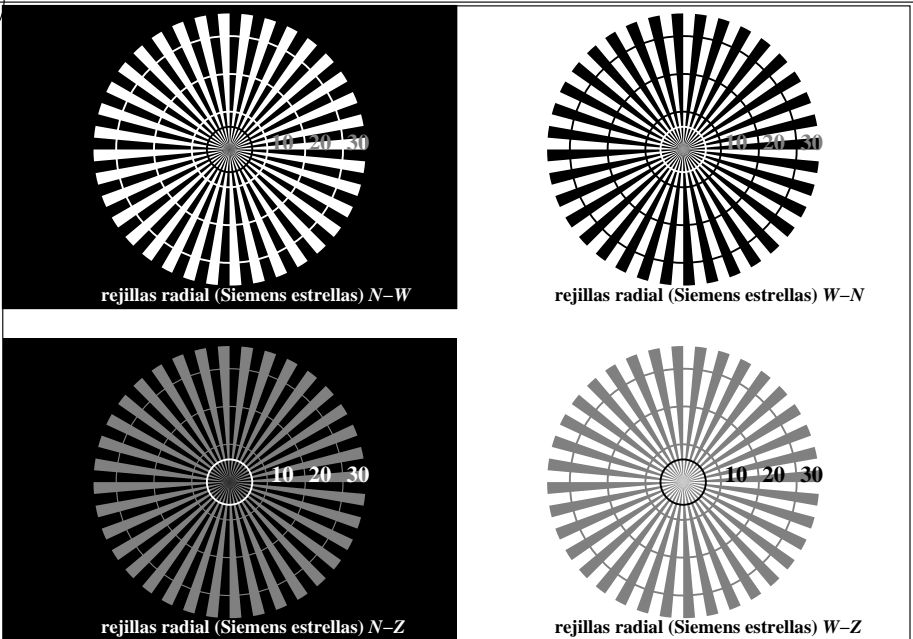


http://130.149.60.45/~farbmetrik/TS71/TS71L0NA.TXT /.PS; comience salida N: ninguna 3D-linealización (OL) en archivo (F) o PS-startup (S), página 1/18

vea archivos semejantes: http://130.149.60.45/~farbmetrik/TS71/TS71.HTM información técnica: http://www.ps.bam.de o http://130.149.60.45/~farbmetrik

TUB matrícula: 20150901-TS71/TS71L0NA.TXT /.PS aplicación para la medida de display output

TUB material: code=rh4ta



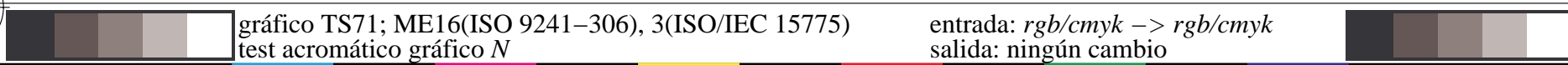
TS710-3, Fig. C1W-: Elemento A: rejillas radial N-W, W-N, N-Z y W-Z; PS operator: rgb/cmy0

$L^*/Y_{entrada}$	18.0/2.5	37.3/9.7	56.7/24.6	76.1/49.9	95.4/88.6	N_0 (min.)	W_I (max.)	
(absoluta)								
$w^* = l^*_{CIE LAB, r}$								
(relativa)	$w^*_{entrada}$	0,000	0,250	0,500	0,750	1,000	N_0 (min.)	W_I (max.)

TS710-5, Fig. C2W-: Elemento B: 5 equidistante L^* pasos de gris + N_0 + W_I ; PS operator: rgb/cmy0

$L^*/Y_{entrada}$	18.0/2.5	23.2/3.8	28.3/5.6	33.5/7.8	38.6/10.5	43.8/13.7	49.0/17.6	54.1/22.1	59.3/27.3	64.4/33.3	69.6/40.2	74.8/47.9	79.9/56.5	85.1/66.2	90.2/76.8	95.4/88.6	
(absoluta)																	
NO y código Hex	00;F	01;E	02;D	03;C	04;B	05;A	06;9	07;8	08;7	09;6	10;5	11;4	12;3	13;2	14;1	15;0	
$w^* = l^*_{CIE LAB, r}$																	
(relativa)	$w^*_{entrada}$	0,000	0,067	0,133	0,200	0,267	0,333	0,400	0,467	0,533	0,600	0,667	0,733	0,800	0,867	0,933	1,000

TS710-7, Fig. C3W-: Elemento C: 16 equidistante L^* pasos de gris; PS operator: rgb/cmy0



paso fondo	0		1	paso del anillo	0-1
Código Hexadecimal	7		8	Código Hexadecimal	7-8
E		F	E-F	E-F	
2		0	2-0	2-0	
8		6	8-6	8-6	
F		D	F-D	F-D	

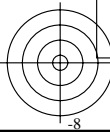
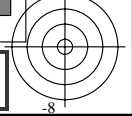
TS711-1, Fig. C4W-: Elemento D: anillos de Landolt W-N; PS operator: rgb/cmy0

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	

TS711-3, Fig. C5W-: Elemento E: Trama línea menores de 45° (o 135°) grados; PS operator: rgb/cmy0

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	

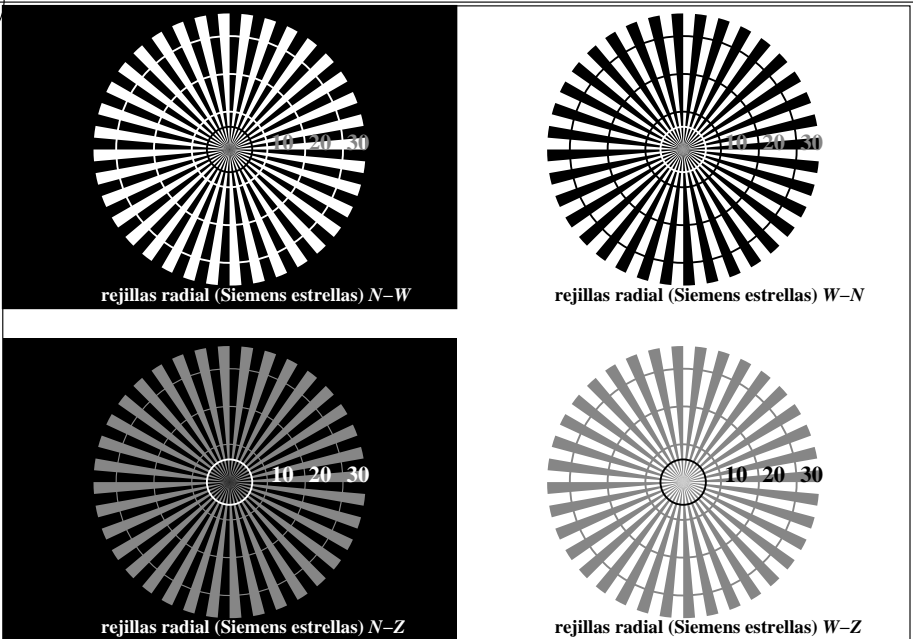
TS711-5, Fig. C6W-: Elemento F: Trama línea menores de 90° (o 0°) grados; PS operator: rgb/cmy0



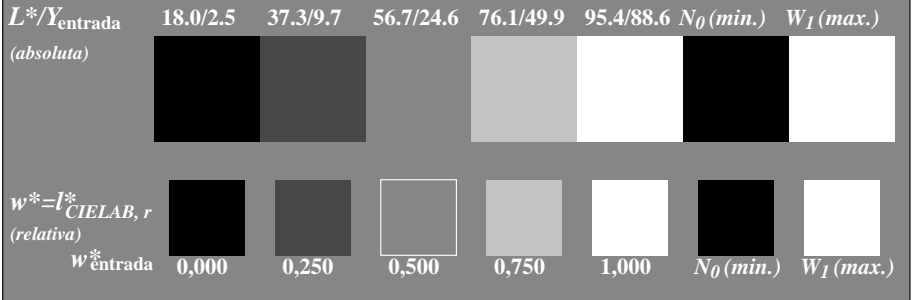
vea archivos semejantes: http://130.149.60.45/~farbmetrik/TS71/TS71.HTM
información técnica: http://www.ps.bam.de o http://130.149.60.45/~farbmetrik

TUB matrícula: 20150901-TS71/TS71L0NA.TXT /.PS
aplicación para la medida de display output, ninguna separación

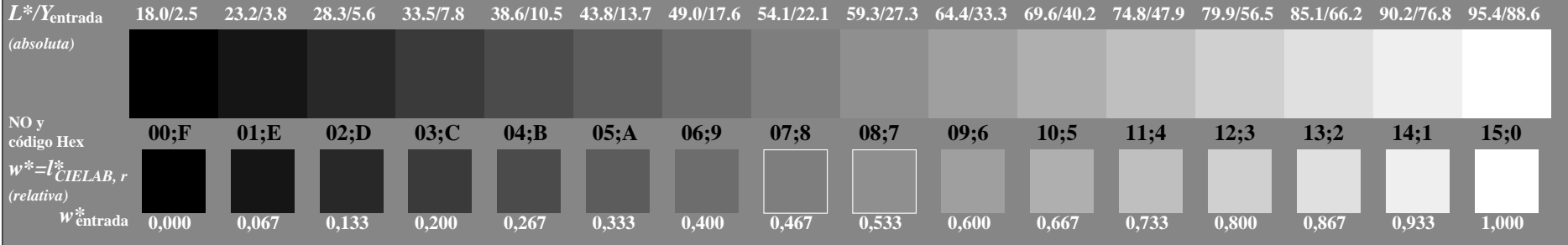
TUB material: code=rh4ta



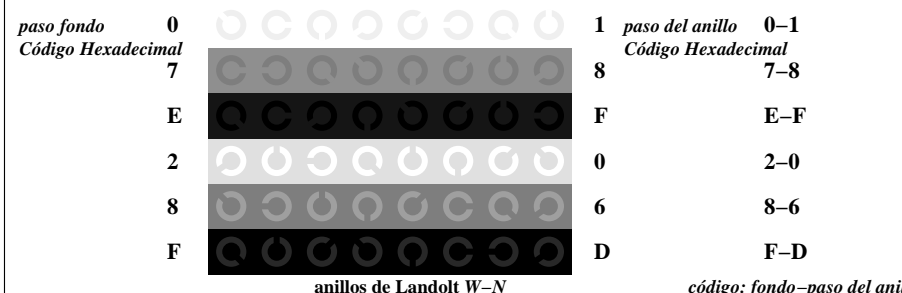
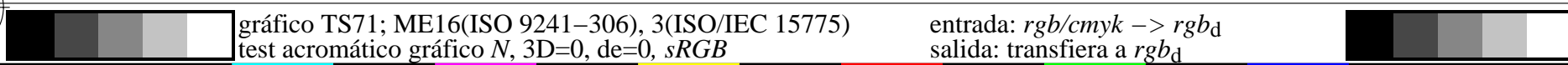
TS710-3, Fig. C1Wd: Elemento A: rejillas radial N-W, W-N, N-Z y W-Z; PS operator: rgb/cmy0



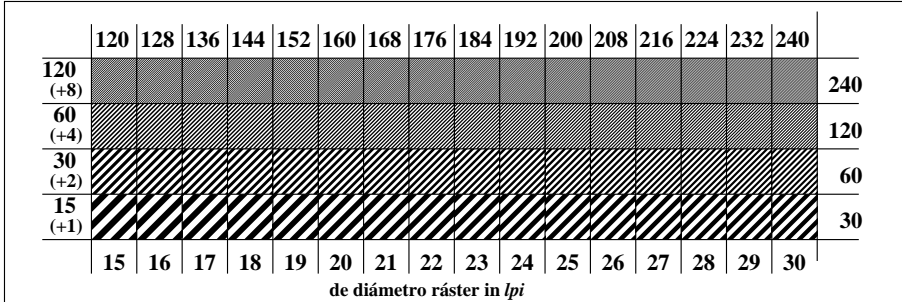
TS710-5, Fig. C2Wd: Elemento B: 5 equidistante L^* pasos de gris + N_0 + W_1 ; PS operator: rgb/cmy0



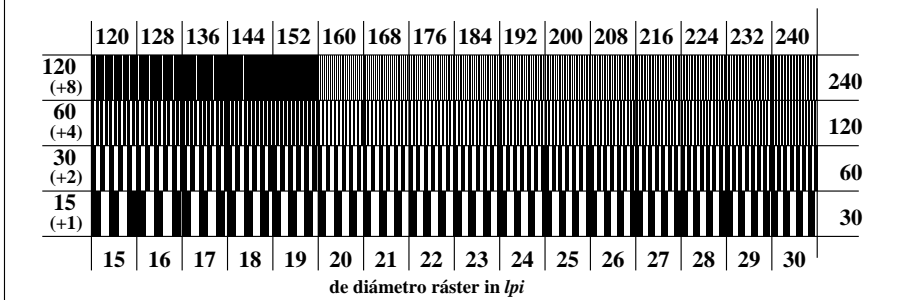
TS710-7, Fig. C3Wd: Elemento C: 16 equidistante L^* pasos de gris; PS operator: rgb/cmy0



TS711-1, Fig. C4Wd: Elemento D: anillos de Landolt W-N; PS operator: rgb/cmy0



TS711-3, Fig. C5Wd: Elemento E: Trama línea menores de 45° (o 135°) grados; PS operator: rgb/cmy0



TS711-5, Fig. C6Wd: Elemento F: Trama línea menores de 90° (o 0°) grados; PS operator: rgb/cmy0

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

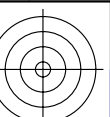
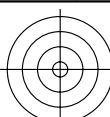
nj	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md
0/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5 100.4 39.9 0.0	389	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0	389
1/657	R13Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	51.4 74.1 64.9 98.5 41.2	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 98.5 41.2	36
2/666	R25Y_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	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 94.4 44.2	42
3/675	R38Y_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	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 88.1 50.3	51
4/684	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	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	59
5/693	R63Y_100_100a	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 79.4 71.8	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 79.4 71.8	68
6/702	R75Y_100_100a	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 81.0 84.4	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 81.0 84.4	77
7/711	R88Y_100_100a	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 85.8 94.4	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 85.8 94.4	83
8/720	Y00G_100_100a	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 93.0 102.8	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 93.0 102.8	89
9/639	Y13G_100_100a	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 94.0 110.0	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 94.0 110.0	96
10/558	Y25G_100_100a	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 96.5 116.6	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 96.5 116.6	102
11/477	Y38G_100_100a	0.625 1.0 0.0	1.0 1.0 0.5	112	0.633 1.0 0.0	87.0 -55.0 84.1 105.1 123.2	0.625 1.0 0.0	86.9 -55.7 83.9 105.1 123.6 0.7	111	0.633 1.0 0.0	87.0 -55.0 84.1 105.1 123.2	111
12/396	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 82.4 100.5 128.3	0.5 1.0 0.0	85.7 -65.2 82.4 100.7 128.3 0.0	119	0.5 1.0 0.0	85.7 -65.2 82.4 100.5 128.3	119
13/315	Y63G_100_100a	0.375 1.0 0.0	1.0 1.0 0.5	128	0.366 1.0 0.0	84.7 -73.2 81.2 109.3 132.0	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 81.2 109.3 132.0	128
14/234	Y75G_100_100a	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	84.0 -78.7 80.4 112.5 134.3	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 80.4 112.5 134.3	137
15/153	Y88G_100_100a	0.125 1.0 0.0	1.0 1.0 0.5	143	0.116 1.0 0.0	83.7 -81.5 80.0 114.2 135.5	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 80.0 114.2 135.5	143
16/72	G00C_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	83.6 -82.7 79.8 115.0 136.0	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 79.8 115.0 136.0	149
17/73	G13C_100_100a	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.116	83.6 -82.1 76.8 112.5 136.9	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 76.8 112.5 136.9	156
18/74	G25C_100_100a	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.233	83.7 -80.8 70.1 106.9 139.0	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 70.1 106.9 139.0	162
19/75	G38C_100_100a	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 97.7 142.9	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 97.7 142.9	171
20/76	G50C_100_100a	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 86.4 148.6	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 86.4 148.6	180
21/77	G63C_100_100a	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 74.3 156.5	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 74.3 156.5	188
22/78	G75C_100_100a	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 62.8 167.3	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 62.8 167.3	197
23/79	G88C_100_100a	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 54.1 180.0	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 54.1 180.0	203
24/80	C00B_100_100a	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 48.1 196.3	210
25/71	C13B_100_100a	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 42.5 218.2	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 42.5 218.2	216
26/62	C25B_100_100a	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 43.9 243.6	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 43.9 243.6	222
27/53	C38B_100_100a	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 53.9 268.3	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 53.9 268.3	231
28/44	C50B_100_100a	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 70.7 285.0	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 70.7 285.0	240
29/35	C63B_100_100a	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 90.7 295.3	0.0 0.375 1.0	43.8 37.6 -82.0 89.5 294.8 1.4	248	0.0 0.366 1.0	43.4 38.7 -82.0 90.7 295.3	248
30/26	C75B_100_100a	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.233 1.0	36.5 57.6 -93.4 109.7 301.6	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 -93.4 109.7 301.6	257
31/17	C88B_100_100a	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.116 1.0	32.3 70.0 -100.3 122.3 304.9	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 -100.3 122.3 304.9	263
32/8	B00M_100_100a	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	270
33/89	B13M_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	276
34/170	B25M_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	282
35/251	B38M_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	291
36/332	B50M_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	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 -89.7 120.0 311.6	300
37/413	B63M_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	308
38/494	B75M_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	47.9 86.4 -74.0 113.8 319.4	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 -74.0 113.8 319.4	317
39/575	B88M_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	0.883 0.0 1.0	52.5 90.1 -66.3 111.9 323.6	323
40/656	M00R_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	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 -58.4 110.9 328.2	330
41/655	M13R_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	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 -44.8 101.1 333.6	336
42/654	M25R_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	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 92.5 340.6	342
43/653	M38R_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	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 85.0 350.7	351
44/652	M50R_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	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	360
45/651	M63R_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	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 82.5 16.0	368
46/650	M75R_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	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 88.2 27.8	377
47/649	M88R_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	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 95.1 35.7	383
48/648	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 40.0	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	389
49/0	NW_000a	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	360	0.0 0.0 0.0	0.0 0.0 0.0 0.0 0.0	360
50/91	NW_013a	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	0.125 0.125 0.125	11.0 0.0 0.0 0.0 0.				

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

Table with columns: n/j, HIC*Fa, rgb_Fa, iet_Fa, hsi_Fa, rgb*Fa, LabCh*Fa, DE*Fa, hsi_Md, rgb*Md, LabCh*Md. It contains multiple rows of color and density data for various color patches and printing conditions.

delta E* = 6.5

TUB matrícula: 20150901-TS71/TS71LONA.TXT /.PS
aplicación para la medida de display output, ninguna separación
TUB material: code=rh4ta



vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71LONA.TXT> / .PS
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20150901-TS71/TS71LONA.TXT / .PS
 aplicación para la medida de display output, ninguna separación
 TUB material: code=rh4ta

n=j	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	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	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1	BOOR_012_012a	0.0 0.0 0.125	0.125 0.125 0.125	0.062 0.062 0.062	0.0 0.0 0.125	3.7 9.5	-12.9 16.0 306.2	0.0 0.0 0.125	0.8 5.8	-15.5 16.6	290.4 5.4 270	30.3 76.0
2	BOOR_025_025a	0.0 0.0 0.25	0.25 0.25 0.25	0.125 0.125 0.125	0.0 0.0 0.25	7.5 19.0	-25.8 32.1 306.2	0.0 0.0 0.25	2.9 20.6	-35.3 40.9	300.2 10.6 270	30.3 76.0
3	BOOR_037_037a	0.0 0.0 0.375	0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.0 0.375	11.3 28.5	-38.8 48.1 306.2	0.0 0.0 0.375	6.7 36.7	-50.3 62.3	306.1 14.9 270	30.3 76.0
4	BOOR_050_050a	0.0 0.0 0.5	0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.0 0.5	15.1 38.0	-51.7 64.2 306.2	0.0 0.0 0.5	11.7 45.5	-61.9 76.8	306.2 13.0 270	30.3 76.0
5	BOOR_062_062a	0.0 0.0 0.625	0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.0 0.625	18.9 47.5	-64.7 80.3 306.2	0.0 0.0 0.625	16.6 55.5	-72.9 90.4	306.2 10.3 270	30.3 76.0
6	BOOR_075_075a	0.0 0.0 0.75	0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.0 0.75	22.7 57.0	-77.6 96.3 306.2	0.0 0.0 0.75	21.3 61.2	-83.4 103.5	306.2 7.2 270	30.3 76.0
7	BOOR_087_087a	0.0 0.0 0.875	0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.0 0.875	26.5 66.5	-90.6 112.4 306.2	0.0 0.0 0.875	25.9 68.7	-93.6 116.1	306.2 3.7 270	30.3 76.0
8	BOOR_100_100a	0.0 0.0 1.0	1.0 1.0 1.0	0.5 0.5 0.5	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	30.3 76.0
9	GOOB_012_012a	0.0 0.125	0.125 0.125 0.125	0.062 0.062 0.062	0.0 0.125	10.4	-10.3 9.9	14.3 136.0	0.0 0.125	0.0 8.2	-16.7 11.9	20.6 144.4
10	G50B_012_012a	0.0 0.125	0.125 0.125 0.125	0.062 0.062 0.062	0.0 0.125	10.8	-5.7 -1.6	6.0 196.3	0.0 0.125	0.125 8.9	-10.7 -3.3	11.2 197.0
11	G75B_025_025a	0.0 0.125	0.25 0.25 0.25	0.125 0.125 0.125	0.0 0.125	12.9	4.5 -17.0	17.6 285.0	0.0 0.125	0.25 10.8	3.0 -22.5	22.7 277.6
12	G84B_037_037a	0.0 0.125	0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.118	0.375	15.2 17.1	-32.5 36.7 297.8	0.0 0.125	0.375 13.5	17.5 -39.0	42.8 299.1
13	G88B_050_050a	0.0 0.125	0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.116	0.5	18.2 28.8	-46.7 54.8 301.6	0.0 0.125	0.5 16.8	30.3 -53.4	61.4 294.6
14	G90B_062_062a	0.0 0.125	0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.114	0.625	21.6 39.4	-60.3 72.1 303.1	0.0 0.125	0.625 20.5	41.6 -66.3	78.3 302.1
15	G92B_075_075a	0.0 0.125	0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.112	0.75	25.0 50.0	-70.9 89.3 304.0	0.0 0.125	0.75 24.4	51.7 -78.2	93.8 303.5
16	G93B_087_087a	0.0 0.125	0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.116	0.875	28.7 60.0	-87.1 105.8 304.5	0.0 0.125	0.875 28.4	61.0 -89.4	108.2 304.3
17	G94B_100_100a	0.0 0.125	1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.116	1.0	32.3 70.0	-100.3 122.3 304.9	0.0 0.125	1.0 32.4	69.6 -100.0	121.9 304.8
18	GOOB_025_025a	0.0 0.25	0.0 0.25 0.25 0.25	0.125 0.125 0.125	0.0 0.25	0.0	20.9 -18.6	19.9 28.7 136.0	0.0 0.25	0.0 20.9	-30.6 28.3	41.7 137.2
19	G25B_025_025a	0.0 0.25	0.125 0.125 0.125 0.125	0.062 0.062 0.062	0.0 0.25	0.125	21.0 -20.4	11.2 21.6 148.6	0.0 0.25	0.125 21.2	-26.3 13.8	29.7 152.3
20	G50B_025_025a	0.0 0.25	0.25 0.25 0.25 0.25	0.125 0.125 0.125	0.0 0.25	0.25	21.7 -11.5	-3.3 12.0 196.3	0.0 0.25	0.25 22.1	-1.1 -5.0	17.8 196.3
21	G65B_037_037a	0.0 0.25	0.375 0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.256	0.375	24.1 -3.4	-18.3 18.6 259.3	0.0 0.25	0.375 23.5	-4.6 -22.9	23.4 258.4
22	G75B_050_050a	0.0 0.25	0.5 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.25	0.5	25.8 9.1	-34.1 35.3 285.0	0.0 0.25	0.5 25.5	8.7 -39.2	40.2 282.5
23	G80B_062_062a	0.0 0.25	0.625 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.239	0.625	27.7 22.6	-50.3 55.1 294.2	0.0 0.25	0.625 27.9	21.8 -54.1	58.3 291.9
24	G84B_075_075a	0.0 0.25	0.75 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.237	0.75	30.3 34.3	-65.0 73.5 297.8	0.0 0.25	0.75 30.7	34.0 -67.7	75.8 296.6
25	G86B_087_087a	0.0 0.25	0.875 0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.233	0.875	33.2 46.6	-79.6 92.3 300.8	0.0 0.25	0.875 33.8	45.4 -80.4	92.3 299.4
26	G88B_100_100a	0.0 0.25	1.0 1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.233	1.0	36.5 57.6	-93.9 109.7 301.6	0.0 0.25	1.0 37.0	54.9 -92.3	107.9 301.1
27	GOOB_037_037a	0.0 0.375	0.0 0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.375	0.0	31.3 -31.0	29.9 43.1 136.0	0.0 0.375	0.0 32.5	-40.3 38.9	56.1 136.0
28	G15B_037_037a	0.0 0.375	0.125 0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.375	0.118	31.4 -29.7	23.6 38.0 141.4	0.0 0.375	0.125 32.7	-37.7 27.7	46.9 143.6
29	G34B_037_037a	0.0 0.375	0.25 0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.375	0.256	31.8 -24.7	8.7 26.2 160.4	0.0 0.375	0.25 33.2	-31.7 11.0	33.6 160.8
30	G50B_037_037a	0.0 0.375	0.375 0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.375	0.375	32.5 -17.3	-5.0 18.0 196.3	0.0 0.375	0.375 34.1	-22.5 -6.6	23.4 196.3
31	G61B_050_050a	0.0 0.375	0.5 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.383	0.5	35.1 -9.7	-19.6 21.9 243.6	0.0 0.375	0.5 35.4	-11.1 -23.5	26.0 244.6
32	G69B_062_062a	0.0 0.375	0.625 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.385	0.625	37.3 0.5	-34.8 34.8 270.8	0.0 0.375	0.625 37.0	1.1 -39.4	39.4 271.7
33	G75B_075_075a	0.0 0.375	0.75 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.375	0.75	38.8 13.7	-51.2 53.0 285.0	0.0 0.375	0.75 39.0	13.7 -54.2	56.0 284.1
34	G79B_087_087a	0.0 0.375	0.875 0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.364	0.875	40.6 26.8	-67.7 72.8 291.5	0.0 0.375	0.875 41.3	25.9 -68.1	72.9 290.8
35	G81B_100_100a	0.0 0.375	1.0 1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.366	1.0	43.4 38.7	-82.0 90.7 295.3	0.0 0.375	1.0 43.8	37.6 -81.2	89.5 294.8
36	GOOB_050_050a	0.0 0.5	0.0 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.5	0.0	41.8 -41.3	39.9 57.5 136.0	0.0 0.5	0.0 43.5	-49.5 47.7	68.8 136.0
37	G11B_050_050a	0.0 0.5	0.125 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.5	0.116	41.8 -40.4	35.0 53.4 139.0	0.0 0.5	0.125 43.7	-47.7 39.5	62.0 140.3
38	G25B_050_050a	0.0 0.5	0.25 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.5	0.25	42.1 -36.8	22.4 43.2 148.6	0.0 0.5	0.25 44.0	-43.5 25.2	50.3 149.9
39	G38B_050_050a	0.0 0.5	0.375 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.5	0.383	42.7 -36.0	6.8 31.4 167.3	0.0 0.5	0.375 44.6	-36.7 8.6	37.7 166.7
40	G50B_050_050a	0.0 0.5	0.5 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.5	0.5	43.4 -23.0	-6.7 24.0 196.3	0.0 0.5	0.5 45.5	-27.6 -8.1	28.7 196.3
41	G59B_062_062a	0.0 0.5	0.625 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.51	0.625	46.1 -16.3	-21.4 26.9 232.6	0.0 0.5	0.625 46.6	-16.9 -24.3	29.6 235.0
42	G65B_075_075a	0.0 0.5	0.75 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.512	0.75	48.3 -6.9	-36.6 37.3 259.3	0.0 0.5	0.75 48.1	-5.4 -39.7	40.1 262.2
43	G70B_087_087a	0.0 0.5	0.875 0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.51	0.875	50.4 4.4	-52.3 52.4 274.9	0.0 0.5	0.875 49.8	6.4 -54.4	54.8 276.7
44	G75B_100_100a	0.0 0.5	1.0 1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.5	1.0	51.7 18.3	-68.3 70.7 285.0	0.0 0.5	1.0 51.7	18.3 -68.3	70.7 285.0
45	GOOB_062_062a	0.0 0.625	0.0 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.0	52.2 -51.7	49.9 71.9 136.0	0.0 0.625	0.0 54.1	-58.2 56.2	80.9 136.0
46	G09B_062_062a	0.0 0.625	0.125 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.114	52.3 -50.9	45.6 68.4 138.1	0.0 0.625	0.125 54.2	-56.9 49.9	75.7 138.7
47	G19B_062_062a	0.0 0.625	0.25 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.239	52.5 -48.5	35.7 60.2 143.5	0.0 0.625	0.25 54.4	-53.8 37.8	65.8 144.9
48	G30B_062_062a	0.0 0.625	0.375 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.385	52.9 -43.0	19.7 47.3 155.4	0.0 0.625	0.375 54.8	-48.8 22.6	53.6 155.0
49	G40B_062_062a	0.0 0.625	0.5 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.51	53.5 -36.5	4.6 36.8 172.6	0.0 0.625	0.5 55.5	-41.3 6.5	41.8 171.0
50	G50B_062_062a	0.0 0.625	0.625 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.625	54.2 -28.8	-8.4 30.0 196.3	0.0 0.625	0.625 56.3	-32.4 -9.5	33.8 196.3
51	G57B_075_075a	0.0 0.625	0.75 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.637	0.75	57.1 -22.4	-22.6 31.9 225.3	0.0 0.625	0.75 57.4	-22.3 -25.1	33.6 228.3
52	G63B_087_087a	0.0 0.625	0.875 0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.641	0.875	59.4 -13.4	-37.5 39.8 250.2	0.0 0.625	0.875 58.7	-11.4 -40.2	41.8 254.0
53	G68B_100_100a	0.0 0.625	1.0 1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.633	1.0	60.9 -1.5	-53.9 53.9 268.3	0.0 0.625	1.0 60.3	-0.1 -54.6	54.6 269.8
54	GOOB_075_075a	0.0 0.75	0.0 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.75	0.0	62.7 -62.0	59.9 86.2 136.0	0.0 0.75	0.0 64.2	-66.6 64.3	92.6 136.0
55	G07B_075_075a	0.0 0.75	0.125 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.75	0.112	62.7 -61.4	56.3 83.3 137.4	0.0 0.75	0.125 64.3	-65.6 59.9	88.5 137.8
56	G15B_075_075a	0.0 0.75	0.25 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.75	0.237	62.9 -59.4	47.3 76.0 141.4	0.0 0.75	0.25 64.5	-63.2 49.1	80.1 141.4
57	G25B_075_075a	0.0 0.75	0.375 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.75	0.375	63.2 -55.3	33.7 64.8 148.6	0.0 0.75	0.375 64.8	-59.1 35.4	68.9 149.0
58	G34B_075_075a	0.0 0.75	0.5 0.75 0.75 0.75									

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

Table with columns for various color channels (HIC*Fa, rgb*Fa, icf*Fa, hsi*Fa, LabCh*Fa, DE*Fa, hsiMd, rgb*Ma, LabCh*Ma) and rows for different color patches (e.g., R00Y, B50R, G25B, etc.).

delta E* = 8.3

TUB matrícula: 20150901-TS71/TS71LONA.TXT /.PS
aplicación para la medida de display output, ninguna separación
TUB material: code=rh4ta

http://130.149.60.45/~farbmetrik/TS71/TS71LONA.TXT /.PS; salida de transferencia
 N: ninguna 3D-linealización (OL) en archivo (F) o PS-startup (S), página 7/18

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71LONA.TXT> / .PS
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	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 25.1 4.0	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 2.0	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 29.0
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	0.683 0.0 1.0	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 -59.9	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	123.0 309.3
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	125.6 307.8
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	0.233 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_012a	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 1.0	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	0.0 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	80.2 306.2	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 0.25 0.125	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	G25B_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 -19.3 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	G50B_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 -17.5 -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	G65B_062_037a	0.25 0.5 0.625	0.625 0.375 0.437	229	0.25 0.506 0.625	48.0 -3.4 -18.3	18.6 259.3	0.25 0.5 0.625	47.9 -10.2 -22.3	24.5 245.3 7.8	228	0.0 0.683 1.0	64.4 -9.2	-48.8 49.7 259.3
204	G75B_075_050a	0.25 0.5 0.75	0.75 0.5 0.5	240	0.25 0.5 0.75	49.7 9.1 -34.1	35.3 285.0	0.25 0.5 0.75	49.3 0.1 -37.8	37.8 270.1 9.7	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
205	G80B_087_062a	0.25 0.5 0.875	0.875 0.625 0.562	247	0.25 0.489 0.875	51.5 22.6 -50.3	55.1 294.2	0.25 0.5 0.875	50.9 10.9 -52.5	53.6 281.7 11.9	247	0.0 0.383 1.0	44.3 36.2 -80.5	88.2 294.2
206	G84B_100_075a	0.25 0.5 1.0	1.0 0.75 0.625	251	0.25 0.487 1.0	54.4 34.3 -60.5	73.5 297.8	0.25 0.5 1.0	52.8 21.9 -66.5	70.0 288.2 12.5	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
207	Y61G_062_062a	0.25 0.625 0.0	0.625 0.625 0.312	127	0.239 0.625 0.0	53.0 -45.2 50.8	68.0 131.6	0.25 0.625 0.0	55.1 -49.5 57.4	75.8 130.7 8.1	127	0.383 1.0 0.0	84.8 -72.3	81.3 108.8 131.6
208	Y76G_062_050a	0.25 0.625 0.125	0.625 0.5 0.375	136	0.241 0.625 0.125	53.9 -39.3 40.2	56.2 134.3	0.25 0.625 0.125	55.2 -48.4 51.2	70.5 133.3 14.3	137	0.233 1.0 0.0	84.0 -78.7	80.4 112.5 134.3
209	G00B_062_037a	0.25 0.625 0.25	0.625 0.375 0.437	150	0.25 0.625 0.25	55.2 -31.0 29.9	43.1 136.0	0.25 0.625 0.25	55.4 -45.7 39.2	60.2 139.3 17.3	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
210	G15B_062_037a													

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71LONA.TXT> / .PS
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	icr_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	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	26.5 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	100.4 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 25.0 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 0.5 0.375	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	31.1 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	30.4 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 -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	0.375 0.375 0.0	36.9 -10.0	44.2 45.3 10.2	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	0.375 0.375 0.125	37.1 -8.7	33.8 34.9 10.4	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	0.375 0.375 0.25	37.5 -5.4	17.5 18.3 10.7	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 0.0 0.0	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	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	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	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	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	79.8 115.0 136.0
283	G50B_050_012a	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.5	46.6 -5.7	-1.6 6.0	0.375 0.5 0.5	48.4 -10.7	-3.5 11.3	198.2 5.6	210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3
284	G75B_062_025a	0.375 0.5 0.625	0.625 0.25 0.5	240	0.375 0.5 0.625	48.7 4.5 -17.0	17.6 285.0	0.375 0.5 0.625	49.4 -2.7	-19.8 20.0	262.1 7.8	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
285	G84B_075_037a	0.375 0.5 0.75	0.75 0.375 0.562	251	0.375 0.493 0.75	51.0 17.1 -32.5	36.7 297.8	0.375 0.5 0.75	50.7 6.3 -35.4	35.9 280.2 11.1	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8	
286	G88B_087_050a	0.375 0.5 0.875	0.875 0.5 0.625	256	0.375 0.491 0.875	54.0 28.8 -46.7	54.8 301.6	0.375 0.5 0.875	52.3 16.1 -50.2	52.7 287.8 13.2	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6	
287	G90B_100_062a	0.375 0.5 1.0	1.0 0.625 0.687	259	0.375 0.489 1.0	57.4 39.4 -60.3	72.1 303.1	0.375 0.5 1.0	54.1 26.2 -64.3	69.4 292.1 14.1	260	0.0 0.183 1.0	34.6 63.0 -96.6	115.3 303.1	
288	Y38G_062_062a	0.375 0.625 0.0	0.625 0.625 0.312	113	0.385 0.625 0.0	54.2 -35.2	52.4 63.1	0.375 0.625 0.0	56.3 -39.9	58.9 71.2	124.1 8.3	112	0.616 1.0 0.0	86.8 -56.4	83.8 101.0 123.9
289	Y50G_062_050a	0.375 0.625 0.125	0.625 0.5 0.375	120	0.375 0.625 0.125	54.7 -32.6	41.1 52.5	0.375 0.625 0.125	56.4 -39.0	52.8 65.7	126.4 13.4	119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3
290	Y68G_062_037a	0.375 0.625 0.25	0.625 0.375 0.437	131	0.368 0.625 0.25	55.5 -28.2	30.3 41.4	0.375 0.625 0.25	56.6 -36.6	40.9 54.9	131.8 13.5	131	0.316 1.0 0.0		

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71LONA.TXT> / .PS
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

Table with columns: n, HIC*Fa, rgb_Fa, icf_Fa, hsi_Fa, rgb*Fa, LabCh*Fa, DE*Fa, hsiMd, rgb*Md, LabCh*Md. It contains a large grid of numerical data representing color transfer characteristics for various color patches (e.g., 324, 325, 326, etc.) across different color channels and measurement methods.

delta E** = 10.1

gráfico TS71; ME16(ISO 9241-306), 3(ISO/IEC 15775)
colores y diferencia en color, ΔE*, 3D=0, de=0, sRGB

entrada: rgb/cmyk -> rgbd
salida: transfiera a rgbd

TUB matrícula: 20150901-TS71/TS71LONA.TXT / .PS
aplicación para la medida de display output, ninguna separación
TUB material: code=rh4ta

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	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 0.1125 0.114	31.0 54.7 30.0	62.4 28.7 6.0	380	1.0 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 0.383	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 35.2	0.625 0.0 0.375	32.4 58.6	-7.7 59.1 35.2	6.4 352	1.0 0.0 0.616	52.9 83.6	-11.4 84.3 35.2
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 0.816	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 69.3 328.2	0.625 0.0 0.625	35.5 66.4	-41.1 78.1 328.2	8.7 330	1.0 0.0 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 0.437	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 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 0.233	50.8 78.0	41.2 88.2 27.8
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 0.5	52.0 81.1	4.1 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 0.766	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 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 66.3	-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 0.233 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 0.383 0.0	57.7 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 0.316	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	16.0 349	1.0 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 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 90.0 314.6	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	-89.7 120.0 314.6
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 3.9	40.3 40.5 84.4	0.625 0.5 0.125	52.0 2.6	50.5 50.6 86.9	10.3 77	1.0 0.766 0.0	73.2 78.8	80.6 81.0 84.4
443	R68Y_062_037a	0.625 0.5 0.25	0.625 0.375 0.437	71	0.625 0.506 0.25	51.3 6.9	29.1 29.9 76.5	0.625 0.5 0.25	52.3 4.4	37.1 37.4 83.2	8.4 71	1.0 0.683 0.0	73.4 18.5	77.6 79.8 76.5
444	R50Y_062_025a	0.625 0.5 0.375	0.625 0.25 0.5	60	0.625 0.5 0.375	51.6 10.3	17.7 20.5 59.7	0.625 0.5 0.375	52.8 7.4	21.1 22.3 70.5	4.5 59	1.0 0.5 0.0	63.6 41.3	71.0 82.2 59.7
445	R00Y_062_012a	0.625 0.5 0.5	0.625 0.125 0.562	390	0.625 0.5 0.5	54.0 9.6	8.0 12.5 40.0	0.625 0.5 0.5	53.4 11.7	4.4 12.6 20.7	4.2 389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0
446	B50R_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	54.8 11.7	-7.3 13.8 328.2	0.625 0.5 0.625	54.4 17.2	-11.8 20.9 325.5	7.1 330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
447	B25R_075_025a	0.625 0.5 0.75	0.75 0.25 0.625	300	0.625 0.5 0.75	57.3 19.9	-22.4 30.0 311.6	0.625 0.5 0.75	55.5 23.7	-27.6 36.4 310.7	6.6 300	0.5 0.0 1.0	38.5 79.8	-89.7 120.0 311.6
448	B15R_087_037a	0.625 0.5 0.875	0.875 0.375 0.687	289	0.618 0.5 0.875	60.4 29.0	-36.5 46.7 308.4	0.625 0.5 0.875	56.9 31.0	-42.7 52.8 305.9	7.3 288	0.316 0.0 1.0	33.9 77.4	-97.5 124.5 308.4
449	B11R_100_050a	0.625 0.5 1.0	1.0 0.5 0.75 284	284	0.616 0.5 1.0	63.9 38.3	-50.0 63.1 307.4	0.625 0.5 1.0	58.5 38.8	-57.1 69.0 304.2	8.8 282	0.233 0.0 1.0	32.3 76.7	-100.1 126.2 307.4
450	Y00G_062_062a	0.625 0.625 0.0	0.625 0.625 0.312	90	0.625 0.625 0.0	57.9	-12.9 56.7 58.1 102.8	0.625 0.625 0.0	60.4	-14.5 63.8 65.4 102.8	7.7 89	1.0 1.0 0.0	92.6	-20.7 90.7 93.0 102.8
451	Y00G_062_050a	0.625 0.625 0.125	0.625 0.5 0.375	90	0.625 0.625 0.125	58.2	-10.0 45.3 46.5 102.8	0.625 0.625 0.125	60.5	-13.9 58.1 59.7 103.4 13.4 89	1.0 1.0 0.0	92.6	-20.7 90.7 93.0 102.8	
452	Y00G_062_037a	0.625 0.625 0.25	0.625 0.375 0.437	90	0.625 0.625 0.25	58.5	-7.7 34.0 34.9 102.8	0.625 0.625 0.25	60.7	-12.2 46.6 48.2 104.7 13.5 89	1.0 1			

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	rgb*Md	LabCh*Md	
486	R00Y_075_075a	0.75	0.0	0.0	0.75	0.375	390	0.75	0.0	0.0	0.75	0.375	390
487	R35Y_075_075a	0.75	0.0	0.125	0.75	0.375	381	0.75	0.0	0.125	0.75	0.375	381
488	R18Y_075_075a	0.75	0.0	0.25	0.75	0.375	371	0.75	0.0	0.25	0.75	0.375	371
489	R00Y_075_075a	0.75	0.0	0.375	0.75	0.375	360	0.75	0.0	0.375	0.75	0.375	360
490	B65R_075_075a	0.75	0.0	0.5	0.75	0.375	349	0.75	0.0	0.5	0.75	0.375	349
491	B57R_075_075a	0.75	0.0	0.625	0.75	0.375	339	0.75	0.0	0.625	0.75	0.375	339
492	B50R_075_075a	0.75	0.0	0.75	0.75	0.375	330	0.75	0.0	0.75	0.75	0.375	330
493	B43R_087_087a	0.75	0.0	0.875	0.875	0.437	322	0.758	0.0	0.875	0.875	0.437	322
494	B38R_100_100a	0.75	0.0	1.0	1.0	0.5	316	0.766	0.0	1.0	1.0	0.5	316
495	R15Y_075_075a	0.75	0.125	0.0	0.75	0.375	390	0.75	0.125	0.0	0.75	0.375	390
496	R00Y_075_062a	0.75	0.125	0.125	0.75	0.625	437	0.75	0.125	0.125	0.75	0.625	437
497	R31Y_075_062a	0.75	0.125	0.25	0.75	0.625	437	0.75	0.125	0.25	0.75	0.625	437
498	R11Y_075_062a	0.75	0.125	0.375	0.75	0.625	437	0.75	0.125	0.375	0.75	0.625	437
499	B69R_075_062a	0.75	0.125	0.5	0.75	0.625	437	0.75	0.125	0.5	0.75	0.625	437
500	B59R_075_062a	0.75	0.125	0.625	0.75	0.625	437	0.75	0.125	0.625	0.75	0.625	437
501	B50R_075_062a	0.75	0.125	0.75	0.75	0.625	437	0.75	0.125	0.75	0.75	0.625	437
502	B42R_087_075a	0.75	0.125	0.875	0.875	0.5	321	0.762	0.125	0.875	0.875	0.5	321
503	B36R_100_087a	0.75	0.125	1.0	1.0	0.875	314	0.766	0.125	1.0	1.0	0.875	314
504	R31Y_075_075a	0.75	0.25	0.0	0.75	0.375	49	0.75	0.25	0.0	0.75	0.375	49
505	R18Y_075_062a	0.75	0.25	0.125	0.75	0.625	437	0.75	0.25	0.125	0.75	0.625	437
506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	390	0.75	0.25	0.25	0.75	0.5	390
507	R26Y_075_050a	0.75	0.25	0.375	0.75	0.5	376	0.75	0.25	0.375	0.75	0.5	376
508	R00Y_075_050a	0.75	0.25	0.5	0.75	0.5	360	0.75	0.25	0.5	0.75	0.5	360
509	B61R_075_050a	0.75	0.25	0.625	0.75	0.5	344	0.75	0.25	0.625	0.75	0.5	344
510	B50R_075_050a	0.75	0.25	0.75	0.75	0.5	330	0.75	0.25	0.75	0.75	0.5	330
511	B40R_087_062a	0.75	0.25	0.875	0.875	0.625	319	0.766	0.25	0.875	0.875	0.625	319
512	B34R_100_075a	0.75	0.25	1.0	1.0	0.75	311	0.762	0.25	1.0	1.0	0.75	311
513	R50Y_075_075a	0.75	0.375	0.0	0.75	0.375	60	0.75	0.375	0.0	0.75	0.375	60
514	R38Y_075_062a	0.75	0.375	0.125	0.75	0.625	437	0.75	0.375	0.125	0.75	0.625	437
515	R23Y_075_050a	0.75	0.375	0.25	0.75	0.5	44	0.75	0.375	0.25	0.75	0.5	44
516	R00Y_075_037a	0.75	0.375	0.375	0.75	0.375	390	0.75	0.375	0.375	0.75	0.375	390
517	R18Y_075_037a	0.75	0.375	0.5	0.75	0.375	371	0.75	0.375	0.5	0.75	0.375	371
518	B65R_075_037a	0.75	0.375	0.625	0.75	0.375	362	0.75	0.375	0.625	0.75	0.375	362
519	B50R_075_037a	0.75	0.375	0.75	0.75	0.375	362	0.75	0.375	0.75	0.75	0.375	362
520	B38R_087_050a	0.75	0.375	0.875	0.875	0.5	316	0.758	0.375	0.875	0.875	0.5	316
521	B30R_100_062a	0.75	0.375	1.0	1.0	0.625	307	0.76	0.375	1.0	1.0	0.625	307
522	R68Y_075_075a	0.75	0.5	0.0	0.75	0.375	71	0.75	0.5	0.0	0.75	0.375	71
523	R61Y_075_062a	0.75	0.5	0.125	0.75	0.625	437	0.75	0.5	0.125	0.75	0.625	437
524	R50Y_075_050a	0.75	0.5	0.25	0.75	0.5	60	0.75	0.5	0.25	0.75	0.5	60
525	R31Y_075_037a	0.75	0.5	0.375	0.75	0.375	390	0.75	0.5	0.375	0.75	0.375	390
526	R00Y_075_025a	0.75	0.5	0.5	0.75	0.25	390	0.75	0.5	0.5	0.75	0.25	390
527	R00Y_075_025a	0.75	0.5	0.625	0.75	0.25	360	0.75	0.5	0.625	0.75	0.25	360
528	B50R_075_025a	0.75	0.5	0.75	0.75	0.25	330	0.75	0.5	0.75	0.75	0.25	330
529	B34R_087_037a	0.75	0.5	0.875	0.875	0.375	287	0.756	0.5	0.875	0.875	0.375	287
530	B25R_100_050a	0.75	0.5	1.0	1.0	0.5	300	0.75	0.5	1.0	1.0	0.5	300
531	R85Y_075_075a	0.75	0.625	0.0	0.75	0.375	81	0.75	0.625	0.0	0.75	0.375	81
532	R81Y_075_062a	0.75	0.625	0.125	0.75	0.625	437	0.75	0.625	0.125	0.75	0.625	437
533	R76Y_075_050a	0.75	0.625	0.25	0.75	0.5	76	0.75	0.625	0.25	0.75	0.5	76
534	R68Y_075_037a	0.75	0.625	0.375	0.75	0.375	371	0.75	0.625	0.375	0.75	0.375	371
535	R50Y_075_025a	0.75	0.625	0.5	0.75	0.25	360	0.75	0.625	0.5	0.75	0.25	360
536	R00Y_075_012a	0.75	0.625	0.625	0.75	0.125	687	0.75	0.625	0.625	0.75	0.125	687
537	B50R_075_012a	0.75	0.625	0.75	0.75	0.125	687	0.75	0.625	0.75	0.75	0.125	687
538	B25R_087_025a	0.75	0.625	0.875	0.875	0.25	300	0.75	0.625	0.875	0.875	0.25	300
539	B15R_100_037a	0.75	0.625	1.0	1.0	0.375	289	0.743	0.625	1.0	1.0	0.375	289
540	Y00G_075_075a	0.75	0.75	0.0	0.75	0.375	90	0.75	0.75	0.0	0.75	0.375	90
541	Y00G_075_062a	0.75	0.75	0.125	0.75	0.625	437	0.75	0.75	0.125	0.75	0.625	437
542	Y00G_075_050a	0.75	0.75	0.25	0.75	0.5	90	0.75	0.75	0.25	0.75	0.5	90
543	Y00G_075_037a	0.75	0.75	0.375	0.75	0.375	390	0.75	0.75	0.375	0.75	0.375	390
544	Y00G_075_025a	0.75	0.75	0.5	0.75	0.25	687	0.75	0.75	0.5	0.75	0.25	687
545	Y00G_075_012a	0.75	0.75	0.625	0.75	0.125	687	0.75	0.75	0.625	0.75	0.125	687
546	NW_075a	0.75	0.75	0.75	0.75	0.0	360	0.75	0.75	0.75	0.75	0.0	360
547	B00R_087_012a	0.75	0.75	0.875	0.875	0.125	812	0.75	0.75	0.875	0.875	0.125	812
548	B00R_100_025a	0.75	0.75	1.0	1.0	0.25	887	0.75	0.75	1.0	1.0	0.25	887
549	Y13G_087_087a	0.75	0.875	0.0	0.875	0.875	437	0.758	0.875	0.0	0.875	0.875	437
550	Y15G_087_075a	0.75	0.875	0.125	0.875	0.5	99	0.762	0.875	0.125	0.875	0.5	99
551	Y18G_087_062a	0.75	0.875	0.25	0.875	0.625	562	0.76	0.875	0.25	0.875	0.625	562
552	Y23G_087_050a	0.75	0.875	0.375	0.875	0.5	104	0.758	0.875	0.375	0.875	0.5	104
553	Y31G_087_037a	0.75	0.875	0.5	0.875	0.375	687	0.756	0.875	0.5	0.875	0.375	687
554	Y50G_087_025a	0.75	0.875	0.625	0.875	0.25	175	0.75	0.875	0.625	0.875	0.25	175
555	G00B_087_012a	0.75	0.875	0.75	0.875	0.125	812	0.75	0.875	0.75	0.875	0.125	812
556	G50B_087_012a	0.75	0.875	0.875	0.875	0.125	812	0.75	0.875	0.875	0.875	0.125	812
557	G75B_100_025a	0.75	0.875	1.0	1.0	0.25	875	0.75	0.875	1.0	1.0	0.25	875
558	Y23G_100_100a	0.75	1.0	0.0	1.0	0.5	104	0.766	1.0	0.0	1.0	0.5	104
559	Y26G_100_087a	0.75	1.0	0.125	1.0	0.875	562	0.766	1.0	0.125	1.0	0.875	562
560	Y31G_100_075a	0.75	1.0	0.25	1.0	0.75	687	0.762	1.0	0.25	1.0	0.75	687
561	Y38G_100_062a	0.75	1.0	0.375	1.0	0.625	687	0.76	1.0	0.375	1.0	0.625	687
562	Y50G_100_050a	0.75	1.0	0.5	1.0	0.5	120	0.75	1.0	0.5	1.0	0.5	120
563	Y68G_100_037a	0.75	1.0	0.625	1.0	0.375	812	0.743	1.0	0.625	1.0	0.375	812
564	G00B_100_025a	0.75	1.0	0.75	1.0	0.25	875	0.75	1.0	0.75	1.0	0.25	875
565	G25B_100_025a	0.75	1.0	0.875	1.0	0.25	875	0.75	1.0	0.875	1.0	0.25	875
566	G50B_100_025a	0.75	1.0	1.0	1.0	0.25	875	0.75	1.0	1.0	1.0	0.25	875

2-0031030-F0

TS710-7N, 11/18-F

gráfico TS71; ME16(ISO 9241-306), 3(ISO/IEC 15775)
 colores y diferencia en color, ΔE^* , 3D=0, de=0, sRGB

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

delta E** = 9.4

2-0031030-F0

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71LONA.TXT> / .PS
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20150901-TS71/TS71LONA.TXT / .PS
 aplicación para la medida de display output, ninguna separación
 TUB material: code=rh4ta

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMa	rgb*Ma	LabCh*Ma			
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	90.8 39.9	2.9 389	1.0 0.0 0.0	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	1.0 0.0 0.133	50.6 77.3 53.9	94.4 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	1.0 0.0 0.266	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	1.0 0.0 0.416	51.5 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	1.0 0.0 0.583	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	1.0 0.0 0.733	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	1.0 0.0 0.866	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 31.1	97.1 328.2	0.875 0.0 0.875	50.2 85.3 32.0	100.3 328.2	3.2 330	1.0 0.0 1.0	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	0.883 0.0 1.0	52.5 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	1.0 0.133 0.0	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	1.0 0.0 0.0	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	1.0 0.0 0.15	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	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6	
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	1.0 0.0 0.5	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	1.0 0.0 0.683	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	1.0 0.0 0.85	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	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2	
584	B43R_100_087a	0.875 0.125 1.0	1.0 1.0 0.875	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	106.7 323.0	11.3 322	0.866 0.0 1.0	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	47.8 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	1.0 0.266 0.0	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	1.0 0.15 0.0	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	1.0 0.0 0.0	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	1.0 0.0 0.183	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	1.0 0.0 0.383	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	52.7 352.1	0.875 0.25 0.625	51.1 66.2 16.0	68.1 346.3	17.5 352	1.0 0.0 0.616	52.9 83.6 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	1.0 0.0 0.816	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	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2	
593	B42R_100_075a	0.875 0.25 1.0	1.0 1.0 0.75	325	0.887 0.25 1.0	62.2 66.8 51.4	84.3 324.4	0.875 0.25 1.0	55.6 79.8 67.3	100.7 322.4	17.6 322	0.85 0.0 1.0	51.2 89.1 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	1.0 0.416 0.0	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	1.0 0.316 0.0	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	1.0 0.183 0.0	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	1.0 0.0 0.0	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	1.0 0.0 0.233	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 35.0	18.7 360	1.0 0.0 0.5	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	1.0 0.0 0.766	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	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2	
602	B40R_100_062a	0.875 0.375 1.0	1.0 1.0 0.625	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	0.816 0.0 1.0	49.8 88.1 70.7	113.0 321.2	
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	1.0 0.583 0.0	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	6.9 52	1.0 0.5 0.0	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 56.7	5.7 59	1.0 0.383 0.0	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	1.0 0.233 0.0	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	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
608	R18Y_087_037a	0.875 0.5 0.625	0.875 0.375 0.687	371	0.875 0.5 0.618	66.8 29.6 11.1	31.7 20.6	0.875 0.5 0.625	61.3 39.0 0.7	39.0 358.9	16.0 371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6	
609	B63R_087_037a	0.875 0.5 0.75	0.875 0.375 0.687	349	0.875 0.5 0.756	67.7 32.0 7.4	32.9 346.8	0.875 0.5 0.75	62.3 43.5 16.6	46.5 339.0	15.6 348	1.0 0.0 0.683	53.5 85.4 19.9	87.7 346.8	
610	B50R_087_037a	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	69.1 35.3 21.9	41.6 328.2	0.875 0.5 0.875	63.5 48.6 31.9	58.2 326.7	17.6 330	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2	
611	B38R_100_050a	0.875 0.5 1.0	1.0 0.5 0.75	316	0.883 0.5 1.0	71.6 43.2 37.0	56.9 319.4	0.875 0.5 1.0	64.8 54.4 46.6	71.7 319.3	16.3 317	0.766 0.0 1.0	47.9 86.4 74.0	113.8 319.4	
612	R73Y_087_087a	0.875 0.625 0.0	0.875 0.875 0.437	74	0.875 0.641 0.0	66.7 10.5 69.4	70.2 81.3	0.875 0.625 0.0	66.5 12.7 70.9	72.0 79.7	2.7 75	1.0 0.733 0.0	76.2 12.0 79.3	80.2 81.3	
613	R68Y_087_075a	0.875 0.625 0.125	0.875 0.75 0.5	71	0.875 0.637 0.125	67.0 13.8 58.2	59.8 76.5	0.875 0.625 0.125	66.5 13.2 65.6	66.9 78.6	7.4 71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5	
614	R61Y_087_062a	0.875 0.625 0.25	0.875 0.625 0.562	67	0.875 0.635 0.25	67.4 16.7 46.8	49.7 70.7	0.875 0.625 0.25	66.7 14.3 54.8	56.6 75.3	8.4 67	1.0 0.616 0.0	69.6 26.8 74.8	79.5 70.2	
615	R50Y_087_050a	0.875 0.625 0.375	0.875 0.5 0.625	60	0.875 0.625 0.375	67.6 20.6 35.5	41.1 59.2	0.875 0.625							

n	HIC*Fa	rgb_Fa	ief_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	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	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	54.9	94.8	35.4
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
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
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
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
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
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
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
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
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
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
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
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	13.2	71.6	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
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
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	-35.1	87.3	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	-57.1	97.1	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
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
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
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
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
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
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
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	-20.5	73.9	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	68.8	70.7	-43.8	83.2	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
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
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
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
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
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
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
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
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
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
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
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
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
688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	390	1.0	0.5	0.5	72.9	38.4	32.2	50.2	40.0
689	R26Y_100_050a	1.0	0.5	0.625	1.0	0.5	0.75	376	1.0	0.5	0.616	73.1	39.0	20.6	44.1	27.8
690	R00Y_100_050a	1.0	0.5	0.75	1.0	0.5	0.75	360	1.0	0.5	0.75	73.7	40.5	20.0	40.6	2.9
691	B61R_100_050a	1.0	0.5	0.875	1.0	0.5	0.75	344	1.0	0.5	0.883	74.9	43.6	-15.3	46.2	340.6
692	B50R_100_050a	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
693	R63Y_100_100a	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	79.4	71.8
694	R58Y_100_087a	1.0	0.625	0.125	1.0	0.875	0.562	65	1.0	0.635	0.125	71.3	27.1	64.8	70.2	67.2
695	R50Y_100_075a	1.0	0.625	0.25	1.0	0.75	0.625	60	1.0	0.625	0.25	71.6	31.0	53.2	61.6	59.7
696	R38Y_100_062a	1.0	0.625	0.375	1.0	0.625	0.687	53	1.0	0.614	0.375	72.3	34.0	42.6	54.6	51.3
697	R23Y_100_050a	1.0	0.625	0.5	1.0	0.5	0.75	44	1.0	0.616	0.5	74.5	33.8	32.9	47.2	44.2
698	R00Y_100_037a	1.0	0.625	0.625	1.0	0.375	0.812	390	1.0	0.625	0.625	78.5	28.8	24.2	37.6	40.0
699	R18Y_100_037a	1.0	0.625	0.75	1.0	0.375	0.812	371	1.0	0.625	0.743	78.8	29.6	11.1	31.7	20.6
700	B65R_100_037a	1.0	0.625	0.875	1.0	0.375	0.812	349	1.0	0.625	0.881	79.7	32.0	-7.4	32.9	346.8
701	B50R_100_037a	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
702	R76Y_100_100a	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	81.0	84.4
703	R73Y_100_087a	1.0	0.75	0.125	1.0	0.875	0.562	74	1.0	0.766	0.125	78.6	10.5	69.4	70.2	81.3
704	R68Y_100_075a	1.0	0.75	0.25	1.0	0.75	0.625	71	1.0	0.762	0.25	78.9	13.8	58.2	59.8	76.5
705	R61Y_100_062a	1.0	0.75	0.375	1.0	0.625	0.687	67	1.0	0.76	0.375	79.3	16.7	46.8	49.7	70.2
706	R50Y_100_050a	1.0	0.75	0.5	1.0	0.5	0.75	60	1.0	0.75	0.5	79.5	20.6	35.5	41.1	59.7
707	R31Y_100_037a	1.0	0.75	0.625	1.0	0.375	0.812	49	1.0	0.743	0.625	80.7	22.7	25.2	33.9	47.9
708	R00Y_100_025a	1.0	0.75	0.75	1.0	0.25	0.875	390	1.0	0.75	0.75	84.1	19.2	16.1	25.1	40.0
709	R00Y_100_025a	1.0	0.75	0.875	1.0	0.25	0.875	360	1.0	0.75	0.875	84.5	20.2	1.0	20.3	2.9
710	B50R_100_025a	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
711	R88Y_100_100a	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	85.8	94.4
712	R86Y_100_087a	1.0	0.875	0.125	1.0	0.875	0.562	81	1.0	0.883	0.125	85.6	-4.0	74.2	74.3	91.1
713	R85Y_100_075a	1.0	0.875	0.25	1.0	0.75	0.625	82	1.0	0.887	0.25	86.3	-1.8	63.2	63.2	93.7
714	R81Y_100_062a	1.0	0.875	0.375	1.0	0.625	0.687	79	1.0	0.885	0.375	86.5	1.0	51.8	51.8	88.7
715	R76Y_100_050a	1.0	0.875	0.5	1.0	0.5	0.75	76	1.0	0.883	0.5	86.8	3.9	40.3	40.5	84.4
716	R68Y_100_037a	1.0	0.875	0.625	1.0	0.375	0.812	71	1.0	0.881	0.625	87.1	6.9	29.1	29.9	76.5
717	R50Y_100_025a	1.0	0.875	0.75	1.0	0.25	0.875	60	1.0	0.875	0.75	87.4	10.3	17.7	20.5	59.7
718	R00Y_100_012a	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
719	B50R_100_012a	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
720	Y00G_100_100a	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	93.0	102.8
721	Y00G_100_087a	1.0	1.0	0.125	1.0	0.875	0.562	90	1.0	1.0	0.125	93.0	-18.1	79.4	81.4	102.8
722	Y00G_100_075a	1.0	1.0	0.25	1.0	0.75	0.625	90	1.0	1.0						

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71.LONA.TXT> / .PS
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

Table with columns: n, HIC*Fa, rgb_Fa, icf_Fa, hsi_Fa, rgb*Fa, LabCh*Fa, hsi_Ma, rgb*Ma, LabCh*Ma, DE*Fa, hsi_Ma, rgb*Ma, LabCh*Ma. It contains a large grid of numerical data for various color patches and colorimetric parameters.

delta E** = 7.3

gráfico TS71; ME16(ISO 9241-306), 3(ISO/IEC 15775)
colores y diferencia en color, ΔE*, 3D=0, de=0, sRGB

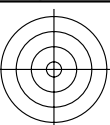
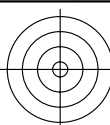
entrada: rgb/cmyk -> rgbd
salida: transfiera a rgbd

TUB matrícula: 20150901-TS71/TS71LONA.TXT / .PS
aplicación para la medida de display output, ninguna separación
TUB material: code=rha4ta

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20150901-TS71/TS71LONA.TXT /.PS
 aplicación para la medida de display output, ninguna separación
 TUB material: code=rh4ta

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md	
810	NW_100a	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	BOOR_100_012a	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
812	BOOR_100_025a	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
813	BOOR_100_037a	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
814	BOOR_100_050a	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
815	BOOR_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 44.9	-76.1 88.2	300.3 14.2 270	0.0 0.0 1.0	30.3 76.0
816	BOOR_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
817	BOOR_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
818	BOOR_100_100a	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 0.0 1.0	30.3 76.0	
819	YOOG_100_012a	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
820	NW_087a	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
821	BOOR_087_012a	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
822	BOOR_087_025a	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
823	BOOR_087_037a	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
824	BOOR_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
825	BOOR_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
826	BOOR_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
827	BOOR_087_087a	0.0 0.0 0.875	0.875 0.875 0.437	270	0.0 0.0 0.875	26.5 66.5	-90.6 112.4 306.2	0.0 0.0 0.875	25.9 68.7	-93.6 116.1	306.2 3.7 270	0.0 0.0 1.0	30.3 76.0
828	YOOG_100_025a	1.0 1.0 0.75	1.0 0.25 0.875	90	1.0 1.0 0.75	94.7 -5.1	22.6 23.2 102.8	1.0 1.0 0.75	94.1 -9.3	29.3 30.8	107.7 7.9 89	1.0 1.0 0.0	92.6 -20.7
829	YOOG_087_012a	0.875 0.875 0.75	0.875 0.125 0.812	90	0.875 0.875 0.75	83.1 -2.5	11.3 11.6 102.8	0.875 0.875 0.75	84.0 -5.1	15.0 15.8	108.7 4.5 89	1.0 1.0 0.0	92.6 -20.7
830	NW_075a	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	71.5 0.0	0.0 0.0 0.0	0.75 0.75 0.75	73.7 0.0	0.0 0.0	325.2 2.1 360	1.0 1.0 1.0	95.4 0.0
831	BOOR_075_012a	0.625 0.625 0.75	0.75 0.125 0.687	270	0.625 0.625 0.75	63.4 9.5	-12.9 16.0 306.2	0.625 0.625 0.75	63.3 6.3	-15.7 16.9	292.0 4.1 270	0.0 0.0 1.0	30.3 76.0
832	BOOR_075_025a	0.5 0.5 0.75	0.75 0.25 0.625	270	0.5 0.5 0.75	55.3 19.0	-25.8 32.1 306.2	0.5 0.5 0.75	52.8 14.4	-31.9 35.1	294.3 7.9 270	0.0 0.0 1.0	30.3 76.0
833	BOOR_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
834	BOOR_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
835	BOOR_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
836	BOOR_075_075a	0.0 0.0 0.75	0.5 0.75 0.375	270	0.0 0.0 0.75	22.7 57.0	-77.6 96.3 306.2	0.0 0.0 0.75	21.3 61.2	-83.4 103.5	306.2 14.2 270	0.0 0.0 1.0	30.3 76.0
837	YOOG_100_037a	1.0 1.0 0.625	1.0 0.375 0.812	90	1.0 1.0 0.625	94.3 -7.7	34.0 34.9 102.8	1.0 1.0 0.625	93.6 -13.0	43.8 45.7	106.5 11.1 89	1.0 1.0 0.0	92.6 -20.7
838	YOOG_087_025a	0.875 0.875 0.625	0.875 0.25 0.75	90	0.875 0.875 0.625	82.7 -5.1	22.6 23.2 102.8	0.875 0.875 0.625	83.4 -9.4	30.0 31.5	107.3 8.5 89	1.0 1.0 0.0	92.6 -20.7
839	YOOG_075_012a	0.75 0.75 0.625	0.75 0.125 0.687	90	0.75 0.75 0.625	71.2 -2.5	11.3 11.6 102.8	0.75 0.75 0.625	73.0 -5.1	15.4 16.3	108.5 5.2 89	1.0 1.0 0.0	92.6 -20.7
840	NW_062a	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	59.6 0.0	0.0 0.0 0.0	0.625 0.625 0.625	62.4 0.0	0.0 0.0	325.2 2.7 360	1.0 1.0 1.0	95.4 0.0
841	BOOR_062_012a	0.5 0.5 0.625	0.625 0.125 0.562	270	0.5 0.5 0.625	51.5 9.5	-12.9 16.0 306.2	0.5 0.5 0.625	51.6 6.7	-16.3 17.6	292.4 4.3 270	0.0 0.0 1.0	30.3 76.0
842	BOOR_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
843	BOOR_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
844	BOOR_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
845	BOOR_062_062a	0.0 0.0 0.625	0.625 0.625 0.312	270	0.0 0.0 0.625	18.9 47.5	-64.7 80.3 306.2	0.0 0.0 0.625	16.6 53.5	-72.9 90.4	306.2 10.3 270	0.0 0.0 1.0	30.3 76.0
846	YOOG_100_050a	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	94.0 -10.3	45.3 46.5 102.8	1.0 1.0 0.5	93.2 -15.9	57.8 59.9	105.3 13.6 89	1.0 1.0 0.0	92.6 -20.7
847	YOOG_087_037a	0.875 0.875 0.5	0.875 0.375 0.687	90	0.875 0.875 0.5	82.4 -7.7	34.0 34.9 102.8	0.875 0.875 0.5	82.9 -12.9	44.8 46.6	106.0 11.9 89	1.0 1.0 0.0	92.6 -20.7
848	YOOG_075_025a	0.75 0.75 0.5	0.75 0.25 0.625	90	0.75 0.75 0.5	70.8 -5.1	22.6 23.2 102.8	0.75 0.75 0.5	72.4 -9.4	30.9 32.3	106.9 9.4 89	1.0 1.0 0.0	92.6 -20.7
849	YOOG_062_012a	0.625 0.625 0.5	0.625 0.125 0.562	90	0.625 0.625 0.5	59.2 -2.5	11.3 11.6 102.8	0.625 0.625 0.5	61.6 -5.2	16.0 16.8	108.2 5.8 89	1.0 1.0 0.0	92.6 -20.7
850	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	47.7 0.0	0.0 0.0 0.0	0.5 0.5 0.5	50.6 0.0	0.0 0.0	325.3 2.9 360	1.0 1.0 1.0	95.4 0.0
851	BOOR_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
852	BOOR_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.25 0.25 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
853	BOOR_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.125 0.125 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
854	BOOR_050_050a	0.0 0.0 0.5	0.5 0.5 0.25	270	0.0 0.0 0.5	15.1 38.0	-51.7 64.2 306.2	0.0 0.0 0.5	11.7 45.5	-61.9 76.8	306.2 13.0 270	0.0 0.0 1.0	30.3 76.0
855	YOOG_100_062a	1.0 1.0 0.375	1.0 0.625 0.687	90	1.0 1.0 0.375	93.6 -12.9	56.7 58.1 102.8	1.0 1.0 0.375	92.9 -18.0	70.4 72.7	104.3 14.7 89	1.0 1.0 0.0	92.6 -20.7
856	YOOG_087_050a	0.875 0.875 0.375	0.875 0.5 0.625	90	0.875 0.875 0.375	82.1 -10.3	45.3 46.5 102.8	0.875 0.875 0.375	82.6 -15.5	58.6 60.6	104.8 14.2 89	1.0 1.0 0.0	92.6 -20.7
857	YOOG_075_037a	0.75 0.75 0.375	0.75 0.375 0.562	90	0.75 0.75 0.375	70.5 -7.7	34.0 34.9 102.8	0.75 0.75 0.375	72.0 -12.6	45.8 47.5	105.4 12.8 89	1.0 1.0 0.0	92.6 -20.7
858	YOOG_062_025a	0.625 0.625 0.375	0.625 0.25 0.5	90	0.625 0.625 0.375	58.9 -5.1	22.6 23.2 102.8	0.625 0.625 0.375	61.1 -9.3	31.9 33.2	106.3 10.3 89	1.0 1.0 0.0	92.6 -20.7
859	YOOG_050_012a	0.5 0.5 0.375	0.5 0.125 0.437	90	0.5 0.5 0.375	47.3 -2.5	11.3 11.6 102.8	0.5 0.5 0.375	49.8 -5.3	16.6 17.5	107.8 6.5 89	1.0 1.0 0.0	92.6 -20.7
860	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
861	BOOR_037_012a	0.25 0.25 0.375	0.375 0.125 0.312	270	0.25 0.25 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
862	BOOR_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.125 0.125 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



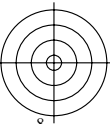
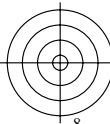
vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20150901-TS71/TS71LONA.TXT /.PS
aplicación para la medida de display output, ninguna separación

TUB material: code=rha4ta

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	rgb*Md	LabCh*Md
972	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
973	NW_012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
974	NW_025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
975	NW_037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
976	NW_050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
977	NW_062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
978	NW_075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
979	NW_087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
980	NW_100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
981	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
982	NW_012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
983	NW_025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
984	NW_037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
985	NW_050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
986	NW_062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
987	NW_075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
988	NW_087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
989	NW_100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
990	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
991	NW_012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
992	NW_025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
993	NW_037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
994	NW_050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
995	NW_062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
996	NW_075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
997	NW_087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
998	NW_100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
999	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1000	NW_012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
1001	NW_025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
1002	NW_037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
1003	NW_050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
1004	NW_062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
1005	NW_075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
1006	NW_087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
1007	NW_100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1008	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1009	NW_006a	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066
1010	NW_013a	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133
1011	NW_020a	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2
1012	NW_026a	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266
1013	NW_033a	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333
1014	NW_040a	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4
1015	NW_046a	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466
1016	NW_053a	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533
1017	NW_060a	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6
1018	NW_066a	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666
1019	NW_073a	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734
1020	NW_080a	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8
1021	NW_086a	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866
1022	NW_093a	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933
1023	NW_100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1024	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1025	NW_006a	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066
1026	NW_013a	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133
1027	NW_020a	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2
1028	NW_026a	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266
1029	NW_033a	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333
1030	NW_040a	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4
1031	NW_046a	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466
1032	NW_053a	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533
1033	NW_060a	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6
1034	NW_066a	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666
1035	NW_073a	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734
1036	NW_080a	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8
1037	NW_086a	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866
1038	NW_093a	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933
1039	NW_100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1040	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1041	NW_006a	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066
1042	NW_013a	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133
1043	NW_020a	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2
1044	NW_026a	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266
1045	NW_033a	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333
1046	NW_040a	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4
1047	NW_046a	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466
1048	NW_053a	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533
1049	NW_060a	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6
1050	NW_066a	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666
1051	NW_073a	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734
1052	NW_080a	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8

delta E* = 1.6



vea archivos semejantes: <http://130.149.60.45/~farbmetrik/TS71/TS71.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	ief_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md
1053	NW_086a	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 0.0 0.0	0.866 0.866 0.866	83.9 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1054	NW_093a	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 0.0 0.0	0.933 0.933 0.933	89.7 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1055	NW_100a	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 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1056	NW_000a	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	1.0 1.0 1.0	95.4 0.0 0.0
1057	NW_006a	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 0.0 0.0	0.066 0.066 0.066	4.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1058	NW_013a	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 0.0 0.0	0.133 0.133 0.133	12.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1059	NW_020a	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 0.0 0.0	0.2 0.2 0.2	19.7 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1060	NW_026a	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 0.0 0.0	0.266 0.266 0.266	27.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1061	NW_033a	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 0.0 0.0	0.333 0.333 0.333	34.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1062	NW_040a	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 0.0 0.0	0.4 0.4 0.4	40.8 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1063	NW_046a	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 0.0 0.0	0.466 0.466 0.466	47.3 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1064	NW_053a	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 0.0 0.0	0.533 0.533 0.533	53.7 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1065	NW_060a	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 0.0 0.0	0.6 0.6 0.6	60.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1066	NW_066a	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 0.0 0.0	0.666 0.666 0.666	66.1 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1067	NW_073a	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 0.0 0.0	0.734 0.734 0.734	72.3 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1068	NW_080a	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 0.0 0.0	0.8 0.8 0.8	78.1 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1069	NW_086a	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 0.0 0.0	0.866 0.866 0.866	83.9 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1070	NW_093a	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 0.0 0.0	0.933 0.933 0.933	89.7 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1071	NW_100a	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 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1072	NW_000a	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	1.0 1.0 1.0	95.4 0.0 0.0
1073	NW_100a	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 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1074	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 40.0	0.0 0.0 0.0	1.0 0.0 0.0	50.4 76.9 64.5	100.4 39.9	0.0 50.4 76.9 64.5
1075	G50B_100_100a	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 0.0 0.0	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3	0.0 86.8 -46.1 -13.5
1076	Y00G_100_100a	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	93.0 102.8	0.0 0.0 0.0	1.0 1.0 0.0	92.6 -20.6 90.7	93.0 102.8	0.0 92.6 -20.7 90.7
1077	B00R_100_100a	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 0.0	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	0.0 0.0 0.0
1078	G00B_100_100a	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	115.0 136.0	0.0 0.0 0.0	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0	0.0 0.0 0.0
1079	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 0.0 0.0	1.0 0.0 1.0	57.2 94.3 -58.4	111.0 328.2	0.0 1.0 0.0

delta E* = 1.0

TUB matrícula: 20150901-TS71/TS71LONA.TXT /.PS
 aplicación para la medida de display output, ninguna separación
 TUB material: code=rh4ta

