours only possible for offset print and colour printers
with at least four basic colours, either *CMYK* or *CMY0*

Series:
metameric
m
A

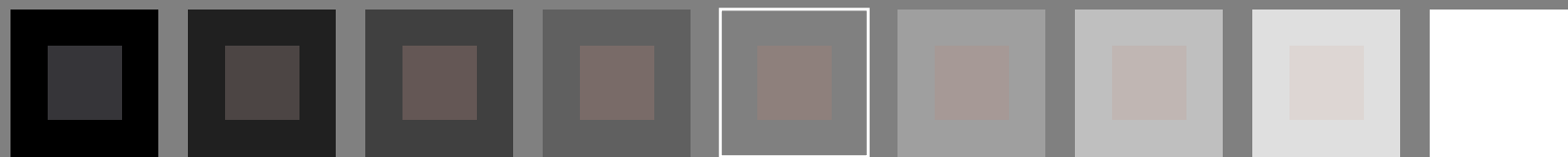
central
z
A/P4000

metameric
m
P4000



28: NW_000_ 29: NW_013_ 30: NW_025_ 31: NW_038_ 32: NW_050_ 33: NW_063_ 34: NW_075_ 35: NW_088_ 36: NW_100_

metameric
m
A



37: NW_000_ 38: NW_013_ 39: NW_025_ 40: NW_038_ 41: NW_050_ 42: NW_063_ 43: NW_075_ 44: NW_088_ 45: NW_100_

grey
g
A/P4000

metameric
m
P4000

metameric colours only possible for offset print and colour printers
with at least four basic colours, either *CMYK* or *CMY0*

2-003030-L0 PS310-7N

gráfico TUB-PS31; reproducción en color; sRGB

54 colores; metameric para A&P4000; tecnología de imagen salida: ningún cambio

entrada: *rgb/cmyk* -> *rgb/cmyk*

TUB matrícula: 20130201-PS31/PS31L0NP.PDF /.PS
aplicación para la medida de display output

TUB material: code=rha4ta

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS31/PS31L0NP.PDF /.PS
aplicación para la medida de display output, ninguna separación

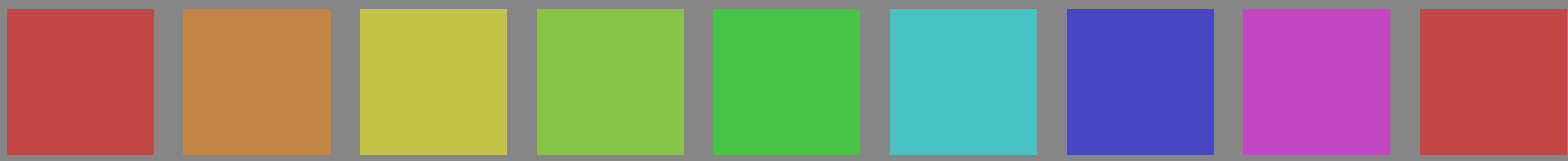
TUB material: code=rha4ta

Test chart 3 for color rendering: metameric colours A and P4000



01: R00Y_075_050_d 02: R50Y_075_050_d 03: Y00G_075_050_d 04: Y50G_075_050_d 05: G00B_075_050_d 06: G50B_075_050_d 07: B00R_075_050_d 08: B50R_075_050_d 09=10: R00Y_075_050_d

Series:
metameric
m
A

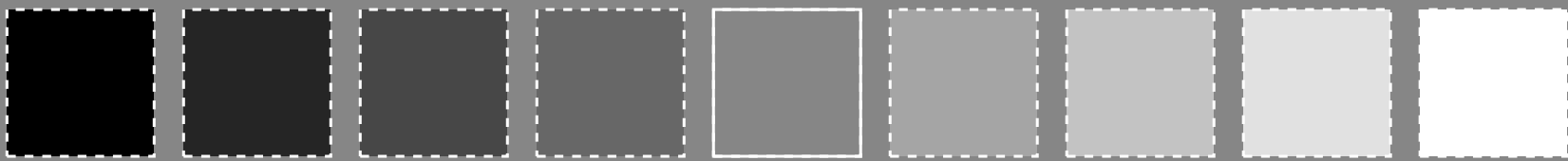


10: R00Y_075_050_d 11: R50Y_075_050_d 12: Y00G_075_050_d 13: Y50G_075_050_d 14: G00B_075_050_d 15: G50B_075_050_d 16: B00R_075_050_d 17: B50R_075_050_d 18=01: R00Y_075_050_d

central
z
A/P4000

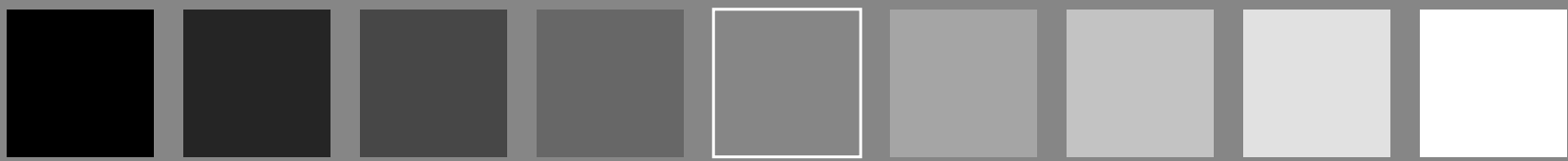
metameric colours only possible for offset print and colour printers
with at least four basic colours, either *CMYK* or *CMY0*

metameric
m
P4000



28: NW_000_d 29: NW_013_d 30: NW_025_d 31: NW_038_d 32: NW_050_d 33: NW_063_d 34: NW_075_d 35: NW_088_d 36=28: NW_100_d

metameric
m
A
AB*N=0.0, 0.0, 0.0
Lab*W=95.4, 0.0, 0.0



37: NW_000_d 38: NW_013_d 39: NW_025_d 40: NW_038_d 41: NW_050_d 42: NW_063_d 43: NW_075_d 44: NW_088_d 45=37: NW_100_d

grey
g
A/P4000

metameric colours only possible for offset print and colour printers
with at least four basic colours, either *CMYK* or *CMY0*

metameric
m
P4000



vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n/f	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsi_Md	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	50.4 76.9 64.5	1.0 0.0 0.0	50.4 76.9 64.5	100.4 39.9	0.0 389	1.0 0.0 0.0	50.4 76.9 64.5
1/657	R13Y_100_100d	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	51.4 74.1 64.9	1.0 0.125 0.0	51.5 73.9 64.9	98.3 41.3	0.2 36	1.0 0.116 0.0	51.4 74.1 64.9
2/666	R25Y_100_100d	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.7 67.6 65.8	1.0 0.25 0.0	54.0 66.7 65.9	93.8 44.6	1.0 42	1.0 0.233 0.0	53.7 67.6 65.8
3/675	R38Y_100_100d	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	57.9 56.2 67.9	1.0 0.375 0.0	58.2 55.4 67.9	87.7 50.7	0.7 51	1.0 0.366 0.0	57.9 56.2 67.9
4/684	R50Y_100_100d	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	63.6 41.3 71.0	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7	0.0 59	1.0 0.5 0.0	63.6 41.3 71.0
5/693	R63Y_100_100d	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	70.5 24.7 75.4	1.0 0.625 0.0	70.1 25.8 75.0	79.3 71.0	1.2 68	1.0 0.633 0.0	70.5 24.7 75.4
6/702	R75Y_100_100d	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	78.2 7.8 80.6	1.0 0.75 0.0	77.2 9.8 79.7	80.3 82.9	2.3 77	1.0 0.766 0.0	78.2 7.8 80.6
7/711	R88Y_100_100d	1.0 0.875 0.0	1.0 1.0 0.5	83	1.0 0.883 0.0	85.3 -6.7 85.5	1.0 0.875 0.0	84.8 -5.7 85.0	85.2 93.8	1.1 83	1.0 0.883 0.0	85.3 -6.7 85.5
8/720	Y00G_100_100d	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	92.6 -20.7 90.7	1.0 1.0 0.0	92.6 -20.6 90.7	93.0 102.8	0.0 89	1.0 1.0 0.0	92.6 -20.7 90.7
9/639	Y13G_100_100d	0.875 1.0 0.0	1.0 1.0 0.5	97	0.883 1.0 0.0	90.5 -32.2 88.3	0.875 1.0 0.0	90.4 -33.0 88.1	94.1 110.5	0.8 96	0.883 1.0 0.0	90.5 -32.2 88.3
10/558	Y25G_100_100d	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	88.7 -43.3 86.2	0.75 1.0 0.0	88.5 -44.9 85.8	96.8 117.6	1.6 102	0.766 1.0 0.0	88.7 -43.3 86.2
11/477	Y38G_100_100d	0.625 1.0 0.0	1.0 1.0 0.5	112	0.633 1.0 0.0	87.0 -55.0 84.1	0.625 1.0 0.0	86.9 -55.7 83.9	100.7 123.6	0.7 111	0.633 1.0 0.0	87.0 -55.0 84.1
12/396	Y50G_100_100d	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	85.7 -65.2 82.4	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3	0.0 119	0.5 1.0 0.0	85.7 -65.2 82.4
13/315	Y63G_100_100d	0.375 1.0 0.0	1.0 1.0 0.5	128	0.366 1.0 0.0	84.7 -73.2 81.2	0.375 1.0 0.0	84.7 -72.8 81.2	109.1 131.8	0.3 128	0.366 1.0 0.0	84.7 -73.2 81.2
14/234	Y75G_100_100d	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	84.0 -78.7 80.4	0.25 1.0 0.0	84.1 -78.2 80.4	112.2 134.1	0.4 137	0.233 1.0 0.0	84.0 -78.7 80.4
15/153	Y88G_100_100d	0.125 1.0 0.0	1.0 1.0 0.5	143	0.116 1.0 0.0	83.7 -81.5 80.0	0.125 1.0 0.0	83.7 -81.4 80.0	114.2 135.5	0.1 143	0.116 1.0 0.0	83.7 -81.5 80.0
16/72	G00C_100_100d	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	83.6 -82.7 79.8	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0	0.0 149	0.0 1.0 0.0	83.6 -82.7 79.8
17/73	G13C_100_100d	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.116	83.6 -82.1 76.8	0.0 1.0 0.125	83.6 -82.1 76.5	112.3 137.0	0.2 156	0.0 1.0 0.116	83.6 -82.1 76.8
18/74	G25C_100_100d	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.233	83.7 -80.8 70.1	0.0 1.0 0.25	83.8 -80.5 69.1	106.1 139.3	1.0 162	0.0 1.0 0.233	83.7 -80.8 70.1
19/75	G38C_100_100d	0.0 1.0 0.375	1.0 1.0 0.5	172	0.0 1.0 0.366	84.0 -78.0 58.8	0.0 1.0 0.375	84.0 -77.7 58.1	97.1 143.2	0.7 171	0.0 1.0 0.366	84.0 -78.0 58.8
20/76	G50C_100_100d	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	84.3 -73.7 44.9	0.0 1.0 0.5	84.3 -73.7 44.9	86.3 148.6	0.0 180	0.0 1.0 0.5	84.3 -73.7 44.9
21/77	G63C_100_100d	0.0 1.0 0.625	1.0 1.0 0.5	188	0.0 1.0 0.633	84.8 -68.1 29.5	0.0 1.0 0.625	84.7 -68.5 30.6	75.0 155.9	1.1 188	0.0 1.0 0.633	84.8 -68.1 29.5
22/78	G75C_100_100d	0.0 1.0 0.75	1.0 1.0 0.5	196	0.0 1.0 0.766	85.4 -61.2 13.7	0.0 1.0 0.75	85.3 -62.0 15.8	64.0 165.6	2.3 197	0.0 1.0 0.766	85.4 -61.2 13.7
23/79	G88C_100_100d	0.0 1.0 0.875	1.0 1.0 0.5	203	0.0 1.0 0.883	86.1 -54.1 0.0	0.0 1.0 0.875	86.0 -54.5 1.0	54.5 178.8	1.1 203	0.0 1.0 0.883	86.1 -54.1 0.0
24/80	C00B_100_100d	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	86.8 -46.1 -13.5	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3	0.0 210	0.0 1.0 1.0	86.8 -46.1 -13.5
25/71	C13B_100_100d	0.0 0.875 1.0	1.0 1.0 0.5	217	0.0 0.883 1.0	78.5 -33.4 -26.3	0.0 0.875 1.0	77.9 -32.3 -27.0	42.1 219.8	1.3 216	0.0 0.883 1.0	78.5 -33.4 -26.3
26/62	C25B_100_100d	0.0 0.75 1.0	1.0 1.0 0.5	224	0.0 0.766 1.0	70.2 -19.5 -39.3	0.0 0.75 1.0	69.1 -17.0 -40.7	44.1 247.2	3.0 222	0.0 0.766 1.0	70.2 -19.5 -39.3
27/53	C38B_100_100d	0.0 0.625 1.0	1.0 1.0 0.5	232	0.0 0.633 1.0	60.9 -1.5 -53.9	0.0 0.625 1.0	60.3 -0.1 -54.6	54.6 269.8	1.7 231	0.0 0.633 1.0	60.9 -1.5 -53.9
28/44	C50B_100_100d	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	51.7 18.3 -68.3	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0	0.0 240	0.0 0.5 1.0	51.7 18.3 -68.3
29/35	C63B_100_100d	0.0 0.375 1.0	1.0 1.0 0.5	248	0.0 0.366 1.0	43.4 38.7 -82.0	0.0 0.375 1.0	43.8 37.6 -81.2	89.5 294.8	1.4 247	0.0 0.366 1.0	43.4 38.7 -82.0
30/26	C75B_100_100d	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.233 1.0	36.5 57.6 -93.4	0.0 0.25 1.0	37.1 55.9 -92.3	107.9 301.1	2.1 257	0.0 0.233 1.0	36.5 57.6 -93.4
31/17	C88B_100_100d	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.116 1.0	32.3 70.0 -100.3	0.0 0.125 1.0	32.4 69.6 -100.0	121.9 304.8	0.5 263	0.0 0.116 1.0	32.3 70.0 -100.3
32/8	B00M_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	30.3 76.0 -103.5	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	0.0 270	0.0 0.0 1.0	30.3 76.0 -103.5
33/89	B13M_100_100d	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	30.9 76.2 -102.5	0.125 0.0 1.0	31.0 76.2 -102.5	127.7 306.6	0.0 276	0.116 0.0 1.0	30.9 76.2 -102.5
34/170	B25M_100_100d	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	32.3 76.7 -100.1	0.25 0.0 1.0	32.6 76.8 -99.8	125.9 307.5	0.4 282	0.233 0.0 1.0	32.3 76.7 -100.1
35/251	B38M_100_100d	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	34.9 77.9 -95.7	0.375 0.0 1.0	35.1 77.9 -95.5	123.3 309.2	0.3 291	0.366 0.0 1.0	34.9 77.9 -95.7
36/332	B50M_100_100d	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	38.5 79.8 -89.7	0.5 0.0 1.0	38.5 79.8 -89.7	120.1 311.6	0.0 300	0.5 0.0 1.0	38.5 79.8 -89.7
37/413	B63M_100_100d	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	43.0 82.7 -82.2	0.625 0.0 1.0	42.7 82.5 -82.8	116.8 314.8	0.6 308	0.633 0.0 1.0	43.0 82.7 -82.2
38/494	B75M_100_100d	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	47.9 86.4 -74.0	0.75 0.0 1.0	47.2 85.8 -75.1	114.1 318.8	1.3 317	0.766 0.0 1.0	47.9 86.4 -74.0
39/575	B88M_100_100d	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	52.5 90.1 -66.3	0.875 0.0 1.0	52.1 89.8 -66.9	112.0 323.3	0.7 323	0.883 0.0 1.0	52.5 90.1 -66.3
40/656	M00R_100_100d	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	57.2 94.3 -58.4	1.0 0.0 1.0	57.2 94.3 -58.4	111.0 328.2	0.0 330	1.0 0.0 1.0	57.2 94.3 -58.4
41/655	M13R_100_100d	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	55.7 90.6 -44.8	1.0 0.0 0.875	55.6 90.3 -43.9	100.4 334.0	0.9 336	1.0 0.0 0.883	55.7 90.6 -44.8
42/654	M25R_100_100d	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	54.4 87.3 -30.6	1.0 0.0 0.75	54.2 86.7 -28.6	91.3 341.6	2.0 342	1.0 0.0 0.766	54.4 87.3 -30.6
43/653	M38R_100_100d	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	53.0 83.9 -13.6	1.0 0.0 0.625	53.0 83.6 -12.6	84.6 351.4	1.0 351	1.0 0.0 0.633	53.0 83.9 -13.6
44/652	M50R_100_100d	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	52.0 81.1 4.1	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	0.0 360	1.0 0.0 0.5	52.0 81.1 4.1
45/651	M63R_100_100d	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	51.3 79.3 22.7	1.0 0.0 0.375	51.3 79.2 21.6	82.1 15.2	1.1 368	1.0 0.0 0.366	51.3 79.3 22.7
46/650	M75R_100_100d	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	50.8 78.0 41.2	1.0 0.0 0.25	50.8 77.9 39.2	87.2 26.6	2.0 377	1.0 0.0 0.233	50.8 78.0 41.2
47/649	M88R_100_100d	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	50.5 77.2 55.6	1.0 0.0 0.125	50.6 77.2 54.9	94.8 35.4	0.6 383	1.0 0.0 0.116	50.5 77.2 55.6
48/648	R00Y_100_100d	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5	1.0 0.0 0.0	50.4 76.9 64.5	100.4 39.9	0.0 389	1.0 0.0 0.0	50.4 76.9 64.5
49/0	NW_000d	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0	0.0 360	1.0 1.0 1.0	95.4 0.0 0.

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS31/PS31L0NP.PDF /.PS
aplicación para la medida de display output, ninguna separación
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*Ma
0/648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.0	50.4	76.9	64.5	100.4	40.0
1/666	R25Y_100_100a	1.0	0.25	0.0	1.0	1.0	0.233	53.7	67.6	65.8	94.4	44.2
2/684	R50Y_100_100a	1.0	0.5	0.0	1.0	1.0	0.5	63.6	41.3	71.0	82.2	59.7
3/702	R75Y_100_100a	1.0	0.75	0.0	1.0	1.0	0.766	78.2	9.8	80.6	81.0	84.4
4/720	Y00G_100_100a	1.0	1.0	0.0	1.0	1.0	1.0	92.6	-20.6	90.7	93.0	102.8
5/558	Y25G_100_100a	0.75	1.0	0.0	1.0	1.0	0.5	88.5	-44.9	85.8	96.8	116.6
6/396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	85.7	-65.2	82.4	105.1	128.3
7/234	Y75G_100_100a	0.25	1.0	0.0	1.0	1.0	0.5	84.1	-78.2	80.4	112.2	134.1
8/72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	1.0	83.6	-82.7	79.8	115.0	136.0
9/72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.0	83.6	-82.7	79.8	115.0	136.0
10/76	G25B_100_100a	0.0	1.0	0.5	1.0	1.0	0.5	84.3	-73.7	44.9	86.4	148.6
11/80	G50B_100_100a	0.0	1.0	1.0	1.0	1.0	1.0	86.8	-46.1	-13.5	48.1	196.3
12/44	G75B_100_100a	0.0	0.5	1.0	1.0	1.0	0.5	51.7	18.3	-68.3	70.7	285.0
13/8	B00M_100_100a	0.0	0.0	1.0	1.0	1.0	0.0	30.3	76.0	-103.5	128.5	306.2
14/332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	38.5	79.8	-89.7	120.1	311.6
15/656	B50R_100_100a	1.0	0.0	1.0	1.0	1.0	0.5	57.2	94.3	-58.4	110.9	328.2
16/652	B75R_100_100a	1.0	0.0	0.5	1.0	1.0	0.5	81.1	81.1	4.1	81.2	2.9
17/648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.0	50.4	76.9	64.5	100.4	40.0
18/688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	72.9	38.4	32.2	50.2	40.0
19/706	R50Y_100_050a	1.0	0.75	0.5	1.0	0.5	0.75	79.5	20.6	35.5	41.1	59.7
20/724	Y00G_100_050a	1.0	1.0	0.5	1.0	0.5	0.75	94.0	-10.3	45.3	46.5	102.8
21/562	Y50G_100_050a	0.75	1.0	0.5	1.0	0.5	0.75	90.5	-32.6	41.2	52.5	128.3
22/400	G00B_100_050a	0.5	1.0	0.5	1.0	0.5	0.75	89.5	-41.3	39.9	57.5	136.0
23/404	G50B_100_050a	0.5	1.0	1.0	1.0	0.5	0.75	91.1	-23.0	-6.7	24.0	196.3
24/368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	62.8	38.0	-51.7	64.2	306.2
25/692	B50R_100_050a	1.0	0.5	1.0	1.0	0.5	0.75	76.3	47.1	-29.2	55.4	328.2
26/688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	72.9	38.4	32.2	50.2	40.0
27/506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	49.0	38.4	32.2	50.2	40.0
28/524	R50Y_075_050a	0.75	0.5	0.25	0.75	0.5	0.5	55.6	20.6	35.5	41.1	59.7
29/542	Y00G_075_050a	0.75	0.75	0.25	0.75	0.5	0.5	70.1	-10.3	45.3	46.5	102.8
30/380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	66.7	-32.6	41.2	52.5	128.3
31/218	G00B_075_050a	0.25	0.75	0.25	0.75	0.5	0.5	65.6	-41.3	39.9	57.5	136.0
32/222	G50B_075_050a	0.25	0.75	0.75	0.75	0.5	0.5	67.2	-23.0	-6.7	24.0	196.3
33/186	B00R_075_050a	0.25	0.25	0.75	0.75	0.5	0.5	62.8	38.0	-51.7	64.2	306.2
34/510	B50R_075_050a	0.75	0.25	0.75	0.75	0.5	0.5	52.5	47.1	-29.2	55.4	328.2
35/506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	49.0	38.4	32.2	50.2	40.0
36/324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.25	25.2	38.4	32.2	50.2	40.0
37/342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	31.8	20.6	35.5	41.1	59.7
38/360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	46.3	-10.3	45.3	46.5	102.8
39/198	Y50G_050_050a	0.25	0.5	0.0	0.5	0.5	0.25	42.8	-32.6	41.2	52.5	128.3
40/36	G00B_050_050a	0.0	0.5	0.0	0.5	0.5	0.25	41.8	-41.3	39.9	57.5	136.0
41/40	G50B_050_050a	0.0	0.5	0.5	0.5	0.5	0.25	43.4	-23.0	-6.7	24.0	196.3
42/4	B00R_050_050a	0.0	0.0	0.5	0.5	0.5	0.25	15.1	38.0	-51.7	64.2	306.2
43/328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.25	28.6	47.1	-29.2	55.4	328.2
44/324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.25	25.2	38.4	32.2	50.2	40.0
45/0	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
46/91	NW_013a	0.125	0.125	0.125	0.125	0.0	0.125	11.9	0.0	0.0	0.0	0.0
47/182	NW_025a	0.25	0.25	0.25	0.25	0.0	0.25	23.8	0.0	0.0	0.0	0.0
48/273	NW_038a	0.375	0.375	0.375	0.375	0.0	0.375	35.7	0.0	0.0	0.0	0.0
49/364	NW_050a	0.5	0.5	0.5	0.5	0.0	0.5	47.7	0.0	0.0	0.0	0.0
50/455	NW_063a	0.625	0.625	0.625	0.625	0.0	0.625	59.6	0.0	0.0	0.0	0.0
51/546	NW_075a	0.75	0.75	0.75	0.75	0.0	0.75	71.5	0.0	0.0	0.0	0.0
52/637	NW_088a	0.875	0.875	0.875	0.875	0.0	0.875	83.4	0.0	0.0	0.0	0.0
53/728	NW_100a	1.0	1.0	1.0	1.0	0.0	1.0	95.4	0.0	0.0	0.0	0.0

delta E* = 6.5

gráfico TUB-PS31; reproducción en color; sRGB
colores y diferencia en color, ΔE^* , 3D=0, de=0, sRGB

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

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31.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	0.125
12	G84B_037_037a	0.0	0.125	0.375	0.375	0.375	0.187	0.187	0.0	0.125	0.375	0.187
13	G88B_050_050a	0.0	0.125	0.5	0.5	0.5	0.25	0.25	0.0	0.125	0.5	0.25
14	G90B_062_062a	0.0	0.125	0.625	0.625	0.625	0.312	0.312	0.0	0.125	0.625	0.312
15	G92B_075_075a	0.0	0.125	0.75	0.75	0.75	0.375	0.375	0.0	0.125	0.75	0.375
16	G93B_087_087a	0.0	0.125	0.875	0.875	0.875	0.437	0.437	0.0	0.125	0.875	0.437
17	G94B_100_100a	0.0	0.125	1.0	1.0	1.0	0.5	0.5	0.0	0.125	1.0	0.5
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	0.125
21	G65B_037_037a	0.0	0.25	0.375	0.375	0.375	0.187	0.187	0.0	0.25	0.375	0.187
22	G75B_050_050a	0.0	0.25	0.5	0.5	0.5	0.25	0.25	0.0	0.25	0.5	0.25
23	G80B_062_062a	0.0	0.25	0.625	0.625	0.625	0.312	0.312	0.0	0.25	0.625	0.312
24	G84B_075_075a	0.0	0.25	0.75	0.75	0.75	0.375	0.375	0.0	0.25	0.75	0.375
25	G86B_087_087a	0.0	0.25	0.875	0.875	0.875	0.437	0.437	0.0	0.25	0.875	0.437
26	G88B_100_100a	0.0	0.25	1.0	1.0	1.0	0.5	0.5	0.0	0.25	1.0	0.5
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	0.187
29	G34B_037_037a	0.0	0.375	0.25	0.375	0.375	0.187	0.187	0.0	0.375	0.25	0.187
30	G50B_037_037a	0.0	0.375	0.375	0.375	0.375	0.187	0.187	0.0	0.375	0.375	0.187
31	G61B_050_050a	0.0	0.375	0.5	0.5	0.5	0.25	0.25	0.0	0.375	0.5	0.25
32	G69B_062_062a	0.0	0.375	0.625	0.625	0.625	0.312	0.312	0.0	0.375	0.625	0.312
33	G75B_075_075a	0.0	0.375	0.75	0.75	0.75	0.375	0.375	0.0	0.375	0.75	0.375
34	G79B_087_087a	0.0	0.375	0.875	0.875	0.875	0.437	0.437	0.0	0.375	0.875	0.437
35	G81B_100_100a	0.0	0.375	1.0	1.0	1.0	0.5	0.5	0.0	0.375	1.0	0.5
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	0.25
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	0.25
40	G50B_050_050a	0.0	0.5	0.5	0.5	0.5	0.25	0.25	0.0	0.5	0.5	0.25
41	G59B_062_062a	0.0	0.5	0.625	0.625	0.625	0.312	0.312	0.0	0.5	0.625	0.312
42	G65B_075_075a	0.0	0.5	0.75	0.75	0.75	0.375	0.375	0.0	0.5	0.75	0.375
43	G70B_087_087a	0.0	0.5	0.875	0.875	0.875	0.437	0.437	0.0	0.5	0.875	0.437
44	G75B_100_100a	0.0	0.5	1.0	1.0	1.0	0.5	0.5	0.0	0.5	1.0	0.5
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	0.312
47	G19B_062_062a	0.0	0.625	0.25	0.625	0.625	0.312	0.312	0.0	0.625	0.25	0.312
48	G30B_062_062a	0.0	0.625	0.375	0.625	0.625	0.312	0.312	0.0	0.625	0.375	0.312
49	G40B_062_062a	0.0	0.625	0.5	0.625	0.625	0.312	0.312	0.0	0.625	0.5	0.312
50	G50B_062_062a	0.0	0.625	0.625	0.625	0.625	0.312	0.312	0.0	0.625	0.625	0.312
51	G57B_075_075a	0.0	0.625	0.75	0.75	0.75	0.375	0.375	0.0	0.625	0.75	0.375
52	G63B_087_087a	0.0	0.625	0.875	0.875	0.875	0.437	0.437	0.0	0.625	0.875	0.437
53	G68B_100_100a	0.0	0.625	1.0	1.0	1.0	0.5	0.5	0.0	0.625	1.0	0.5
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	0.375
56	G15B_075_075a	0.0	0.75	0.25	0.75	0.75	0.375	0.375	0.0	0.75	0.25	0.375
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	0.375
59	G42B_075_075a	0.0	0.75	0.625	0.75	0.75	0.375	0.375	0.0	0.75	0.625	0.375
60	G50B_075_075a	0.0	0.75	0.75	0.75	0.75	0.375	0.375	0.0	0.75	0.75	0.375
61	G56B_087_087a	0.0	0.75	0.875	0.875	0.875	0.437	0.437	0.0	0.75	0.875	0.437
62	G61B_100_100a	0.0	0.75	1.0	1.0	1.0	0.5	0.5	0.0	0.75	1.0	0.5
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	0.437
65	G13B_087_087a	0.0	0.875	0.25	0.875	0.875	0.437	0.437	0.0	0.875	0.25	0.437
66	G20B_087_087a	0.0	0.875	0.375	0.875	0.875	0.437	0.437	0.0	0.875	0.375	0.437
67	G29B_087_087a	0.0	0.875	0.5	0.875	0.875	0.437	0.437	0.0	0.875	0.5	0.437
68	G36B_087_087a	0.0	0.875	0.625	0.875	0.875	0.437	0.437	0.0	0.875	0.625	0.437
69	G43B_087_087a	0.0	0.875	0.75	0.875	0.875	0.437	0.437	0.0	0.875	0.75	0.437
70	G50B_087_087a	0.0	0.875	0.875	0.875	0.875	0.437	0.437	0.0	0.875	0.875	0.437
71	G55B_100_100a	0.0	0.875	1.0	1.0	1.0	0.5	0.5	0.0	0.875	1.0	0.5
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	0.5
78	G38B_100_100a	0.0	1.0	0.75	1.0	1.0	0.5	0.5	0.0	1.0	0.75	0.5
79	G44B_100_100a	0.0	1.0	0.875	1.0	1.0	0.5	0.5	0.0	1.0	0.875	0.5
80	G50B_100_100a	0.0	1.0	1.0	1.0	1.0	0.5	0.5	0.0	1.0	1.0	0.5

delta E** = 4.6

gráfico TUB-PS31; reproducción en color; sRGB
colores y diferencia en color, ΔE^* , 3D=0, de=0, sRGB

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

TUB matrícula: 20130201-PS31/PS31L0NP.PDF /.PS
aplicación para la medida de display output, ninguna separación
TUB material: code=rha4ta

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31.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		
81	R00Y_012_012a	0.125 0.0 0.0	0.125 0.125 0.062	390	0.125 0.0 0.0	6.3 9.6 8.0	12.5 40.0	0.125 0.0 0.0	2.4 10.9 3.8	11.6 19.4 5.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 100.4 40.0
82	B50R_012_012a	0.125 0.0 0.125	0.125 0.125 0.062	330	0.125 0.0 0.125	7.1 11.7 -7.3	13.8 328.2	0.125 0.0 0.125	3.2 16.7 -11.6	20.4 325.1 7.6	330	1.0 0.0 0.0	57.2 94.3 -58.4	110.9 328.2
83	B25R_025_025a	0.125 0.0 0.25	0.25 0.25 0.125	300	0.125 0.0 0.25	9.6 19.9 -22.4	30.0 311.6	0.125 0.0 0.25	5.3 28.5 -31.2	42.3 312.3 13.0	300	0.5 0.0 0.0	38.5 79.8 -89.7	120.0 311.6
84	B15R_037_037a	0.125 0.0 0.375	0.375 0.375 0.187	289	0.118 0.0 0.375	12.7 29.0 -36.5	46.7 308.4	0.125 0.0 0.375	9.0 38.1 -46.3	60.0 309.4 13.8	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
85	B11R_050_050a	0.125 0.0 0.5	0.5 0.5 0.25	284	0.116 0.0 0.5	16.1 38.3 -50.0	63.1 307.4	0.125 0.0 0.5	13.4 46.1 -59.0	74.9 307.9 12.1	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
86	B09R_062_062a	0.125 0.0 0.625	0.625 0.625 0.312	281	0.114 0.0 0.625	19.8 47.8 -63.2	79.3 307.0	0.125 0.0 0.625	17.9 53.9 -70.7	88.9 307.3 9.8	279	0.183 0.0 1.0	31.7 76.5 -101.2	126.9 307.0
87	B07R_075_075a	0.125 0.0 0.75	0.75 0.75 0.375	279	0.112 0.0 0.75	23.5 57.2 -76.4	95.5 306.8	0.125 0.0 0.75	22.3 61.5 -81.7	102.3 306.9 6.9	278	0.15 0.0 1.0	31.3 76.3 -101.9	127.4 306.8
88	B06R_087_087a	0.125 0.0 0.875	0.875 0.875 0.437	278	0.110 0.0 0.875	27.2 66.7 -89.5	111.6 306.7	0.125 0.0 0.875	26.7 69.0 -92.3	115.2 306.7 3.6	277	0.133 0.0 1.0	31.1 76.3 -102.3	127.6 306.7
89	B05R_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	30.9 76.2 -102.5	127.8 306.6	0.125 0.0 1.0	31.0 76.2 -102.5	127.7 306.6 0.0	276	0.116 0.0 1.0	30.9 76.2 -102.5	127.8 306.6
90	Y00G_012_012a	0.125 0.125 0.0	0.125 0.125 0.062	90	0.125 0.125 0.0	11.5 -25.5 11.3	11.6 102.8	0.125 0.125 0.0	10.4 -5.0 15.4	16.2 108.0 4.8	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
91	NW_012a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	11.9 0.0 0.0	0.0 0.0	0.0 0.125 0.125	11.0 0.0 0.0	0.0 325.7 0.8	360	1.0 1.0 1.0	95.4 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 0.124 0.25	15.7 9.5 -12.9	16.0 306.2	0.125 0.125 0.25	12.6 9.6 -19.5	21.8 296.2 7.3	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
93	B00R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.124 0.375	19.5 19.0 -25.8	32.1 306.2	0.125 0.125 0.375	15.0 21.1 -36.5	42.1 300.0 11.6	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
94	B00R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.124 0.5	23.3 28.5 -38.8	48.1 306.2	0.125 0.125 0.5	18.1 32.4 -51.3	60.6 302.2 14.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
95	B00R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	27.1 38.0 -51.7	64.2 306.2	0.125 0.125 0.625	21.6 42.8 -64.6	77.5 303.5 14.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
96	B00R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	30.9 47.5 -64.7	80.3 306.2	0.125 0.125 0.75	25.3 52.5 -76.8	93.0 304.3 14.2	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
97	B00R_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	34.7 57.0 -77.6	96.3 306.2	0.125 0.125 0.875	29.1 61.5 -88.2	107.5 304.8 12.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
98	B00R_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	38.5 66.5 -90.6	112.4 306.2	0.125 0.125 1.0	33.0 69.9 -99.0	121.3 305.2 10.6	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
99	Y50G_025_025a	0.125 0.25 0.0	0.25 0.25 0.125	120	0.125 0.25 0.0	21.4 -16.3 20.6	26.2 128.3	0.125 0.25 0.0	21.9 -22.3 29.7	37.2 126.9 10.9	119	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
100	G00B_025_012a	0.125 0.25 0.125	0.25 0.125 0.187	150	0.124 0.25 0.124	22.3 -10.3 9.9	14.3 136.0	0.125 0.25 0.125	22.2 -18.8 15.2	24.2 141.0 10.0	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
101	G50B_025_012a	0.125 0.25 0.25	0.25 0.125 0.187	210	0.124 0.25 0.25	22.7 -5.7 -1.6	6.0 196.3	0.125 0.25 0.25	23.0 -11.2 -3.5	11.7 197.3 5.7	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
102	G75B_037_025a	0.125 0.25 0.375	0.375 0.25 0.25	240	0.124 0.25 0.375	24.8 4.5 -17.0	17.6 285.0	0.125 0.25 0.375	24.4 -0.5 -21.5	21.5 268.6 6.7	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
103	G84B_050_037a	0.125 0.25 0.5	0.5 0.375 0.312	251	0.124 0.243 0.5	27.2 17.1 -32.5	36.7 297.8	0.125 0.25 0.5	26.3 11.5 -37.9	39.6 286.9 7.8	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
104	G88B_062_050a	0.125 0.25 0.625	0.625 0.5 0.375	256	0.125 0.241 0.625	30.1 28.8 -46.7	54.8 301.6	0.125 0.25 0.625	28.7 23.7 -52.9	58.0 294.1 8.1	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6
105	G90B_075_062a	0.125 0.25 0.75	0.75 0.625 0.437	259	0.125 0.239 0.75	33.5 39.4 -60.3	72.1 303.1	0.125 0.25 0.75	31.4 35.4 -66.7	75.5 297.9 7.7	260	0.0 0.183 1.0	34.6 63.0 -96.6	115.3 303.1
106	G92B_087_075a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.237 0.875	36.9 50.0 -73.9	89.3 304.0	0.125 0.25 0.875	34.4 46.3 -79.5	92.0 300.2 7.1	262	0.0 0.15 1.0	33.4 66.7 -98.6	119.1 304.0
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.241 1.0	40.6 60.0 -87.1	105.8 304.5	0.125 0.25 1.0	37.6 56.5 -94.4	107.5 301.7 6.3	262	0.0 0.133 1.0	32.8 68.6 -99.6	120.9 304.5
108	Y68G_037_037a	0.125 0.375 0.0	0.375 0.375 0.187	131	0.118 0.375 0.0	31.6 -28.2 30.3	41.4 132.9	0.125 0.375 0.0	33.1 -35.2 39.6	53.0 131.5 11.7	131	0.316 1.0 0.0	84.4 -75.3 80.9	110.6 132.9
109	G00B_037_025a	0.125 0.375 0.125	0.375 0.25 0.25	150	0.124 0.375 0.124	32.8 -20.6 19.9	28.7 136.0	0.125 0.375 0.125	33.3 -32.9 28.6	43.6 138.9 14.9	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
110	G25B_037_025a	0.125 0.375 0.25	0.375 0.25 0.25	180	0.124 0.375 0.25	33.0 -18.4 11.2	21.6 148.6	0.125 0.375 0.25	33.8 -27.4 11.9	29.9 156.5 9.0	180	0.0 1.0 0.5	84.3 -73.7 44.9	86.4 148.6
111	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	33.6 -11.5 -3.3	12.0 196.3	0.125 0.375 0.375	34.7 -18.9 -5.7	19.8 196.8 7.8	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
112	G65B_050_037a	0.125 0.375 0.5	0.5 0.375 0.312	229	0.124 0.381 0.5	36.0 -3.4 -18.3	18.6 259.3	0.125 0.375 0.5	35.9 -8.3 -22.7	24.1 249.7 6.5	228	0.0 0.683 1.0	64.4 -9.2 -48.8	49.7 259.3
113	G75B_062_050a	0.125 0.375 0.625	0.625 0.5 0.375	240	0.125 0.375 0.625	37.8 9.1 -34.1	35.3 285.0	0.125 0.375 0.625	37.5 3.3 -38.6	38.7 274.9 7.3	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.364 0.75	39.6 22.6 -50.3	55.1 294.2	0.125 0.375 0.75	39.5 15.3 -53.5	55.6 285.9 7.9	247	0.0 0.383 1.0	44.3 36.2 -80.5	88.2 294.2
115	G84B_087_075a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.362 0.875	42.4 34.3 -65.0	73.5 297.8	0.125 0.375 0.875	41.7 27.1 -67.4	72.7 291.9 7.5	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
116	G86B_100_087a	0.125 0.375 1.0	1.0 0.875 0.562	254	0.125 0.358 1.0	45.2 46.6 -79.6	92.2 300.3	0.125 0.375 1.0	44.2 38.6 -80.5	89.3 295.5 8.1	255	0.0 0.266 1.0	38.0 53.3 -91.0	105.4 300.3
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 -39.3 40.2	56.2 134.3	0.125 0.5 0.0	43.9 -45.9 48.2	66.6 133.6 10.6	137	0.233 1.0 0.0	84.0 -78.7 80.4	112.5 134.3
118	G00B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.124	43.2 -31.0 29.9	43.1 136.0	0.125 0.5 0.125	44.1 -44.3 40.1	59.8 137.8 16.7	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.243	43.3 -29.7 23.6	38.0 141.4	0.125 0.5 0.25	44.4 -40.3 25.7	47.9 147.4 10.9	168	0.0 1.0 0.316	83.9 -79.2 63.1	101.3 141.4
120	G34B_050_037a	0.125 0.5 0.375	0.5 0.375 0.312	191	0.124 0.5 0.381	43.8 -24.7 8.7	26.2 160.4	0.125 0.5 0.375	45.0 -33.8 9.2	35.1 164.7 9.2	191	0.0 1.0 0.683	85.0 -65.8 23.4	69.9 160.4
121														

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31.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	
162	R00Y_025_025a	0.25 0.0 0.0	0.25 0.25 0.125	390	0.25 0.0 0.0	12.6 19.2 16.1	25.1 40.0	0.25 0.0 0.0	8.6 28.5 13.6	31.6 25.5 10.4	389 1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
163	R00Y_025_025a	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.125	13.0 20.2 1.0	20.3 2.9	0.25 0.0 0.125	9.4 30.5 -1.8	30.6 356.5 11.2	360 1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
164	B50R_025_025a	0.25 0.0 0.25	0.25 0.25 0.125	330	0.25 0.0 0.25	14.3 23.5 -14.6	27.7 328.2	0.25 0.0 0.25	11.1 34.9 -21.6	41.1 328.2 13.7	330 1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
165	B34R_037_037a	0.25 0.0 0.375	0.375 0.375 0.187	311	0.256 0.0 0.375	16.8 31.5 -29.7	43.3 316.7	0.25 0.0 0.375	13.8 41.1 -38.3	56.2 316.9 13.2	311 1.0 0.683 0.0	44.8 84.1 -79.2	115.5 316.7
166	B25R_050_050a	0.25 0.0 0.5	0.5 0.5 0.25	300	0.25 0.0 0.5	19.2 39.9 -44.8	60.0 311.6	0.25 0.0 0.5	17.1 48.0 -52.8	71.4 312.2 11.6	300 0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
167	B19R_062_062a	0.25 0.0 0.625	0.625 0.625 0.312	293	0.239 0.0 0.625	22.1 48.8 -59.4	76.9 309.3	0.25 0.0 0.625	20.7 55.2 -65.9	86.0 309.9 9.2	292 0.383 0.0 1.0	35.3 78.1 -95.1	120.3 309.3
168	B15R_075_075a	0.25 0.0 0.75	0.75 0.75 0.375	289	0.237 0.0 0.75	25.4 58.1 -73.1	93.4 308.4	0.25 0.0 0.75	24.6 62.5 -77.8	99.8 308.7 6.5	288 0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
169	B13R_087_087a	0.25 0.0 0.875	0.875 0.875 0.437	286	0.233 0.0 0.875	28.8 67.3 -86.8	109.9 307.8	0.25 0.0 0.875	28.6 69.7 -89.1	113.1 308.0 3.2	284 0.266 0.0 1.0	32.9 77.0 -99.2	125.6 307.0
170	B11R_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4	0.25 0.0 1.0	32.6 76.8 -99.8	125.9 307.5 0.4	282 0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
171	R50Y_025_025a	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.125 0.0	15.9 10.3 17.7	20.5 59.7	0.25 0.125 0.0	14.7 12.2 22.0	25.2 60.9 4.8	59 1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
172	R00Y_025_012a	0.25 0.125 0.125	0.25 0.125 0.187	390	0.25 0.124 0.124	18.2 9.6 8.0	12.5 40.0	0.25 0.125 0.125	15.2 14.7 6.5	16.1 23.9 6.1	389 1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
173	B50R_025_012a	0.25 0.125 0.25	0.25 0.125 0.187	330	0.25 0.124 0.25	19.0 11.7 -7.3	13.8 328.2	0.25 0.125 0.25	16.4 20.2 -13.2	24.2 326.7 10.6	330 1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
174	B25R_037_025a	0.25 0.125 0.375	0.375 0.25 0.25	300	0.25 0.124 0.375	21.5 19.9 -22.4	30.0 311.6	0.25 0.125 0.375	18.4 28.0 -30.9	41.7 312.1 12.1	300 0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
175	B15R_050_037a	0.25 0.125 0.5	0.5 0.375 0.312	289	0.243 0.124 0.5	24.6 29.0 -36.5	46.7 308.4	0.25 0.125 0.5	20.9 36.7 -46.5	59.3 308.3 13.1	288 0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
176	B11R_062_050a	0.25 0.125 0.625	0.625 0.5 0.375	284	0.241 0.125 0.625	28.1 38.3 -50.0	63.1 307.4	0.25 0.125 0.625	23.9 45.7 -60.5	75.9 307.0 13.4	282 0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
177	B09R_075_062a	0.25 0.125 0.75	0.75 0.625 0.437	281	0.239 0.125 0.75	31.7 47.8 -63.2	79.3 307.0	0.25 0.125 0.75	27.3 54.4 -73.4	91.4 306.5 12.9	279 0.183 0.0 1.0	31.7 76.5 -101.2	126.9 307.0
178	B07R_087_075a	0.25 0.125 0.875	0.875 0.75 0.5	279	0.237 0.125 0.875	35.4 57.2 -76.4	95.5 306.8	0.25 0.125 0.875	30.8 62.8 -85.3	106.0 306.3 11.4	278 0.15 0.0 1.0	31.3 76.3 -101.9	127.4 306.8
179	B06R_100_087a	0.25 0.125 1.0	1.0 0.875 0.562	278	0.241 0.125 1.0	39.1 66.7 -89.5	111.6 306.7	0.25 0.125 1.0	34.5 70.9 -96.6	119.8 306.2 9.4	277 0.133 0.0 1.0	31.1 76.3 -102.3	127.6 306.7
180	Y00G_025_025a	0.25 0.25 0.0	0.25 0.25 0.125	90	0.25 0.25 0.0	23.1 -5.1 22.6	23.2 102.8	0.25 0.25 0.0	24.2 -7.6 32.9	33.7 103.1 10.5	89 1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
181	Y00G_025_012a	0.25 0.25 0.125	0.25 0.125 0.187	90	0.25 0.25 0.124	23.5 -2.5 11.3	11.6 102.8	0.25 0.25 0.125	24.5 -5.3 18.6	19.4 105.9 7.8	89 1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	23.8 0.0 0.0	0.0 0.0	0.25 0.25 0.25	25.2 0.0 0.0	0.0 325.5 1.4	360 1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0
183	B00R_037_012a	0.25 0.25 0.375	0.375 0.125 0.312	270	0.249 0.249 0.375	27.6 9.5 -12.9	16.0 306.2	0.25 0.25 0.375	26.5 8.0 -18.0	19.8 294.0 5.4	270 0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
184	B00R_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.249 0.5	31.4 19.0 -25.8	32.1 306.2	0.25 0.25 0.5	28.2 17.7 -34.7	39.0 297.0 9.5	270 0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
185	B00R_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	35.2 28.5 -38.8	48.1 306.2	0.25 0.25 0.625	30.4 28.1 -50.0	57.4 299.3 12.2	270 0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	39.0 38.0 -51.7	64.2 306.2	0.25 0.25 0.75	32.9 38.5 -64.1	74.8 301.0 13.7	270 0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
187	B00R_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.8 47.5 -64.7	80.3 306.2	0.25 0.25 0.875	35.8 48.6 -77.1	91.2 302.1 14.3	270 0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
188	B00R_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	46.6 57.0 -77.6	96.3 306.2	0.25 0.25 1.0	38.8 58.2 -89.4	106.7 303.0 14.1	270 0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
189	Y31G_037_037a	0.25 0.375 0.0	0.375 0.375 0.187	109	0.256 0.375 0.0	32.8 -19.0 31.8	37.1 120.8	0.25 0.375 0.0	34.6 -24.3 41.4	48.0 120.4 11.0	108 0.683 1.0 0.0	87.6 -50.7 84.9	98.9 120.8
190	Y50G_037_025a	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	33.3 -16.3 20.6	26.2 128.3	0.25 0.375 0.125	34.8 -22.5 30.5	38.0 126.3 11.8	119 0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
191	G00B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	34.3 -10.3 9.9	14.3 136.0	0.25 0.375 0.25	35.2 -18.1 14.0	22.9 142.2 8.8	149 0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
192	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	34.7 -5.7 -1.6	6.0 196.3	0.25 0.375 0.375	36.0 -11.0 -3.5	11.6 197.8 5.8	210 0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
193	G75B_050_025a	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.375 0.5	36.7 4.5 -17.0	17.6 285.0	0.25 0.375 0.5	37.2 -2.0 -20.5	20.6 264.3 7.4	240 0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
194	G84B_062_037a	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.368 0.625	39.1 17.1 -32.5	36.7 297.8	0.25 0.375 0.625	38.7 8.2 -36.6	37.5 282.7 9.7	251 0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
195	G88B_075_050a	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.366 0.75	42.1 28.8 -46.7	54.8 301.6	0.25 0.375 0.75	40.6 19.1 -51.6	55.0 290.3 10.9	257 0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6
196	G90B_087_062a	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.364 0.875	45.5 39.4 -60.3	72.1 303.1	0.25 0.375 0.875	42.8 30.1 -65.7	72.2 294.6 11.0	260 0.0 0.183 1.0	34.6 63.0 -96.6	115.3 303.1
197	G92B_100_075a	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.362 1.0	48.9 50.0 -73.9	89.3 304.0	0.25 0.375 1.0	45.2 40.8 -78.9	88.9 297.3 11.1	262 0.0 0.15 1.0	33.4 66.7 -98.6	119.1 304.0
198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	42.8 -32.6 41.2	52.5 128.3	0.25 0.5 0.0	44.9 -37.9 49.4	62.3 127.5 10.0	119 0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
199	Y68G_050_037a	0.25 0.5 0.125	0.5 0.375 0.312	131	0.243 0.5 0.124	43.6 -28.2 30.3	41.4 132.9	0.25 0.5 0.125	45.0 -36.5 41.4	55.2 131.4 13.9	131 0.316 1.0 0.0	84.4 -75.3 80.9	110.6 132.9
200	G00B_050_025a	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.249	44.7 -20.6 19.9	28.7 136.0	0.25 0.5 0.25	45.4 -33.0 27.2	42.8 140.5 14.3	149 0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
201	C25B_050_025a	0.25 0.5 0.375	0.5 0.25 0.375	180	0.249 0.5 0.375	44.9 -18.4 11.2	21.6 148.6	0.25 0.5 0.375	45.9 -27.3 10.6	29.3 158.6 8.9	180 0.0 1.0 0.5	84.3 -73.7 44.9	86.4 148.6
202	C50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	45.5 -11.5 -3.3	12.0 196.3	0.25 0.5 0.5	46.8 -19.5 -6.0	20.4 197.2 8.5	210 0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
203	C65B_062_037a	0.25 0.5											

	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	18.9 28.8 24.2	37.6 40.0	0.375 0.0 0.0	16.4 37.5 25.4	45.3 34.1 9.1	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
244	R18Y_037_037a	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.118	19.1 29.6 11.1	31.7 20.6	0.375 0.0 0.125	16.8 38.7 9.7	39.9 14.1 9.4	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6
245	B65R_037_037a	0.375 0.0 0.25	0.375 0.375 0.187	349	0.375 0.0 0.256	20.0 32.0 -7.4	32.9 346.8	0.375 0.0 0.25	17.9 41.5 -10.4	42.8 345.8 10.2	348	1.0 0.0 0.683	53.5 85.4 -19.9	87.7 346.8
246	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	21.4 35.3 -21.9	41.6 328.2	0.375 0.0 0.375	19.7 46.0 -28.5	54.1 328.2 12.6	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
247	B38R_050_050a	0.375 0.0 0.5	0.5 0.5 0.25	316	0.383 0.0 0.5	23.9 43.2 -37.0	56.9 319.4	0.375 0.0 0.5	22.1 51.5 -44.4	68.1 319.2 11.3	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4
248	B30R_062_062a	0.375 0.0 0.625	0.625 0.625 0.312	307	0.385 0.0 0.625	25.5 51.4 -52.0	73.1 314.6	0.375 0.0 0.625	24.9 57.8 -58.7	82.4 314.5 9.4	307	0.616 0.0 1.0	42.4 82.3 -83.2	117.0 314.6
249	B25R_075_075a	0.375 0.0 0.75	0.75 0.75 0.375	300	0.375 0.0 0.75	28.9 59.8 -67.2	90.0 311.6	0.375 0.0 0.75	28.1 64.4 -71.9	96.5 311.8 6.5	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
250	B20R_087_087a	0.375 0.0 0.875	0.875 0.875 0.437	295	0.364 0.0 0.875	31.7 68.8 -81.8	106.9 310.0	0.375 0.0 0.875	31.6 71.2 -84.0	110.1 310.2 3.2	294	0.416 0.0 1.0	36.3 78.6 -93.5	122.2 310.0
251	B18R_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	34.9 77.9 -95.7	123.4 309.1	0.375 0.0 1.0	35.1 77.9 -95.5	123.3 309.2 0.3	291	0.366 0.0 1.0	34.9 77.9 -95.7	123.4 309.1
252	R31Y_037_037a	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.118 0.0	21.1 22.7 25.2	33.9 47.9	0.375 0.125 0.0	20.4 26.4 30.1	40.1 48.7 6.2	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9
253	R00Y_037_025a	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.124	24.5 19.2 16.1	25.1 40.0	0.375 0.125 0.125	20.7 27.8 14.8	31.5 28.0 9.5	389	1.0 0.0 0.0	50.4 76.9 64.5	100.0 40.0
254	R00Y_037_025a	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.25	24.9 20.2 1.0	20.3 2.9	0.375 0.125 0.25	21.6 31.1 -4.9	31.5 351.0 12.8	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
255	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.124 0.375	26.2 23.5 -14.6	27.7 328.2	0.375 0.125 0.375	23.1 36.3 -23.1	43.0 327.5 15.6	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
256	B34R_050_037a	0.375 0.125 0.5	0.5 0.375 0.312	311	0.381 0.124 0.5	28.7 31.5 -29.7	43.3 316.7	0.375 0.125 0.5	25.1 42.8 -39.5	58.3 317.2 15.3	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7
257	B25R_062_050a	0.375 0.125 0.625	0.625 0.5 0.375	300	0.375 0.125 0.625	31.2 39.9 -44.8	60.0 311.6	0.375 0.125 0.625	27.6 50.0 -54.4	73.9 312.5 14.4	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
258	B19R_075_062a	0.375 0.125 0.75	0.75 0.625 0.437	293	0.364 0.125 0.75	34.0 48.8 -59.4	76.9 309.3	0.375 0.125 0.75	34.0 57.5 -68.1	89.1 310.2 12.8	292	0.383 0.0 1.0	35.3 78.1 -95.1	123.0 309.3
259	B15R_087_075a	0.375 0.125 0.875	0.875 0.75 0.5	289	0.362 0.125 0.875	37.4 58.1 -73.1	93.4 308.4	0.375 0.125 0.875	33.6 65.1 -80.7	103.7 308.9 11.0	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
260	B13R_100_087a	0.375 0.125 1.0	1.0 0.875 0.562	286	0.358 0.125 1.0	40.7 67.3 -86.8	109.9 307.8	0.375 0.125 1.0	36.9 72.6 -92.6	117.7 308.1 8.7	284	0.266 0.0 1.0	32.9 77.0 -99.2	125.6 307.8
261	R68Y_037_037a	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.256 0.0	27.5 6.9 29.1	29.9 76.5	0.375 0.25 0.0	27.8 8.3 37.5	38.4 77.4 8.5	71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5
262	R50Y_037_025a	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.25 0.124	27.8 10.3 17.7	20.5 59.7	0.375 0.25 0.125	28.1 9.8 23.7	25.7 67.5 6.0	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
263	R00Y_037_012a	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.249	30.1 9.6 8.0	12.5 40.0	0.375 0.25 0.25	28.7 13.3 5.4	14.4 22.0 4.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
264	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.249 0.375	31.0 11.7 -7.3	13.8 328.2	0.375 0.25 0.375	29.7 19.0 -12.7	22.9 326.1 9.1	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
265	B25R_050_025a	0.375 0.25 0.5	0.5 0.25 0.375	300	0.375 0.249 0.5	33.5 19.9 -22.4	30.0 311.6	0.375 0.25 0.5	31.2 26.3 -29.7	39.7 311.5 9.9	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
266	B15R_062_037a	0.375 0.25 0.625	0.625 0.375 0.437	289	0.368 0.25 0.625	36.5 29.0 -36.5	46.7 308.4	0.375 0.25 0.625	33.2 34.6 -45.4	57.0 307.3 10.9	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
267	B11R_075_050a	0.375 0.25 0.75	0.75 0.5 0.5	284	0.366 0.25 0.75	40.0 38.3 -50.0	63.1 307.4	0.375 0.25 0.75	35.4 43.3 -59.8	73.9 305.9 11.8	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
268	B09R_087_062a	0.375 0.25 0.875	0.875 0.625 0.562	281	0.364 0.25 0.875	43.7 47.8 -63.2	79.3 307.0	0.375 0.25 0.875	38.0 52.2 -73.3	90.0 305.4 12.3	279	0.183 0.0 1.0	31.7 76.5 -101.2	126.9 307.0
269	B07R_100_075a	0.375 0.25 1.0	1.0 0.75 0.625	279	0.362 0.25 1.0	47.3 57.2 -76.4	95.5 306.8	0.375 0.25 1.0	40.9 60.9 -86.0	105.4 305.3 12.0	278	0.15 0.0 1.0	31.3 76.3 -101.9	127.4 306.8
270	Y00G_037_037a	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.375 0.0	34.7 -7.7 34.0	34.9 102.8	0.375 0.375 0.0	36.9 -10.0 44.2	45.3 102.8 10.7	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
271	Y00G_037_025a	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.375 0.124	35.0 -5.1 22.6	23.2 102.8	0.375 0.375 0.125	37.1 -8.7 33.8	34.9 104.4 11.8	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
272	Y00G_037_012a	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.375 0.249	35.4 -2.5 11.3	11.6 102.8	0.375 0.375 0.25	37.5 -5.4 17.5	18.3 107.1 7.1	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
273	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	35.7 0.0 0.0	0.0 0.0	0.375 0.375 0.375	38.3 0.0 0.0	0.0 325.3 2.5	360	1.0 1.0 1.0	95.4 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	39.5 9.5 -12.9	16.0 306.2	0.375 0.375 0.5	39.4 7.2 -17.0	18.5 292.9 4.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
275	B00R_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	43.3 19.0 -25.8	32.1 306.2	0.375 0.375 0.625	40.8 15.7 -33.2	36.8 295.4 8.4	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
276	B00R_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	47.1 28.5 -38.8	48.1 306.2	0.375 0.375 0.75	42.5 25.1 -48.4	54.5 297.4 11.1	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
277	B00R_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	50.9 38.0 -51.7	64.2 306.2	0.375 0.375 0.875	44.6 34.8 -62.7	71.7 299.0 13.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
278	B00R_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	54.7 47.5 -64.7	80.3 306.2	0.375 0.375 1.0	46.8 44.5 -76.1	88.2 300.3 14.2	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
279	Y23G_050_050a	0.375 0.5 0.0	0.5 0.5 0.25	104	0.383 0.5 0.0	44.3 -21.6 43.1	48.2 116.6	0.375 0.5 0.0	46.6 -26.1 51.4	57.7 116.9 9.7	102	0.766 1.0 0.0	88.7 -43.3 86.2	96.5 116.6
280	Y31G_050_037a	0.375 0.5 0.125	0.5 0.375 0.312	109	0.381 0.5 0.124	44.8 -19.0 31.8	37.1 120.8	0.375 0.5 0.125	46.7 -25.0 43.6	50.2 119.8 13.3	108	0.683 1.0 0.0	87.6 -50.7 84.9	98.9 120.8
281	Y50G_050_025a	0.375 0.5 0.25	0.5 0.25 0.375	120	0.375 0.5 0.249	45.2 -16.3 20.6	26.2 128.3	0.375 0.5 0.25	47.0 -22.1 29.6	36.9 126.8 10.8	119	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
282	G00B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	46.2 -10.3 9.9	14.3 136.0	0.375 0.5 0.375	47.6 -17.3 13.1	21.8 142.8 7.8	149	0.0 1.0 0.0	83.6 -82.7 7	

http://130.149.60.45/~farbmetrik/PS31/PS31L0NP.PDF /.PS; salida de transferencia
N: ninguna 3D-linealización (OL) en archivo (F) o PS-startup (S), página 9/18

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	0.25	390	0.5	0.0	0.0	50.4	76.9	64.5	100.4	40.0	
325	R26Y_050_050a	0.5	0.0	0.125	0.5	0.5	0.25	376	1.0	0.0	0.233	50.8	78.0	41.2	88.2	27.8	
326	R00Y_050_050a	0.5	0.0	0.25	0.5	0.5	0.25	360	1.0	0.0	0.5	52.0	81.1	4.1	81.2	2.9	
327	B61R_050_050a	0.5	0.0	0.375	0.5	0.5	0.25	344	1.0	0.0	0.766	54.4	87.3	-30.6	92.5	340.6	
328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.25	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	
329	B40R_062_062a	0.5	0.0	0.625	0.625	0.625	0.312	319	0.816	0.0	1.0	49.8	88.1	-70.7	113.0	321.2	
330	B34R_075_075a	0.5	0.0	0.75	0.75	0.75	0.375	311	0.683	0.0	1.0	44.8	84.1	-79.2	115.5	316.7	
331	B29R_087_087a	0.5	0.0	0.875	0.875	0.875	0.437	305	0.583	0.0	1.0	41.3	81.6	-85.1	117.9	313.8	
332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	300	0.5	0.0	1.0	38.5	79.8	-89.7	120.0	311.6	
333	R23Y_050_050a	0.5	0.125	0.0	0.5	0.5	0.25	44	1.0	0.233	0.0	53.7	67.6	65.8	94.4	44.2	
334	R00Y_050_037a	0.5	0.125	0.125	0.5	0.375	0.312	390	1.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0	
335	R18Y_050_037a	0.5	0.125	0.25	0.5	0.375	0.312	371	1.0	0.0	0.316	51.1	79.1	29.7	84.5	20.6	
336	B65R_050_037a	0.5	0.125	0.375	0.5	0.375	0.312	349	1.0	0.0	0.683	53.5	85.4	-19.9	87.7	346.8	
337	B50R_050_037a	0.5	0.125	0.5	0.5	0.375	0.312	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	
338	B38R_062_050a	0.5	0.125	0.625	0.625	0.5	0.375	316	0.766	0.0	1.0	47.9	86.4	-74.0	113.8	319.4	
339	B30R_075_062a	0.5	0.125	0.75	0.75	0.625	0.437	307	0.616	0.0	1.0	42.4	82.3	-83.2	117.0	314.6	
340	B25R_087_075a	0.5	0.125	0.875	0.875	0.75	0.5	300	0.5	0.0	1.0	38.5	79.8	-89.7	120.0	311.6	
341	B20R_100_087a	0.5	0.125	1.0	1.0	0.875	0.562	295	0.416	0.0	1.0	36.3	78.6	-93.5	122.2	310.0	
342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	60	1.0	0.5	0.0	63.6	41.3	71.0	82.2	59.7	
343	R31Y_050_037a	0.5	0.25	0.125	0.5	0.375	0.312	49	1.0	0.316	0.0	56.2	60.6	67.2	90.5	47.9	
344	R00Y_050_025a	0.5	0.25	0.25	0.5	0.25	0.375	390	1.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0	
345	R00Y_050_025a	0.5	0.25	0.375	0.5	0.25	0.375	360	1.0	0.0	0.5	52.0	81.1	4.1	81.2	2.9	
346	B50R_050_025a	0.5	0.25	0.5	0.5	0.25	0.375	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	
347	B34R_062_037a	0.5	0.25	0.625	0.625	0.375	0.437	311	0.683	0.0	1.0	44.8	84.1	-79.2	115.5	316.7	
348	B25R_075_050a	0.5	0.25	0.75	0.75	0.5	0.5	300	0.5	0.0	1.0	38.5	79.8	-89.7	120.0	311.6	
349	B19R_087_062a	0.5	0.25	0.875	0.875	0.625	0.562	293	0.489	0.0	1.0	35.3	78.1	-95.1	123.0	309.3	
350	B15R_100_075a	0.5	0.25	1.0	1.0	0.75	0.625	289	0.487	0.0	1.0	33.9	77.4	-97.5	124.5	308.4	
351	R76Y_050_050a	0.5	0.375	0.0	0.5	0.5	0.25	76	0.5	0.383	0.0	40.2	4.9	48.0	84.2	78.7	
352	R68Y_050_037a	0.5	0.375	0.125	0.5	0.375	0.312	71	0.7	0.666	0.0	78.2	7.8	80.6	81.0	84.4	
353	R50Y_050_025a	0.5	0.375	0.25	0.5	0.25	0.375	60	0.5	0.5	0.0	63.6	41.3	71.0	82.2	59.7	
354	R00Y_050_012a	0.5	0.375	0.375	0.5	0.125	0.437	390	1.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0	
355	B50R_050_012a	0.5	0.375	0.5	0.5	0.125	0.437	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	
356	B25R_062_025a	0.5	0.375	0.625	0.625	0.25	0.5	300	0.5	0.0	1.0	38.5	79.8	-89.7	120.0	311.6	
357	B15R_075_037a	0.5	0.375	0.75	0.75	0.375	0.562	289	0.316	0.0	1.0	33.9	77.4	-97.5	124.5	308.4	
358	B11R_087_050a	0.5	0.375	0.875	0.875	0.5	0.625	284	0.233	0.0	1.0	32.3	76.7	-100.1	126.2	307.4	
359	B09R_100_062a	0.5	0.375	1.0	1.0	0.625	0.687	281	0.183	0.0	1.0	31.7	76.5	-101.2	126.9	307.0	
360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	90	1.0	1.0	0.0	92.6	-20.7	90.7	93.0	102.8	
361	Y00G_050_037a	0.5	0.5	0.125	0.5	0.375	0.312	90	1.0	1.0	0.0	92.6	-20.7	90.7	93.0	102.8	
362	Y00G_050_025a	0.5	0.5	0.25	0.5	0.25	0.375	90	1.0	1.0	0.0	92.6	-20.7	90.7	93.0	102.8	
363	Y00G_050_012a	0.5	0.5	0.375	0.5	0.125	0.437	90	1.0	1.0	0.0	92.6	-20.7	90.7	93.0	102.8	
364	NW_050a	0.5	0.5	0.5	0.5	0.0	0.5	360	0.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0	
365	B00R_062_012a	0.5	0.625	0.625	0.625	0.125	0.562	270	0.5	0.5	0.625	51.6	6.7	-16.3	17.6	292.4	4.3
366	B00R_075_025a	0.5	0.5	0.75	0.75	0.25	0.625	270	0.0	0.0	1.0	30.3	76.0	-103.5	128.5	306.2	
367	B00R_087_037a	0.5	0.5	0.875	0.875	0.375	0.687	270	0.0	0.0	1.0	30.3	76.0	-103.5	128.5	306.2	
368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	270	0.0	0.0	1.0	30.3	76.0	-103.5	128.5	306.2	
369	Y18G_062_062a	0.5	0.625	0.0	0.625	0.625	0.312	101	0.816	1.0	0.0	89.5	-38.6	87.2	95.4	113.8	
370	Y23G_062_050a	0.5	0.625	0.125	0.625	0.5	0.375	104	0.766	1.0	0.0	88.7	-43.3	86.2	96.5	116.6	
371	Y31G_062_037a	0.5	0.625	0.25	0.625	0.375	0.437	109	0.683	1.0	0.0	87.6	-50.7	84.9	98.9	120.8	
372	Y50G_062_025a	0.5	0.625	0.375	0.625	0.25	0.5	120	0.5	1.0	0.0	85.7	-65.2	82.4	105.1	128.3	
373	G00B_062_012a	0.5	0.625	0.5	0.625	0.125	0.562	150	0.0	1.0	0.0	83.6	-82.7	79.8	115.0	136.0	
374	G50B_062_012a	0.5	0.625	0.625	0.625	0.125	0.562	210	0.0	1.0	1.0	86.8	-46.1	-13.5	48.1	196.3	
375	G75B_075_025a	0.5	0.625	0.75	0.75	0.25	0.625	240	0.0	0.5	1.0	51.7	18.3	-68.3	70.7	285.0	
376	G84B_087_037a	0.5	0.625	0.875	0.875	0.375	0.687	251	0.0	0.316	1.0	40.7	45.8	-86.7	98.1	297.8	
377	G88B_100_050a	0.5	0.625	1.0	1.0	0.5	0.75	256	0.5	0.616	1.0	36.3	14.0	-48.9	50.9	286.0	
378	Y31G_075_075a	0.5	0.75	0.0	0.75	0.75	0.375	109	0.637	1.0	0.0	63.7	-41.7	68.1	79.9	121.4	
379	Y38G_075_062a	0.5	0.75	0.125	0.75	0.625	0.437	113	0.5	0.75	0.125	67.4	-41.0	63.4	75.5	122.9	
380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	120	0.5	0.75	0.25	67.6	-38.2	53.4	66.3	126.3	
381	Y68G_075_037a	0.5	0.75	0.375	0.75	0.375	0.562	131	0.316	1.0	0.0	84.4	-75.3	80.9	110.6	132.9	
382	G00B_075_025a	0.5	0.75	0.5	0.75	0.25	0.625	150	0.5	0.75	0.5	68.3	-31.8	24.8	40.4	142.0	
383	G25B_075_025a	0.5	0.75	0.625	0.75	0.25	0.625	180	0.5	0.75	0.625	69.0	-26.2	9.2	27.8	160.5	
384	G50B_075_025a	0.5	0.75	0.75	0.75	0.25	0.625	210	0.5	0.75	0.75	69.8	-19.4	-6.2	20.3	197.8	
385	G65B_087_037a	0.5	0.75	0.875	0.875	0.375	0.687	229	0.5	0.75	0.875	70.7	-11.6	-21.4	24.3	214.8	
386	G75B_100_050a	0.5	0.75	1.0	1.0	0.5	0.75	240	0.5	0.75	1.0	71.9	-3.1	-36.1	36.3	265.0	
387	Y41G_087_087a	0.5	0.875	0.0	0.875	0.875	0.437	115	0.5	0.875	0.0	76.5	-54.0	75.2	92.6	125.6	
388	Y50G_087_075a	0.5	0.875	0.125	0.875	0.75	0.5	120	0.5	0.875	0.125	76.6	-53.4	71.4	89.1	126.8	
389	Y61G_087_062a	0.5	0.875	0.25	0.875	0.625	0.562	127	0.489	0.5	0.875	76.7	-51.9	62.8	81.5	129.5	
390	Y76G_087_050a	0.5	0.875	0.375	0.875	0.5	0.625	136	0.491	0.875	0.375	77.0	-49.2	50.7	70.7	134.1	
391	G00B_087_037a	0.5	0.875	0.5	0.875	0.375	0.687	150	0.5	0.875	0.5	77.4	-45.4	36.6	58.3	141.0	
392	G15B_087_037a	0.5	0.875	0.625	0.875	0.375	0.687	169	0.5	0.875	0.625	77.9	-40.3	21.7	45.8	151.7	
393	G34B_087_037a	0.5	0.875	0.75	0.875	0.375	0.687	191	0.5	0.875	0.75	78.5	-34.2	6.5	34.8	169.1	
394	G50B_087_037a	0.5	0.875	0.875	0.875	0.375	0.687	210	0.5	0.875	0.875	79.3	-27.1	-8.5	28.4	197.4	
395	G61B_100_050a	0.5	0.875	1.0	1.0	0.5	0.75	224	0.5	0.875	1.0	80.3	-19.1	-23.2	30.1	230.5	
396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	120	0.5	1.0	0.0	85.7	-65.2	82.4	105.1	128.3	
397	Y58G_100_087a	0.5	1.0	0.125	1.0	0.875	0.562	125	0.489	1.0	0.125	85.7	-64.7	79.1	102.2	129.2	
398	Y68G_100_075a	0.5	1.0	0.25	1.0	0.75	0.625	131	0.487	1.0	0.25						

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31.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
405	R00Y_062_062a	0.625 0.0 0.0	0.625 0.625 0.312	390	0.625 0.0 0.0	31.5 48.0 40.3	62.7 40.0	0.625 0.0 0.0	30.7 54.1 44.5	70.1 39.4 7.4	389	1.0 0.0 0.0 50.4 76.9 64.5 100.4 40.0
406	R31Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.114	31.7 48.7 29.7	57.0 31.3	0.625 0.0 0.125	31.0 54.7 30.0	62.4 28.7 6.0	380	1.0 0.0 0.0 0.183 50.7 77.9 47.5 91.2 31.3
407	R11Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.239	32.1 49.6 12.8	51.3 14.4	0.625 0.0 0.25	31.5 56.2 10.9	57.2 11.0 6.7	367	1.0 0.0 0.0 0.383 51.4 79.5 20.4 82.1 14.4
408	B69R_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.385	33.0 52.2 -7.1	52.7 352.1	0.625 0.0 0.375	32.4 58.6 -7.7	59.1 352.5 6.4	352	1.0 0.0 0.0 0.616 52.9 83.6 -11.4 84.3 352.1
409	B59R_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.625 0.0 0.51	34.3 55.5 -22.8	60.1 337.6	0.625 0.0 0.5	33.8 62.1 -25.0	67.0 338.0 6.9	339	1.0 0.0 0.0 0.816 54.9 88.9 -36.6 96.2 337.6
410	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	35.8 58.9 -36.5	63.9 328.2	0.625 0.0 0.625	35.5 66.4 -41.1	78.1 328.2 8.7	330	1.0 0.0 0.0 1.0 57.2 94.3 -58.4 110.9 328.2
411	B42R_075_075a	0.625 0.0 0.75	0.75 0.75 0.375	321	0.637 0.0 0.75	38.4 66.8 -51.4	84.3 322.4	0.625 0.0 0.75	37.6 71.3 -55.9	90.6 321.8 6.4	322	0.85 0.0 1.0 51.2 89.1 -68.5 112.4 322.4
412	B36R_087_087a	0.625 0.0 0.875	0.875 0.875 0.375	314	0.641 0.0 0.875	40.8 74.7 -66.6	100.1 318.3	0.625 0.0 0.875	40.0 76.7 -69.8	103.7 317.7 3.8	315	0.733 0.0 1.0 46.6 85.4 -76.1 114.4 318.3
413	B31R_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	43.0 82.7 -82.2	116.6 315.1	0.625 0.0 1.0	42.7 82.5 -82.8	116.8 314.8 0.6	308	0.633 0.0 1.0 43.0 82.7 -82.2 116.6 315.1
414	R18Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	41	0.625 0.114 0.0	32.9 44.0 40.9	60.1 42.8	0.625 0.125 0.0	32.8 48.2 45.9	66.6 34.6 6.5	39	1.0 0.183 0.0 52.7 70.5 65.5 96.2 42.8
415	R00Y_062_050a	0.625 0.125 0.125	0.625 0.5 0.375	390	0.625 0.125 0.125	37.1 38.4 32.2	50.2 40.0	0.625 0.125 0.125	33.0 48.8 32.2	58.5 33.3 11.1	389	1.0 0.0 0.0 50.4 76.9 64.5 100.4 40.0
416	R26Y_062_050a	0.625 0.125 0.25	0.625 0.5 0.375	376	0.625 0.125 0.241	37.3 39.0 20.6	44.1 27.8	0.625 0.125 0.25	33.5 50.4 13.6	52.2 15.1 13.9	377	1.0 0.0 0.0 0.233 50.7 77.9 47.5 91.2 31.3
417	R00Y_062_050a	0.625 0.125 0.375	0.625 0.5 0.375	360	0.625 0.125 0.375	37.9 40.5 2.0	40.6 2.9	0.625 0.125 0.375	34.4 53.1 -4.8	53.3 354.8 14.7	360	1.0 0.0 0.0 0.5 52.0 81.1 41.1 81.2 2.9
418	B61R_062_050a	0.625 0.125 0.5	0.625 0.5 0.375	344	0.625 0.125 0.508	39.1 43.6 -15.3	46.2 340.6	0.625 0.125 0.5	35.6 56.7 -22.2	60.9 338.6 15.2	342	1.0 0.0 0.0 0.766 54.4 87.3 -30.6 92.5 340.6
419	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	40.5 47.1 -29.2	55.4 328.2	0.625 0.125 0.625	37.3 61.3 -38.3	72.3 327.9 17.2	330	1.0 0.0 0.0 1.0 57.2 94.3 -58.4 110.9 328.2
420	B40R_075_062a	0.625 0.125 0.75	0.75 0.625 0.437	319	0.635 0.125 0.75	43.1 55.0 -44.2	70.6 321.2	0.625 0.125 0.75	39.2 66.6 -53.4	85.3 321.2 15.2	320	0.816 0.0 1.0 49.8 88.1 -70.7 113.0 321.2
421	B34R_087_075a	0.625 0.125 0.875	0.875 0.75 0.5	311	0.637 0.125 0.875	45.5 63.1 -59.4	86.6 316.7	0.625 0.125 0.875	41.5 72.3 -67.4	98.9 317.0 12.9	311	0.683 0.0 1.0 44.8 84.1 -79.2 115.5 316.7
422	B29R_100_087a	0.625 0.125 1.0	1.0 0.875 0.562	305	0.635 0.125 1.0	48.0 71.4 -74.4	103.2 313.8	0.625 0.125 1.0	44.0 78.4 -80.5	112.4 314.2 10.1	305	0.583 0.0 1.0 41.3 81.6 -85.1 117.9 313.8
423	R38Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	53	0.625 0.239 0.0	36.6 34.0 42.6	54.6 51.3	0.625 0.25 0.0	37.4 35.7 48.5	60.2 53.5 6.1	52	1.0 0.383 0.0 58.5 54.5 68.2 87.3 51.3
424	R23Y_062_050a	0.625 0.25 0.125	0.625 0.5 0.375	44	0.625 0.241 0.125	38.8 33.8 32.9	47.2 44.2	0.625 0.25 0.125	37.6 36.4 36.8	51.8 45.2 4.8	42	1.0 0.233 0.0 53.7 67.6 65.8 94.4 44.2
425	R00Y_062_037a	0.625 0.25 0.25	0.625 0.375 0.437	390	0.625 0.25 0.25	42.7 28.8 24.2	37.6 40.0	0.625 0.25 0.25	38.0 38.2 19.6	42.9 27.1 11.4	389	1.0 0.0 0.0 50.4 76.9 64.5 100.4 40.0
426	R18Y_062_037a	0.625 0.25 0.375	0.625 0.375 0.437	371	0.625 0.25 0.368	43.0 29.6 11.1	31.7 20.6	0.625 0.25 0.375	38.7 41.1 1.5	41.1 21.1 15.5	371	1.0 0.0 0.0 0.316 51.1 79.1 29.7 84.5 20.6
427	B65R_062_037a	0.625 0.25 0.5	0.625 0.375 0.437	349	0.625 0.25 0.506	43.9 32.0 -7.4	32.9 346.8	0.625 0.25 0.5	39.8 45.1 -15.7	47.8 340.7 16.0	348	1.0 0.0 0.0 0.683 53.5 85.4 -19.9 87.7 346.8
428	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	45.3 35.3 -21.9	41.6 328.2	0.625 0.25 0.625	41.2 50.2 -32.1	59.6 327.4 18.5	330	1.0 0.0 0.0 1.0 57.2 94.3 -58.4 110.9 328.2
429	B38R_075_050a	0.625 0.25 0.75	0.75 0.5 0.5	316	0.633 0.25 0.75	47.8 43.2 -37.0	56.9 319.4	0.625 0.25 0.75	42.9 56.0 -47.4	73.4 319.7 17.2	317	0.766 0.0 1.0 47.9 86.4 -74.0 113.8 319.4
430	B30R_087_062a	0.625 0.25 0.875	0.875 0.625 0.562	307	0.635 0.25 0.875	50.3 51.4 -52.0	73.1 314.6	0.625 0.25 0.875	44.9 62.4 -61.8	87.9 315.2 15.7	307	0.616 0.0 1.0 42.4 82.3 -83.2 117.0 314.6
431	B25R_100_075a	0.625 0.25 1.0	1.0 0.75 0.625	300	0.625 0.25 1.0	52.8 59.8 -67.2	90.0 311.6	0.625 0.25 1.0	47.2 69.2 -75.4	102.3 312.5 13.5	300	0.5 0.0 1.0 38.5 79.8 -89.7 120.0 311.6
432	R61Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	67	0.625 0.385 0.0	43.5 16.7 46.8	49.7 70.2	0.625 0.375 0.0	44.1 19.3 52.4	55.9 69.7 6.2	67	1.0 0.616 0.0 69.6 26.8 74.8 79.5 70.2
433	R50Y_062_050a	0.625 0.375 0.125	0.625 0.5 0.375	60	0.625 0.375 0.125	43.7 20.6 35.5	41.1 59.7	0.625 0.375 0.125	44.2 20.0 43.2	47.6 65.1 7.7	59	1.0 0.5 0.0 63.6 41.3 71.0 82.2 59.7
434	R31Y_062_037a	0.625 0.375 0.25	0.625 0.375 0.437	49	0.625 0.368 0.25	44.9 22.7 25.2	33.9 47.9	0.625 0.375 0.25	44.5 21.8 27.8	35.4 51.9 2.8	48	1.0 0.316 0.0 56.2 60.6 67.2 90.5 47.9
435	R00Y_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	48.4 19.2 16.1	25.1 40.0	0.625 0.375 0.375	45.1 24.9 10.6	27.0 23.1 8.5	389	1.0 0.0 0.0 50.4 76.9 64.5 100.4 40.0
436	R00Y_062_025a	0.625 0.375 0.5	0.625 0.25 0.5	360	0.625 0.375 0.5	48.7 20.2 1.0	20.3 2.9	0.625 0.375 0.5	46.0 29.2 -6.4	29.9 347.5 11.9	360	1.0 0.0 0.5 52.0 81.1 4.1 81.2 2.9
437	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	50.1 23.5 -14.6	27.7 328.2	0.625 0.375 0.625	47.1 34.6 -22.9	41.5 326.5 14.1	330	1.0 0.0 1.0 57.2 94.3 -58.4 110.9 328.2
438	B34R_075_037a	0.625 0.375 0.75	0.75 0.375 0.562	311	0.631 0.375 0.75	52.5 31.5 -29.7	43.3 316.7	0.625 0.375 0.75	48.5 40.9 -38.5	56.2 316.7 13.5	311	0.683 0.0 1.0 44.8 84.1 -79.2 115.5 316.7
439	B25R_087_050a	0.625 0.375 0.875	0.875 0.5 0.625	300	0.625 0.375 0.875	55.0 39.9 -44.8	60.0 311.6	0.625 0.375 0.875	50.2 47.9 -53.3	71.7 311.9 12.6	300	0.5 0.0 1.0 38.5 79.8 -89.7 120.0 311.6
440	B19R_100_062a	0.625 0.375 1.0	1.0 0.625 0.687	293	0.614 0.375 1.0	57.8 48.8 -59.4	76.9 309.3	0.625 0.375 1.0	52.1 55.3 -67.3	87.1 309.4 11.7	292	0.383 0.0 1.0 35.3 78.1 -95.1 123.0 309.3
441	R81Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	79	0.625 0.51 0.0	50.8 1.0 51.8	51.8 88.7	0.625 0.5 0.0	51.9 1.9 57.7	57.8 88.0 6.0	80	1.0 0.816 0.0 81.2 1.7 82.9 83.0 88.7
442	R76Y_062_050a	0.625 0.5 0.125	0.625 0.5 0.375	76	0.625 0.508 0.125	51.0						

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31L0NP.PDF /.PS>
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	0.75	0.75	5.5	389	1.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0
487	R35Y_075_075a	0.75	0.0	0.125	0.75	0.75	0.75	0.75	4.2	382	1.0	0.0	0.15	50.6	77.6	51.7	93.3	33.6
488	R18Y_075_075a	0.75	0.0	0.25	0.75	0.75	0.75	0.75	3.1	371	1.0	0.0	0.316	51.1	79.1	29.7	84.5	20.6
489	R00Y_075_075a	0.75	0.0	0.375	0.75	0.75	0.75	0.75	2.4	360	1.0	0.0	0.5	52.0	81.1	4.1	81.2	2.9
490	B65R_075_075a	0.75	0.0	0.5	0.75	0.75	0.75	0.75	1.8	349	1.0	0.0	0.683	53.5	85.4	-19.9	87.7	346.8
491	B57R_075_075a	0.75	0.0	0.625	0.75	0.75	0.75	0.75	1.3	338	1.0	0.0	0.85	55.3	89.8	-40.7	98.6	335.5
492	B50R_075_075a	0.75	0.0	0.75	0.75	0.75	0.75	0.75	1.0	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2
493	B43R_087_087a	0.75	0.0	0.875	0.875	0.875	0.875	0.875	0.8	322	0.866	0.0	1.0	51.8	89.6	-67.4	112.1	323.0
494	B38R_100_100a	0.75	0.0	1.0	1.0	1.0	1.0	1.0	0.7	317	0.766	0.0	1.0	47.9	86.4	-74.0	113.8	319.4
495	R15Y_075_075a	0.75	0.125	0.0	0.75	0.75	0.375	39	0.75	0.112	0.0	0.0	0.75	51.2	81.2	65.2	97.4	41.9
496	R00Y_075_062a	0.75	0.125	0.125	0.75	0.625	0.437	390	0.75	0.125	0.125	39.3	57.8	40.4	70.6	34.9	10.0	50.4
497	R31Y_075_062a	0.75	0.125	0.25	0.75	0.625	0.437	379	0.75	0.125	0.25	39.0	57.8	40.4	70.6	34.9	10.0	50.4
498	R11Y_075_062a	0.75	0.125	0.375	0.75	0.625	0.437	367	0.75	0.125	0.375	40.2	61.0	4.6	61.2	14.5	367	0.0
499	B69R_075_062a	0.75	0.125	0.5	0.75	0.625	0.437	353	0.75	0.125	0.5	41.4	64.0	-12.9	65.2	34.85	13.5	352
500	B59R_075_062a	0.75	0.125	0.625	0.75	0.625	0.437	341	0.75	0.125	0.625	42.7	67.7	-29.4	73.8	36.4	14.2	339
501	B50R_075_062a	0.75	0.125	0.75	0.75	0.625	0.437	330	0.75	0.125	0.75	44.3	72.1	-44.9	84.9	32.80	15.9	330
502	B42R_087_075a	0.75	0.125	0.875	0.875	0.75	0.5	321	0.762	0.125	0.875	50.3	66.8	-51.4	84.3	32.24	13.6	322
503	B36R_100_087a	0.75	0.125	1.0	1.0	0.875	0.562	314	0.766	0.125	1.0	52.7	74.7	-66.6	100.1	31.83	10.9	315
504	R31Y_075_075a	0.75	0.25	0.0	0.75	0.75	0.375	49	0.75	0.237	0.0	42.2	45.5	50.4	67.9	47.9	0.75	0.237
505	R18Y_075_062a	0.75	0.25	0.125	0.75	0.625	0.437	41	0.75	0.239	0.125	44.8	44.0	40.9	60.1	42.8	0.75	0.239
506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	390	0.75	0.25	0.25	49.0	38.4	32.2	50.2	40.0	0.75	0.25
507	R26Y_075_050a	0.75	0.25	0.375	0.75	0.5	0.5	376	0.75	0.25	0.366	49.2	39.0	20.6	44.1	27.8	0.75	0.25
508	R00Y_075_050a	0.75	0.25	0.5	0.75	0.5	0.5	360	0.75	0.25	0.5	49.8	40.5	2.0	46.6	2.9	0.75	0.25
509	B61R_075_050a	0.75	0.25	0.625	0.75	0.5	0.5	344	0.75	0.25	0.633	51.0	43.6	-15.3	46.2	340.6	0.75	0.25
510	B50R_075_050a	0.75	0.25	0.75	0.75	0.5	0.5	330	0.75	0.25	0.75	52.5	47.1	-29.2	55.4	328.2	0.75	0.25
511	B40R_087_062a	0.75	0.25	0.875	0.875	0.625	0.562	319	0.76	0.25	0.875	55.0	-44.2	70.6	32.17	0.75	0.25	
512	B34R_100_075a	0.75	0.25	1.0	1.0	0.75	0.625	311	0.762	0.25	1.0	57.4	63.1	-59.4	86.6	316.2	0.75	0.25
513	R50Y_075_075a	0.75	0.375	0.0	0.75	0.75	0.375	60	0.75	0.375	0.0	47.7	31.0	53.2	61.6	59.7	0.75	0.375
514	R38Y_075_062a	0.75	0.375	0.125	0.75	0.625	0.437	53	0.75	0.364	0.125	48.5	34.0	42.6	54.6	51.3	0.75	0.375
515	R23Y_075_050a	0.75	0.375	0.25	0.75	0.5	0.5	44	0.75	0.366	0.25	50.7	33.8	32.9	47.2	44.2	0.75	0.375
516	R00Y_075_037a	0.75	0.375	0.375	0.75	0.375	0.562	390	0.75	0.375	0.375	54.7	28.8	24.2	37.6	40.0	0.75	0.375
517	R18Y_075_037a	0.75	0.375	0.5	0.75	0.375	0.562	371	0.75	0.375	0.493	54.9	29.6	11.1	31.7	20.6	0.75	0.375
518	B65R_075_037a	0.75	0.375	0.625	0.75	0.375	0.562	349	0.75	0.375	0.631	55.8	32.0	-7.4	32.9	346.8	0.75	0.375
519	B50R_075_037a	0.75	0.375	0.75	0.75	0.375	0.562	330	0.75	0.375	0.75	57.2	35.3	-21.9	41.6	328.2	0.75	0.375
520	B38R_087_050a	0.75	0.375	0.875	0.875	0.5	0.625	316	0.758	0.375	0.875	59.7	43.2	-37.0	56.9	319.4	0.75	0.375
521	B30R_100_062a	0.75	0.375	1.0	1.0	0.625	0.687	307	0.76	0.375	1.0	62.2	51.4	-52.0	73.1	314.6	0.75	0.375
522	R68Y_075_075a	0.75	0.5	0.0	0.75	0.75	0.375	71	0.75	0.512	0.0	55.0	13.8	58.2	59.8	76.5	0.75	0.5
523	R61Y_075_062a	0.75	0.5	0.125	0.75	0.625	0.437	67	0.75	0.51	0.125	55.4	16.7	46.8	49.7	70.2	0.75	0.5
524	R50Y_075_050a	0.75	0.5	0.25	0.75	0.5	0.5	60	0.75	0.5	0.25	55.6	20.6	35.5	41.1	59.7	0.75	0.5
525	R31Y_075_037a	0.75	0.5	0.375	0.75	0.375	0.562	49	0.75	0.493	0.375	56.8	22.7	25.2	33.9	47.9	0.75	0.5
526	R00Y_075_025a	0.75	0.5	0.5	0.75	0.25	0.625	390	0.75	0.5	0.5	56.8	19.2	16.1	25.1	40.0	0.75	0.5
527	R00Y_075_025a	0.75	0.5	0.625	0.75	0.25	0.625	360	0.75	0.5	0.625	60.7	20.2	1.0	20.3	2.9	0.75	0.5
528	B50R_075_025a	0.75	0.5	0.75	0.75	0.25	0.625	330	0.75	0.5	0.75	62.0	23.5	-14.6	27.7	328.2	0.75	0.5
529	B34R_087_037a	0.75	0.5	0.875	0.875	0.375	0.687	311	0.756	0.5	0.875	64.5	31.5	-29.7	43.3	316.7	0.75	0.5
530	B25R_100_050a	0.75	0.5	1.0	1.0	0.5	0.75	300	0.75	0.5	1.0	67.0	39.9	-44.8	60.0	311.6	0.75	0.5
531	R85Y_075_075a	0.75	0.625	0.0	0.75	0.75	0.375	81	0.75	0.637	0.0	62.4	-1.8	63.2	63.2	91.7	0.75	0.625
532	R81Y_075_062a	0.75	0.625	0.125	0.75	0.625	0.437	79	0.75	0.635	0.125	62.7	1.0	51.8	51.8	88.7	0.75	0.625
533	R76Y_075_050a	0.75	0.625	0.25	0.75	0.5	0.5	76	0.75	0.633	0.25	62.9	3.9	40.3	40.5	84.4	0.75	0.625
534	R68Y_075_037a	0.75	0.625	0.375	0.75	0.375	0.562	71	0.75	0.631	0.375	63.3	6.9	29.1	29.9	76.5	0.75	0.625
535	R50Y_075_025a	0.75	0.625	0.5	0.75	0.25	0.625	60	0.75	0.625	0.5	63.6	10.3	17.7	20.5	59.7	0.75	0.625
536	R00Y_075_012a	0.75	0.625	0.625	0.75	0.125	0.687	390	0.75	0.625	0.625	65.9	9.6	8.0	12.5	40.0	0.75	0.625
537	B50R_075_012a	0.75	0.625	0.75	0.75	0.125	0.687	330	0.75	0.625	0.75	66.7	11.7	-7.3	13.8	328.2	0.75	0.625
538	B25R_087_025a	0.75	0.625	0.875	0.875	0.25	0.75	300	0.75	0.625	0.875	69.2	19.9	-22.4	30.0	311.6	0.75	0.625
539	B15R_100_037a	0.75	0.625	1.0	1.0	0.375	0.812	289	0.743	0.625	1.0	72.3	29.0	-36.5	46.7	308.4	0.75	0.625
540	Y00G_075_075a	0.75	0.75	0.0	0.75	0.75												

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31.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
567	R00Y_087_087a	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.0	44.1 67.3 56.4	87.8 40.0	0.875 0.0 0.0	44.1 69.5 58.3	39.9 2.9	389	50.4 76.9 64.5 100.4 40.0
568	R36Y_087_087a	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.116	44.2 67.7 47.1	82.5 34.8	0.875 0.0 0.125	44.2 69.9 47.2	84.3 34.0 2.2	382	50.6 77.3 53.9 94.3 34.8
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 68.5 32.2	75.7 25.1	0.875 0.0 0.25	44.5 70.8 30.2	77.0 23.1 2.9	375	50.9 78.3 36.8 86.6 25.1
570	R08Y_087_087a	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.364	45.1 70.2 13.8	71.6 11.1	0.875 0.0 0.375	45.1 72.4 12.2	73.4 9.5 2.6	365	51.0 80.3 15.8 81.8 11.1
571	B70R_087_087a	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.51	46.1 72.8 -6.0	73.0 355.2	0.875 0.0 0.5	46.0 74.6 -5.3	74.8 355.8 1.9	354	52.7 83.2 -6.9 83.5 355.2
572	B63R_087_087a	0.875 0.0 0.625	0.875 0.875 0.437	346	0.875 0.0 0.641	47.2 75.6 -23.1	79.1 342.9	0.875 0.0 0.625	47.1 77.6 -22.1	80.7 344.0 2.2	344	54.0 86.5 -26.4 90.4 342.9
573	B56R_087_087a	0.875 0.0 0.75	0.875 0.875 0.437	338	0.875 0.0 0.758	48.6 78.8 -37.5	87.3 334.5	0.875 0.0 0.75	48.5 81.2 -37.9	89.6 334.9 2.3	337	55.5 90.1 -42.8 99.8 334.5
574	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	50.1 82.5 -51.1	97.1 328.2	0.875 0.0 0.875	50.2 85.3 -52.8	100.3 328.2 3.2	330	57.2 94.3 -58.4 110.9 328.2
575	B44R_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	52.5 90.1 -66.3	111.9 323.6	0.875 0.0 1.0	52.1 89.8 -66.9	112.0 323.3 0.7	323	58.3 90.1 -66.3 111.9 323.6
576	R13Y_087_087a	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.116 0.0	45.2 64.2 56.9	85.8 41.5	0.875 0.125 0.0	45.3 65.8 58.8	88.3 41.7 2.5	37	51.7 73.4 65.0 98.0 41.5
577	R00Y_087_075a	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.125	49.7 57.7 48.4	75.3 40.0	0.875 0.125 0.125	45.5 66.2 48.3	81.9 36.0 9.5	389	50.4 76.9 64.5 100.4 40.0
578	R35Y_087_075a	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.237	49.9 58.2 38.8	69.9 33.6	0.875 0.125 0.25	45.8 67.1 31.8	74.3 25.3 12.0	382	50.6 77.6 51.7 93.3 33.6
579	R18Y_087_075a	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.362	50.2 59.3 22.3	63.4 20.6	0.875 0.125 0.375	46.4 68.8 13.9	70.2 11.4 13.2	371	51.0 80.3 15.8 81.8 11.1
580	R00Y_087_075a	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.5	50.9 60.8 3.1	60.9 2.9	0.875 0.125 0.5	47.2 71.1 -3.6	71.2 357.1 12.8	360	52.0 81.1 4.1 81.2 2.9
581	B65R_087_075a	0.875 0.125 0.625	0.875 0.75 0.5	349	0.875 0.125 0.637	52.1 64.1 -14.9	65.8 346.8	0.875 0.125 0.625	48.3 74.2 -20.3	76.9 344.6 12.0	348	53.5 85.4 -19.9 87.7 346.8
582	B57R_087_075a	0.875 0.125 0.75	0.875 0.75 0.5	339	0.875 0.125 0.762	53.4 67.3 -30.5	73.9 335.5	0.875 0.125 0.75	49.6 77.9 -36.1	85.9 335.1 12.5	337	55.3 89.8 -40.7 98.6 335.5
583	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	54.9 70.7 -43.8	83.2 328.2	0.875 0.125 0.875	51.3 82.1 -51.1	96.7 328.1 14.0	330	57.2 94.3 -58.4 110.9 328.2
584	B43R_100_087a	0.875 0.125 1.0	1.0 0.875 0.562	322	0.883 0.125 1.0	57.3 78.4 -59.0	98.1 323.0	0.875 0.125 1.0	53.1 86.9 -65.3	108.7 323.0 11.3	322	51.8 89.6 -67.4 112.1 323.0
585	R26Y_087_087a	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.233 0.0	47.0 57.0 58.0	81.3 45.5	0.875 0.25 0.0	48.4 57.3 60.1	83.0 46.3 2.1	44	54.6 65.1 66.3 93.0 45.5
586	R15Y_087_075a	0.875 0.25 0.125	0.875 0.75 0.5	39	0.875 0.237 0.125	50.9 54.3 48.9	73.1 41.9	0.875 0.25 0.125	48.5 57.7 50.8	76.9 41.3 4.5	37	52.0 72.4 65.2 97.4 41.9
587	R00Y_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	55.4 48.0 40.3	62.7 40.0	0.875 0.25 0.25	48.8 58.7 35.3	68.5 31.0 13.4	389	50.4 76.9 64.5 100.4 40.0
588	R31Y_087_062a	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.364	55.5 48.7 29.7	57.0 31.3	0.875 0.25 0.375	49.3 60.4 17.9	63.0 16.5 17.7	380	50.7 77.9 47.5 91.2 31.3
589	R11Y_087_062a	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.489	55.9 49.6 12.8	51.3 14.4	0.875 0.25 0.5	50.1 63.0 0.6	63.0 0.5 18.9	367	51.4 79.5 20.4 82.1 14.4
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 52.2 -7.1	52.7 35.1	0.875 0.25 0.625	51.1 66.2 -16.0	68.1 346.3 17.5	352	53.6 -11.4 84.3 352.1
591	B59R_087_062a	0.875 0.25 0.75	0.875 0.625 0.562	341	0.875 0.25 0.76	58.2 55.5 -22.8	60.1 337.6	0.875 0.25 0.75	52.3 70.2 -31.9	77.1 335.5 18.1	339	54.9 88.9 -36.6 96.2 337.6
592	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	59.6 58.9 -36.5	69.3 328.2	0.875 0.25 0.875	53.8 74.7 -47.0	88.3 327.8 19.8	330	57.2 94.3 -58.4 110.9 328.2
593	B42R_100_075a	0.875 0.25 1.0	1.0 0.75 0.625	321	0.887 0.25 1.0	62.2 66.8 -51.4	84.3 322.4	0.875 0.25 1.0	55.6 79.8 -61.3	100.7 322.4 17.6	322	58.5 90.1 -68.5 112.4 322.4
594	R41Y_087_087a	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.364 0.0	52.5 44.4 60.6	75.1 53.7	0.875 0.375 0.0	53.2 44.5 62.6	76.8 54.5 2.1	54	60.0 50.7 69.3 85.9 53.7
595	R31Y_087_075a	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.362 0.125	54.1 45.5 50.4	67.9 47.9	0.875 0.375 0.125	53.3 44.9 54.7	70.8 50.6 4.4	48	56.2 60.6 67.2 90.5 47.9
596	R18Y_087_062a	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.364 0.25	56.8 44.0 40.9	60.1 42.8	0.875 0.375 0.25	53.5 45.9 40.7	61.4 41.5 3.7	39	52.7 70.5 65.5 96.2 42.8
597	R00Y_087_050a	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	61.0 38.4 32.2	50.2 40.0	0.875 0.375 0.375	54.0 47.8 24.1	53.6 26.8 14.2	389	50.4 76.9 64.5 100.4 40.0
598	R26Y_087_050a	0.875 0.375 0.5	0.875 0.5 0.625	376	0.875 0.375 0.491	61.1 39.0 20.6	44.1 27.8	0.875 0.375 0.5	54.6 50.5 7.2	51.0 8.1 18.8	377	50.8 78.0 41.2 88.2 27.8
599	R00Y_087_050a	0.875 0.375 0.625	0.875 0.5 0.625	360	0.875 0.375 0.625	61.8 40.5 2.0	40.6 2.9	0.875 0.375 0.625	55.5 54.0 -9.3	54.8 350.2 18.7	360	52.0 81.1 4.1 81.2 2.9
600	B61R_087_050a	0.875 0.375 0.75	0.875 0.5 0.625	344	0.875 0.375 0.758	62.9 43.6 -15.3	46.2 340.6	0.875 0.375 0.75	56.6 58.3 -25.2	63.5 336.5 18.7	342	54.4 87.3 -30.6 92.5 340.6
601	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	64.4 47.1 -29.2	55.4 328.2	0.875 0.375 0.875	58.0 63.2 -40.5	75.0 327.3 20.6	330	57.2 94.3 -58.4 110.9 328.2
602	B40R_100_062a	0.875 0.375 1.0	1.0 0.625 0.687	319	0.885 0.375 1.0	66.9 55.0 -44.2	70.6 321.2	0.875 0.375 1.0	59.5 68.6 -55.0	88.0 321.2 18.8	320	58.1 90.1 -70.7 110.3 321.2
603	R58Y_087_087a	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.51 0.0	59.4 27.1 64.8	70.2 67.2	0.875 0.5 0.0	59.4 29.0 66.2	72.3 66.2 2.4	65	67.9 31.0 74.0 80.3 67.2
604	R50Y_087_075a	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.5 0.125	59.6 31.0 53.2	61.6 59.7	0.875 0.5 0.125	59.4 29.5 59.8	66.7 63.7 6.7	59	63.6 41.3 71.0 82.2 59.7
605	R38Y_087_062a	0.875 0.5 0.25	0.875 0.625 0.562	53	0.875 0.489 0.25	60.4 34.0 42.6	54.6 51.3	0.875 0.5 0.25	59.7 30.6 47.4	56.4 57.1 5.9	52	58.5 54.5 68.2 87.3 51.3
606	R23Y_087_050a	0.875 0.5 0.375	0.875 0.5 0.625	44	0.875 0.491 0.375	62.6 33.8 32.9	47.2 44.2	0.875 0.5 0.375	60.0 32.5 31.9	45.6 44.4 3.0	42	53.7 67.6 65.8 94.4 44.2
607	R00Y_087_037a	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.5	66.6 28.8 24.2	37.6 40.0	0.875 0.5 0.5	60.6 35.3 15.5	38.6 23.7 12.3	389</	

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31.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													
648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	50.4	76.9	64.5	100.4	39.9	0.0	389	1.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0
649	R38Y_100_100a	1.0	0.0	0.125	1.0	1.0	0.5	383	1.0	0.0	0.116	50.5	77.2	55.6	95.1	35.7	0.0	383	1.0	0.0	0.116	50.5	77.2	55.6	95.1	35.7
650	R26Y_100_100a	1.0	0.0	0.25	1.0	1.0	0.5	376	1.0	0.0	0.233	50.8	78.0	41.2	88.2	27.8	0.0	377	1.0	0.0	0.233	50.8	78.0	41.2	88.2	27.8
651	R13Y_100_100a	1.0	0.0	0.375	1.0	1.0	0.5	368	1.0	0.0	0.366	51.3	79.3	22.7	82.5	16.0	0.0	368	1.0	0.0	0.366	51.3	79.3	22.7	82.5	16.0
652	R00Y_100_100a	1.0	0.0	0.5	1.0	1.0	0.5	360	1.0	0.0	0.5	52.0	81.1	4.1	81.2	2.9	0.0	360	1.0	0.0	0.5	52.0	81.1	4.1	81.2	2.9
653	B68R_100_100a	1.0	0.0	0.625	1.0	1.0	0.5	352	1.0	0.0	0.633	53.0	83.9	-13.6	85.0	350.7	0.0	351	1.0	0.0	0.633	53.0	83.9	-13.6	85.0	350.7
654	B61R_100_100a	1.0	0.0	0.75	1.0	1.0	0.5	344	1.0	0.0	0.766	54.4	87.3	-30.6	92.5	340.6	0.0	342	1.0	0.0	0.766	54.4	87.3	-30.6	92.5	340.6
655	B55R_100_100a	1.0	0.0	0.875	1.0	1.0	0.5	337	1.0	0.0	0.883	55.7	90.6	-44.8	101.1	333.6	0.0	336	1.0	0.0	0.883	55.7	90.6	-44.8	101.1	333.6
656	B50R_100_100a	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	0.0	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2
657	R11Y_100_100a	1.0	0.125	0.0	1.0	1.0	0.5	37	1.0	0.116	0.0	51.4	74.1	64.9	98.5	41.2	0.0	37	1.0	0.116	0.0	51.4	74.1	64.9	98.5	41.2
658	R00Y_100_087a	1.0	0.125	0.125	1.0	0.875	0.562	390	1.0	0.125	0.125	56.1	67.3	56.4	87.8	40.0	0.0	389	1.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0
659	R36Y_100_087a	1.0	0.125	0.25	1.0	0.875	0.562	382	1.0	0.125	0.241	56.2	67.7	47.1	82.5	34.8	0.0	382	1.0	0.0	0.133	50.6	77.3	53.9	94.3	34.8
660	R23Y_100_087a	1.0	0.125	0.375	1.0	0.875	0.562	374	1.0	0.125	0.358	56.4	68.5	32.2	75.7	25.1	0.0	375	1.0	0.0	0.266	50.9	78.3	36.8	86.6	25.1
661	R08Y_100_087a	1.0	0.125	0.5	1.0	0.875	0.562	365	1.0	0.125	0.489	57.0	70.2	13.8	71.6	11.1	0.0	365	1.0	0.0	0.416	51.5	80.3	15.8	81.8	11.1
662	B70R_100_087a	1.0	0.125	0.625	1.0	0.875	0.562	355	1.0	0.125	0.635	58.0	72.8	-6.0	73.0	355.2	0.0	354	1.0	0.0	0.583	52.7	83.2	-6.9	83.5	355.2
663	B63R_100_087a	1.0	0.125	0.75	1.0	0.875	0.562	346	1.0	0.125	0.766	59.2	75.6	-23.1	79.1	342.9	0.0	344	1.0	0.0	0.733	54.0	86.5	-26.4	90.4	342.9
664	B56R_100_087a	1.0	0.125	0.875	1.0	0.875	0.562	338	1.0	0.125	0.883	60.5	78.8	-37.5	87.3	334.5	0.0	337	1.0	0.0	0.866	55.5	90.1	-42.8	99.8	334.5
665	B50R_100_087a	1.0	0.125	1.0	1.0	0.875	0.562	330	1.0	0.125	1.0	62.0	82.5	-51.1	97.1	328.2	0.0	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2
666	R23Y_100_100a	1.0	0.25	0.0	1.0	1.0	0.5	44	1.0	0.233	0.0	53.7	67.6	65.8	94.4	44.2	0.0	42	1.0	0.233	0.0	53.7	67.6	65.8	94.4	44.2
667	R13Y_100_087a	1.0	0.25	0.125	1.0	0.875	0.562	38	1.0	0.241	0.125	57.1	64.2	56.9	85.8	41.5	0.0	37	1.0	0.133	0.0	51.7	73.4	65.0	98.0	41.5
668	R00Y_100_075a	1.0	0.25	0.25	1.0	0.75	0.625	390	1.0	0.25	0.25	61.7	57.7	48.4	75.3	40.0	0.0	389	1.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0
669	R35Y_100_075a	1.0	0.25	0.375	1.0	0.75	0.625	381	1.0	0.25	0.362	61.8	58.2	38.8	69.9	33.6	0.0	382	1.0	0.0	0.15	50.6	77.6	51.7	93.3	33.6
670	R18Y_100_075a	1.0	0.25	0.5	1.0	0.75	0.625	371	1.0	0.25	0.487	62.1	59.3	22.3	63.4	20.6	0.0	371	1.0	0.0	0.316	51.1	79.1	29.7	84.5	20.6
671	R00Y_100_075a	1.0	0.25	0.625	1.0	0.75	0.625	360	1.0	0.25	0.625	62.9	60.8	3.1	60.9	2.9	0.0	360	1.0	0.0	0.5	52.0	81.1	4.1	81.2	2.9
672	B65R_100_075a	1.0	0.25	0.75	1.0	0.75	0.625	349	1.0	0.25	0.762	64.0	64.1	-14.9	65.8	346.8	0.0	347	1.0	0.0	0.683	53.5	85.4	-19.9	87.7	346.8
673	B57R_100_075a	1.0	0.25	0.875	1.0	0.75	0.625	339	1.0	0.25	0.887	65.3	67.3	-30.5	73.9	335.5	0.0	337	1.0	0.0	0.85	53.8	89.8	-40.7	98.6	335.5
674	B50R_100_075a	1.0	0.25	1.0	1.0	0.75	0.625	330	1.0	0.25	1.0	66.8	70.7	-43.8	83.2	328.2	0.0	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2
675	R36Y_100_100a	1.0	0.375	0.0	1.0	1.0	0.5	52	1.0	0.366	0.0	57.9	56.2	67.9	88.1	50.3	0.0	51	1.0	0.366	0.0	57.9	56.2	67.9	88.1	50.3
676	R26Y_100_087a	1.0	0.375	0.125	1.0	0.875	0.562	46	1.0	0.358	0.125	59.7	57.0	58.0	81.3	45.5	0.0	44	1.0	0.266	0.0	54.6	65.1	66.3	93.0	45.5
677	R15Y_100_075a	1.0	0.375	0.25	1.0	0.75	0.625	39	1.0	0.362	0.25	62.8	54.3	48.9	73.1	41.9	0.0	37	1.0	0.15	0.0	52.0	72.4	65.2	97.4	41.9
678	R00Y_100_062a	1.0	0.375	0.375	1.0	0.625	0.687	390	1.0	0.375	0.375	67.3	48.0	40.3	62.7	40.0	0.0	389	1.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0
679	R31Y_100_062a	1.0	0.375	0.5	1.0	0.625	0.687	379	1.0	0.375	0.489	67.4	48.7	29.7	57.0	31.3	0.0	380	1.0	0.0	0.183	50.7	77.9	47.5	91.2	31.3
680	R11Y_100_062a	1.0	0.375	0.625	1.0	0.625	0.687	367	1.0	0.375	0.614	67.9	49.6	12.8	51.3	14.4	0.0	367	1.0	0.0	0.383	51.4	79.5	20.4	82.1	14.4
681	B69R_100_062a	1.0	0.375	0.75	1.0	0.625	0.687	353	1.0	0.375	0.76	68.8	52.2	-7.1	52.7	352.1	0.0	352	1.0	0.0	0.616	52.9	83.6	-11.4	84.3	352.1
682	B59R_100_062a	1.0	0.375	0.875	1.0	0.625	0.687	341	1.0	0.375	0.885	70.1	55.5	-22.8	60.1	337.6	0.0	349	1.0	0.0	0.816	54.9	88.9	-36.6	96.2	337.6
683	B50R_100_062a	1.0	0.375	1.0	1.0	0.625	0.687	330	1.0	0.375	1.0	71.5	58.9	-36.5	69.3	328.2	0.0	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2
684	R50Y_100_100a	1.0	0.5	0.0	1.0	1.0	0.5	60	1.0	0.5	0.0	63.6	41.3	71.0	82.2	59.7	0.0	59	1.0	0.5	0.0	63.6	41.3	71.0	82.2	59.7
685	R41Y_100_087a	1.0	0.5	0.125	1.0	0.875	0.562	55	1.0	0.489	0.125	64.4	44.4	60.6	75.1	53.7	0.0	54	1.0	0.416	0.0	60.0	50.7	69.3	85.9	53.7
686	R31Y_100_075a	1.0	0.5	0.25	1.0	0.75	0.625	49	1.0	0.487	0.25	66.0	45.5	50.4	67.9	47.9	0.0	48	1.0	0.316	0.0	56.2	60.6	67.2	90.5	47.9
687	R18Y_100_062a	1.0	0.5	0.375	1.0	0.625	0.687	41	1.0	0.489	0.375	68.7	44.0	40.9	60.1	42.8	0.0	40	1.0	0.183	0.0	52.7	70.5	65.5	96.2	42.8
688	R00Y_100_050a	1.0	0.5	0.5																						

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS31/PS31.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	95.4 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	360	1.0 1.0 1.0	95.4 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	94.3 -5.7 -1.6	6.0 196.3	0.875 1.0 1.0	93.3 -9.7 -3.3	10.3 198.8 4.4	210 0.0 1.0 1.0	86.8 -46.1 -13.5
731	G50B_100_025d	0.75 1.0 1.0	1.0 0.25 0.875	210	0.75 1.0 1.0	93.2 -11.5 -3.3	12.0 196.3	0.75 1.0 1.0	91.5 -18.9 -6.2	19.9 198.1 8.0	210 0.0 1.0 1.0	86.8 -46.1 -13.5
732	G50B_100_037d	0.625 1.0 1.0	1.0 0.375 0.812	210	0.625 1.0 1.0	92.2 -17.3 -5.0	18.0 196.3	0.625 1.0 1.0	90.0 -27.0 -8.5	28.3 197.6 10.5	210 0.0 1.0 1.0	86.8 -46.1 -13.5
733	G50B_100_050d	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	91.1 -23.0 -6.7	24.0 196.3	0.5 1.0 1.0	88.8 -33.9 -10.4	35.4 197.1 11.6	210 0.0 1.0 1.0	86.8 -46.1 -13.5
734	G50B_100_062d	0.375 1.0 1.0	1.0 0.625 0.687	210	0.375 1.0 1.0	90.0 -28.8 -8.4	30.0 196.3	0.375 1.0 1.0	87.9 -39.3 -11.8	41.0 196.8 11.1	210 0.0 1.0 1.0	86.8 -46.1 -13.5
735	G50B_100_075d	0.25 1.0 1.0	1.0 0.75 0.625	210	0.25 1.0 1.0	89.0 -34.6 -10.1	36.1 196.3	0.25 1.0 1.0	87.3 -43.0 -12.8	44.9 196.5 9.0	210 0.0 1.0 1.0	86.8 -46.1 -13.5
736	G50B_100_087d	0.125 1.0 1.0	1.0 0.875 0.562	210	0.125 1.0 1.0	87.9 -40.4 -11.8	42.1 196.3	0.125 1.0 1.0	87.0 -45.2 -13.3	47.2 196.4 5.1	210 0.0 1.0 1.0	86.8 -46.1 -13.5
737	G50B_100_100d	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3 0.0	210 0.0 1.0 1.0	86.8 -46.1 -13.5
738	RO0Y_100_012d	1.0 0.875 0.875	1.0 0.125 0.937	390	1.0 0.875 0.875	89.7 9.6 8.0	12.5 40.0	1.0 0.875 0.875	87.1 10.5 3.8	11.2 20.1 5.0	389 1.0 0.0 0.0	50.4 76.9 64.5
739	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	83.4 0.0 0.0	0.0 0.0	0.875 0.875 0.875	84.7 0.0 0.0	0.0 325.2 1.2	360 1.0 1.0 1.0	95.4 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	82.4 -5.7 -1.6	6.0 196.3	0.75 0.875 0.875	80.7 0.0 0.0	0.0 325.2 1.2	360 1.0 1.0 1.0	95.4 0.0 0.0
741	G50B_087_025d	0.625 0.875 0.875	0.875 0.25 0.75	210	0.625 0.875 0.875	81.3 -11.5 -3.3	12.0 196.3	0.625 0.875 0.875	80.7 -19.1 -6.2	20.1 197.9 8.1	210 0.0 1.0 1.0	86.8 -46.1 -13.5
742	G50B_087_037d	0.5 0.875 0.875	0.875 0.375 0.687	210	0.5 0.875 0.875	80.2 -17.3 -5.0	18.0 196.3	0.5 0.875 0.875	79.3 -27.1 -8.5	28.4 197.4 10.4	210 0.0 1.0 1.0	86.8 -46.1 -13.5
743	G50B_087_050d	0.375 0.875 0.875	0.875 0.5 0.625	210	0.375 0.875 0.875	79.2 -23.0 -6.7	24.0 196.3	0.375 0.875 0.875	78.3 -33.4 -10.2	34.9 196.9 10.9	210 0.0 1.0 1.0	86.8 -46.1 -13.5
744	G50B_087_062d	0.25 0.875 0.875	0.87									

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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	1.0 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	325.2 0.0 360	1.0 1.0 1.0	95.4 0.0 0.0	
811	B00R_100_012d	0.875 0.875 1.0	1.0 0.125 0.937	270	0.875 0.875 1.0	87.2 9.5 -12.9	16.0 306.2	0.875 0.875 1.0	85.5 5.8 -14.8	15.9 291.5 4.4	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
812	B00R_100_025d	0.75 0.75 1.0	1.0 0.25 0.875	270	0.75 0.75 1.0	79.1 19.0 -25.8	32.1 306.2	0.75 0.75 1.0	75.6 12.8 -30.0	32.7 293.1 8.2	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
813	B00R_100_037d	0.625 0.625 1.0	1.0 0.375 0.812	270	0.625 0.625 1.0	71.0 28.5 -38.8	48.1 306.2	0.625 0.625 1.0	65.7 21.4 -45.6	50.4 295.1 11.1	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
814	B00R_100_050d	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	62.8 38.0 -51.7	64.2 306.2	0.5 0.5 1.0	56.0 31.9 -61.1	69.0 297.5 13.0	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
815	B00R_100_062d	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	54.7 47.5 -64.7	80.3 306.2	0.375 0.375 1.0	46.8 44.5 -76.1	88.2 300.3 14.2	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
816	B00R_100_075d	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	46.6 57.0 -77.6	96.3 306.2	0.25 0.25 1.0	38.8 58.2 -89.4	106.7 303.0 14.1	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
817	B00R_100_087d	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	38.5 66.5 -90.6	112.4 306.2	0.125 0.125 1.0	33.0 69.9 -99.0	121.3 305.2 10.6	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
818	B00R_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2 0.0	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
819	Y00G_100_012d	1.0 1.0 0.875	1.0 0.125 0.937	90	1.0 1.0 0.875	95.0 -2.5 11.3	11.6 102.8	1.0 1.0 0.875	94.7 -5.0 14.6	15.4 108.9 4.1	89	1.0 1.0 0.0	92.6 -20.7 90.7 93.0 102.8
820	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	83.4 0.0 0.0	0.0 0.0	0.875 0.875 0.875	84.7 0.0 0.0	325.2 1.2 360	1.0 1.0 1.0	95.4 0.0 0.0	
821	B00R_087_012d	0.75 0.75 0.875	0.875 0.125 0.812	270	0.75 0.75 0.875	75.3 9.5 -12.9	16.0 306.2	0.75 0.75 0.875	74.6 6.0 -15.2	16.4 291.7 4.1	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
822	B00R_087_025d	0.625 0.625 0.875	0.875 0.25 0.75	270	0.625 0.625 0.875	67.2 19.0 -25.8	32.1 306.2	0.625 0.625 0.875	64.4 13.5 -30.9	33.8 293.6 7.9	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
823	B00R_087_037d	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.5 0.875	59.1 28.5 -38.8	48.1 306.2	0.5 0.5 0.875	54.3 23.0 -46.9	52.2 296.1 10.8	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
824	B00R_087_050d	0.375 0.375 0.875</											

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		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	95.4	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.4	0.0	0.0	0.0	1.0	1.0	1.0	95.4	0.0	0.0	325.2	0.0	360	1.0	1.0	1.0	95.4	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	90.6	11.7	-7.3	13.8	328.2	1.0	0.875	1.0	87.9	15.7	-10.9	19.1	325.1	5.9	330	1.0	0.875	1.0	87.2	94.3	-58.4	110.9	328.2	1.0	0.875	1.0	87.2	94.3	-58.4	110.9	328.2							
893	B50R_100_025d	1.0	0.75	1.0	1.0	0.25	0.875	330	1.0	0.75	1.0	85.8	23.5	-14.6	27.7	328.2	1.0	0.75	1.0	80.9	31.7	-21.5	38.4	325.8	11.8	330	1.0	0.75	1.0	57.2	94.3	-58.4	110.9	328.2	1.0	0.75	1.0	57.2	94.3	-58.4	110.9	328.2							
894	B50R_100_037d	1.0	0.625	1.0	1.0	0.375	0.812	330	1.0	0.625	1.0	81.1	35.3	-21.9	41.6	328.2	1.0	0.625	1.0	74.3	47.6	-31.5	57.1	326.4	17.0	330	1.0	0.625	1.0	57.2	94.3	-58.4	110.9	328.2	1.0	0.625	1.0	57.2	94.3	-58.4	110.9	328.2							
895	B50R_100_050d	1.0	0.5	1.0	1.0	0.5	0.75	330	1.0	0.5	1.0	76.3	47.1	-29.2	55.4	328.2	1.0	0.5	1.0	68.6	62.6	-40.5	74.6	327.0	20.6	330	1.0	0.5	1.0	57.2	94.3	-58.4	110.9	328.2	1.0	0.5	1.0	57.2	94.3	-58.4	110.9	328.2							
896	B50R_100_062d	1.0	0.375	1.0	1.0	0.625	0.687	330	1.0	0.375	1.0	71.5	58.9	-36.5	69.3	328.2	1.0	0.375	1.0	63.8	75.6	-48.0	89.6	327.5	21.6	330	1.0	0.375	1.0	57.2	94.3	-58.4	110.9	328.2	1.0	0.375	1.0	57.2	94.3	-58.4	110.9	328.2							
897	B50R_100_075d	1.0	0.25	1.0	1.0	0.75	0.625	330	1.0	0.25	1.0	66.8	70.7	-43.8	83.2	328.2	1.0	0.25	1.0	60.2	85.6	-53.6	101.0	327.9	18.9	330	1.0	0.25	1.0	57.2	94.3	-58.4	110.9	328.2	1.0	0.25	1.0	57.2	94.3	-58.4	110.9	328.2							
898	B50R_100_087d	1.0	0.125	1.0	1.0	0.875	0.562	330	1.0	0.125	1.0	62.0	82.5	-51.1	97.1	328.2	1.0	0.125	1.0	58.1	91.8	-57.0	108.0	328.1	11.6	330	1.0	0.125	1.0	57.2	94.3	-58.4	110.9	328.2	1.0	0.125	1.0	57.2	94.3	-58.4	110.9	328.2							
899	B50R_100_100d	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	1.0	0.0	1.0	57.2	94.3	-58.4	111.0	328.2	0.0																								

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		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	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.

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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	82.6	0.0	0.0	0.0	0.0	0.0	0.0	0.866	0.866	0.866	83.9	0.0	0.0	0.0	0.0	325.2	1.3	360	1.0	1.0	1.0	95.4	0.0	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	89.0	0.0	0.0	0.0	0.0	0.0	0.0	0.933	0.933	0.933	89.7	0.0	0.0	0.0	0.0	325.2	0.6	360	1.0	1.0	1.0	95.4	0.0	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	95.4	0.0	0.0	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0	325.2	0.0	360	1.0	1.0	1.0	95.4	0.0	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	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.4	0.0	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	6.2	0.0	0.0	0.0	0.0	0.0	0.0	0.066	0.066	0.066	4.4	0.0	0.0	0.0	0.0	326.3	1.8	360	1.0	1.0	1.0	95.4	0.0	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	12.6	0.0	0.0	0.0	0.0	0.0	0.0	0.133	0.133	0.133	12.0	0.0	0.0	0.0	0.0	325.6	0.6	360	1.0	1.0	1.0	95.4	0.0	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	19.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2	0.2	0.2	19.7	0.0	0.0	0.0	0.0	325.5	0.6	360	1.0	1.0	1.0	95.4	0.0	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	25.3	0.0	0.0	0.0	0.0	0.0	0.0	0.266	0.266	0.266	27.0	0.0	0.0	0.0	0.0	325.4	1.6	360	1.0	1.0	1.0	95.4	0.0	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	31.7	0.0	0.0	0.0	0.0	0.0	0.0	0.333	0.333	0.333	34.0	0.0	0.0	0.0	0.0	325.3	2.2	360	1.0	1.0	1.0	95.4	0.0	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	38.1	0.0	0.0	0.0	0.0	0.0	0.0	0.4	0.4	0.4	40.8	0.0	0.0	0.0	0.0	325.3	2.6	360	1.0	1.0	1.0	95.4	0.0	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	44.4	0.0	0.0	0.0	0.0	0.0	0.0	0.466	0.466	0.466	47.3	0.0	0.0	0.0	0.0	325.4	2.8	360	1.0	1.0	1.0	95.4	0.0	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	50.8	0.0	0.0	0.0	0.0	0.0	0.0	0.533	0.533	0.533	53.7	0.0	0.0	0.0	0.0	325.3	2.9	360	1.0	1.0	1.0	95.4	0.0	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	57.2	0.0	0.0	0.0	0.0	0.0	0.0	0.6	0.6	0.6	60.0	0.0	0.0	0.0	0.0	325.3	2.8	360	1.0	1.0	1.0	95.4	0.0	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	63.5	0.0	0.0	0.0	0.0	0.0	0.0	0.666	0.666	0.666	66.1	0.0	0.0	0.0	0.0	325.2	2.6	360	1.0	1.0	1.0	95.4	0.0	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	70.0	0.0	0.0	0.0	0.0	0.0	0.0	0.734	0.734	0.734	72.3	0.0	0.0	0.0	0.0	325.2	2.2	360	1.0	1.0	1.0	95.4	0.0	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	76.3	0.0	0.0	0.0	0.0	0.0	0.0	0.8	0.8	0.8	78.1	0.0	0.0	0.0	0.0	325.2	1.									

delta E* = 1.0

gráfico TUB-PS31; reproducción en color; sRGB
colores y diferencia en color, ΔE^* , 3D=0, de=0, sRGB

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