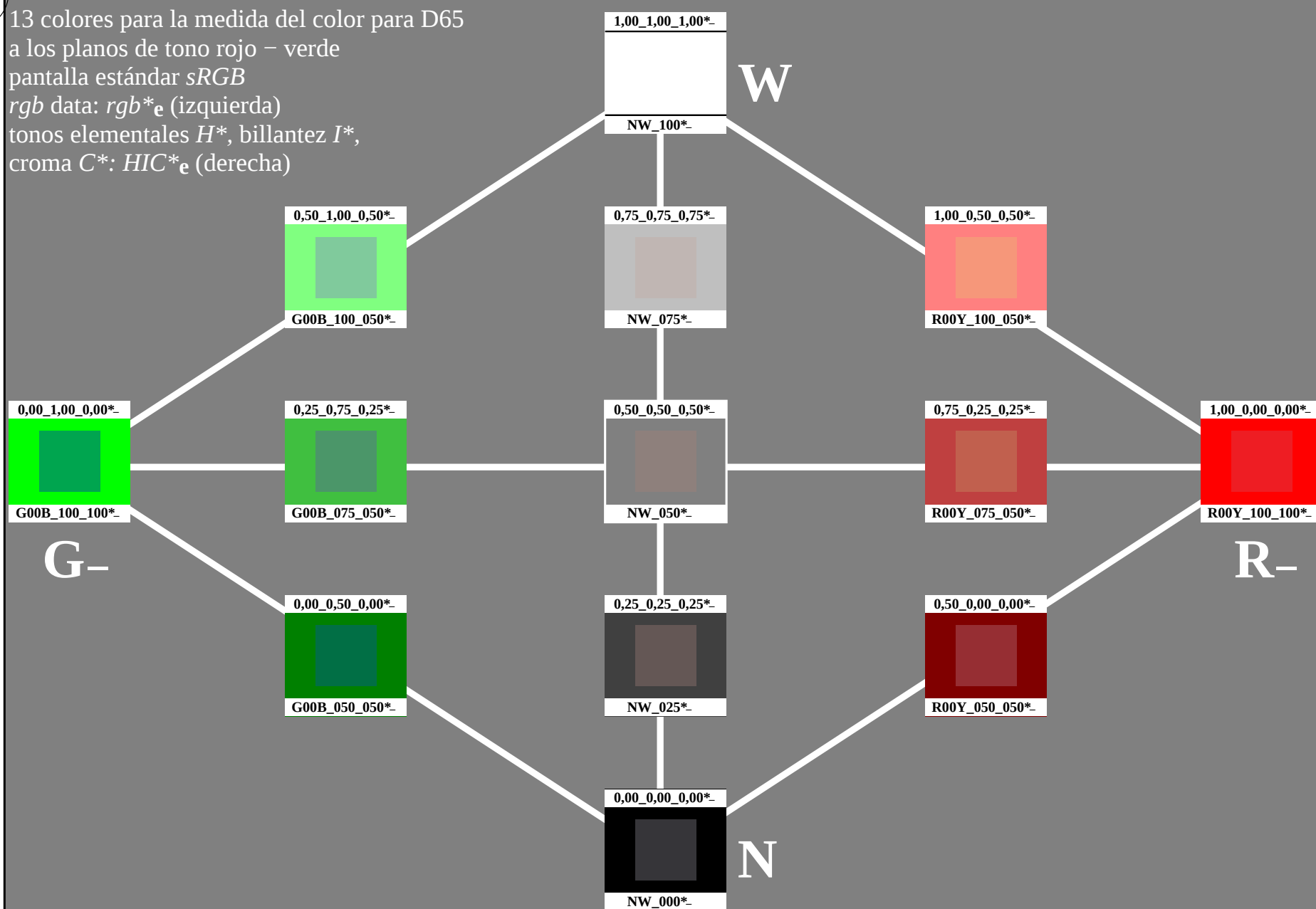


13 colores para la medida del color para D65
a los planos de tono rojo - verde
pantalla estándar sRGB
rgb data: *rgb**_e (izquierda)
tonos elementales *H**, billantez *I**,
croma *C**: *HIC**_e (derecha)



2-003031-L0

PS570-7N

gráfico TUB-PS57; tonos rojo - verde
13 colores del estándar para D65

entrada: *rgb/cmyk* -> *rgb/cmyk*
salida: ningún cambio

TUB matrícula: 20130201-PS57/PS57L0NA.TXT /.PS
aplicación para la medida salida en la impresión offset

TUB material: code=rha4ta

13 colores para la medida del color para D65
a los planos de tono rojo - verde
pantalla estándar *sRGB*
rgb data: rgb^*_e (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : HIC^*_e (derecha)

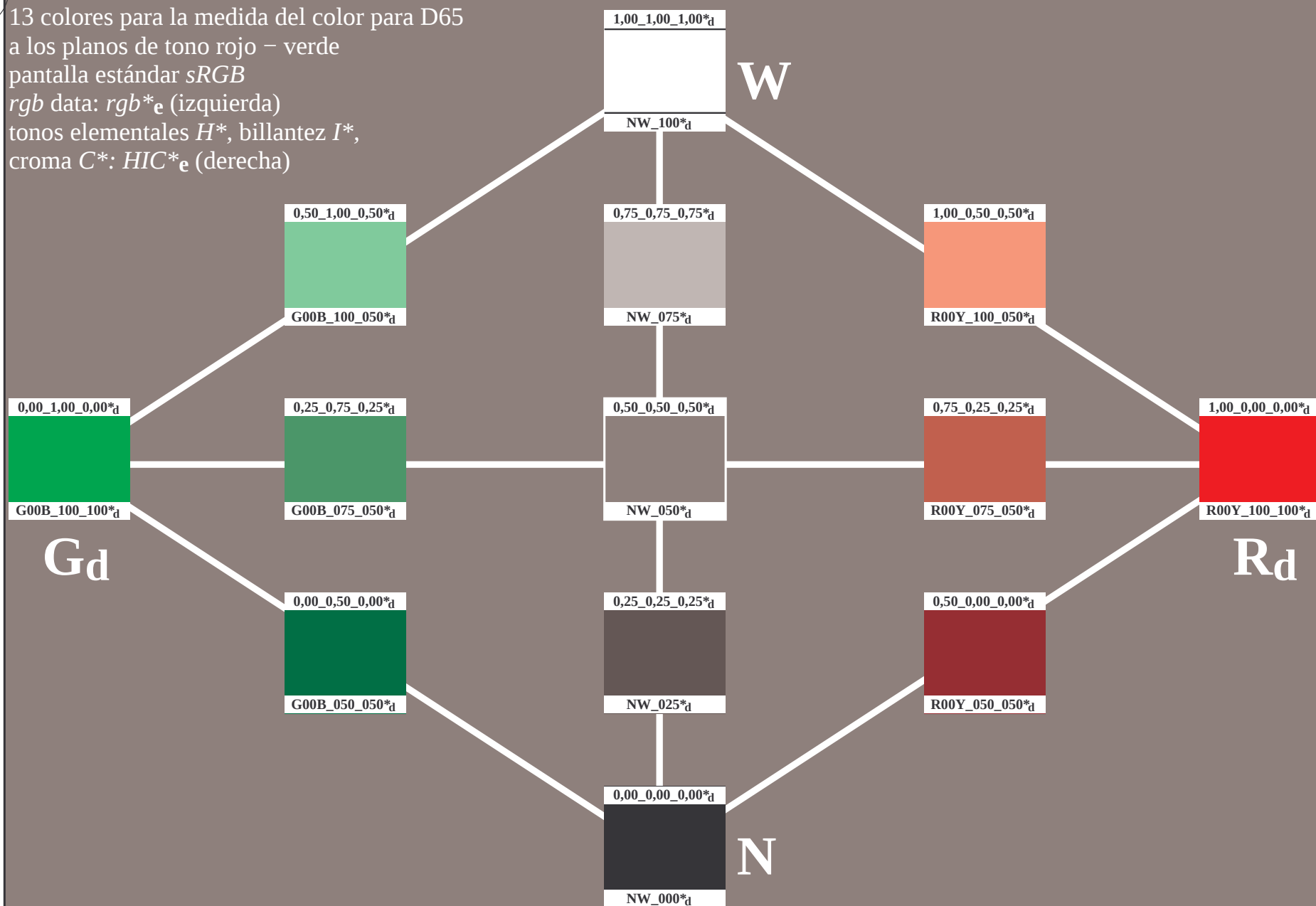
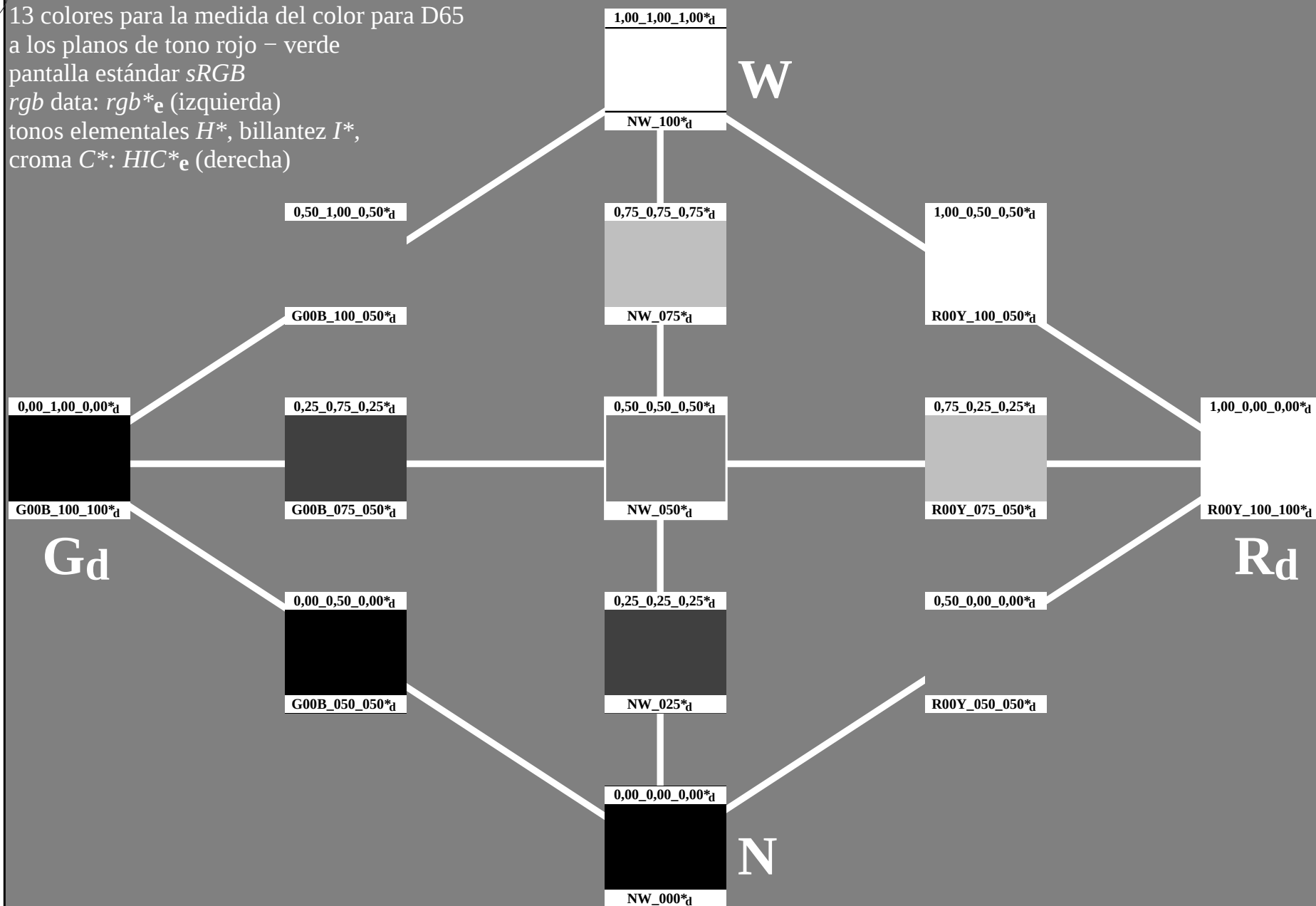


gráfico TUB-PS57; tonos rojo - verde
13 colores del estándar para D65, 3D=0, de=0, *cmy0*

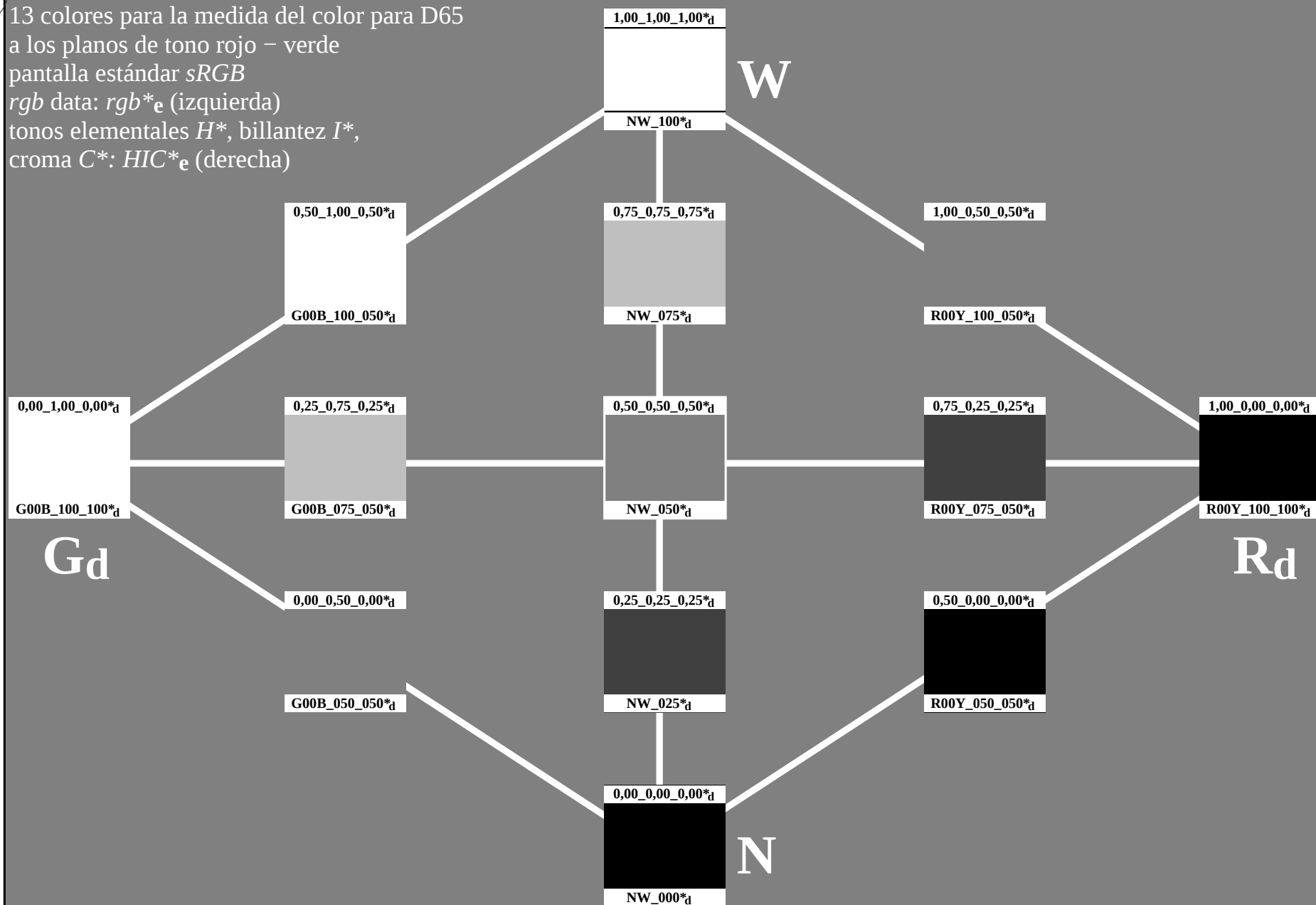
entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmy0_d$

PE4600L_120830.TXT, 1080 colors, Separation *cmy0**

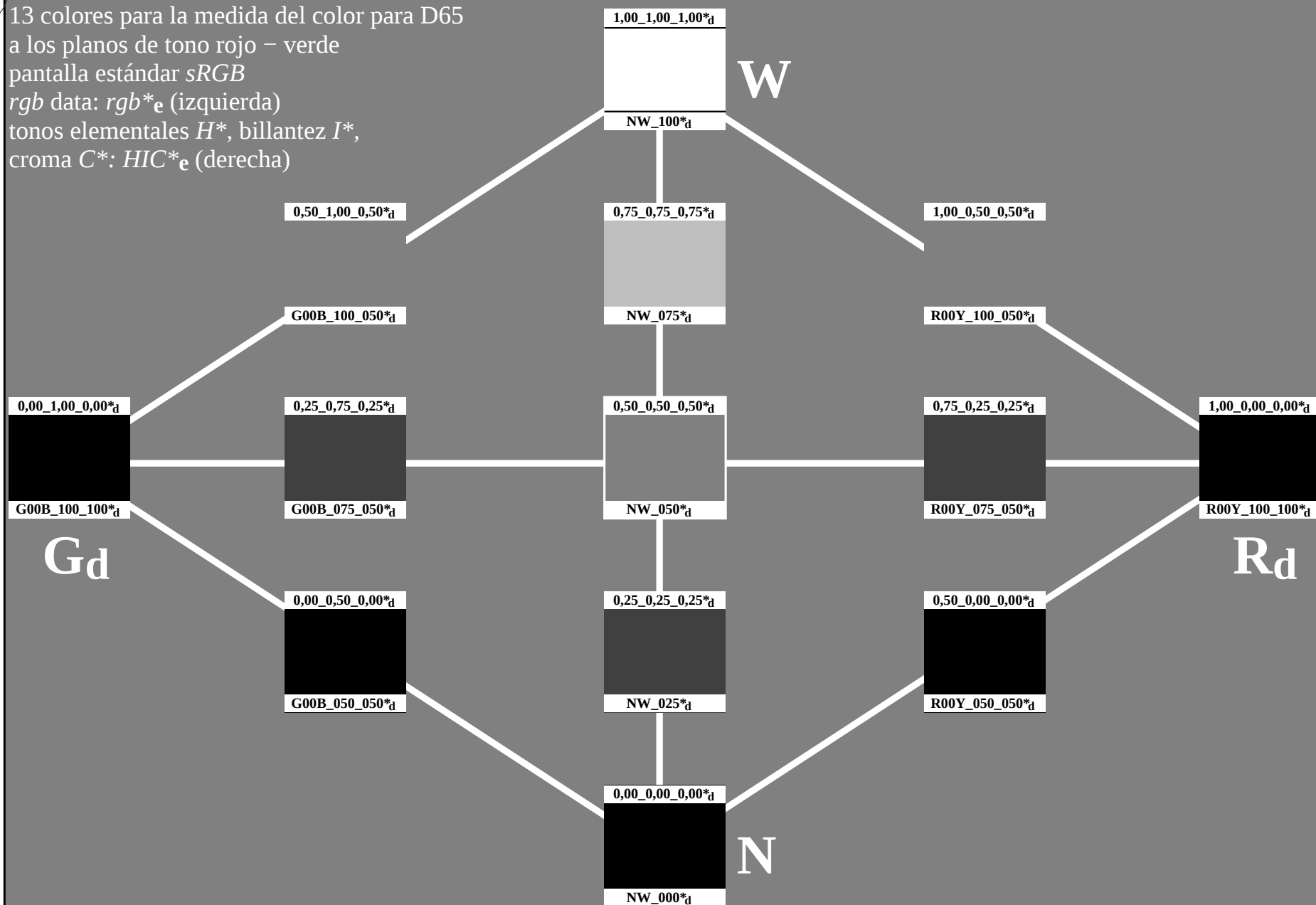
13 colores para la medida del color para D65
a los planos de tono rojo - verde
pantalla estándar *sRGB*
rgb data: $rgb*_e$ (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : $HIC*_e$ (derecha)

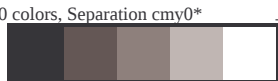


13 colores para la medida del color para D65
a los planos de tono rojo – verde
pantalla estándar *sRGB*
rgb data: $rgb*_e$ (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : $HIC*_e$ (derecha)



13 colores para la medida del color para D65
a los planos de tono rojo - verde
pantalla estándar *sRGB*
rgb data: rgb^*_e (izquierda)
tonos elementales H^* , billantez I^* ,
croma C^* : HIC^*_e (derecha)

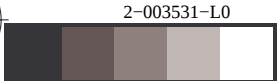




PE4600L_120830.TXT, 1080 colors, Separation cmy0*

entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmy0_d$

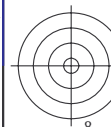
gráfico TUB-PS57; tonos rojo – verde
13 colores del estándar para D65, 3D=0, de=0, $cmy0$



2-003531-E0

PS570-70

2-003531-L0



<http://130.149.60.45/~farbmetrik/PS57/PS57L0NA.TXT> /.PS; salida de transferencia
N: ninguna 3D-linealización (OL) en archivo (F) o PS-startup (S), página 6/26



13 colores para la medida del color para D65

a los planos de tono rojo - verde
pantalla estándar *sRGB*

rgb data: *rgb***e* (izquierda)

tonos elementales *H**, billantez *I**,

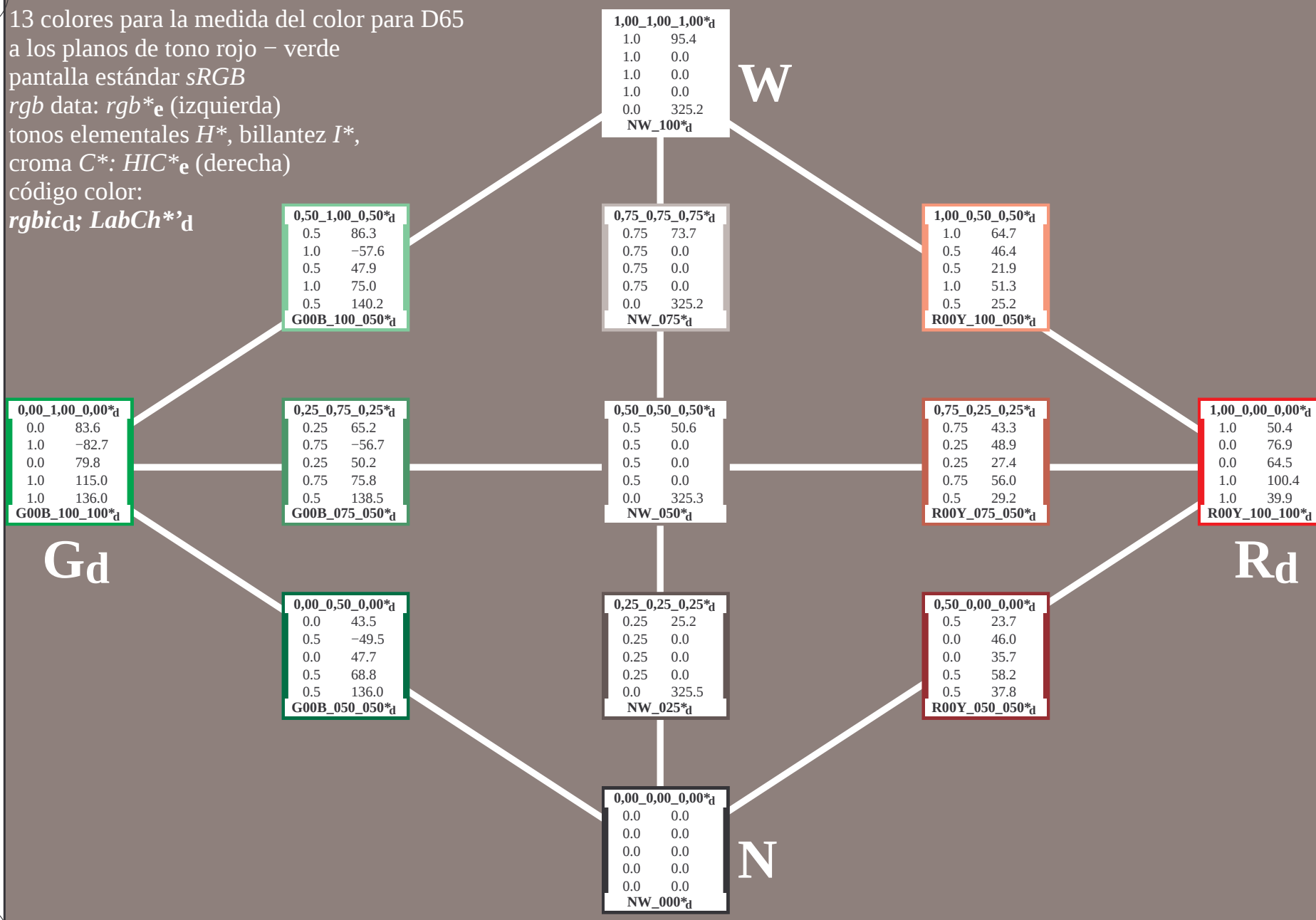
croma *C**: *HIC***e* (derecha)

código color:

rgbicd; *LabCh***'d*

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS57/PS57L0NA.TXT /.PS
aplicación para la medida salida en la impresión offset, separación *cmY0* (CMY0)
TUB material: code=rha4ta



2-003631-L0

PS570-70

PE4600L_120830.TXT, 1080 colors, Separation *cmY0**

gráfico TUB-PS57; tonos rojo - verde
13 colores del estándar para D65, 3D=0, de=0, *cmY0*

entrada: *rgb/cmyk* -> *rgb*_d
salida: transfiera a *cmY0*_d

13 colores para la medida del color para D65

a los planos de tono rojo - verde

pantalla estándar *sRGB*

rgb data: $rgb * e$ (izquierda)

tonos elementales H^* , billantez I^* ,

croma C^* : $HIC * e$ (derecha)

código color:

$rgbic' * d$; $LabCh * d$

0,50_1,00_0,50*d		
0.5	72.8	
1.0	-32.5	
0.5	14.8	
1.0	35.7	
0.5	155.5	
G00B_100_050*d		

1,00_1,00_1,00*d		
1.0	95.6	
1.0	0.0	
1.0	0.0	
1.0	0.0	
0.0	0.0	
NW_100*d		

W

0,75_0,75_0,75*d		
0.75	77.8	
0.75	0.0	
0.75	0.0	
0.75	0.0	
0.0	0.0	
NW_075*d		

1,00_0,50_0,50*d		
1.0	70.5	
0.5	35.4	
0.5	22.4	
1.0	41.9	
0.5	32.3	
R00Y_100_050*d		

0,00_1,00_0,00*d		
0.0	50.0	
1.0	-65.0	
0.0	29.6	
1.0	71.4	
1.0	155.5	
G00B_100_100*d		

0,25_0,75_0,25*d		
0.25	55.0	
0.75	-32.5	
0.25	14.8	
0.75	35.7	
0.5	155.5	
G00B_075_050*d		

0,50_0,50_0,50*d		
0.5	60.0	
0.5	0.0	
0.5	0.0	
0.5	0.0	
0.0	0.0	
NW_050*d		

0,75_0,25_0,25*d		
0.75	52.7	
0.25	35.4	
0.25	22.4	
0.75	41.9	
0.5	32.3	
R00Y_075_050*d		

1,00_0,00_0,00*d		
1.0	45.4	
0.0	70.9	
0.0	44.8	
1.0	83.9	
1.0	32.3	
R00Y_100_100*d		

Gd

0,00_0,50_0,00*d		
0.0	37.2	
0.5	-32.5	
0.0	14.8	
0.5	35.7	
0.5	155.5	
G00B_050_050*d		

0,25_0,25_0,25*d		
0.25	42.1	
0.25	0.0	
0.25	0.0	
0.25	0.0	
0.0	0.0	
NW_025*d		

0,50_0,00_0,00*d		
0.5	34.9	
0.0	35.4	
0.0	22.4	
0.5	41.9	
0.5	32.3	
R00Y_050_050*d		

Rd

0,00_0,00_0,00*d		
0.0	24.3	
0.0	0.0	
0.0	0.0	
0.0	0.0	
0.0	0.0	
NW_000*d		

N

2-003731-L0

PS570-70

PE4600L_120830.TXT, 1080 colors, Separation cmy0*

gráfico TUB-PS57; tonos rojo - verde
13 colores del estándar para D65, 3D=0, de=0, *cmy0*

entrada: *rgb/cmyk* -> rgb_d
salida: transfiera a $cmy0_d$

13 colores para la medida del color para D65

a los planos de tono rojo - verde

pantalla estándar *sRGB*

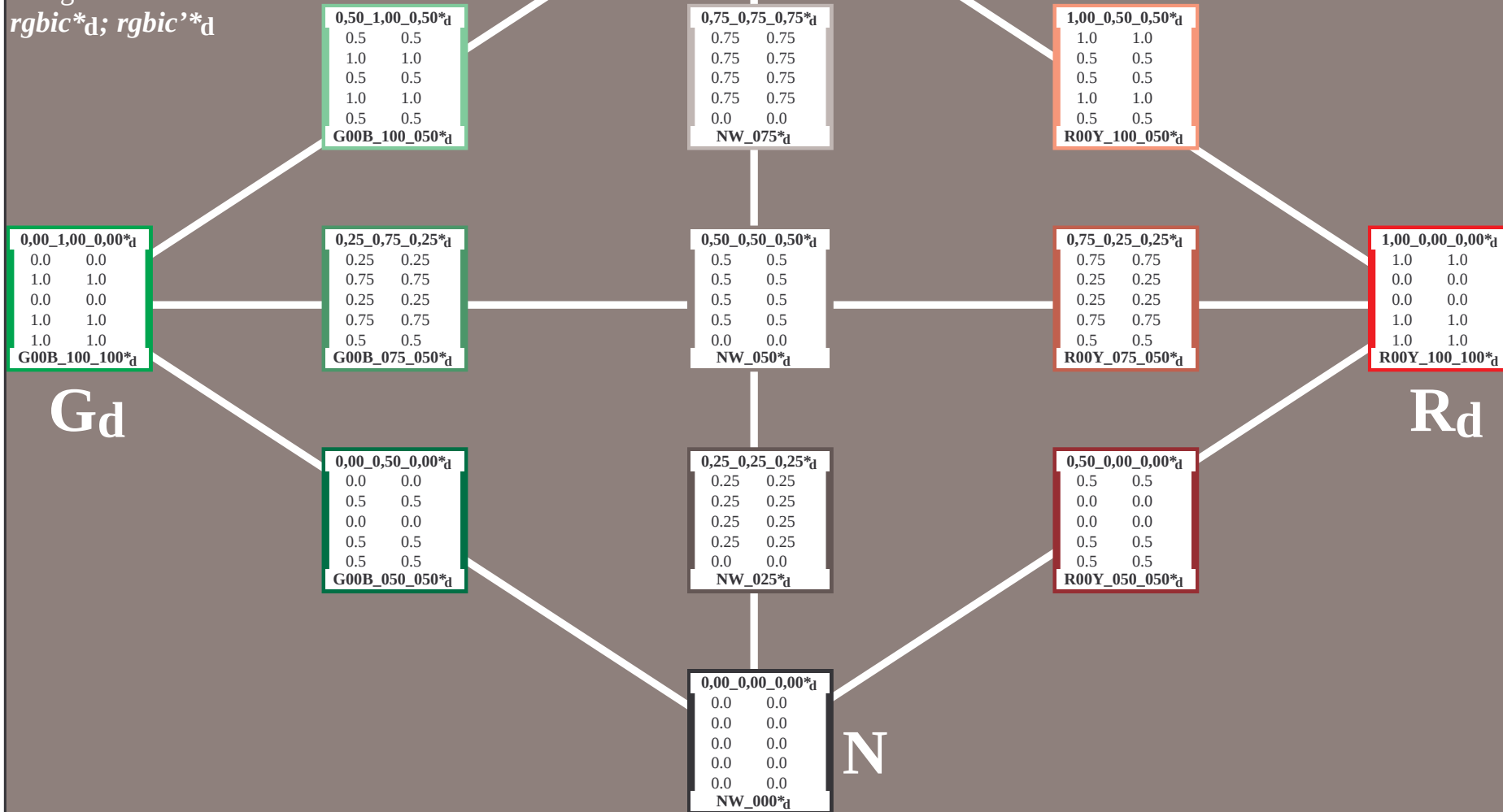
rgb data: $rgb * e$ (izquierda)

tonos elementales H^* , billantez I^* ,

croma C^* : $HIC * e$ (derecha)

código color:

$rgbic * d$; $rgbic' * d$



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PS570-70

PE4600L_120830.TXT, 1080 colors, Separation cmy0*

gráfico TUB-PS57; tonos rojo - verde
13 colores del estándar para D65, 3D=0, de=0, *cmy0*

entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmy0_d$

13 colores para la medida del color para D65

a los planos de tono rojo - verde

pantalla estándar *sRGB*

rgb data: $rgb*_e$ (izquierda)

tonos elementales H^* , billantez I^* ,

croma C^* : $HIC*_e$ (derecha)

código color:

$LabCh*_d$; $Lab^*/DE^*/h^*_d$

$0,50_1,00_0,50*_d$

86.3	72.8
-57.6	-32.5
47.9	14.8
75.0	43.7
140.2	155.5

$G00B_100_050*_d$

$1,00_1,00_1,00*_d$

95.4	95.6
0.0	0.0
0.0	0.0
0.0	0.2
325.2	0.0

NW_100*_d

W

$0,75_0,75_0,75*_d$

73.7	77.8
0.0	0.0
0.0	0.0
0.0	4.0
325.2	0.0

NW_075*_d

$1,00_0,50_0,50*_d$

64.7	70.5
46.4	35.4
21.9	22.4
51.3	12.4
25.2	32.3

$R00Y_100_050*_d$

$0,00_1,00_0,00*_d$

83.6	50.0
-82.7	-65.0
79.8	29.6
115.0	63.0
136.0	155.5

$G00B_100_100*_d$

$0,25_0,75_0,25*_d$

65.2	55.0
-56.7	-32.5
50.2	14.8
75.8	44.1
138.5	155.5

$G00B_075_050*_d$

$0,50_0,50_0,50*_d$

50.6	60.0
0.0	0.0
0.0	0.0
0.0	9.3
325.3	0.0

NW_050*_d

$0,75_0,25_0,25*_d$

43.3	52.7
48.9	35.4
27.4	22.4
56.0	17.1
29.2	32.3

$R00Y_075_050*_d$

$1,00_0,00_0,00*_d$

50.4	45.4
76.9	70.9
64.5	44.8
100.4	21.1
39.9	32.3

$R00Y_100_100*_d$

Gd

Rd

$0,00_0,50_0,00*_d$

43.5	37.2
-49.5	-32.5
47.7	14.8
68.8	37.6
136.0	155.5

$G00B_050_050*_d$

$0,25_0,25_0,25*_d$

25.2	42.1
0.0	0.0
0.0	0.0
0.0	16.9
325.5	0.0

NW_025*_d

$0,50_0,00_0,00*_d$

23.7	34.9
46.0	35.4
35.7	22.4
58.2	20.3
37.8	32.3

$R00Y_050_050*_d$

$0,00_0,00_0,00*_d$

0.0	24.3
0.0	0.0
0.0	0.0
0.0	24.3
0.0	0.0

NW_000*_d

N

2-003931-L0

PS570-70

PE4600L_120830.TXT, 1080 colors, Separation cmy0*

gráfico TUB-PS57; tonos rojo - verde
13 colores del estándar para D65, 3D=0, de=0, *cmy0*

entrada: *rgb/cmyk* -> rgb_d
salida: transfiera a $cmy0_d$

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS57/PS57L0NA.TXT /.PS TUB material: code=rh4ta
aplicación para la medida salida en la impresión offset, separación cmy0 (CMY0)

n/f	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiM,d	rgb*Md	LabCh*Md
0/648	R00Y_100_100d	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	0.0 0.0	389	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3
1/657	R13Y_100_100d	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	48.6 63.3 49.1 80.2 37.7	1.0 0.125 0.0	48.9 62.8 49.4 79.9 38.1	0.6 0.6	36	1.0 0.116 0.0	48.6 63.3 49.1 80.2 37.7
2/666	R25Y_100_100d	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.0 53.4 54.8 76.5 45.7	1.0 0.25 0.0	53.6 51.9 55.5 76.0 46.8	1.7 4.2	42	1.0 0.233 0.0	53.0 53.4 54.8 76.5 45.7
3/675	R38Y_100_100d	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	58.8 41.1 61.7 74.1 56.3	1.0 0.375 0.0	59.1 40.3 62.0 74.0 56.9	0.9 51	51	1.0 0.366 0.0	58.8 41.1 61.7 74.1 56.3
4/684	R50Y_100_100d	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	64.9 28.9 68.6 74.5 67.1	1.0 0.5 0.0	64.9 28.9 68.6 74.5 67.1	0.0 59	59	1.0 0.5 0.0	64.9 28.9 68.6 74.5 67.1
5/693	R63Y_100_100d	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	72.5 14.8 77.6 79.0 79.1	1.0 0.625 0.0	72.1 15.4 77.1 78.6 78.6	0.8 68	68	1.0 0.633 0.0	72.5 14.8 77.6 79.0 79.1
6/702	R75Y_100_100d	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	78.6 4.3 84.7 84.8 87.0	1.0 0.75 0.0	77.9 5.4 83.8 84.0 86.2	1.6 77	77	1.0 0.766 0.0	78.6 4.3 84.7 84.8 87.0
7/711	R88Y_100_100d	1.0 0.875 0.0	1.0 1.0 0.5	83	1.0 0.883 0.0	83.7 -3.8 90.5 90.6 92.4	1.0 0.875 0.0	83.4 -3.4 90.2 90.2 92.1	0.6 83	83	1.0 0.883 0.0	83.7 -3.8 90.5 90.6 92.4
8/720	Y00G_100_100d	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	87.8 -10.2 95.4 96.0 96.1	1.0 1.0 0.0	87.8 -10.2 95.4 96.0 96.1	0.0 89	89	1.0 1.0 0.0	87.8 -10.2 95.4 96.0 96.1
9/639	Y13G_100_100d	0.875 1.0 0.0	1.0 1.0 0.5	97	0.883 1.0 0.0	84.5 -13.6 89.7 90.7 98.6	0.875 1.0 0.0	84.3 -13.9 89.2 90.3 98.8	0.5 96	96	0.883 1.0 0.0	84.5 -13.6 89.7 90.7 98.6
10/558	Y25G_100_100d	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	81.2 -17.0 84.3 86.0 101.4	0.75 1.0 0.0	80.7 -17.5 83.5 85.3 101.8	1.0 102	102	0.766 1.0 0.0	81.2 -17.0 84.3 86.0 101.4
11/477	Y38G_100_100d	0.625 1.0 0.0	1.0 1.0 0.5	112	0.633 1.0 0.0	75.6 -23.6 76.2 79.8 107.2	0.625 1.0 0.0	75.3 -24.0 75.7 79.4 107.6	0.7 111	111	0.625 1.0 0.0	75.6 -23.6 76.2 79.8 107.2
12/396	Y50G_100_100d	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	70.6 -29.7 66.5 72.8 114.0	0.5 1.0 0.0	70.6 -29.7 66.5 72.8 114.0	0.0 119	119	0.5 1.0 0.0	70.6 -29.7 66.5 72.8 114.0
13/315	Y63G_100_100d	0.375 1.0 0.0	1.0 1.0 0.5	128	0.366 1.0 0.0	65.2 -36.4 57.6 68.2 122.3	0.375 1.0 0.0	65.7 -35.6 58.3 68.3 121.4	1.2 128	128	0.366 1.0 0.0	65.2 -36.4 57.6 68.2 122.3
14/234	Y75G_100_100d	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	57.9 -48.3 45.8 66.5 136.5	0.25 1.0 0.0	58.4 -47.3 46.8 66.6 135.3	1.4 137	137	0.233 1.0 0.0	57.9 -48.3 45.8 66.5 136.5
15/153	Y88G_100_100d	0.125 1.0 0.0	1.0 1.0 0.5	143	0.116 1.0 0.0	54.4 -54.7 38.0 66.6 145.1	0.125 1.0 0.0	54.7 -53.9 38.5 66.3 144.4	0.9 143	143	0.116 1.0 0.0	54.4 -54.7 38.0 66.6 145.1
16/72	G00C_100_100d	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	0.0 149	149	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5
17/73	G13C_100_100d	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.116	50.5 -62.9 22.4 66.8 160.4	0.0 1.0 0.125	50.5 -62.8 21.9 66.5 160.7	0.5 156	156	0.0 1.0 0.116	50.5 -62.9 22.4 66.8 160.4
18/74	G25C_100_100d	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.233	51.1 -59.5 13.9 61.1 166.8	0.0 1.0 0.25 0.5	51.2 -58.9 12.7 60.3 167.7	1.2 162	162	0.0 1.0 0.233	51.1 -59.5 13.9 61.1 166.8
19/75	G38C_100_100d	0.0 1.0 0.375	1.0 1.0 0.5	172	0.0 1.0 0.366	51.9 -54.9 3.7 55.0 176.1	0.0 1.0 0.375 0.5	52.0 -54.5 3.1 54.6 176.7	0.6 171	171	0.0 1.0 0.366	51.9 -54.9 3.7 55.0 176.1
20/76	G50C_100_100d	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	52.9 -48.6 -8.0 49.3 189.3	0.0 1.0 0.5 0.5	52.9 -48.6 -8.0 49.3 189.3	0.0 180	180	0.0 1.0 0.5	52.9 -48.6 -8.0 49.3 189.3
21/77	G63C_100_100d	0.0 1.0 0.625	1.0 1.0 0.5	188	0.0 1.0 0.633	54.1 -42.0 -18.8 46.0 204.1	0.0 1.0 0.625 0.5	54.0 -42.3 -18.1 46.1 203.2	0.7 188	188	0.0 1.0 0.633	54.1 -42.0 -18.8 46.0 204.1
22/78	G75C_100_100d	0.0 1.0 0.75	1.0 1.0 0.5	196	0.0 1.0 0.766	55.1 -35.4 -28.4 45.4 218.7	0.0 1.0 0.75 0.5	55.0 -36.0 -27.4 45.3 217.2	1.1 197	197	0.0 1.0 0.766	55.1 -35.4 -28.4 45.4 218.7
23/79	G88C_100_100d	0.0 1.0 0.875	1.0 1.0 0.5	203	0.0 1.0 0.883	55.9 -30.4 -35.0 46.3 229.0	0.0 1.0 0.875 0.5	55.8 -30.7 -34.5 46.2 228.3	0.5 203	203	0.0 1.0 0.883	55.9 -30.4 -35.0 46.3 229.0
24/80	C00B_100_100d	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5 48.7 238.4	0.0 1.0 1.0	56.8 -25.5 -41.5 48.7 238.4	0.0 210	210	0.0 1.0 1.0	56.8 -25.5 -41.5 48.7 238.4
25/71	C13B_100_100d	0.0 0.875 1.0	1.0 1.0 0.5	217	0.0 0.883 1.0	54.3 -21.4 -41.4 46.6 242.6	0.0 0.875 1.0	54.1 -21.1 -41.3 46.4 242.9	0.3 216	216	0.0 0.883 1.0	54.3 -21.4 -41.4 46.6 242.6
26/62	C25B_100_100d	0.0 0.75 1.0	1.0 1.0 0.5	224	0.0 0.766 1.0	50.9 -16.2 -41.2 44.2 248.4	0.0 0.75 1.0	50.4 -15.5 -41.1 43.9 249.3	0.8 222	222	0.0 0.766 1.0	50.9 -16.2 -41.2 44.2 248.4
27/53	C38B_100_100d	0.0 0.625 1.0	1.0 1.0 0.5	232	0.0 0.633 1.0	46.8 -9.8 -40.9 42.1 256.4	0.0 0.625 1.0	46.5 -9.4 -40.8 41.9 256.9	0.4 231	231	0.0 0.633 1.0	46.8 -9.8 -40.9 42.1 256.4
28/44	C50B_100_100d	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	41.7 -1.2 -40.6 40.6 268.2	0.0 0.5 1.0	41.7 -1.2 -40.6 40.6 268.2	0.0 240	240	0.0 0.5 1.0	41.7 -1.2 -40.6 40.6 268.2
29/35	C63B_100_100d	0.0 0.375 1.0	1.0 1.0 0.5	248	0.0 0.366 1.0	37.0 6.6 -40.2 40.8 279.3	0.0 0.375 1.0	37.3 6.1 -40.2 40.7 278.6	0.6 248	248	0.0 0.366 1.0	37.0 6.6 -40.2 40.8 279.3
30/26	C75B_100_100d	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.233 1.0	32.2 15.3 -40.3 43.1 290.8	0.0 0.25 1.0	32.8 14.3 -40.2 42.7 289.6	1.1 257	257	0.0 0.233 1.0	32.2 15.3 -40.3 43.1 290.8
31/17	C88B_100_100d	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.116 1.0	28.4 22.8 -40.3 46.3 299.5	0.0 0.125 1.0	28.6 22.4 -40.2 46.1 299.0	0.5 263	263	0.0 0.116 1.0	28.4 22.8 -40.3 46.3 299.5
32/8	B00M_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	25.0 29.5 -40.4 50.0 306.2	0.0 0.0 1.0	25.0 29.5 -40.4 50.0 306.2	0.0 270	270	0.0 0.0 1.0	25.0 29.5 -40.4 50.0 306.2
33/89	B13M_100_100d	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	27.7 35.6 -36.7 51.1 314.1	0.125 0.0 1.0	27.9 36.0 -36.4 51.2 314.7	0.5 276	276	0.116 0.0 1.0	27.7 35.6 -36.7 51.1 314.1
34/170	B25M_100_100d	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	28.7 41.2 -33.1 52.9 321.1	0.25 0.0 1.0	28.8 41.9 -32.5 53.1 322.1	0.9 282	282	0.233 0.0 1.0	28.7 41.2 -33.1 52.9 321.1
35/251	B38M_100_100d	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	32.5 51.2 -26.5 57.7 332.6	0.375 0.0 1.0	32.7 51.8 -26.0 58.0 333.3	0.8 291	291	0.366 0.0 1.0	32.5 51.2 -26.5 57.7 332.6
36/332	B50M_100_100d	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	35.6 58.6 -20.7 62.1 340.5	0.5 0.0 1.0	35.6 58.6 -20.7 62.1 340.5	0.0 300	300	0.5 0.0 1.0	35.6 58.6 -20.7 62.1 340.5
37/413	B63M_100_100d	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	38.3 65.8 -13.7 67.2 348.2	0.625 0.0 1.0	38.1 65.4 -14.0 66.9 347.9	0.5 308	308	0.633 0.0 1.0	38.3 65.8 -13.7 67.2 348.2
38/494	B75M_100_100d	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	42.1 71.6 -8.7 72.1 353.0	0.75 0.0 1.0	41.8 71.0 -9.2 71.6 352.5	0.8 317	317	0.766 0.0 1.0	42.1 71.6 -8.7 72.1 353.0
39/575	B88M_100_100d	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	44.3 75.4 -4.7 75.6 356.3	0.875 0.0 1.0	44.2 75.2 -5.0 75.3 356.1	0.4 323	323	0.883 0.0 1.0	44.3 75.4 -4.7 75.6 356.3
40/656	M00R_100_100d	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2 79.3 359.8	1.0 0.0 1.0	46.1 79.3 -0.2 79.3 359.8	0.0 330	330	1.0 0.0 1.0	46.1 79.3 -0.2 79.3 359.8
41/655	M13R_100_100d	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	45.9 78.3 3.8 78.4 2.8	1.0 0.0 0.875 0.5	45.9 78.2 4.1 78.3 363.0	0.2 336	336	1.0 0.0 0.883	45.9 78.3 3.8 78.4 2.8
42/654	M25R_100_100d	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	45.9 77.3 8.0 77.7 5.9	1.0 0.0 0.75 0.5	45.9 77.1 8.6 77.6 366.4	0.6 342	342	1.0 0.0 0.766	45.9 77.3 8.0 77.7 5.9
43/653	M38R_100_100d	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	46.0 75.7 14.4 77.1 10.8	1.0 0.0 0.625 0.6	46.0 75.6 14.8 77.0 371.1	0.4 351	351	1.0 0.0 0.633	46.0 75.7 14.4 77.1 10.8
44/652	M50R_100_100d	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	45.9 74.2 21.1 77.1 15.9	1.0 0.0 0.5 0.5	45.9 74.2 21.1 77.1 375.9	0.0 360	360	1.0 0.0 0.5	45.9 74.2 21.1

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS57/PS57L0NA.TXT /.PS TUB material: code=rha4ta
aplicación para la medida salida en la impresión offset, separación cmy0 (CMY0)

n/f	HIC*Fd	rgb_Fd			ict_Fd			hsi_Fd	rgb*Fd			LabCh*Fd			rgb**Fd			LabCh**Fd			DE**Fd			hsiMd	rgb*Md			LabCh*Md						
0/648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3		
1/666	R25Y_100_100a	1.0	0.25	0.0	1.0	1.0	0.5	44	1.0	0.233	0.0	53.0	53.4	54.8	76.5	45.7	1.0	0.25	0.0	53.6	51.9	55.5	76.0	46.8	1.7	42	1.0	0.233	0.0	53.0	53.4	54.8	76.5	45.7
2/684	R50Y_100_100a	1.0	0.5	0.0	1.0	1.0	0.5	60	1.0	0.5	0.0	64.9	28.9	68.6	74.5	67.1	1.0	0.5	0.0	64.9	28.9	68.6	74.5	67.1	1.0	0.5	0.0	64.9	28.9	68.6	74.5	67.1		
3/702	R75Y_100_100a	1.0	0.75	0.0	1.0	1.0	0.5	76	1.0	0.766	0.0	78.6	4.3	84.7	84.8	87.0	1.0	0.75	0.0	77.9	5.4	83.8	84.0	86.2	1.6	77	1.0	0.766	0.0	78.6	4.3	84.7	84.8	87.0
4/720	Y00G_100_100a	1.0	1.0	0.0	1.0	1.0	0.5	90	1.0	1.0	0.0	87.8	-10.2	95.4	96.0	96.1	1.0	1.0	0.0	87.8	-10.2	95.4	96.0	96.1	0.0	89	1.0	1.0	0.0	87.8	-10.2	95.4	96.0	96.1
5/558	Y25G_100_100a	0.75	1.0	0.0	1.0	1.0	0.5	104	0.766	1.0	0.0	81.2	-17.0	84.3	86.0	101.4	0.75	1.0	0.0	80.7	-17.5	83.5	85.3	101.8	1.0	102	0.766	1.0	0.0	81.2	-17.0	84.3	86.0	101.4
6/396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	120	0.5	1.0	0.0	70.6	-29.7	66.5	72.8	114.0	0.5	1.0	0.0	70.6	-29.7	66.5	72.8	114.0	0.0	119	0.5	1.0	0.0	70.6	-29.7	66.5	72.8	114.0
7/234	Y75G_100_100a	0.25	1.0	0.0	1.0	1.0	0.5	136	0.233	1.0	0.0	57.9	-48.3	45.8	66.5	136.5	0.25	1.0	0.0	58.4	-47.3	46.8	66.6	135.3	1.4	137	0.233	1.0	0.0	57.9	-48.3	45.8	66.5	136.5
8/72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5	0.0	149	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5
9/72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5	0.0	149	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5
10/76	G25B_100_100a	0.0	1.0	0.5	1.0	1.0	0.5	180	0.0	1.0	0.5	52.9	-48.6	-8.0	49.3	189.3	0.0	1.0	0.5	52.9	-48.6	-8.0	49.3	189.3	0.0	180	0.0	1.0	0.5	52.9	-48.6	-8.0	49.3	189.3
11/80	G50B_100_100a	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	1.0	56.8	-25.5	-41.5	48.7	238.4	0.0	1.0	1.0	56.8	-25.5	-41.5	48.7	238.4	0.0	210	0.0	1.0	1.0	56.8	-25.5	-41.5	48.7	238.4
12/44	G75B_100_100a	0.0	0.5	1.0	1.0	1.0	0.5	240	0.0	0.5	1.0	41.7	-1.2	-40.6	40.6	268.2	0.0	0.5	1.0	41.7	-1.2	-40.6	40.6	268.2	0.0	240	0.0	0.5	1.0	41.7	-1.2	-40.6	40.6	268.2
13/8	B00R_100_100a	0.0	0.0	1.0	1.0	1.0	0.5	270	0.0	0.0	1.0	25.0	29.5	-40.4	50.0	306.2	0.0	0.0	1.0	25.0	29.5	-40.4	50.0	306.2	0.0	270	0.0	0.0	1.0	25.0	29.5	-40.4	50.0	306.2
14/332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	300	0.5	0.0	1.0	35.6	58.6	-20.7	62.1	340.5	0.5	0.0	1.0	35.6	58.6	-20.7	62.1	340.5	0.0	300	0.5	0.0	1.0	35.6	58.6	-20.7	62.1	340.5
15/656	B50R_100_100a	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8	0.0	330	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8
16/652	B75R_100_100a	1.0	0.0	0.5	1.0	1.0	0.5	360	1.0	0.0	0.5	45.9	74.2	21.1	77.1	15.9	1.0	0.0	0.5	45.9	74.2	21.1	77.1	15.9	0.0	360	1.0	0.0	0.5	45.9	74.2	21.1	77.1	15.9
17/648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3	0.0	389	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3
18/688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	390	1.0	0.5	0.5	70.5	35.4	22.4	41.9	32.3	1.0	0.5	0.5	68.0	29.9	28.7	41.5	43.8	8.7	389	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3
19/706	R50Y_100_050a	1.0	0.75	0.5	1.0	0.5	0.75	60	1.0	0.75	0.5	80.2	14.4	34.3	37.2	67.1	1.0	0.75	0.5	80.4	9.0	35.3	36.5	75.5	5.4	59	1.0	0.5	0.0	64.9	28.9	68.6	74.5	67.1
20/724	Y00G_100_050a	1.0	1.0	0.5	1.0	0.5	0.75	90	1.0	1.0	0.5	91.7	-5.1	47.7	48.0	96.1	1.0	1.0	0.5	91.4	-7.7	42.5	43.2	100.3	5.8	89	1.0	1.0	0.0	87.8	-10.2	95.4	96.0	96.1
21/562	Y50G_100_050a	0.75	1.0	0.5	1.0	0.5	0.75	120	0.75	1.0	0.5	83.1	-14.8	33.2	36.4	114.0	0.75	1.0	0.5	84.2	-14.1	31.5	34.5	114.0	2.1	119	0.5	1.0	0.0	70.6	-29.7	66.5	72.8	114.0
22/490	G00B_100_050a	0.5	1.0	0.5	1.0	0.5	0.75	150	0.5	1.0	0.5	72.8	-32.5	14.8	35.7	155.5	0.5	1.0	0.5	73.9	-23.7	19.9	31.0	140.0	10.1	149	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5
23/404	G50B_100_050a	0.5	1.0	1.0	1.0	0.5	0.75	210	0.5	1.0	1.0	76.2	-12.7	-20.7	24.3	238.4	0.5	1.0	1.0	78.7	-11.6	-18.3	21.7	237.6	3.6	210	0.0	1.0	1.0	56.8	-25.5	-41.5	48.7	238.4
24/368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	270	0.5	0.5	1.0	60.3	14.7	-20.2	25.0	306.2	0.5	0.5	1.0	57.9	18.3	-20.7	27.7	311.4	4.3	270	0.0	1.0	1.0	25.0	29.5	-40.4	50.0	306.2
25/692	B50R_100_050a	1.0	0.5	1.0	1.0	0.5	0.75	330	1.0	0.5	1.0	70.8	39.6	-0.1	39.6	359.8	1.0	0.5	1.0	70.7	35.2	-3.7	35.4	353.9	5.7	330	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8
26/688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	390	1.0	0.5	0.5	70.5	35.4	22.4	41.9	32.3	1.0	0.5	0.5	68.0	29.9	28.7	41.5	43.8	8.7	389	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3
27/506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	390	0.75	0.25	0.25	52.7	35.4	22.4	41.9	32.3	0.75	0.25	0.25	50.4	39.4	31.9	50.7	38.9	10.5	389	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3
28/524	R50Y_075_050a	0.75	0.5	0.25	0.75	0.5	0.5	60	0.75	0.5	0.25	62.4	14.4	34.3	37.2	67.1	0.75	0.5	0.25	61.2	18.1	39.5	43.4	65.3	6.4	59	1.0	0.5	0.0	64.9	28.9	68.6	74.5	67.1
29/542	Y00G_075_050a	0.75	0.75	0.25	0.75	0.5	0.5	90	0.75	0.75	0.25	73.9	-5.1	47.7	48.0	96.1	0.75	0.75	0.25	72.4	-1.4	48.0	48.0	91.7	3.9	89	1.0	1.0	0.0	87.8	-10.2	95.4	96.0	96.1
30/380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	120	0.5	0.75	0.25	65.3	-14.8	33.2	36.4	114.0	0.5	0.75	0.25	63.2	-12.6	35.5	37.7	109.6	3.7	119	0.5	1.0	0.0	70.6	-29.7	66.5	72.8	114.0
31/218	G00B_075_050a	0.25	0.75	0.25	0.75	0.5	0.5	150	0.25	0.75	0.25	55.0	-32.5	14.8	35.7	155.5	0.25	0.75	0.25	53.0	-27.9	21.7	35.3	142.0	8.5	149	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5
32/222	G50B_075_050a	0.25	0.75	0.75	0.75	0.5	0.5	210	0.25	0.75	0.75	58.4	-12.7	-20.7	24.3	238.4	0.25	0.75	0.75	55.9	-14.3	-16.3	21.7	228.6	5.3	210	0.0	1.0	1.0	56.8	-25.5	-41.5	48.7	238.4
33/186	B00R_075_050a	0.25	0.25	0.75	0.75	0.5	0.5	270	0.25	0.25	0.75	42.5	14.7	-20.2	25.0	306.2	0.25	0.25	0.75	37.5	18.9	-20.4	27.9	312.8	6.5	270	0.0	0.0	1.0	25.0	29.5	-40.4	50.0	306.2
34/510	B50R_075_050a																																	

delta E* = 5.0

2-0

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

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

delta E* = 4.2

gráfico TUB-PS57; tonos rojo - verde
colores y diferencia en color, ΔE^* , 3D=0, de=0, *cmY0*

entrada: *rgb/cmyk* -> *rgb*_d
salida: transfiera a *cmY0*_d

TUB matrícula: 20130201-PS57/PS57L0NA.TXT /.PS
aplicación para la medida salida en la impresión offset, separación cmY0 (CMY0)
TUB material: code=rh4ta

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.L0NA.TXT> / .PS
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md			
81	R00Y_012_012a	0.125 0.0 0.0	0.125 0.125 0.062	390	0.125 0.0 0.0	27.0 8.8 5.6	10.4 32.3	0.125 0.0 0.0	26.6 14.6 4.2	15.2 16.1 5.9	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	
82	B50R_012_012a	0.125 0.0 0.125	0.125 0.125 0.062	330	0.125 0.0 0.125	27.0 9.9 0.0	9.9 359.8	0.125 0.0 0.125	26.7 15.8 0.3	15.8 1.1 5.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	
83	B25R_025_025a	0.125 0.0 0.25	0.25 0.25 0.125	300	0.125 0.0 0.25	27.1 14.6 -5.1	15.5 340.5	0.125 0.0 0.25	26.9 17.8 -4.5	18.4 345.8 3.2	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5	
84	B15R_037_037a	0.125 0.0 0.375	0.375 0.375 0.187	289	0.118 0.0 0.375	26.8 17.7 -11.0	20.9 328.1	0.125 0.0 0.375	26.6 19.3 -9.3	21.5 334.2 2.3	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1	
85	B11R_050_050a	0.125 0.0 0.5	0.5 0.5 0.25	284	0.116 0.0 0.5	26.5 20.6 -16.5	26.4 321.1	0.125 0.0 0.5	27.0 21.7 -15.4	26.6 324.6 1.7	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1	
86	B09R_062_062a	0.125 0.0 0.625	0.625 0.625 0.312	281	0.114 0.0 0.625	26.8 24.2 -21.7	32.5 318.2	0.125 0.0 0.625	27.1 25.2 -21.3	33.1 319.7 1.0	279	0.183 0.0 1.0	28.3 38.8 -34.7	52.1 318.2	
87	B07R_075_075a	0.125 0.0 0.75	0.75 0.75 0.375	279	0.112 0.0 0.75	27.1 27.9 -26.8	38.7 316.2	0.125 0.0 0.75	27.4 29.1 -26.9	39.7 317.2 1.2	278	0.15 0.0 1.0	28.1 37.2 -35.7	51.6 316.2	
88	B06R_087_087a	0.125 0.0 0.875	0.875 0.875 0.437	278	0.116 0.0 0.875	27.5 31.9 -31.6	44.9 315.2	0.125 0.0 0.875	27.4 33.0 -32.0	46.0 315.8 1.1	277	0.133 0.0 1.0	27.9 36.4 -36.2	51.3 315.2	
89	B05R_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	27.7 35.6 -36.7	51.1 314.1	0.125 0.0 1.0	27.9 36.0 -36.4	51.2 314.7 0.5	276	0.116 0.0 1.0	27.7 35.6 -36.7	51.1 314.1	
90	Y00G_012_012a	0.125 0.125 0.0	0.125 0.125 0.062	90	0.125 0.125 0.0	32.3 -1.2 11.9	12.0 96.1	0.125 0.125 0.0	29.6 5.9 7.7	9.7 52.8 8.6	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0	96.1
91	NW_012a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	33.2 0.0 0.0	0.0 0.0	0.125 0.125 0.125	29.8 7.2 3.6	8.1 26.3 8.7	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	
92	B00R_025_012a	0.125 0.125 0.25	0.25 0.125 0.187	270	0.124 0.124 0.25	33.3 3.6 -5.0	6.2 306.2	0.125 0.125 0.25	30.0 8.9 -1.7	9.1 349.1 7.0	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
93	B00R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.124 0.375	33.4 7.3 -10.1	12.5 306.2	0.125 0.125 0.375	30.4 11.8 -7.5	14.0 327.5 5.9	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
94	B00R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.124 0.5	33.5 11.0 -15.1	18.7 306.2	0.125 0.125 0.5	30.5 14.5 -14.1	20.3 315.8 4.7	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
95	B00R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	33.6 14.7 -20.2	25.0 306.2	0.125 0.125 0.625	30.9 17.9 -20.2	27.0 311.4 4.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
96	B00R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	33.7 18.4 -25.2	31.3 306.2	0.125 0.125 0.75	31.5 21.1 -26.2	33.7 308.7 3.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
97	B00R_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	33.8 22.1 -30.3	37.5 306.2	0.125 0.125 0.875	31.5 25.0 -31.5	40.2 308.4 3.8	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
98	B00R_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	33.9 25.8 -35.3	43.8 306.2	0.125 0.125 1.0	32.0 28.2 -36.3	46.0 307.8 3.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
99	Y50G_025_025a	0.125 0.25 0.0	0.25 0.25 0.125	120	0.125 0.25 0.0	35.9 -7.4 16.6	18.2 114.0	0.125 0.25 0.0	33.7 -4.5 12.9	13.6 109.2 5.2	119	0.5 1.0 0.0	70.6 -29.7	66.5 72.8	114.0
100	G00B_025_012a	0.125 0.25 0.125	0.25 0.125 0.187	150	0.124 0.25 0.124	36.4 -8.1 3.7	8.9 155.5	0.125 0.25 0.125	33.9 -3.6 8.3	9.1 113.6 6.9	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4	155.5
101	G50B_025_012a	0.125 0.25 0.25	0.25 0.125 0.187	210	0.124 0.25 0.25	37.3 -3.1 -5.1	6.0 238.4	0.125 0.25 0.25	34.4 -1.1 1.6	2.0 124.6 7.7	210	0.0 1.0 1.0	56.8 -25.5	-41.5 48.7	238.4
102	G75B_037_025a	0.125 0.25 0.375	0.375 0.25 0.25	240	0.124 0.25 0.375	37.6 -0.3 -10.1	10.1 268.2	0.125 0.25 0.375	34.7 1.3 -4.5	4.7 286.1 6.5	240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6	268.2
103	G84B_050_037a	0.125 0.25 0.5	0.5 0.375 0.312	251	0.124 0.243 0.5	37.3 3.7 -15.1	15.6 283.7	0.125 0.25 0.5	35.0 4.5 -11.8	12.7 291.0 4.1	251	0.0 0.316 1.0	35.2 9.9	-40.4 41.6	283.7
104	G88B_062_050a	0.125 0.25 0.625	0.625 0.5 0.375	256	0.125 0.241 0.625	37.2 7.6 -20.1	21.5 290.8	0.125 0.25 0.625	35.2 8.5 -18.0	20.0 295.3 2.9	257	0.0 0.233 1.0	32.2 15.3	-40.3 43.1	290.8
105	G90B_075_062a	0.125 0.25 0.75	0.75 0.625 0.437	259	0.125 0.239 0.75	37.1 11.6 -25.2	27.8 294.6	0.125 0.25 0.75	35.7 12.5 -24.8	27.8 296.7 1.7	260	0.0 0.183 1.0	30.6 18.5	-40.4 44.5	294.6
106	G92B_087_075a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.237 0.875	37.1 15.5 -30.3	34.0 297.1	0.125 0.25 0.875	36.1 16.4 -30.6	34.8 298.2 1.3	262	0.0 0.15 1.0	29.5 20.7	-40.4 45.4	297.1
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.241 1.0	37.2 19.1 -35.2	40.1 298.4	0.125 0.25 1.0	36.4 19.7 -35.8	40.8 298.8 1.1	262	0.0 0.133 1.0	28.9 21.8	-40.3 45.8	298.4
108	Y68G_037_037a	0.125 0.375 0.0	0.375 0.375 0.187	131	0.118 0.375 0.0	38.6 -15.5 19.9	25.3 127.8	0.125 0.375 0.0	37.4 -15.0 17.0	22.7 131.3 3.1	131	0.316 1.0 0.0	62.3 -41.4	53.2 67.5	127.8
109	G00B_037_025a	0.125 0.375 0.125	0.375 0.25 0.25	150	0.124 0.375 0.124	39.7 -16.2 7.4	17.8 155.5	0.125 0.375 0.125	37.6 -12.8 11.7	17.3 137.3 5.9	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4	155.5
110	G25B_037_025a	0.125 0.375 0.25	0.375 0.25 0.25	180	0.124 0.375 0.25	40.4 -12.1 -2.0	12.3 189.3	0.125 0.375 0.25	38.4 -10.8 5.2	12.0 154.3 7.6	180	0.0 1.0 0.5	52.9 -48.6	-8.0 49.3	189.3
111	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	41.3 -6.3 -10.3	12.1 238.4	0.125 0.375 0.375	38.8 -7.8 -2.3	8.2 196.2 8.6	210	0.0 1.0 1.0	56.8 -25.5	-41.5 48.7	238.4
112	G65B_050_037a	0.125 0.375 0.5	0.5 0.375 0.312	229	0.124 0.381 0.5	42.2 -4.6 -15.4	16.0 253.3	0.125 0.375 0.5	39.7 -5.2 -9.5	10.8 241.1 6.4	228	0.0 0.683 1.0	48.3 -12.2	-41.1 42.9	253.3
113	G75B_062_050a	0.125 0.375 0.625	0.625 0.5 0.375	240	0.125 0.375 0.625	41.9 -0.6 -20.3	20.3 268.2	0.125 0.375 0.625	39.7 -0.9 -16.6	16.6 266.8 4.2	240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6	268.2
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.364 0.75	41.5 3.5 -25.1	25.4 277.9	0.125 0.375 0.75	39.8 4.0 -24.0	24.4 279.5 2.0	247	0.0 0.383 1.0	37.6 5.6	-40.3 40.7	277.9
115	G84B_087_075a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.362 0.875	41.4 7.4 -30.3	31.2 283.7	0.125 0.375 0.875	40.3 8.1 -30.2	31.3 285.1 1.3	251	0.0 0.316 1.0	35.2 9.9	-40.4 41.6	283.7
116	G86B_100_087a	0.125 0.375 1.0	1.0 0.875 0.562	254	0.125 0.358 1.0	41.2 11.5 -35.2	37.1 288.1	0.125 0.375 1.0	40.4 12.6 -35.8	37.9 289.4 1.4	255	0.0 0.266 1.0	33.4 13.2	-40.3 42.4	288.1
117	Y76G_050_050a	0.125 0.5 0.0	0.5 0.5 0.25	136	0.116 0.5 0.0	41.1 -24.1 22.9	33.2 136.5	0.125 0.5 0.0	41.0 -23.7 21.5	32.0 137.7 1.4	137	0.233 1.0 0.0	57.9 -48.3	45.8 66.5	136.5
118	G00B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.124	42.9 -24.3 11.1	26.7 155.5	0.125 0.5 0.125	41.5 -21.6 15.4	26.6 144.4 5.3	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4	155.5
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.243	43.5 -21.3 2.7	21.4 172.5	0.125 0.5 0.25	42.1 -19.2 8.0	20.8 157.3 5.8	168	0.0 1.0 0.316	51.6 -56.8	7.4 57.3	172.5
120	G34B_050_037a	0.125 0.5 0.375	0.5 0.375 0.312	191	0.124 0.5 0.381	44.									

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

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
162	R00Y_025_025a	0.25 0.0 0.0	0.25 0.25 0.125	390	0.25 0.0 0.0	29.6 17.7 11.2	20.9 32.3	0.25 0.0 0.0	28.1 24.0 7.8	25.2 18.0 7.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
163	R00Y_025_025a	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.125	29.7 18.5 5.2	19.2 15.9	0.25 0.0 0.125	28.1 25.5 4.4	25.9 9.7 7.2	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
164	B50R_025_025a	0.25 0.0 0.25	0.25 0.25 0.125	330	0.25 0.0 0.25	29.8 19.8 0.0	19.8 359.8	0.25 0.0 0.25	28.3 27.3 -0.1	27.3 359.7 7.6	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
165	B34R_037_037a	0.25 0.0 0.375	0.375 0.375 0.187	311	0.256 0.0 0.375	30.1 25.5 -4.4	25.9 350.0	0.25 0.0 0.375	28.5 29.3 -4.3	29.6 351.6 4.1	311	1.0 0.683 0.0	39.8 68.1 -11.9	69.1 350.0
166	B25R_050_050a	0.25 0.0 0.5	0.5 0.5 0.25	300	0.25 0.0 0.5	29.9 29.3 -10.3	31.0 340.5	0.25 0.0 0.5	28.5 30.6 -10.4	32.3 341.1 1.9	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
167	B19R_062_062a	0.25 0.0 0.625	0.625 0.625 0.312	293	0.239 0.0 0.625	29.7 32.7 -16.0	36.4 333.8	0.25 0.0 0.625	28.5 32.6 -17.0	36.8 332.3 1.5	292	0.383 0.0 1.0	32.9 52.3 -25.7	58.3 333.8
168	B15R_075_075a	0.25 0.0 0.75	0.75 0.75 0.375	289	0.237 0.0 0.75	29.3 35.5 -22.0	41.8 328.1	0.25 0.0 0.75	28.7 36.0 -23.1	42.8 327.3 1.3	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
169	B13R_087_087a	0.25 0.0 0.875	0.875 0.875 0.437	286	0.233 0.0 0.875	28.7 37.9 -27.8	47.0 323.6	0.25 0.0 0.875	28.6 39.2 -28.1	48.2 324.3 1.3	284	0.266 0.0 1.0	29.4 43.3 -31.8	53.8 323.6
170	B11R_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1	0.25 0.0 1.0	28.8 41.9 -32.5	53.1 322.1 0.9	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
171	R50Y_025_025a	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.125 0.0	34.5 7.2 17.1	18.6 67.1	0.25 0.125 0.0	31.0 16.0 11.8	19.9 36.4 10.8	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
172	R00Y_025_012a	0.25 0.125 0.125	0.25 0.125 0.187	390	0.25 0.124 0.124	35.9 8.8 5.6	10.4 32.3	0.25 0.125 0.125	31.0 16.8 8.0	18.6 25.3 9.7	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
173	B50R_025_012a	0.25 0.125 0.25	0.25 0.125 0.187	330	0.25 0.124 0.25	36.0 9.9 0.0	9.9 359.8	0.25 0.125 0.25	31.5 18.6 3.2	18.9 9.7 10.3	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
174	B25R_037_025a	0.25 0.125 0.375	0.375 0.25 0.25	300	0.25 0.124 0.375	36.0 14.6 -5.1	15.5 340.5	0.25 0.125 0.375	31.7 20.5 -2.2	20.6 353.7 7.9	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
175	B15R_050_037a	0.25 0.125 0.5	0.5 0.375 0.312	289	0.243 0.124 0.5	35.7 17.7 -11.0	20.9 328.1	0.25 0.125 0.5	31.9 22.3 -9.9	24.4 359.6 6.0	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
176	B11R_062_050a	0.25 0.125 0.625	0.625 0.5 0.375	284	0.241 0.125 0.625	35.4 20.6 -16.5	26.4 321.1	0.25 0.125 0.625	32.1 24.7 -16.1	29.5 326.9 5.3	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
177	B09R_075_062a	0.25 0.125 0.75	0.75 0.625 0.437	281	0.239 0.125 0.75	35.7 24.2 -21.7	32.5 318.2	0.25 0.125 0.75	32.4 28.2 -22.2	35.9 321.8 5.1	279	0.183 0.0 1.0	28.3 38.8 -34.7	52.1 318.2
178	B07R_087_075a	0.25 0.125 0.875	0.875 0.75 0.5	279	0.237 0.125 0.875	36.0 27.9 -26.8	38.7 316.2	0.25 0.125 0.875	32.8 30.9 -27.8	41.6 318.0 4.5	278	0.15 0.0 1.0	28.1 37.2 -35.7	51.6 316.2
179	B06R_100_087a	0.25 0.125 1.0	1.0 0.875 0.562	278	0.241 0.125 1.0	36.4 31.9 -31.6	44.9 315.2	0.25 0.125 1.0	33.1 33.9 -32.5	47.0 316.2 3.9	277	0.133 0.0 1.0	27.9 36.4 -36.2	51.3 315.2
180	Y00G_025_025a	0.25 0.25 0.0	0.25 0.25 0.125	90	0.25 0.25 0.0	40.2 -2.5 23.8	24.0 96.1	0.25 0.25 0.0	35.1 5.0 17.5	18.2 74.0 11.1	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
181	Y00G_025_012a	0.25 0.25 0.125	0.25 0.125 0.187	90	0.25 0.25 0.124	41.2 -1.2 11.9	12.0 96.1	0.25 0.25 0.125	35.4 5.7 13.1	14.3 66.3 9.1	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	42.1 0.0 0.0	0.0 0.0	0.25 0.25 0.25	35.7 7.5 7.1	10.4 43.4 12.2	360	1.0 1.0 1.0	95.6 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	42.2 3.6 -5.0	6.2 306.2	0.25 0.25 0.375	36.3 9.3 0.7	9.4 4.5 10.0	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
184	B00R_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.249 0.5	42.3 7.3 -10.1	12.5 306.2	0.25 0.25 0.5	36.4 12.3 -6.8	14.1 330.9 8.3	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
185	B00R_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	42.4 11.0 -15.1	18.7 306.2	0.25 0.25 0.625	37.0 15.5 -14.0	20.9 318.0 7.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	42.5 14.7 -20.2	25.0 306.2	0.25 0.25 0.75	37.5 18.9 -20.4	27.9 312.8 6.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
187	B00R_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.6 18.4 -25.2	31.3 306.2	0.25 0.25 0.875	38.0 22.0 -26.5	34.4 309.7 5.9	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
188	B00R_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	42.7 22.1 -30.3	37.5 306.2	0.25 0.25 1.0	38.2 25.3 -31.6	40.5 308.7 5.6	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
189	Y31G_037_037a	0.25 0.375 0.0	0.375 0.375 0.187	109	0.256 0.375 0.0	44.4 -7.9 29.8	30.8 104.9	0.25 0.375 0.0	39.4 -4.2 23.2	23.6 100.4 9.0	108	0.683 1.0 0.0	77.8 -21.1	79.4 82.2 104.9
190	Y50G_037_025a	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	44.8 -7.4 16.6	18.2 114.0	0.25 0.375 0.125	39.4 -3.5 17.7	18.1 101.1 6.7	119	0.5 1.0 0.0	70.6 -29.7	66.5 72.8 114.0
191	G00B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	45.4 -8.1 3.7	8.9 155.5	0.25 0.375 0.25	40.0 -1.6 10.9	11.0 98.5 11.0	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
192	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	46.2 -3.1 -5.1	6.0 238.4	0.25 0.375 0.375	40.5 0.2 4.0	4.0 87.0 11.3	210	0.0 1.0 1.0	56.8 -25.5	-41.5 48.7 238.4
193	G75B_050_025a	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.375 0.5	46.5 -0.3 -10.1	10.1 268.2	0.25 0.375 0.5	40.8 3.5 -4.5	5.8 307.8 8.8	240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6 268.2
194	G84B_062_037a	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.368 0.625	46.2 3.7 -15.1	15.6 283.7	0.25 0.375 0.625	40.7 7.2 -11.9	13.9 301.0 7.2	251	0.0 0.316 1.0	35.2 9.9 -40.4	41.6 283.7
195	G88B_075_050a	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.366 0.75	46.1 7.6 -20.1	21.5 290.8	0.25 0.375 0.75	41.7 10.7 -19.4	22.2 299.0 5.4	257	0.0 0.233 1.0	32.2 15.3 -40.3	43.1 290.8
196	G90B_087_062a	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.364 0.875	46.0 11.6 -25.2	27.8 294.6	0.25 0.375 0.875	41.7 14.8 -26.1	30.1 299.6 5.5	260	0.0 0.183 1.0	30.6 18.5 -40.4	44.5 294.6
197	G92B_100_075a	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.362 1.0	46.0 15.5 -30.3	34.0 297.1	0.25 0.375 1.0	42.1 18.3 -31.8	36.7 299.9 5.0	262	0.0 0.15 1.0	29.5 20.7 -40.4	45.4 297.1
198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	47.4 -14.8 33.2	36.4 114.0	0.25 0.5 0.0	43.1 -14.1 28.4	31.7 116.4 6.5	119	0.5 1.0 0.0	70.6 -29.7	66.5 72.8 114.0
199	Y68G_050_037a	0.25 0.5 0.125	0.5 0.375 0.312	131	0.243 0.5 0.124	47.5 -15.5 19.9	25.3 127.8	0.25 0.5 0.125	43.2 -13.5 21.7	25.6 121.9 4.9	131	0.316 1.0 0.0	62.3 -41.4	53.2 67.5 127.8
200	G00B_050_025a	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.249	48.6 -16.2 7.4	17.8 155.5	0.25 0.5 0.25	44.2 -10.9 14.5	18.2 126.7 9.9	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
201	G25B_050_025a	0.25 0.5 0.375	0.5 0.25 0.375	180	0.249 0.5 0.375	49.3 -12.1 -2.0	12.3 189.3	0.25 0.5 0.375	44.9 -8.5 6.8	10.9 141.0 10.5	180	0.0 1.0 0.5	52.9 -48.6	-8.0 49.3 189.3
202	G50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	50.2 -6.3 -10.3	12.1 238.4	0.25 0.5 0.5	45.1 -5.3 -2.5	5.9 205.3 9.3	210	0.0 1.		

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

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md		
243	R00Y_037_037a	0.375 0.0 0.0	0.375 0.375 0.187	390	0.375 0.0 0.0	32.2 26.6 16.8	31.4 32.3	0.375 0.0 0.0	31.7 36.2 17.7	40.3 26.1 9.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
244	R18Y_037_037a	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.118	32.3 27.2 11.7	29.6 23.2	0.375 0.0 0.125	31.6 36.7 13.2	39.0 19.8 9.6	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
245	B65R_037_037a	0.375 0.0 0.25	0.375 0.375 0.187	349	0.375 0.0 0.256	32.4 28.6 4.4	29.0 8.9	0.375 0.0 0.25	31.7 38.5 8.1	39.3 11.9 10.5	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
246	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	32.5 29.7 0.0	29.7 359.8	0.375 0.0 0.375	31.7 39.8 3.0	39.9 4.3 10.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
247	B38R_050_050a	0.375 0.0 0.5	0.5 0.5 0.25	316	0.383 0.0 0.5	33.2 35.8 -4.3	36.0 353.0	0.375 0.0 0.5	32.2 42.9 -3.3	43.0 355.5 7.3	317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0
248	B30R_062_062a	0.375 0.0 0.625	0.625 0.625 0.312	307	0.385 0.0 0.625	32.8 40.6 -9.0	41.6 347.4	0.375 0.0 0.625	32.4 45.1 -9.5	46.1 348.0 4.5	307	0.616 0.0 1.0	37.9 65.0 -14.5	66.6 347.4
249	B25R_075_075a	0.375 0.0 0.75	0.75 0.75 0.375	300	0.375 0.0 0.75	32.7 43.9 -15.5	46.6 340.5	0.375 0.0 0.75	32.5 47.1 -15.8	49.6 341.4 3.1	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
250	B20R_087_087a	0.375 0.0 0.875	0.875 0.875 0.437	295	0.364 0.0 0.875	32.5 47.4 -21.3	51.9 335.7	0.375 0.0 0.875	32.6 49.3 -21.4	53.8 336.5 1.9	294	0.416 0.0 1.0	33.7 54.1 -24.4	59.4 335.7
251	B18R_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	32.5 51.2 -26.5	57.7 332.6	0.375 0.0 1.0	32.7 51.8 -26.0	58.0 333.3 0.8	291	0.366 0.0 1.0	32.5 51.2 -26.5	57.7 332.6
252	R31Y_037_037a	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.118 0.0	36.4 17.1 22.2	28.1 52.2	0.375 0.125 0.0	34.8 28.0 21.3	35.2 37.3 10.9	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
253	R00Y_037_025a	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.124	38.5 17.7 11.2	20.9 32.3	0.375 0.125 0.125	35.1 28.3 16.7	32.9 30.6 12.4	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
254	R00Y_037_025a	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.25	38.6 18.5 5.2	19.2 15.9	0.375 0.125 0.25	35.3 29.6 10.7	31.5 19.8 12.7	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
255	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.124 0.375	38.7 19.8 0.0	19.8 359.8	0.375 0.125 0.375	35.5 31.2 5.0	31.6 9.2 12.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
256	B34R_050_037a	0.375 0.125 0.5	0.5 0.375 0.312	311	0.381 0.124 0.5	39.0 25.5 -4.4	25.9 350.0	0.375 0.125 0.5	36.2 33.7 -2.3	33.7 355.9 8.9	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
257	B25R_062_050a	0.375 0.125 0.625	0.625 0.5 0.375	300	0.375 0.125 0.625	38.8 29.3 -10.3	31.0 340.5	0.375 0.125 0.625	36.2 35.2 -9.0	36.3 345.6 6.6	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
258	B19R_075_062a	0.375 0.125 0.75	0.75 0.625 0.437	293	0.364 0.125 0.75	38.6 32.7 -16.0	36.4 333.8	0.375 0.125 0.75	36.6 37.1 -15.7	40.3 337.0 4.8	292	0.383 0.0 1.0	32.9 52.3 -25.7	58.3 333.8
259	B15R_087_075a	0.375 0.125 0.875	0.875 0.75 0.5	289	0.362 0.125 0.875	38.2 35.5 -22.0	41.8 328.1	0.375 0.125 0.875	36.9 39.8 -21.4	45.2 331.6 4.5	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
260	B13R_100_087a	0.375 0.125 1.0	1.0 0.875 0.562	286	0.358 0.125 1.0	37.6 37.9 -27.8	47.0 323.6	0.375 0.125 1.0	36.8 42.2 -26.6	49.9 327.7 4.5	284	0.266 0.0 1.0	29.4 43.3 -31.8	53.8 323.6
261	R68Y_037_037a	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.256 0.0	43.2 4.1 30.1	30.4 82.1	0.375 0.25 0.0	39.9 16.0 27.6	31.9 59.7 12.6	71	1.0 0.683 0.0	74.8 11.0 80.4	81.1 82.1
262	R50Y_037_025a	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.25 0.124	43.4 7.2 17.1	18.6 67.1	0.375 0.25 0.125	39.9 17.1 21.7	27.7 51.6 11.5	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
263	R00Y_037_012a	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.249	44.8 8.8 5.6	10.4 32.3	0.375 0.25 0.25	40.0 18.4 15.1	23.9 39.3 14.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
264	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.249 0.375	44.9 9.9 0.0	9.9 359.8	0.375 0.25 0.375	40.7 19.7 8.1	21.3 22.2 13.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
265	B25R_050_025a	0.375 0.25 0.5	0.5 0.25 0.375	300	0.375 0.249 0.5	44.9 14.6 -5.1	15.5 340.5	0.375 0.25 0.5	41.2 22.1 -0.1	22.1 359.5 9.7	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
266	B15R_062_037a	0.375 0.25 0.625	0.625 0.375 0.437	289	0.368 0.25 0.625	44.6 17.7 -11.0	20.9 328.1	0.375 0.25 0.625	41.6 23.9 -7.1	25.0 343.2 7.9	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
267	B11R_075_050a	0.375 0.25 0.75	0.75 0.5 0.5	284	0.366 0.25 0.75	44.3 20.6 -16.5	26.4 321.1	0.375 0.25 0.75	42.1 26.2 -14.0	29.7 331.7 6.5	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
268	B09R_087_062a	0.375 0.25 0.875	0.875 0.625 0.562	283	0.364 0.25 0.875	44.6 24.2 -21.7	32.5 318.2	0.375 0.25 0.875	42.9 28.9 -20.3	35.3 324.8 5.1	279	0.183 0.0 1.0	28.3 38.8 -34.7	52.1 318.2
269	B07R_100_075a	0.375 0.25 1.0	1.0 0.75 0.625	279	0.362 0.25 1.0	44.9 27.9 -26.8	38.7 316.2	0.375 0.25 1.0	43.1 31.3 -26.0	40.7 320.3 3.9	278	0.15 0.0 1.0	28.1 37.2 -35.7	51.6 316.2
270	Y00G_037_037a	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.375 0.0	48.1 -3.8 35.8	36.0 96.1	0.375 0.375 0.0	44.1 6.7 33.2	33.8 78.5 11.5	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
271	Y00G_037_025a	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.375 0.124	49.1 -2.5 23.8	24.0 96.1	0.375 0.375 0.125	44.5 7.0 26.3	27.2 75.0 10.9	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
272	Y00G_037_012a	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.375 0.249	50.1 -1.2 11.9	12.0 96.1	0.375 0.375 0.25	44.7 8.5 18.5	20.4 65.3 12.9	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
273	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	51.0 0.0 0.0	0.0 0.0	0.375 0.375 0.375	45.3 10.0 11.0	14.9 47.8 16.0	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0
274	B00R_050_012a	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.375 0.5	51.1 3.6 -5.0	6.2 306.2	0.375 0.375 0.5	46.1 12.2 2.1	12.3 10.0 12.2	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
275	B00R_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	51.2 7.3 -10.1	12.5 306.2	0.375 0.375 0.625	46.7 14.8 -5.3	15.7 340.2 9.9	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
276	B00R_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	51.3 11.0 -15.1	18.7 306.2	0.375 0.375 0.75	47.4 17.2 -12.5	21.3 323.8 7.7	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
277	B00R_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	51.4 14.7 -20.2	25.0 306.2	0.375 0.375 0.875	48.1 19.9 -19.3	27.7 315.9 6.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
278	B00R_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	51.5 18.4 -25.2	31.3 306.2	0.375 0.375 1.0	48.4 23.0 -25.3	34.2 312.3 5.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
279	Y23G_050_050a	0.375 0.5 0.0	0.5 0.5 0.25	104	0.383 0.5 0.0	52.8 -8.5 42.1	43.0 101.4	0.375 0.5 0.0	49.1 -2.0 38.9	38.9 92.9 8.1	102	0.766 1.0 0.0	81.2 -17.0 84.3	86.0 101.4
280	Y31G_050_037a	0.375 0.5 0.125	0.5 0.375 0.312	109	0.381 0.5 0.124	53.3 -7.9 29.8	30.8 104.9	0.375 0.5 0.125	49.5 -1.7 31.0	31.0 93.2 7.3	108	0.683 1.0 0.0	77.8 -21.1 79.4	82.2 104.9
281	Y50G_050_025a	0.375 0.5 0.25	0.5 0.25 0.375	120	0.375 0.5 0.249	53.7 -7.4 16.6	18.2 114.0	0.375 0.5 0.25	49.7 -0.9 22.3	22.3 95.5 9.4	119	0.5 1.0 0.0	70.6 -2	

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md				
324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.25	390	0.5	0.0	0.0	45.4	70.9	44.8	83.9	32.3
325	R26Y_050_050a	0.5	0.0	0.125	0.5	0.5	0.25	376	0.5	0.0	0.125	34.7	45.7	18.0	49.1	26.1
326	R00Y_050_050a	0.5	0.0	0.25	0.5	0.5	0.25	360	0.5	0.0	0.25	34.8	46.7	12.4	48.3	14.9
327	B61R_050_050a	0.5	0.0	0.375	0.5	0.5	0.25	344	0.5	0.0	0.375	34.8	48.4	6.7	48.9	7.8
328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.25	330	0.5	0.0	0.5	35.0	49.8	0.6	49.8	0.7
329	B40R_062_062a	0.5	0.0	0.625	0.625	0.625	0.312	319	0.51	0.0	0.625	35.3	52.5	-4.7	52.7	354.8
330	B34R_075_075a	0.5	0.0	0.75	0.75	0.75	0.375	311	0.512	0.0	0.75	35.7	54.4	-10.3	55.4	349.2
331	B29R_087_087a	0.5	0.0	0.875	0.875	0.875	0.437	305	0.51	0.0	0.875	35.8	56.7	-15.7	58.8	344.4
332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	300	0.5	0.0	1.0	35.6	58.6	-20.7	62.1	340.5
333	R23Y_050_050a	0.5	0.125	0.0	0.5	0.5	0.25	44	0.5	0.116	0.0	38.7	26.7	27.4	38.2	45.7
334	R00Y_050_037a	0.5	0.125	0.125	0.5	0.375	0.312	390	0.5	0.124	0.124	41.1	26.6	16.8	31.4	32.3
335	R18Y_050_037a	0.5	0.125	0.25	0.5	0.375	0.312	371	0.5	0.124	0.243	41.2	27.2	11.7	29.6	23.2
336	B65R_050_037a	0.5	0.125	0.375	0.5	0.375	0.312	349	0.5	0.124	0.381	41.3	28.6	4.4	29.0	8.9
337	B50R_050_037a	0.5	0.125	0.5	0.5	0.375	0.312	330	0.5	0.124	0.5	41.4	29.7	0.0	29.7	359.8
338	B38R_062_050a	0.5	0.125	0.625	0.625	0.5	0.375	316	0.508	0.125	0.625	42.1	35.8	-4.3	36.0	353.0
339	B30R_075_062a	0.5	0.125	0.75	0.75	0.625	0.437	307	0.51	0.125	0.75	41.7	40.6	-9.0	41.6	347.4
340	B25R_087_050a	0.5	0.125	0.875	0.875	0.75	0.5	300	0.5	0.125	0.875	41.7	43.9	-15.5	46.6	340.5
341	B20R_100_087a	0.5	0.125	1.0	1.0	0.875	0.562	295	0.489	0.125	1.0	41.4	47.4	-21.3	51.9	335.7
342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	60	0.5	0.25	0.0	44.6	14.4	34.3	37.2	67.1
343	R31Y_050_037a	0.5	0.25	0.125	0.5	0.375	0.312	49	0.5	0.243	0.124	45.3	17.1	22.2	28.1	52.2
344	R00Y_050_025a	0.5	0.25	0.25	0.5	0.25	0.375	390	0.5	0.249	0.249	47.4	17.7	11.2	20.9	32.3
345	R00Y_050_025a	0.5	0.25	0.375	0.5	0.25	0.375	360	0.5	0.249	0.375	47.5	18.5	5.2	19.2	15.9
346	B50R_050_025a	0.5	0.25	0.5	0.5	0.25	0.375	330	0.5	0.249	0.5	47.6	19.8	0.0	19.8	359.8
347	B34R_062_037a	0.5	0.25	0.625	0.625	0.375	0.437	311	0.506	0.25	0.625	47.9	25.5	-4.4	25.9	350.0
348	B25R_075_050a	0.5	0.25	0.75	0.75	0.5	0.5	300	0.5	0.25	0.75	47.9	29.3	-10.3	31.0	340.5
349	B19R_087_062a	0.5	0.25	0.875	0.875	0.625	0.562	293	0.489	0.25	0.875	47.5	32.7	-16.0	36.4	333.8
350	B15R_100_075a	0.5	0.25	1.0	1.0	0.75	0.625	289	0.487	0.25	1.0	47.1	35.5	-22.0	41.8	328.1
351	R76Y_050_050a	0.5	0.375	0.0	0.5	0.5	0.25	76	0.5	0.383	0.0	51.5	2.1	42.3	42.4	87.0
352	R68Y_050_037a	0.5	0.375	0.125	0.5	0.375	0.312	71	0.5	0.381	0.124	52.2	4.1	30.1	30.4	82.1
353	R50Y_050_025a	0.5	0.375	0.25	0.5	0.25	0.375	60	0.5	0.375	0.249	52.3	7.2	17.1	18.6	67.1
354	R00Y_050_012a	0.5	0.375	0.375	0.5	0.125	0.437	390	0.5	0.375	0.375	53.7	8.8	5.6	10.4	32.3
355	B50R_050_012a	0.5	0.375	0.5	0.5	0.125	0.437	330	0.5	0.375	0.5	53.8	9.9	0.0	9.9	359.8
356	B25R_062_025a	0.5	0.375	0.625	0.625	0.25	0.5	300	0.5	0.375	0.625	53.9	14.6	-5.1	15.5	340.5
357	B15R_075_037a	0.5	0.375	0.75	0.75	0.375	0.562	289	0.493	0.375	0.75	53.5	17.7	-11.0	20.9	328.1
358	B11R_087_050a	0.5	0.375	0.875	0.875	0.5	0.625	284	0.491	0.375	0.875	53.2	20.6	-16.5	26.4	321.1
359	B09R_100_062a	0.5	0.375	1.0	1.0	0.625	0.687	281	0.489	0.375	1.0	53.5	24.2	-21.7	32.5	318.2
360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	90	0.5	0.5	0.0	56.1	-5.1	47.7	48.0	96.1
361	Y00G_050_037a	0.5	0.5	0.125	0.5	0.375	0.312	90	0.5	0.5	0.124	57.0	-3.8	35.8	36.0	96.1
362	Y00G_050_025a	0.5	0.5	0.25	0.5	0.25	0.375	90	0.5	0.5	0.249	58.0	-2.5	23.8	24.0	96.1
363	Y00G_050_012a	0.5	0.5	0.375	0.5	0.125	0.437	90	0.5	0.5	0.375	59.0	-1.2	11.9	12.0	96.1
364	NW_050a	0.5	0.5	0.5	0.5	0.0	0.5	360	0.5	0.5	0.5	60.0	0.0	0.0	0.0	0.0
365	B00R_062_012a	0.5	0.5	0.625	0.625	0.125	0.562	270	0.5	0.5	0.625	60.0	3.6	-5.0	6.2	306.2
366	B00R_075_025a	0.5	0.5	0.75	0.75	0.25	0.625	270	0.5	0.5	0.75	60.1	7.3	-10.1	12.5	306.2
367	B00R_087_037a	0.5	0.5	0.875	0.875	0.375	0.687	270	0.5	0.5	0.875	60.2	11.0	-15.1	18.7	306.2
368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	270	0.5	0.5	1.0	60.3	14.7	-20.2	25.0	306.2
369	Y18G_062_062a	0.5	0.625	0.0	0.625	0.625	0.312	101	0.51	0.625	0.0	60.8	-9.7	54.1	55.0	100.2
370	Y23G_062_050a	0.5	0.625	0.125	0.625	0.5	0.375	104	0.508	0.625	0.125	61.7	-8.5	42.1	43.0	101.4
371	Y31G_062_037a	0.5	0.625	0.25	0.625	0.375	0.437	109	0.506	0.625	0.25	62.2	-7.9	29.8	30.8	104.9
372	Y50G_062_025a	0.5	0.625	0.375	0.625	0.25	0.5	120	0.5	0.625	0.375	62.6	-7.4	16.6	18.2	114.0
373	G00B_062_012a	0.5	0.625	0.5	0.625	0.125	0.562	150	0.5	0.625	0.5	63.2	-8.1	3.7	8.9	155.5
374	G50B_062_012a	0.5	0.625	0.625	0.625	0.125	0.562	210	0.5	0.625	0.625	64.0	-3.1	-5.1	6.0	238.4
375	G75B_075_025a	0.5	0.625	0.75	0.75	0.25	0.625	240	0.5	0.625	0.75	64.3	-0.3	-10.1	10.1	268.2
376	G84B_087_037a	0.5	0.625	0.875	0.875	0.375	0.687	251	0.5	0.618	0.875	64.0	3.7	-15.1	15.6	283.7
377	G88B_100_050a	0.5	0.625	1.0	1.0	0.5	0.75	256	0.5	0.616	1.0	63.9	7.6	-20.1	21.5	290.8
378	Y31G_075_075a	0.5	0.75	0.0	0.75	0.75	0.375	109	0.512	0.75	0.0	64.4	-15.8	59.6	61.6	104.9
379	Y38G_075_062a	0.5	0.75	0.125	0.75	0.625	0.437	113	0.51	0.75	0.125	64.9	-15.3	46.9	49.4	108.0
380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	120	0.5	0.75	0.25	65.3	-14.8	33.2	36.4	114.0
381	Y68G_075_037a	0.5	0.75	0.375	0.75	0.375	0.562	131	0.493	0.75	0.375	65.3	-15.5	19.9	25.3	127.8
382	G00B_075_025a	0.5	0.75	0.5	0.75	0.25	0.625	150	0.5	0.75	0.5	66.4	-16.2	7.4	17.8	155.5
383	G25B_075_025a	0.5	0.75	0.625	0.75	0.25	0.625	180	0.5	0.75	0.625	67.1	-12.1	2.0	12.3	189.3
384	G50B_075_025a	0.5	0.75	0.75	0.75	0.25	0.625	210	0.5	0.75	0.75	68.1	-6.3	-10.3	12.1	238.4
385	G65B_087_037a	0.5	0.75	0.875	0.875											

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md		
405	R00Y_062_062a	0.625 0.0 0.0	0.625 0.625 0.312	390	0.625 0.0 0.0	37.5 44.3 28.0	52.4 32.3	0.625 0.0 0.0	37.2 53.3 28.6	60.5 28.2 9.0	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
406	R31Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.114	37.6 44.9 23.4	50.6 27.5	0.625 0.0 0.125	37.4 54.0 24.4	59.3 24.3 9.2	380	1.0 0.0 0.183	45.5 71.8 37.5	81.0 27.5
407	R11Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.239	37.7 45.6 17.4	48.8 20.8	0.625 0.0 0.25	37.3 54.8 19.5	58.2 19.6 9.4	367	1.0 0.0 0.383	45.8 73.0 27.8	78.2 20.8
408	B69R_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.385	37.8 47.2 9.5	48.1 11.4	0.625 0.0 0.375	37.4 56.1 13.0	57.6 13.0 9.5	352	1.0 0.0 0.616	46.0 75.5 15.2	77.1 11.4
409	B59R_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.625 0.0 0.51	37.8 48.6 3.9	48.7 4.6	0.625 0.0 0.5	37.4 57.9 6.5	58.2 6.4 9.6	339	1.0 0.0 0.816	45.9 77.7 6.2	78.0 4.6
410	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	37.9 49.5 -0.1	49.5 359.8	0.625 0.0 0.625	37.4 59.3 1.1	59.3 1.0 9.8	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
411	B42R_075_075a	0.625 0.0 0.75	0.75 0.75 0.375	321	0.637 0.0 0.75	38.9 55.7 -4.4	55.9 355.4	0.625 0.0 0.75	37.9 61.6 -4.2	61.8 356.0 5.9	322	0.85 0.0 1.0	43.7 74.3 -5.9	74.6 355.4
412	B36R_087_087a	0.625 0.0 0.875	0.875 0.875 0.437	314	0.641 0.0 0.875	39.2 61.5 -8.7	62.1 351.9	0.625 0.0 0.875	38.3 64.0 -9.1	64.6 351.8 2.6	315	0.733 0.0 1.0	41.3 70.3 -9.9	71.0 351.9
413	B31R_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	38.3 65.8 -13.7	67.2 348.2	0.625 0.0 1.0	38.1 65.4 -14.0	66.9 347.9 0.5	308	0.633 0.0 1.0	38.3 65.8 -13.7	67.2 348.2
414	R18Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	41	0.625 0.114 0.0	41.1 36.1 32.8	48.8 42.2	0.625 0.125 0.0	40.5 45.1 32.7	55.7 35.9 9.0	39	1.0 0.183 0.0	51.1 57.8 52.5	78.1 42.2
415	R00Y_062_050a	0.625 0.125 0.125	0.625 0.5 0.375	390	0.625 0.125 0.125	43.8 35.4 22.4	41.9 32.3	0.625 0.125 0.125	41.0 44.9 28.0	53.0 31.9 11.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
416	R26Y_062_050a	0.625 0.125 0.25	0.625 0.5 0.375	376	0.625 0.125 0.241	43.9 36.0 17.6	40.1 32.6	0.625 0.125 0.25	41.0 45.8 22.3	51.0 25.9 11.2	377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1
417	R00Y_062_050a	0.625 0.125 0.375	0.625 0.5 0.375	360	0.625 0.125 0.375	44.0 37.1 10.5	38.5 15.9	0.625 0.125 0.375	41.1 47.2 15.5	49.7 18.2 11.6	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
418	B61R_062_050a	0.625 0.125 0.5	0.625 0.5 0.375	344	0.625 0.125 0.508	44.0 38.6 4.0	38.8 5.9	0.625 0.125 0.5	41.4 48.6 7.7	49.3 9.0 11.0	342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
419	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	44.1 39.6 -0.1	39.6 359.8	0.625 0.125 0.625	41.7 50.4 1.6	50.4 1.8 11.1	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
420	B40R_075_062a	0.625 0.125 0.75	0.75 0.625 0.437	319	0.635 0.125 0.75	44.9 45.8 -4.4	46.0 354.4	0.625 0.125 0.75	42.7 52.1 -4.3	52.3 355.2 6.7	320	0.816 0.0 1.0	43.1 73.2 -7.0	73.6 354.4
421	B34R_087_075a	0.625 0.125 0.875	0.875 0.75 0.5	311	0.637 0.125 0.875	44.8 51.0 -8.9	51.8 350.0	0.625 0.125 0.875	42.7 54.6 -10.3	55.5 349.2 4.3	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
422	B29R_100_087a	0.625 0.125 1.0	1.0 0.875 0.562	305	0.635 0.125 1.0	44.5 55.3 -14.3	57.1 345.4	0.625 0.125 1.0	43.0 56.2 -15.1	58.2 344.9 1.9	305	0.583 0.0 1.0	37.2 63.2 -16.4	65.3 345.4
423	R38Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	53	0.625 0.239 0.0	46.3 24.7 39.1	46.2 57.6	0.625 0.25 0.0	45.1 34.1 38.7	51.6 48.5 9.5	52	1.0 0.383 0.0	59.5 39.5 62.5	74.0 57.6
424	R23Y_062_050a	0.625 0.25 0.125	0.625 0.5 0.375	44	0.625 0.241 0.125	47.6 26.7 27.4	38.2 45.7	0.625 0.25 0.125	45.7 34.0 33.2	47.6 44.3 9.6	42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7
425	R00Y_062_037a	0.625 0.25 0.25	0.625 0.375 0.437	390	0.625 0.25 0.25	50.1 26.6 16.8	31.4 32.3	0.625 0.25 0.25	46.1 34.0 26.2	43.0 37.6 12.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
426	R18Y_062_037a	0.625 0.25 0.375	0.625 0.375 0.437	371	0.625 0.25 0.368	50.2 27.2 11.7	29.6 32.6	0.625 0.25 0.375	46.5 35.2 19.1	40.1 28.4 11.4	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
427	B65R_062_037a	0.625 0.25 0.5	0.625 0.375 0.437	349	0.625 0.25 0.506	50.2 28.6 4.4	29.0 8.9	0.625 0.25 0.5	46.9 37.0 10.1	38.4 15.3 10.6	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
428	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	50.3 29.7 0.0	29.7 359.8	0.625 0.25 0.625	47.5 38.1 3.1	38.3 4.7 9.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
429	B38R_075_050a	0.625 0.25 0.75	0.75 0.5 0.5	316	0.633 0.25 0.75	51.0 35.8 -4.3	36.0 353.0	0.625 0.25 0.75	48.6 39.4 -3.6	39.6 354.7 4.4	317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0
430	B30R_087_062a	0.625 0.25 0.875	0.875 0.625 0.562	307	0.635 0.25 0.875	50.6 40.6 -9.0	41.6 347.4	0.625 0.25 0.875	46.0 42.1 -9.7	43.2 346.9 2.2	307	0.616 0.0 1.0	37.9 65.0 -14.5	66.6 347.4
431	B25R_100_075a	0.625 0.25 1.0	1.0 0.75 0.625	300	0.635 0.25 1.0	50.6 49.9 -15.5	46.6 340.5	0.625 0.25 1.0	49.1 43.7 -15.5	46.4 340.3 1.5	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
432	R61Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	67	0.625 0.385 0.0	53.9 10.2 47.9	49.0 77.8	0.625 0.375 0.0	50.8 21.2 46.0	50.6 65.2 11.5	67	1.0 0.616 0.0	71.6 16.4 76.6	78.4 77.8
433	R50Y_062_050a	0.625 0.375 0.125	0.625 0.5 0.375	60	0.625 0.375 0.125	53.5 14.4 34.3	37.2 67.1	0.625 0.375 0.125	50.7 22.7 38.2	44.5 59.2 9.5	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
434	R31Y_062_037a	0.625 0.375 0.25	0.625 0.375 0.437	49	0.625 0.368 0.25	54.2 17.1 22.2	28.1 52.2	0.625 0.375 0.25	50.9 23.8 30.0	38.3 51.5 10.8	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
435	R00Y_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	56.3 17.7 11.2	20.9 32.3	0.625 0.375 0.375	51.6 24.4 22.1	33.0 42.1 13.7	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
436	R00Y_062_025a	0.625 0.375 0.5	0.625 0.25 0.5	360	0.625 0.375 0.5	56.4 18.5 5.2	19.2 15.9	0.625 0.375 0.5	52.0 26.1 13.2	29.2 26.9 11.8	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
437	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	56.5 19.8 0.0	19.8 359.8	0.625 0.375 0.625	52.6 27.8 4.7	28.2 9.6 10.1	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
438	B34R_075_037a	0.625 0.375 0.75	0.75 0.375 0.562	311	0.631 0.375 0.75	56.8 25.5 -4.4	25.9 350.0	0.625 0.375 0.75	53.8 29.6 -2.9	29.8 354.2 5.3	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
439	B25R_087_050a	0.625 0.375 0.875	0.875 0.5 0.625	300	0.625 0.375 0.875	56.7 29.3 -10.3	31.0 340.5	0.625 0.375 0.875	54.2 31.4 -9.8	32.9 342.6 3.3	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
440	B19R_100_062a	0.625 0.375 1.0	1.0 0.625 0.687	293	0.614 0.375 1.0	56.4 32.7 -16.0	36.4 333.8	0.625 0.375 1.0	54.3 32.9 -16.3	36.8 333.5 2.1	292	0.383 0.0 1.0	32.9 52.3 -25.7	58.3 333.8
441	R81Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	79	0.625 0.51 0.0	59.7 0.5 54.6	54.6 89.4	0.625 0.5 0.0	55.7 11.1 52.4	53.6 73.9 11.5	80	1.0 0.816 0.0	80.8 0.8 87.3	87.3 89.4
442	R76Y_062_050a	0.625 0.5 0.125	0.625 0.5 0.375	76	0.625 0.508 0.125	60.4 2.1 42.3	42.4 87.0	0.625 0.5 0.125	56.2 11.5 43.8	45.3 75.3 10.3	77	1.0 0.766 0.0	78.6 4.3 84.7	84.8 87.0
443	R68Y_062_037a	0.625 0.5 0.25	0.625 0.375 0.437	71	0.625 0.506 0.25	61.1 4.1 30.1	30.4 82.1	0.625 0.5 0.25	56.7 12.5 34.7	36.9 70.0 10.5	71	1.0 0.683 0.0	74.8 11.0 80.4	81.1 82.1
444	R50Y_062_025a	0.625 0.5 0.375	0.625 0.25 0.5	60	0.625 0.5 0.375	61.2 7.2 17.1	18.6 67.1	0.625 0.5 0.375	57.0 14.3 25.0	28.8 60.2 11.4	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
445	R00Y_062_012a	0.625 0.5 0.5	0.625 0.125 0.562	3										

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.L0NA.TXT> / .PS
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS57/PS57L0NA.TXT /.PS
aplicación para la medida salida en la impresión offset, separación cmy0 (CMY0)

TUB material: code=rha4ta

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
486	R00Y_075_075a	0.75 0.0 0.0	0.75 0.75 0.375	390	0.75 0.0 0.0	40.2 53.2 33.6	62.9 32.3	0.75 0.0 0.0	40.7 59.2 36.3	69.4 31.5 6.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
487	R35Y_075_075a	0.75 0.0 0.125	0.75 0.75 0.375	381	0.75 0.0 0.112	40.2 53.7 29.2	61.1 28.5	0.75 0.0 0.125	40.6 60.2 31.6	68.0 27.7 6.9	382	1.0 0.0 0.15	45.5 71.6 39.0	81.5 28.5
488	R18Y_075_075a	0.75 0.0 0.25	0.75 0.75 0.375	371	0.75 0.0 0.237	40.4 54.5 23.4	59.3 23.2	0.75 0.0 0.25	40.9 61.1 25.5	66.2 22.6 6.9	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
489	R00Y_075_075a	0.75 0.0 0.375	0.75 0.75 0.375	360	0.75 0.0 0.375	40.5 55.6 15.8	57.8 15.9	0.75 0.0 0.375	41.0 62.2 19.2	65.1 17.1 7.4	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
490	B65R_075_075a	0.75 0.0 0.5	0.75 0.75 0.375	349	0.75 0.0 0.512	40.5 57.3 8.9	58.0 8.9	0.75 0.0 0.5	40.9 64.0 11.4	65.1 10.1 7.2	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
491	B57R_075_075a	0.75 0.0 0.625	0.75 0.75 0.375	339	0.75 0.0 0.637	40.5 58.5 3.7	58.6 3.7	0.75 0.0 0.625	41.1 65.4 5.1	65.6 4.4 7.0	337	1.0 0.0 0.85	45.9 78.0 5.0	78.2 3.7
492	B50R_075_075a	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	40.6 59.4 -0.1	59.4 359.8	0.75 0.0 0.75	41.1 66.9 0.0	66.9 0.0 7.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
493	B43R_087_087a	0.75 0.0 0.875	0.875 0.875 0.375	322	0.758 0.0 0.875	41.6 65.5 -4.6	65.7 355.9	0.75 0.0 0.875	41.4 69.0 -4.7	69.2 356.0 3.4	322	0.866 0.0 1.0	44.0 74.9 -5.3	75.1 355.9
494	B38R_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0	0.75 0.0 1.0	41.8 71.0 -9.2	71.6 352.5 0.8	317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0
495	R15Y_075_075a	0.75 0.125 0.0	0.75 0.75 0.375	39	0.75 0.112 0.0	43.4 45.5 38.0	59.3 39.9	0.75 0.125 0.0	43.9 51.3 40.0	65.1 37.8 6.1	37	1.0 0.15 0.0	49.8 60.7 50.7	79.1 39.9
496	R00Y_075_062a	0.75 0.125 0.125	0.75 0.625 0.437	390	0.75 0.125 0.125	46.4 44.3 28.0	52.4 32.3	0.75 0.125 0.125	44.5 50.6 34.5	61.3 34.3 9.2	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
497	R31Y_075_062a	0.75 0.125 0.25	0.75 0.625 0.437	379	0.75 0.125 0.239	46.5 44.9 23.4	50.6 27.5	0.75 0.125 0.25	44.8 51.4 28.4	58.8 28.9 8.3	380	1.0 0.0 0.183	45.5 71.8 37.5	81.0 27.5
498	R11Y_075_062a	0.75 0.125 0.375	0.75 0.625 0.437	367	0.75 0.125 0.364	46.6 45.6 17.4	48.8 20.8	0.75 0.125 0.375	45.0 52.4 21.2	56.5 22.0 7.9	367	1.0 0.0 0.383	45.8 73.0 27.8	78.2 20.8
499	B69R_075_062a	0.75 0.125 0.5	0.75 0.625 0.437	353	0.75 0.125 0.51	46.8 47.2 9.5	48.1 11.4	0.75 0.125 0.5	45.4 54.0 12.4	55.4 12.9 7.5	352	1.0 0.0 0.616	46.0 75.5 15.2	77.1 11.4
500	B59R_075_062a	0.75 0.125 0.625	0.75 0.625 0.437	341	0.75 0.125 0.635	46.7 48.6 3.9	48.7 4.6	0.75 0.125 0.625	45.8 55.0 5.5	55.3 5.7 6.6	339	1.0 0.0 0.816	45.9 77.7 6.2	78.0 4.6
501	B50R_075_062a	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	46.8 49.5 -0.1	49.5 359.8	0.75 0.125 0.75	45.9 56.5 -0.2	56.5 359.7 7.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
502	B42R_087_075a	0.75 0.125 0.875	0.875 0.75 0.5	321	0.762 0.125 0.875	47.8 55.7 -4.4	55.9 355.4	0.75 0.125 0.875	46.6 58.6 -5.6	58.9 354.5 3.3	322	0.85 0.0 1.0	43.7 74.3 -5.9	74.6 355.4
503	B36R_100_087a	0.75 0.125 1.0	1.0 0.875 0.562	314	0.766 0.125 1.0	48.1 61.5 -8.7	62.1 351.9	0.75 0.125 1.0	47.0 60.4 -10.4	61.3 350.2 2.2	315	0.733 0.0 1.0	41.3 70.3 -9.9	71.0 351.9
504	R31Y_075_075a	0.75 0.25 0.0	0.75 0.75 0.375	49	0.75 0.237 0.0	48.5 34.3 44.4	56.2 52.2	0.75 0.25 0.0	48.9 39.7 46.7	61.3 49.6 5.8	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
505	R18Y_075_062a	0.75 0.25 0.125	0.75 0.625 0.437	41	0.75 0.239 0.125	50.0 36.1 32.8	48.8 42.2	0.75 0.25 0.125	49.3 39.8 39.4	56.1 44.7 7.6	39	1.0 0.183 0.0	51.1 57.8 52.5	78.1 42.2
506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	52.7 35.4 22.4	41.9 32.3	0.75 0.25 0.25	50.4 39.4 31.9	50.7 38.9 10.5	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
507	R26Y_075_050a	0.75 0.25 0.375	0.75 0.5 0.5	376	0.75 0.25 0.366	52.8 36.0 17.6	40.1 26.1	0.75 0.25 0.375	51.0 39.9 24.4	46.8 31.4 8.0	377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1
508	R00Y_075_050a	0.75 0.25 0.5	0.75 0.5 0.5	360	0.75 0.25 0.5	52.9 37.1 10.5	38.5 15.9	0.75 0.25 0.5	51.3 41.4 15.2	44.1 20.2 6.5	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
509	B61R_075_050a	0.75 0.25 0.625	0.75 0.5 0.5	344	0.75 0.25 0.633	52.9 38.6 4.0	38.8 5.9	0.75 0.25 0.625	52.0 42.7 7.1	43.3 9.4 5.1	342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	53.0 39.6 -0.1	39.6 359.8	0.75 0.25 0.75	52.4 44.4 0.5	44.4 0.6 4.8	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
511	B40R_087_062a	0.75 0.25 0.875	0.875 0.625 0.562	319	0.76 0.25 0.875	53.9 45.8 -4.4	46.0 354.4	0.75 0.25 0.875	53.4 46.0 -5.4	46.3 353.2 1.1	320	0.816 0.0 1.0	43.1 73.2 -7.0	73.6 354.4
512	B34R_100_075a	0.75 0.25 1.0	1.0 0.75 0.625	311	0.762 0.25 1.0	53.7 51.0 -8.9	51.8 350.0	0.75 0.25 1.0	53.7 47.7 -10.9	48.9 347.1 3.9	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
513	R50Y_075_075a	0.75 0.375 0.0	0.75 0.75 0.375	60	0.75 0.375 0.0	54.7 21.6 51.5	55.9 67.1	0.75 0.375 0.0	54.3 28.1 53.1	60.1 62.1 6.6	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
514	R38Y_075_062a	0.75 0.375 0.125	0.75 0.625 0.437	53	0.75 0.364 0.125	55.2 24.7 39.1	46.2 57.6	0.75 0.375 0.125	54.7 28.8 44.2	52.8 56.8 6.6	52	1.0 0.383 0.0	59.5 39.5 62.5	74.0 57.6
515	R23Y_075_050a	0.75 0.375 0.25	0.75 0.5 0.5	44	0.75 0.366 0.25	56.5 26.7 27.4	38.2 45.7	0.75 0.375 0.25	55.2 29.4 35.2	45.9 50.0 8.3	42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7
516	R00Y_075_037a	0.75 0.375 0.375	0.75 0.375 0.562	390	0.75 0.375 0.375	59.0 26.6 16.8	31.4 32.3	0.75 0.375 0.375	56.5 29.0 26.5	39.3 42.3 10.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
517	R18Y_075_037a	0.75 0.375 0.5	0.75 0.375 0.562	371	0.75 0.375 0.493	59.1 27.2 11.7	29.6 23.2	0.75 0.375 0.5	56.9 30.5 18.0	35.4 30.6 7.4	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
518	B65R_075_037a	0.75 0.375 0.625	0.75 0.375 0.562	349	0.75									

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

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
567	R00Y_087_087a	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.0	42.8 62.0 39.2	73.4 32.3	0.875 0.0 0.0	43.2 65.4 40.5	76.9 31.8 3.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
568	R36Y_087_087a	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.116	42.9 62.5 34.7	71.6 29.0	0.875 0.0 0.125	43.3 66.0 35.3	74.9 28.1 3.5	382	1.0 0.0 0.133	45.5 71.5 39.7	81.8 29.0
569	R23Y_087_087a	0.875 0.0 0.25	0.875 0.875 0.437	374	0.875 0.0 0.233	43.0 63.2 29.5	68.1 25.0	0.875 0.0 0.25	43.6 66.5 29.6	72.8 23.9 3.3	375	1.0 0.0 0.266	45.6 72.3 33.8	79.8 25.0
570	R08Y_087_087a	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.364	43.1 64.2 22.7	68.1 19.4	0.875 0.0 0.375	43.6 67.7 23.3	71.6 19.0 3.5	365	1.0 0.0 0.416	45.8 73.4 25.9	77.9 19.4
571	B70R_087_087a	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.51	43.2 65.8 14.8	67.4 12.7	0.875 0.0 0.5	43.7 69.3 16.0	71.2 13.0 3.7	354	1.0 0.0 0.583	45.9 75.2 16.9	77.1 12.7
572	B63R_087_087a	0.875 0.0 0.625	0.875 0.875 0.437	346	0.875 0.0 0.641	43.2 67.3 8.3	67.8 7.0	0.875 0.0 0.625	43.8 70.8 9.3	71.4 7.5 3.6	344	1.0 0.0 0.733	45.9 77.0 9.4	77.5 7.0
573	B56R_087_087a	0.875 0.0 0.75	0.875 0.875 0.437	338	0.875 0.0 0.758	43.2 68.4 3.8	68.5 3.2	0.875 0.0 0.75	43.8 72.3 4.2	72.5 3.3 4.0	337	1.0 0.0 0.866	45.9 78.1 4.4	78.3 3.2
574	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	43.4 69.4 -0.1	69.4 359.8	0.875 0.0 0.875	44.0 73.5 -0.8	73.5 359.3 4.2	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
575	B44R_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	44.3 75.4 -4.7	75.6 356.3	0.875 0.0 1.0	44.2 75.2 -5.0	75.3 356.1 0.4	323	0.883 0.0 1.0	44.3 75.4 -4.7	75.6 356.3
576	R13Y_087_087a	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.116 0.0	46.1 54.3 43.6	69.7 38.7	0.875 0.125 0.0	47.3 56.4 44.0	71.5 38.0 2.4	37	1.0 0.133 0.0	49.2 62.1 49.8	79.6 38.7
577	R00Y_087_075a	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.125	49.1 53.2 33.6	62.9 32.3	0.875 0.125 0.125	47.6 56.0 38.5	67.9 34.5 5.8	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
578	R35Y_087_075a	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.237	49.1 53.7 29.2	61.1 28.5	0.875 0.125 0.25	47.9 56.7 32.6	65.4 29.8 4.6	382	1.0 0.0 0.15	45.5 71.6 39.0	81.5 28.5
579	R18Y_087_075a	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.362	49.3 54.5 23.4	59.3 23.2	0.875 0.125 0.375	48.2 57.5 25.3	62.8 23.7 3.7	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
580	R00Y_087_075a	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.5	49.4 55.6 15.8	57.8 15.9	0.875 0.125 0.5	48.4 59.1 16.9	61.5 15.9 3.7	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
581	B65R_087_075a	0.875 0.125 0.625	0.875 0.75 0.5	349	0.875 0.125 0.637	49.4 57.3 8.9	58.0 8.9	0.875 0.125 0.625	48.8 60.3 9.3	61.0 8.8 3.1	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
582	B57R_087_075a	0.875 0.125 0.75	0.875 0.75 0.5	339	0.875 0.125 0.762	49.4 58.5 3.7	58.6 3.7	0.875 0.125 0.75	48.9 62.0 2.9	62.0 2.7 3.6	337	1.0 0.0 0.85	45.9 78.0 5.0	78.2 3.7
583	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	49.5 59.4 -0.1	59.4 359.8	0.875 0.125 0.875	49.3 62.9 -2.0	62.9 358.1 3.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
584	B43R_100_087a	0.875 0.125 1.0	1.0 0.875 0.562	322	0.883 0.125 1.0	50.5 65.5 -4.6	65.7 355.9	0.875 0.125 1.0	49.6 64.5 -6.6	64.9 354.1 2.3	322	0.866 0.0 1.0	44.0 74.9 -5.3	75.1 355.9
585	R26Y_087_087a	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.233 0.0	50.6 44.1 49.4	66.2 48.2	0.875 0.25 0.0	51.7 45.6 50.7	68.2 48.0 2.3	44	1.0 0.266 0.0	54.4 50.4 56.5	75.7 48.2
586	R15Y_087_075a	0.875 0.25 0.125	0.875 0.75 0.5	39	0.875 0.237 0.125	52.4 45.5 38.0	59.3 39.9	0.875 0.25 0.125	52.6 45.0 43.6	62.7 44.1 5.6	37	1.0 0.15 0.0	49.8 60.7 50.7	79.1 39.9
587	R00Y_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	55.3 44.3 28.0	52.4 32.3	0.875 0.25 0.25	53.7 44.1 35.9	56.8 39.1 8.0	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
588	R31Y_087_062a	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.364	55.4 44.9 23.4	50.6 27.5	0.875 0.25 0.375	54.3 44.5 28.2	52.7 32.3 4.8	380	1.0 0.0 0.183	45.5 71.8 37.5	81.0 27.5
589	R11Y_087_062a	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.489	55.6 45.6 17.4	48.8 20.8	0.875 0.25 0.5	54.5 45.9 19.9	50.0 23.4 2.7	367	1.0 0.0 0.383	45.8 73.0 27.8	78.2 20.8
590	B69R_087_062a	0.875 0.25 0.625	0.875 0.625 0.562	353	0.875 0.25 0.635	55.7 47.2 9.5	48.1 11.4	0.875 0.25 0.625	55.1 47.5 10.8	48.7 12.8 1.4	352	1.0 0.0 0.616	46.0 75.5 15.2	77.1 11.4
591	B59R_087_062a	0.875 0.25 0.75	0.875 0.625 0.562	341	0.875 0.25 0.76	55.6 48.6 3.9	48.7 4.6	0.875 0.25 0.75	55.4 48.8 4.0	49.0 4.6 0.3	339	1.0 0.0 0.816	45.9 77.7 6.2	78.0 4.6
592	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	55.7 49.5 -0.1	49.5 359.8	0.875 0.25 0.875	56.0 49.9 -1.8	49.9 357.9 1.7	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
593	B42R_100_075a	0.875 0.25 1.0	1.0 0.75 0.625	321	0.887 0.25 1.0	56.7 55.7 -4.4	55.9 355.4	0.875 0.25 1.0	56.7 51.9 -6.8	52.3 352.4 4.5	322	0.85 0.0 1.0	43.7 74.3 -5.9	74.6 355.4
594	R41Y_087_087a	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.364 0.0	56.5 32.0 56.4	64.9 60.3	0.875 0.375 0.0	57.5 33.5 57.7	66.8 59.8 2.2	54	1.0 0.416 0.0	61.0 36.6 64.5	74.1 60.3
595	R31Y_087_075a	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.362 0.125	57.4 34.3 44.4	56.2 52.2	0.875 0.375 0.125	57.9 33.6 48.9	59.4 55.5 4.6	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
596	R18Y_087_062a	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.364 0.25	58.9 36.1 32.8	48.8 42.2	0.875 0.375 0.25	58.6 34.1 39.3	52.1 49.0 6.8	39	1.0 0.183 0.0	51.1 57.8 52.5	78.1 42.2
597	R00Y_087_050a	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	61.6 35.4 22.4	41.9 32.3	0.875 0.375 0.375	59.7 33.8 30.7	45.6 42.2 8.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
598	R26Y_087_050a	0.875 0.375 0.5	0.875 0.5 0.625	376	0.875 0.375 0.491	61.7 36.0 17.6	40.1 26.1	0.875 0.375 0.5	60.3 34.8 21.9	41.1 32.1 4.6	377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1
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 37.1 10.5	38.5 15.9	0.875 0.375 0.625	61.1 36.1 12.9	38.3 19.7 2.6	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
600	B61R_087_050a	0.875 0.375 0.75	0.875 0.5 0.625	344	0.875 0.375 0.758	61.8 38.6 4.0	38.8 5.9	0.875 0.375 0.75	61.4 37.8 4.6	38.1 7.0 1.0	342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
601	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	61.9 39.6 -0.1	39.6 359.8	0.875 0.375 0.875	62.3 38.7 -1.4	38.7 357.8 1.6	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
602	B40R_100_062a	0.875 0.375 1.0	1.0 0.625 0.687	319	0.885 0.375 1.0	62.8 45.8 -4.4	46.0 354.4	0.875 0.375 1.0	63.0 40.3 -7.2	40.9 349.7 6.1	320	0.816 0.0 1.0	43.1 73.2 -7.0	73.6 354.4
603	R58Y_087_087a	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.51 0.0	64.0 17.7 65.2	67.6 74.8	0.875 0.5 0.0	63.7 21.0 64.7	68.1 72.0 3.3	65	1.0 0.583 0.0	69.7 20.2 74.6	77.3 74.8
604	R50Y_087_075a	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.5 0.125	63.6 21.6 51.5	55.9 67.1	0.875 0.5 0.125	63.9 22.1 53.8	58.2 67.6 2.4	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
605	R38Y_087_062a	0.875 0.5 0.25	0.875 0.625 0.562	53	0.875 0.489 0.25	64.1 24.7 39.1	46.2 57.6	0.875 0.5 0.25	64.0 23.7 43.4	49.4 61.3 4.4	52	1.0 0.383 0.0	59.5 39.5 62.5	74.0 57.6
606	R23Y_087_050a	0.875 0.5 0.375	0.875 0.5 0.625	44	0.875 0.491 0.375	65.4 26.7 27.4	38.2 45.7	0.875 0.5 0.375	64.9 24.1 33.4	41.2 54.1 6.5	42	1.0 0.2		

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3 0.0	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
649	R38Y_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	45.5 71.4 40.4	82.1 29.5	1.0 0.0 0.125	45.5 71.4 40.1	81.9 29.3 0.3	383	1.0 0.0 0.116	45.5 71.4 40.4	82.1 29.5
650	R26Y_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1	1.0 0.0 0.25	45.6 72.1 34.6	80.0 25.6 0.7	377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1
651	R13Y_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	45.8 72.9 28.7	78.4 21.5	1.0 0.0 0.375	45.8 72.9 28.3	78.3 21.2 0.4	368	1.0 0.0 0.366	45.8 72.9 28.7	78.4 21.5
652	R00Y_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9 0.0	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
653	B68R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	46.0 75.7 14.4	77.1 10.8	1.0 0.0 0.625	46.0 75.6 14.8	77.0 11.1 0.4	351	1.0 0.0 0.633	46.0 75.7 14.4	77.1 10.8
654	B61R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9	1.0 0.0 0.75	45.9 77.1 8.6	77.6 6.4 0.6	342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
655	B55R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	45.9 78.3 3.8	78.4 2.8	1.0 0.0 0.875	45.9 78.2 4.1	78.3 3.0 0.2	336	1.0 0.0 0.883	45.9 78.3 3.8	78.4 2.8
656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8 0.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
657	R11Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	48.6 63.3 49.1	80.2 37.7	1.0 0.125 0.0	48.9 62.8 49.4	79.9 38.1 0.6	36	1.0 0.116 0.0	48.6 63.3 49.1	80.2 37.7
658	R00Y_100_087a	1.0 0.125 0.125	1.0 0.875 0.562	390	1.0 0.125 0.125	51.7 62.0 39.2	73.4 32.3	1.0 0.125 0.125	49.6 62.3 43.6	76.1 34.9 4.8	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
659	R36Y_100_087a	1.0 0.125 0.25	1.0 0.875 0.562	382	1.0 0.125 0.241	51.8 62.5 34.7	71.6 29.0	1.0 0.125 0.25	49.6 61.3 36.9	73.1 30.3 3.0	382	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
660	R23Y_100_087a	1.0 0.125 0.375	1.0 0.875 0.562	374	1.0 0.125 0.358	51.9 63.2 29.5	69.8 25.0	1.0 0.125 0.375	50.0 63.5 30.1	70.3 25.3 2.0	375	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
661	R08Y_100_087a	1.0 0.125 0.5	1.0 0.875 0.562	365	1.0 0.125 0.489	52.0 64.2 22.7	68.1 19.4	1.0 0.125 0.5	50.2 64.7 22.4	68.5 19.1 1.9	365	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
662	B70R_100_087a	1.0 0.125 0.625	1.0 0.875 0.562	355	1.0 0.125 0.635	52.1 65.8 14.8	67.4 12.7	1.0 0.125 0.625	50.6 64.3 14.3	63.2 12.7 1.6	354	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
663	B63R_100_087a	1.0 0.125 0.75	1.0 0.875 0.562	346	1.0 0.125 0.766	52.1 67.3 8.3	67.8 7.0	1.0 0.125 0.75	50.9 66.9 7.4	67.3 8.3 1.5	344	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
664	B56R_100_087a	1.0 0.125 0.875	1.0 0.875 0.562	338	1.0 0.125 0.883	52.1 68.4 3.8	68.5 3.2	1.0 0.125 0.875	51.0 68.3 2.4	68.3 2.0 1.9	337	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
665	B50R_100_087a	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	52.3 69.4 -0.1	69.4 359.8	1.0 0.125 1.0	51.1 69.1 -2.3	69.2 358.0 2.4	330	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
666	R23Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7	1.0 0.25 0.0	53.6 51.9 55.5	76.0 46.8 1.7	42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7
667	R13Y_100_087a	1.0 0.25 0.125	1.0 0.875 0.562	38	1.0 0.241 0.125	55.0 54.3 43.6	69.7 38.7	1.0 0.25 0.125	54.4 51.3 48.5	70.6 43.3 5.7	37	1.0 0.133 0.0	49.2 62.1 49.8	79.6 38.7
668	R00Y_100_075a	1.0 0.25 0.25	1.0 0.75 0.625	390	1.0 0.25 0.25	58.0 53.2 33.6	62.9 32.3	1.0 0.25 0.25	55.3 50.6 40.6	64.9 38.7 7.8	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
669	R35Y_100_075a	1.0 0.25 0.375	1.0 0.75 0.625	381	1.0 0.25 0.362	58.0 53.7 29.2	61.1 28.5	1.0 0.25 0.375	55.8 50.9 33.0	60.7 32.9 5.2	382	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
670	R18Y_100_075a	1.0 0.25 0.5	1.0 0.75 0.625	371	1.0 0.25 0.487	58.2 54.5 23.4	59.3 23.2	1.0 0.25 0.5	56.4 51.4 24.6	57.0 25.5 3.6	371	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
671	R00Y_100_075a	1.0 0.25 0.625	1.0 0.75 0.625	360	1.0 0.25 0.625	58.3 55.6 15.8	57.8 15.9	1.0 0.25 0.625	56.8 52.8 15.9	55.2 16.7 3.1	360	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
672	B65R_100_075a	1.0 0.25 0.75	1.0 0.75 0.625	349	1.0 0.25 0.762	58.3 57.3 8.9	58.0 8.9	1.0 0.25 0.75	57.1 54.5 7.8	55.1 8.1 3.2	348	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
673	B57R_100_075a	1.0 0.25 0.875	1.0 0.75 0.625	339	1.0 0.25 0.887	58.3 58.5 3.7	58.6 3.7	1.0 0.25 0.875	57.6 55.4 1.7	55.5 1.7 3.7	337	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
674	B50R_100_075a	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	58.4 59.4 -0.1	59.4 359.8	1.0 0.25 1.0	58.0 56.2 -3.2	56.3 356.6 4.4	330	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
675	R36Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	58.8 41.1 61.7	74.1 56.3	1.0 0.375 0.0	59.1 40.3 62.0	74.0 56.9 0.9	51	1.0 0.366 0.0	58.8 41.1 61.7	74.1 56.3
676	R26Y_100_087a	1.0 0.375 0.125	1.0 0.875 0.562	46	1.0 0.358 0.125	59.5 44.1 49.4	66.2 48.2	1.0 0.375 0.125	59.2 41.2 53.0	67.1 52.1 4.6	44	1.0 0.266 0.0	54.4 50.4 56.5	75.7 48.2
677	R15Y_100_075a	1.0 0.375 0.25	1.0 0.75 0.625	39	1.0 0.362 0.25	61.3 45.5 38.0	59.3 39.9	1.0 0.375 0.25	59.8 41.2 44.0	60.3 46.8 7.4	37	1.0 0.15 0.0	49.8 60.7 50.7	79.1 39.9
678	R00Y_100_062a	1.0 0.375 0.375	1.0 0.625 0.687	390	1.0 0.375 0.375	64.2 44.3 28.0	52.4 32.3	1.0 0.375 0.375	61.2 40.1 35.6	53.7 41.6 9.2	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
679	R31Y_100_062a	1.0 0.375 0.5	1.0 0.625 0.687	379	1.0 0.375 0.489	64.3 44.9 23.4	50.6 27.5	1.0 0.375 0.5	61.7 40.7 27.1	48.9 33.6 6.1	380	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
680	R11Y_100_062a	1.0 0.375 0.625	1.0 0.625 0.687	367	1.0 0.375 0.614	64.5 45.6 17.4	48.8 20.8	1.0 0.375 0.625	62.6 41.7 17.7	45.3 23.0 4.4	367	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
681	B69R_100_062a	1.0 0.375 0.75	1.0 0.625 0.687	353	1.0 0.375 0									

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.L0NA.TXT>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md
729	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.5 0.0 0.0	0.1 360	1.0 1.0 1.0	95.6 0.0 0.0
730	G50B_100_012d	0.875 1.0 1.0	1.0 0.125 0.937	210	0.875 1.0 1.0	90.7 -3.1 -5.1	6.0 238.4	0.875 1.0 1.0	91.9 -2.9 -4.1	5.0 234.3	1.0 1.0 1.0	96.8 -25.5 -41.5
731	G50B_100_025d	0.75 1.0 1.0	1.0 0.25 0.875	210	0.75 1.0 1.0	85.9 -6.3 -10.3	12.1 238.4	0.75 1.0 1.0	87.8 -5.7 -8.6	10.3 236.4	2.7 210	1.0 1.0 1.0
732	G50B_100_037d	0.625 1.0 1.0	1.0 0.375 0.812	210	0.625 1.0 1.0	81.0 -9.5 -15.5	18.2 238.4	0.625 1.0 1.0	83.2 -8.6 -13.4	15.9 237.2	3.2 210	1.0 1.0 1.0
733	G50B_100_050d	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	76.2 -12.7 -20.7	24.3 238.4	0.5 1.0 1.0	77.6 -12.2 -19.4	22.9 237.6	2.0 210	1.0 1.0 1.0
734	G50B_100_062d	0.375 1.0 1.0	1.0 0.625 0.687	210	0.375 1.0 1.0	71.3 -15.9 -25.9	30.4 238.4	0.375 1.0 1.0	72.3 -15.5 -24.9	29.4 238.1	1.4 210	1.0 1.0 1.0
735	G50B_100_075d	0.25 1.0 1.0	1.0 0.75 0.625	210	0.25 1.0 1.0	66.5 -19.1 -31.1	36.5 238.4	0.25 1.0 1.0	66.5 -19.1 -31.2	36.6 238.4	0.0 210	1.0 1.0 1.0
736	G50B_100_087d	0.125 1.0 1.0	1.0 0.875 0.562	210	0.125 1.0 1.0	61.6 -22.3 -36.3	42.6 238.4	0.125 1.0 1.0	61.2 -21.8 -36.5	42.5 239.0	0.6 210	1.0 1.0 1.0
737	G50B_100_100d	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4	0.0 1.0 1.0	55.3 -24.7 -42.3	49.0 239.6	1.7 210	1.0 1.0 1.0
738	RO0Y_100_012d	1.0 0.875 0.875	1.0 0.125 0.937	390	1.0 0.875 0.875	89.8 8.8 5.6	10.4 32.3	1.0 0.875 0.875	89.7 4.4 7.8	9.0 60.1 4.9	389	1.0 0.0 0.0
739	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0	0.875 0.875 0.875	86.1 1.2 3.6	3.8 70.9 3.8	360	1.0 1.0 1.0
740	G50B_087_012d	0.75 0.875 0.875	0.875 0.125 0.812	210	0.75 0.875 0.875	81.8 -3.1 -5.1	6.0 238.4	0.75 0.875 0.875	82.2 -1.9 -0.8	2.1 204.3 4.4	210	1.0 1.0 1.0
741	G50B_087_025d	0.625 0.875 0.875	0.875 0.25 0.75	210	0.625 0.875 0.875	77.0 -6.3 -10.3	12.1 238.4	0.625 0.875 0.875	77.9 -5.4 -5.5	7.8 225.6 4.9	210	1.0 1.0 1.0
742	G50B_087_037d	0.5 0.875 0.875	0.875 0.375 0.687	210	0.5 0.875 0.875	72.1 -9.5 -15.5	18.2 238.4	0.5 0.875 0.875	72.8 -9.5 -11.3	14.8 229.9 4.2	210	1.0 1.0 1.0
743	G50B_087_050d	0.375 0.875 0.875	0.875 0.5 0.625	210	0.375 0.875 0.875	67.3 -12.7 -20.7	24.3 238.4	0.375 0.875 0.875	67.6 -13.7 -16.9	21.8 230.9 3.9	210	1.0 1.0 1.0
744	G50B_087_062d	0.25 0.875 0.875	0.875 0.625 0.562	210	0.25 0.875 0.875	62.4 -15.9 -25.9	30.4 238.4	0.25 0.875 0.875	62.2 -18.3 -23.4	29.8 231.9 3.4	210	1.0 1.0 1.0
745	G50B_087_075d	0.125 0.875 0.875	0.875 0.75 0.5	210	0.125 0.875 0.875	57.6 -19.1 -31.1	36.5 238.4	0.125 0.875 0.875	57.2 -22.1 -28.6	36.1 232.2 3.9	210	1.0 1.0 1.0
746	G50B_087_087d	0.0 0.875 0.875	0.875 0.875 0.437	210	0.0 0.875 0.875	52.7 -22.3 -36.3	42.6 238.4	0.0 0.875 0.875	51.9 -26.3 -34.9	43.7 232.9 4.3	210	1.0 1.0 1.0
747	RO0Y_100_025d	1.0 0.75 0.75	1.0 0.25 0.875	390	1.0 0.75 0.75	83.0 17.7 11.2	20.9 32.3	1.0 0.75 0.75	82.3 11.7 15.1	19.1 52.1 7.1	389	1.0 0.0 0.0
748	RO0Y_087_012d	0.875 0.75 0.75	0.875 0.125 0.812	390	0.875 0.75 0.75	80.4 8.8 5.6	10.4 32.3	0.875 0.75 0.75	79.1 8.0 10.9	13.6 53.6 5.5	389	1.0 0.0 0.0
749	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0	0.75 0.75 0.75	75.6 4.4 6.7	8.0 56.1 8.3	360	1.0 1.0 1.0
750	G50B_075_012d	0.625 0.75 0.75	0.75 0.125 0.687	210	0.625 0.75 0.75	72.9 -3.1 -5.1	6.0 238.4	0.625 0.75 0.75	71.2 0.3 1.9	2.0 79.0 8.2	210	1.0 1.0 1.0
751	G50B_075_025d	0.5 0.75 0.75	0.75 0.25 0.625	210	0.5 0.75 0.75	68.1 -6.3 -10.3	12.1 238.4	0.5 0.75 0.75	66.4 -4.7 -3.8	6.1 219.4 6.9	210	1.0 1.0 1.0
752	G50B_075_037d	0.375 0.75 0.75	0.75 0.375 0.562	210	0.375 0.75 0.75	63.2 -9.5 -15.5	18.2 238.4	0.375 0.75 0.75	61.8 -9.3 -9.6	13.4 225.8 6.0	210	1.0 1.0 1.0
753	G50B_075_050d	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.75	58.4 -12.7 -20.7	24.3 238.4	0.25 0.75 0.75	56.5 -15.2 -16.0	22.1 226.3 5.6	210	1.0 1.0 1.0
754	G50B_075_062d	0.125 0.75 0.75	0.75 0.625 0.437	210	0.125 0.75 0.75	53.5 -15.9 -25.9	30.4 238.4	0.125 0.75 0.75	52.2 -19.8 -21.1	28.9 226.8 6.3	210	1.0 1.0 1.0
755	G50B_075_075d	0.0 0.75 0.75	0.75 0.75 0.375	210	0.0 0.75 0.75	48.7 -19.1 -31.1	36.5 238.4	0.0 0.75 0.75	47.3 -25.7 -27.2	37.5 226.6 7.8	210	1.0 1.0 1.0
756	RO0Y_100_037d	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	76.8 26.6 16.8	31.4 32.3	1.0 0.625 0.625	76.1 18.3 22.9	29.3 51.3 10.2	389	1.0 0.0 0.0
757	RO0Y_087_025d	0.875 0.625 0.625	0.875 0.25 0.75	390	0.875 0.625 0.625	74.1 17.7 11.2	20.9 32.3	0.875 0.625 0.625	73.0 14.4 18.5	23.5 52.0 8.0	389	1.0 0.0 0.0
758	RO0Y_075_012d	0.75 0.625 0.625	0.75 0.125 0.687	390	0.75 0.625 0.625	71.5 8.8 5.6	10.4 32.3	0.75 0.625 0.625	69.8 10.1 14.0	17.3 54.0 8.6	389	1.0 0.0 0.0
759	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0	0.625 0.625 0.625	65.4 5.8 9.1	10.9 57.3 11.4	360	1.0 1.0 1.0
760	G50B_062_012d	0.5 0.625 0.625	0.625 0.125 0.562	210	0.5 0.625 0.625	64.0 -3.1 -5.1	6.0 238.4	0.5 0.625 0.625	61.0 0.4 3.7	3.7 83.2 10.1	210	1.0 1.0 1.0
761	G50B_062_025d	0.375 0.625 0.625	0.625 0.25 0.5	210	0.375 0.625 0.625	59.2 -6.3 -10.3	12.1 238.4	0.375 0.625 0.625	56.7 -5.3 -2.1	5.7 201.6 8.6	210	1.0 1.0 1.0
762	G50B_062_037d	0.25 0.625 0.625	0.625 0.375 0.437	210	0.25 0.625 0.625	54.3 -9.5 -15.5	18.2 238.4	0.25 0.625 0.625	51.9 -12.3 -8.5	14.9 214.7 7.9	210	1.0 1.0 1.0
763	G50B_062_050d	0.125 0.625 0.625	0.625 0.5 0.375	210	0.125 0.625 0.625	49.4 -12.7 -20.7	24.3 238.4	0.125 0.625 0.625	48.0 -18.0 -13.9	22.8 217.6 8.7	210	1.0 1.0 1.0
764	G50B_062_062d	0.0 0.625 0.625	0.625 0.625 0.312	210	0.0 0.625 0.625	44.6 -15.9 -25.9	30.4 238.4	0.0 0.625 0.625	43.3 -25.1 -20.1	32.1 218.6 10.9	210	1.0 1.0 1.0
765	RO0Y_100_050d	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	70.5 35.4 22.4	41.9 32.3	1.0 0.5 0.5	68.2 29.0 29.0	41.1 45.0 9.5	389	1.0 0.0 0.0
766	RO0Y_087_037d	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.5	67.9 26.6 16.8	31.4 32.3	0.875 0.5 0.5	65.3 24.5 25.2	35.1 45.7 9.0	389	1.0 0.0 0.0
767	RO0Y_075_025d	0.75 0.5 0.5	0.75 0.25 0.62									

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md
810	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.6 0.0 0.1	0.1 116.7	0.1 360	1.0 1.0 1.0
811	BOOR_100_012d	0.875 0.875 1.0	1.0 0.125 0.937	270	0.875 0.875 1.0	86.8 3.6 -5.0	6.2 306.2	0.875 0.875 1.0	87.2 3.8 -5.3	6.6 305.3	0.5 270	1.0 0.0 1.0
812	BOOR_100_025d	0.75 0.75 1.0	1.0 0.25 0.875	270	0.75 0.75 1.0	77.9 7.3 -10.1	12.5 306.2	0.75 0.75 1.0	76.6 9.6 -10.6	14.3 312.1	2.6 270	0.0 0.0 1.0
813	BOOR_100_037d	0.625 0.625 1.0	1.0 0.375 0.812	270	0.625 0.625 1.0	69.1 11.0 -15.1	18.7 306.2	0.625 0.625 1.0	67.2 13.6 -15.6	20.8 310.1	3.2 270	0.0 0.0 1.0
814	BOOR_100_050d	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	60.3 14.7 -20.2	25.0 306.2	0.5 0.5 1.0	55.8 19.6 -21.4	29.1 312.4	6.7 270	0.0 0.0 1.0
815	BOOR_100_062d	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	51.5 18.4 -25.2	31.3 306.2	0.375 0.375 1.0	45.8 24.1 -26.3	35.7 312.5	8.1 270	0.0 0.0 1.0
816	BOOR_100_075d	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	42.7 22.1 -30.3	37.5 306.2	0.25 0.25 1.0	37.4 26.6 -31.6	41.3 310.1	6.9 270	0.0 0.0 1.0
817	BOOR_100_087d	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	33.9 25.8 -35.3	43.8 306.2	0.125 0.125 1.0	28.7 31.4 -36.1	47.8 311.0	7.6 270	0.0 0.0 1.0
818	BOOR_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 0.0 1.0	23.4 30.6 -39.6	50.1 307.6	2.0 270	0.0 0.0 1.0
819	Y00G_100_012d	1.0 1.0 0.875	1.0 0.125 0.937	90	1.0 1.0 0.875	94.6 -1.2 11.9	12.0 96.1	1.0 1.0 0.875	94.6 -2.5 9.9	10.2 104.1	2.3 89	1.0 1.0 0.0
820	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0	0.875 0.875 0.875	86.3 1.2 3.7	3.9 71.1	3.9 360	1.0 1.0 1.0
821	BOOR_087_012d	0.75 0.75 0.875	0.875 0.125 0.812	270	0.75 0.75 0.875	77.9 3.6 -5.0	6.2 306.2	0.75 0.75 0.875	76.0 6.9 -5.3	7.3 341.0	4.5 270	0.0 0.0 1.0
822	BOOR_087_025d	0.625 0.625 0.875	0.875 0.25 0.75	270	0.625 0.625 0.875	69.0 7.3 -10.1	12.5 306.2	0.625 0.625 0.875	66.7 11.0 -8.0	13.6 323.8	4.7 270	0.0 0.0 1.0
823	BOOR_087_037d	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.5 0.875	60.2 11.0 -15.1	18.7 306.2	0.5 0.5 0.875	55.5 16.6 -14.6	22.1 318.6	7.2 270	0.0 0.0 1.0
824	BOOR_087_050d	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	51.4 14.7 -20.2	25.0 306.2	0.375 0.375 0.875	45.6 21.0 -20.4	29.2 315.8	8.5 270	0.0 0.0 1.0
825	BOOR_087_062d	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.6 18.4 -25.2	31.3 306.2	0.25 0.25 0.875	37.1 23.2 -26.2	35.0 311.5	7.3 270	0.0 0.0 1.0
826	BOOR_087_075d	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	33.8 22.1 -30.3	37.5 306.2	0.125 0.125 0.875	29.0 26.9 -31.2	41.2 310.8	6.8 270	0.0 0.0 1.0
827	BOOR_087_087d	0.0 0.0 0.875	0.875 0.875 0.437	270	0.0 0.0 0.875	24.9 25.8 -35.3	43.8 306.2	0.0 0.0 0.875	23.4 26.1 -35.1	43.8 306.6	1.6 270	0.0 0.0 1.0
828	Y00G_100_025d	1.0 1.0 0.75	1.0 0.25 0.875	90	1.0 1.0 0.75	93.6 -2.5 23.8	24.0 96.1	1.0 1.0 0.75	93.5 -4.4 20.0	20.4 102.4	4.2 89	1.0 1.0 0.0
829	Y00G_087_012d	0.875 0.875 0.75	0.875 0.125 0.812	90	0.875 0.875 0.75	85.7 -1.2 11.9	12.0 96.1	0.875 0.875 0.75	85.2 -0.7 13.0	13.1 93.4	1.3 89	1.0 1.0 0.0
830	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0	0.75 0.75 0.75	75.1 4.6 6.6	8.1 54.7	8.5 360	1.0 1.0 1.0
831	BOOR_075_012d	0.625 0.625 0.75	0.75 0.125 0.687	270	0.625 0.625 0.75	68.9 3.6 -5.0	6.2 306.2	0.625 0.625 0.75	66.1 8.4 0.2	8.4 1.7	7.7 270	1.0 0.0 1.0
832	BOOR_075_025d	0.5 0.5 0.75	0.75 0.25 0.625	270	0.5 0.5 0.75	60.1 7.3 -10.1	12.5 306.2	0.5 0.5 0.75	54.8 13.8 -6.8	15.4 333.6	8.9 270	0.0 0.0 1.0
833	BOOR_075_037d	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	51.3 11.0 -15.1	18.7 306.2	0.375 0.375 0.75	45.6 17.2 -13.3	21.7 322.1	8.5 270	0.0 0.0 1.0
834	BOOR_075_050d	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	42.5 14.7 -20.2	25.0 306.2	0.25 0.25 0.75	37.2 19.3 -19.7	27.6 314.5	7.0 270	0.0 0.0 1.0
835	BOOR_075_062d	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	33.7 18.4 -25.2	31.3 306.2	0.125 0.125 0.75	29.3 22.6 -25.7	34.2 311.4	6.1 270	0.0 0.0 1.0
836	BOOR_075_075d	0.0 0.0 0.75	0.75 0.75 0.375	270	0.0 0.0 0.75	24.9 22.1 -30.3	37.5 306.2	0.0 0.0 0.75	23.6 21.0 -30.2	36.9 304.8	1.6 270	0.0 0.0 1.0
837	Y00G_100_037d	1.0 1.0 0.625	1.0 0.375 0.812	90	1.0 1.0 0.625	92.6 -3.8 35.8	36.0 96.1	1.0 1.0 0.625	92.4 -6.1 30.9	31.6 101.2	5.3 89	1.0 1.0 0.0
838	Y00G_087_025d	0.875 0.875 0.625	0.875 0.25 0.75	90	0.875 0.875 0.625	84.7 -2.5 23.8	24.0 96.1	0.875 0.875 0.625	84.2 -2.8 23.6	23.8 96.7	0.5 89	1.0 1.0 0.0
839	Y00G_075_012d	0.75 0.75 0.625	0.75 0.125 0.687	90	0.75 0.75 0.625	76.8 -1.2 11.9	12.0 96.1	0.75 0.75 0.625	74.4 2.4 16.3	16.5 81.4	6.2 89	1.0 1.0 0.0
840	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0	0.625 0.625 0.625	65.5 5.9 9.4	11.1 57.6	11.6 360	1.0 1.0 1.0
841	BOOR_062_012d	0.5 0.5 0.625	0.625 0.125 0.562	270	0.5 0.5 0.625	60.0 3.6 -5.0	6.2 306.2	0.5 0.5 0.625	54.5 11.4 1.1	11.4 5.8	11.3 270	1.0 0.0 1.0
842	BOOR_062_025d	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	51.2 7.3 -10.1	12.5 306.2	0.375 0.375 0.625	45.2 14.8 -6.0	16.0 337.7	10.3 270	0.0 0.0 1.0
843	BOOR_062_037d	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	42.4 11.0 -15.1	18.7 306.2	0.25 0.25 0.625	36.9 16.3 -13.2	21.0 320.9	7.8 270	0.0 0.0 1.0
844	BOOR_062_050d	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	33.6 14.7 -20.2	25.0 306.2	0.125 0.125 0.625	29.1 19.3 -19.9	27.7 314.1	6.3 270	0.0 0.0 1.0
845	BOOR_062_062d	0.0 0.0 0.625	0.625 0.625 0.312	270	0.0 0.0 0.625	24.8 18.4 -25.2	31.3 306.2	0.0 0.0 0.625	23.5 16.8 -24.9	30.0 304.0	2.1 270	0.0 0.0 1.0
846	Y00G_100_050d	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	91.7 -5.1 47.7	48.0 96.1	1.0 1.0 0.5	91.2 -7.6 43.4	44.1 100.0	5.0 89	1.0 1.0 0.0
847	Y00G_087_037d	0.875 0.875 0.5	0.875 0.375 0.687	90	0.875 0.875 0.5	83.7 -3.8 35.8	36.0 96.1	0.875 0.875 0.5	83.1 -4.5 35.6	35.8 97.2	0.9 89	1.0 1.0 0.0
848	Y00G_075_025d	0.75 0.75 0.5	0.75 0.25 0.625	90	0.75 0.75 0.5	75.8 -2.5 23.8						

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS57/PS57L0NA.TXT /.PS TUB material: code=rh4ta
aplicación para la medida salida en la impresión offset, separación cmy0 (CMY0)

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md
891	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.6 0.0 0.1	111.4 0.1 360	1.0 1.0 1.0	95.6 0.0 0.0
892	B50R_100_012d	1.0 0.875 1.0	1.0 0.125 0.937	330	1.0 0.875 1.0	89.4 9.9 0.0	9.9 359.8	1.0 0.875 1.0	90.7 6.8 -1.4	6.9 348.2 3.6 330	1.0 0.0 1.0	46.1 79.3 -0.2
893	B50R_100_025d	1.0 0.75 1.0	1.0 0.25 0.875	330	1.0 0.75 1.0	83.2 19.8 0.0	19.8 359.8	1.0 0.75 1.0	84.2 15.6 -2.4	15.8 351.1 4.9 330	1.0 0.0 1.0	46.1 79.3 -0.2
894	B50R_100_037d	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	77.0 29.7 0.0	29.7 359.8	1.0 0.625 1.0	78.5 23.6 -3.2	23.8 352.2 7.0 330	1.0 0.0 1.0	46.1 79.3 -0.2
895	B50R_100_050d	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	70.8 39.6 -0.1	39.6 359.8	1.0 0.5 1.0	70.6 35.6 -3.8	35.8 353.8 5.5 330	1.0 0.0 1.0	46.1 79.3 -0.2
896	B50R_100_062d	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	64.6 49.5 -0.1	49.5 359.8	1.0 0.375 1.0	63.5 46.7 -3.8	46.9 355.3 4.7 330	1.0 0.0 1.0	46.1 79.3 -0.2
897	B50R_100_075d	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	58.4 59.4 -0.1	59.4 359.8	1.0 0.25 1.0	57.0 58.1 -2.9	58.1 357.1 3.4 330	1.0 0.0 1.0	46.1 79.3 -0.2
898	B50R_100_087d	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	52.3 69.4 -0.1	69.4 359.8	1.0 0.125 1.0	50.3 70.4 -1.6	70.4 358.6 2.6 330	1.0 0.0 1.0	46.1 79.3 -0.2
899	B50R_100_100d	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	1.0 0.0 1.0	45.4 79.5 0.0	79.5 0.7 1.4 330	1.0 0.0 1.0	46.1 79.3 -0.2
900	GO0B_100_012d	0.875 1.0 0.875	1.0 0.125 0.937	150	0.875 1.0 0.875	89.9 -8.1 3.7	8.9 155.5	0.875 1.0 0.875	90.9 -5.6 5.6	7.9 135.3 3.2 149	0.0 1.0 0.0	50.0 -65.0
901	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0	0.875 0.875 0.875	86.2 1.2 3.6	3.8 71.0 3.8 360	1.0 1.0 1.0	95.6 0.0 0.0
902	B50R_087_012d	0.875 0.75 0.875	0.875 0.125 0.812	330	0.875 0.75 0.875	80.5 9.9 0.0	9.9 359.8	0.875 0.75 0.875	80.1 10.0 2.1	10.2 11.8 2.1 330	1.0 0.0 1.0	46.1 79.3 -0.2
903	B50R_087_025d	0.875 0.625 0.875	0.875 0.25 0.75	330	0.875 0.625 0.875	74.3 19.8 0.0	19.8 359.8	0.875 0.625 0.875	74.6 18.0 0.9	18.1 2.9 2.0 330	1.0 0.0 1.0	46.1 79.3 -0.2
904	B50R_087_037d	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	68.1 29.7 0.0	29.7 359.8	0.875 0.5 0.875	66.7 30.6 -0.6	30.6 358.7 1.7 330	1.0 0.0 1.0	46.1 79.3 -0.2
905	B50R_087_050d	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	61.9 39.6 -0.1	39.6 359.8	0.875 0.375 0.875	60.5 40.8 -1.0	40.8 358.5 2.0 330	1.0 0.0 1.0	46.1 79.3 -0.2
906	B50R_087_062d	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	55.7 49.5 -0.1	49.5 359.8	0.875 0.25 0.875	54.0 52.3 -1.0	52.3 358.7 3.3 330	1.0 0.0 1.0	46.1 79.3 -0.2
907	B50R_087_075d	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	49.5 59.4 -0.1	59.4 359.8	0.875 0.125 0.875	47.7 64.4 -0.5	64.4 359.4 5.3 330	1.0 0.0 1.0	46.1 79.3 -0.2
908	B50R_087_087d	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	43.4 69.4 -0.1	69.4 359.8	0.875 0.0 0.875	42.9 73.7 1.1	73.7 0.8 4.5 330	1.0 0.0 1.0	46.1 79.3 -0.2
909	GO0B_100_025d	0.75 1.0 0.75	1.0 0.25 0.875	150	0.75 1.0 0.75	84.2 -16.2 7.4	17.8 155.5	0.75 1.0 0.75	85.6 -11.0 10.4	15.2 136.5 6.2 149	0.0 1.0 0.0	50.0 -65.0
910	GO0B_087_012d	0.75 0.875 0.75	0.875 0.125 0.812	150	0.75 0.875 0.75	81.0 -8.1 3.7	8.9 155.5	0.75 0.875 0.75	81.1 -4.3 8.3	9.4 117.5 5.9 149	0.0 1.0 0.0	50.0 -65.0
911	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0	0.75 0.75 0.75	75.6 4.3 6.4	7.8 56.1 8.1 360	1.0 1.0 1.0	95.6 0.0 0.0
912	B50R_075_012d	0.75 0.625 0.75	0.75 0.125 0.687	330	0.75 0.625 0.75	71.6 9.9 0.0	9.9 359.8	0.75 0.625 0.75	70.5 12.2 4.7	13.1 21.4 5.4 330	1.0 0.0 1.0	46.1 79.3 -0.2
913	B50R_075_025d	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	65.4 19.8 0.0	19.8 359.8	0.75 0.5 0.75	63.2 23.9 2.7	24.1 6.6 5.4 330	1.0 0.0 1.0	46.1 79.3 -0.2
914	B50R_075_037d	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	59.2 29.7 0.0	29.7 359.8	0.75 0.375 0.75	57.3 34.4 1.7	34.4 2.9 5.4 330	1.0 0.0 1.0	46.1 79.3 -0.2
915	B50R_075_050d	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	53.0 39.6 -0.1	39.6 359.8	0.75 0.25 0.75	50.7 45.7 0.7	45.8 0.9 6.6 330	1.0 0.0 1.0	46.1 79.3 -0.2
916	B50R_075_062d	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	46.8 49.5 -0.1	49.5 359.8	0.75 0.125 0.75	44.9 57.7 0.1	57.7 0.1 8.4 330	1.0 0.0 1.0	46.1 79.3 -0.2
917	B50R_075_075d	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	40.6 59.4 -0.1	59.4 359.8	0.75 0.0 0.75	40.3 67.0 1.0	67.0 0.8 7.6 330	1.0 0.0 1.0	46.1 79.3 -0.2
918	GO0B_100_037d	0.625 1.0 0.625	1.0 0.375 0.812	150	0.625 1.0 0.625	78.5 -24.3 11.1	26.7 155.5	0.625 1.0 0.625	79.8 -17.2 15.5	23.2 137.8 8.5 149	0.0 1.0 0.0	50.0 -65.0
919	GO0B_087_025d	0.625 0.875 0.625	0.875 0.25 0.75	150	0.625 0.875 0.625	75.3 -16.2 7.4	17.8 155.5	0.625 0.875 0.625	76.0 -10.5 12.9	16.7 129.1 8.0 149	0.0 1.0 0.0	50.0 -65.0
920	GO0B_075_012d	0.625 0.75 0.625	0.75 0.125 0.687	150	0.625 0.75 0.625	72.1 -8.1 3.7	8.9 155.5	0.625 0.75 0.625	70.7 -2.0 10.9	11.1 100.3 9.5 149	0.0 1.0 0.0	50.0 -65.0
921	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0	0.625 0.625 0.625	66.0 5.6 8.9	10.5 57.5 10.9 360	1.0 1.0 1.0	95.6 0.0 0.0
922	B50R_062_012d	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	62.7 9.9 0.0	9.9 359.8	0.625 0.5 0.625	59.5 17.0 6.1	18.1 19.9 9.9 330	1.0 0.0 1.0	46.1 79.3 -0.2
923	B50R_062_025d	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	56.5 19.8 0.0	19.8 359.8	0.625 0.375 0.625	53.7 26.9 4.3	27.3 9.1 8.8 330	1.0 0.0 1.0	46.1 79.3 -0.2
924	B50R_062_037d	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	50.3 29.7 0.0	29.7 359.8	0.625 0.25 0.625	47.9 38.2 2.9	38.3 4.3 9.3 330	1.0 0.0 1.0	46.1 79.3 -0.2
925	B50R_062_050d	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	44.1 39.6 -0.1	39.6 359.8	0.625 0.125 0.625	42.0 50.1 1.3	50.1 1.5 10.7 330	1.0 0.0 1.0	46.1 79.3 -0.2
926	B50R_062_062d	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	37.9 49.5 -0.1	49.5 359.8	0.625 0.0 0.625	37.5 59.5 0.8	59.5 0.7 10.0 330	1.0 0.0 1.0	46.1 79.3 -0.2
927	GO0B_100_050d	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	72.8 -32.5 14.8	35.7 15					

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

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md	
972	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0 0.0	23.1 1.0	-1.6 1.9	302.0 2.2	360	95.6 0.0 0.0 0.0	
973	NW_012a	0.125 0.125 0.125	0.125 0.125 0.125	0.125 360	0.125 0.125 0.125	33.2 0.0 0.0	0.125 0.125 0.125	28.5 8.0 4.0	8.9 26.4 10.1	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
974	NW_025a	0.25 0.25 0.25	0.25 0.25 0.25	0.25 360	0.25 0.25 0.25	42.1 0.0 0.0	0.25 0.25 0.25	36.5 9.3 8.5	12.6 42.5 13.9	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
975	NW_037a	0.375 0.375 0.375	0.375 0.375 0.375	0.375 360	0.375 0.375 0.375	51.0 0.0 0.0	0.375 0.375 0.375	45.3 10.1 10.9	14.8 47.1 15.9	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
976	NW_050a	0.5 0.5 0.5	0.5 0.5 0.5	0.5 360	0.5 0.5 0.5	60.0 0.0 0.0	0.5 0.5 0.5	55.2 8.8 10.0	13.3 48.4 14.2	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
977	NW_062a	0.625 0.625 0.625	0.625 0.625 0.625	0.625 360	0.625 0.625 0.625	68.9 0.0 0.0	0.625 0.625 0.625	66.4 5.6 9.0	10.6 58.3 10.9	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
978	NW_075a	0.75 0.75 0.75	0.75 0.75 0.75	0.75 360	0.75 0.75 0.75	77.8 0.0 0.0	0.75 0.75 0.75	76.2 3.9 6.3	7.5 57.9 7.6	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
979	NW_087a	0.875 0.875 0.875	0.875 0.875 0.875	0.875 360	0.875 0.875 0.875	86.7 0.0 0.0	0.875 0.875 0.875	86.5 1.1 3.3	3.6 70.5 3.6	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
980	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 360	1.0 1.0 1.0	95.6 0.0 0.0	1.0 1.0 1.0	95.6 0.0 0.0	0.1 126.7 0.1	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
981	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0 0.0	22.9 1.2	-0.6 1.4	332.7 2.0	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0
982	NW_012a	0.125 0.125 0.125	0.125 0.125 0.125	0.125 360	0.125 0.125 0.125	33.2 0.0 0.0	0.125 0.125 0.125	28.4 8.3 4.3	9.4 27.2 10.5	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
983	NW_025a	0.25 0.25 0.25	0.25 0.25 0.25	0.25 360	0.25 0.25 0.25	42.1 0.0 0.0	0.25 0.25 0.25	35.9 9.7 9.1	13.3 43.2 14.7	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
984	NW_037a	0.375 0.375 0.375	0.375 0.375 0.375	0.375 360	0.375 0.375 0.375	51.0 0.0 0.0	0.375 0.375 0.375	45.6 9.9 11.0	14.9 47.9 15.8	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
985	NW_050a	0.5 0.5 0.5	0.5 0.5 0.5	0.5 360	0.5 0.5 0.5	60.0 0.0 0.0	0.5 0.5 0.5	55.1 8.6 9.9	13.1 49.1 14.0	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
986	NW_062a	0.625 0.625 0.625	0.625 0.625 0.625	0.625 360	0.625 0.625 0.625	68.9 0.0 0.0	0.625 0.625 0.625	66.2 5.6 9.1	10.7 58.2 11.1	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
987	NW_075a	0.75 0.75 0.75	0.75 0.75 0.75	0.75 360	0.75 0.75 0.75	77.8 0.0 0.0	0.75 0.75 0.75	76.0 4.1 7.4	7.6 56.0 7.6	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
988	NW_087a	0.875 0.875 0.875	0.875 0.875 0.875	0.875 360	0.875 0.875 0.875	86.7 0.0 0.0	0.875 0.875 0.875	86.6 1.2 3.4	3.6 70.8 3.6	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
989	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 360	1.0 1.0 1.0	95.6 0.0 0.0	1.0 1.0 1.0	95.6 0.0 0.0	0.0 133.9 0.1	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
990	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0 0.0	23.0 0.5	-0.7 0.9	307.9 1.6	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0
991	NW_012a	0.125 0.125 0.125	0.125 0.125 0.125	0.125 360	0.125 0.125 0.125	33.2 0.0 0.0	0.125 0.125 0.125	28.1 7.9 4.7	9.2 30.9 10.6	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
992	NW_025a	0.25 0.25 0.25	0.25 0.25 0.25	0.25 360	0.25 0.25 0.25	42.1 0.0 0.0	0.25 0.25 0.25	36.3 9.2 9.2	13.0 45.2 14.3	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
993	NW_037a	0.375 0.375 0.375	0.375 0.375 0.375	0.375 360	0.375 0.375 0.375	51.0 0.0 0.0	0.375 0.375 0.375	44.9 10.0 11.2	15.1 48.2 16.3	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
994	NW_050a	0.5 0.5 0.5	0.5 0.5 0.5	0.5 360	0.5 0.5 0.5	60.0 0.0 0.0	0.5 0.5 0.5	54.7 8.9 9.9	13.3 48.3 14.3	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
995	NW_062a	0.625 0.625 0.625	0.625 0.625 0.625	0.625 360	0.625 0.625 0.625	68.9 0.0 0.0	0.625 0.625 0.625	66.3 5.6 9.3	10.9 59.0 11.2	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
996	NW_075a	0.75 0.75 0.75	0.75 0.75 0.75	0.75 360	0.75 0.75 0.75	77.8 0.0 0.0	0.75 0.75 0.75	75.8 4.1 6.3	7.5 56.9 7.8	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
997	NW_087a	0.875 0.875 0.875	0.875 0.875 0.875	0.875 360	0.875 0.875 0.875	86.7 0.0 0.0	0.875 0.875 0.875	86.3 1.1 3.4	3.6 71.6 3.6	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
998	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 360	1.0 1.0 1.0	95.6 0.0 0.0	1.0 1.0 1.0	95.7 0.0 0.1	120.9 0.2	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
999	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0 0.0	22.8 0.5	-0.5 0.8	317.5 1.7	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0
1000	NW_012a	0.125 0.125 0.125	0.125 0.125 0.125	0.125 360	0.125 0.125 0.125	33.2 0.0 0.0	0.125 0.125 0.125	27.9 8.0 4.4	9.1 28.8 10.5	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1001	NW_025a	0.25 0.25 0.25	0.25 0.25 0.25	0.25 360	0.25 0.25 0.25	42.1 0.0 0.0	0.25 0.25 0.25	35.8 9.1 9.3	13.0 45.5 14.5	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1002	NW_037a	0.375 0.375 0.375	0.375 0.375 0.375	0.375 360	0.375 0.375 0.375	51.0 0.0 0.0	0.375 0.375 0.375	44.9 10.0 11.4	15.2 48.7 16.4	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1003	NW_050a	0.5 0.5 0.5	0.5 0.5 0.5	0.5 360	0.5 0.5 0.5	60.0 0.0 0.0	0.5 0.5 0.5	54.7 9.1 10.4	13.8 48.7 14.8	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1004	NW_062a	0.625 0.625 0.625	0.625 0.625 0.625	0.625 360	0.625 0.625 0.625	68.9 0.0 0.0	0.625 0.625 0.625	66.0 5.6 9.5	11.1 59.3 11.4	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1005	NW_075a	0.75 0.75 0.75	0.75 0.75 0.75	0.75 360	0.75 0.75 0.75	77.8 0.0 0.0	0.75 0.75 0.75	75.7 4.1 6.4	7.6 57.3 7.9	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1006	NW_087a	0.875 0.875 0.875	0.875 0.875 0.875	0.875 360	0.875 0.875 0.875	86.7 0.0 0.0	0.875 0.875 0.875	86.3 1.1 3.5	3.7 71.9 3.8	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1007	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 360	1.0 1.0 1.0	95.6 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	0.0 113.6 0.1	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1008	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0 0.0	23.1 1.4	-1.9 2.4	306.9 2.7	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0
1009	NW_006a	0.066 0.066 0.066	0.066 0.066 0.066	0.066 360	0.066 0.066 0.066	29.0 0.0 0.0	0.066 0.066 0.066	26.0 5.8 0.2	5.8 2.4 6.6	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1010	NW_013a	0.133 0.133 0.133	0.133 0.133 0.133	0.133 360	0.133 0.133 0.133	33.8 0.0 0.0	0.133 0.133 0.133	28.8 8.4 3.0	9.0 19.7 10.3	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1011	NW_020a	0.2 0.2 0.2	0.2 0.2 0.2	0.2 360	0.2 0.2 0.2	38.6 0.0 0.0	0.2 0.2 0.2	32.3 9.7 5.8	11.4 30.8 13.0	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1012	NW_026a	0.266 0.266 0.266	0.266 0.266 0.266	0.266 360	0.266 0.266 0.266	43.3 0.0 0.0	0.266 0.266 0.266	37.0 9.1 8.3	12.3 42.4 13.8	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1013	NW_033a	0.333 0.333 0.333	0.333 0.333 0.333	0.333 360	0.333 0.333 0.333	48.1 0.0 0.0	0.333 0.333 0.333	41.4 10.4 9.4	14.0 42.0 15.5	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1014	NW_040a	0.4 0.4 0.4	0.4 0.4 0.4	0.4 360	0.4 0.4 0.4	52.8 0.0 0.0	0.4 0.4 0.4	47.5 8.9 9.8	13.3 47.7 14.3	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1015	NW_046a	0.466 0.466 0.466	0.466 0.466 0.466	0.466 360	0.466 0.466 0.466	57.5 0.0 0.0	0.466 0.466 0.466	52.0 8.9 10.0	13.4 48.0 14.5	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0	
1016	NW_053a	0.533 0.533 0.533	0.533 0.533 0.533	0.533									

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS57/PS57.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd		ict_Fd		hsi_Fd		rgb**Fd				LabCh**Fd				rgb**Fd				LabCh**Fd				DE**Fd		hsiM,d	rgb**Md		LabCh**Md				
1053	NW_086d	0.866	0.866	0.866	0.866	0.0	0.866	360	0.866	0.866	0.866	86.0	0.0	0.0	0.0	0.0	0.866	0.866	0.866	86.1	1.2	3.4	3.7	69.9	3.7	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1054	NW_093d	0.933	0.933	0.933	0.933	0.0	0.933	360	0.933	0.933	0.933	90.8	0.0	0.0	0.0	0.0	0.933	0.933	0.933	90.8	0.4	1.4	1.5	71.6	1.5	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1055	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.6	0.0	0.1	0.1	114.3	0.1	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1056	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	24.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	23.0	0.7	-0.9	1.1	308.5	1.7	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1057	NW_006d	0.066	0.066	0.066	0.066	0.0	0.066	360	0.066	0.066	0.066	29.0	0.0	0.0	0.0	0.0	0.066	0.066	0.066	25.6	5.5	0.6	5.5	6.7	6.5	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1058	NW_013d	0.133	0.133	0.133	0.133	0.0	0.133	360	0.133	0.133	0.133	33.8	0.0	0.0	0.0	0.0	0.133	0.133	0.133	28.2	8.3	3.4	9.0	22.4	10.6	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1059	NW_020d	0.2	0.2	0.2	0.2	0.0	0.2	360	0.2	0.2	0.2	38.6	0.0	0.0	0.0	0.0	0.2	0.2	0.2	32.0	10.0	5.8	11.6	30.4	13.3	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1060	NW_026d	0.266	0.266	0.266	0.266	0.0	0.266	360	0.266	0.266	0.266	43.3	0.0	0.0	0.0	0.0	0.266	0.266	0.266	36.7	8.8	8.7	12.4	44.7	14.0	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1061	NW_033d	0.333	0.333	0.333	0.333	0.0	0.333	360	0.333	0.333	0.333	48.1	0.0	0.0	0.0	0.0	0.333	0.333	0.333	40.7	10.4	8.9	13.7	40.4	15.5	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1062	NW_040d	0.4	0.4	0.4	0.4	0.0	0.4	360	0.4	0.4	0.4	52.8	0.0	0.0	0.0	0.0	0.4	0.4	0.4	46.8	8.7	10.2	13.4	49.7	14.7	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1063	NW_046d	0.466	0.466	0.466	0.466	0.0	0.466	360	0.466	0.466	0.466	57.5	0.0	0.0	0.0	0.0	0.466	0.466	0.466	51.8	8.8	9.9	13.3	48.4	14.5	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1064	NW_053d	0.533	0.533	0.533	0.533	0.0	0.533	360	0.533	0.533	0.533	62.3	0.0	0.0	0.0	0.0	0.533	0.533	0.533	57.5	7.3	9.2	11.8	51.6	12.7	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1065	NW_060d	0.6	0.6	0.6	0.6	0.0	0.6	360	0.6	0.6	0.6	67.1	0.0	0.0	0.0	0.0	0.6	0.6	0.6	63.6	6.0	9.2	11.0	56.7	11.5	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1066	NW_066d	0.666	0.666	0.666	0.666	0.0	0.666	360	0.666	0.666	0.666	71.8	0.0	0.0	0.0	0.0	0.666	0.666	0.666	69.3	5.2	8.3	9.8	57.5	10.1	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1067	NW_073d	0.734	0.734	0.734	0.734	0.0	0.734	360	0.734	0.734	0.734	76.6	0.0	0.0	0.0	0.0	0.734	0.734	0.734	74.5	4.8	6.5	8.1	53.5	8.3	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1068	NW_080d	0.8	0.8	0.8	0.8	0.0	0.8	360	0.8	0.8	0.8	81.3	0.0	0.0	0.0	0.0	0.8	0.8	0.8	80.5	2.7	5.2	5.9	62.0	5.9	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1069	NW_086d	0.866	0.866	0.866	0.866	0.0	0.866	360	0.866	0.866	0.866	86.0	0.0	0.0	0.0	0.0	0.866	0.866	0.866	86.1	1.2	3.4	3.6	69.4	3.6	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1070	NW_093d	0.933	0.933	0.933	0.933	0.0	0.933	360	0.933	0.933	0.933	90.8	0.0	0.0	0.0	0.0	0.933	0.933	0.933	90.7	0.4	1.4	1.5	71.7	1.5	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1071	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.7	0.0	0.0	0.1	118.4	0.1	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1072	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	24.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	23.3	1.3	-2.4	2.8	299.2	2.9	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1073	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.7	0.0	0.0	0.0	138.7	0.0	360	1.0	1.0	1.0	95.6	0.0	0.0	0.0
1074	R00Y_100_100d	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3	1.0	0.0	0.0	45.4	70.5	45.5	83.9	32.8	0.7	389	1.0	0.0	0.0	45.4	70.9	44.8	83.9
1075	G50B_100_100d	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	1.0	56.8	-25.5	-41.5	48.7	238.4	0.0	1.0	1.0	56.4	-25.2	-41.8	48.8	238.9	0.5	210	0.0	1.0	1.0	56.8	-25.5	-41.5	48.7
1076	Y00G_100_100d	1.0	1.0	0.0	1.0	1.0	0.5	90	1.0	1.0	0.0	87.8	-10.2	95.4	96.0	96.1	1.0	1.0	0.0	87.5	-10.0	95.1	95.7	96.0	0.4	89	1.0	1.0	0.0	87.8	-10.2	95.4	96.0
1077	B00R_100_100d	0.0	0.0	1.0	1.0	1.0	0.5	270	0.0	0.0	1.0	25.0	29.5	-40.4	50.0	306.2	0.0	0.0	1.0	24.7	29.8	-40.1	49.9	306.6	0.5	270	0.0	0.0	1.0	25.0	29.5	-40.4	50.0
1078	G00B_100_100d	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5	0.0	1.0	0.0	49.2	-65.4	28.0	71.2	156.7	1.8	149	0.0	1.0	0.0	50.0	-65.0	29.6	71.4
1079	B50R_100_100d	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8	1.0	0.0	1.0	45.8	79.2	-0.2	79.2	359.8	0.2	330	1.0	0.0	1.0	46.1	79.3	-0.2	79.3

delta E* = 5.8

2-0032531-F0

PS570-7N, 26/26-F

PE4600L_120830.TXT, 1080 colors, Separation cmy0*

gráfico TUB-PS57; tonos rojo - verde
colores y diferencia en color, ΔE^* , 3D=0, de=0, cmy0

entrada: $rgb/cmyk \rightarrow rgb_d$
salida: transfiera a $cmy0_d$