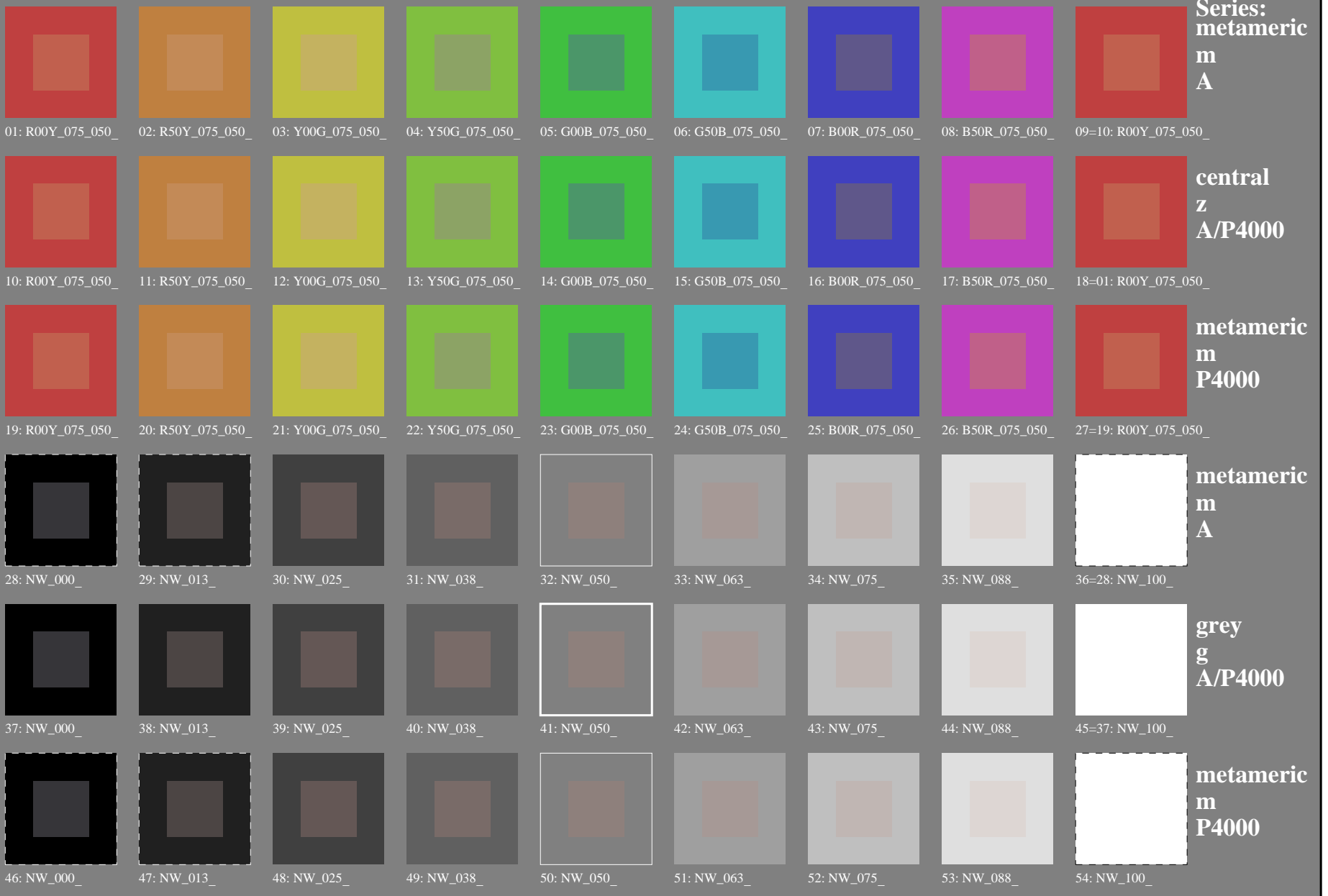


Test chart 3 for color rendering: metameric colours A and P4000



vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS
aplicación para la medida salida en la impresión offset

TUB material: code=rh4ta

2-003030-L0 PS370-7N

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS
aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)
TUB material: code=rh4ta

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

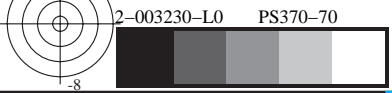
									Series: metameric m A
01: R00Y_075_050_d	02: R50Y_075_050_d	03: Y00G_075_050_d	04: Y50G_075_050_d	05: G00B_075_050_d	06: G50B_075_050_d	07: B00R_075_050_d	08: B50R_075_050_d	09=10: R00Y_075_050_d	
									central z A/P4000
10: R00Y_075_050_d	11: R50Y_075_050_d	12: Y00G_075_050_d	13: Y50G_075_050_d	14: G00B_075_050_d	15: G50B_075_050_d	16: B00R_075_050_d	17: B50R_075_050_d	18=01: R00Y_075_050_d	
									metameric m P4000
19: R00Y_075_050_d	20: R50Y_075_050_d	21: Y00G_075_050_d	22: Y50G_075_050_d	23: G00B_075_050_d	24: G50B_075_050_d	25: B00R_075_050_d	26: B50R_075_050_d	27=19: R00Y_075_050_d	
									metameric m A <i>Lab*N0=17.8, 1.3, 0.7</i> <i>Lab*W0=95.3, 0.3, -4.9</i> <i>Lab*N=23.1, -3.5, -9.1</i> <i>Lab*W=95.4, 0.3, -5.0</i>
28: NW_000_d	29: NW_013_d	30: NW_025_d	31: NW_038_d	32: NW_050_d	33: NW_063_d	34: NW_075_d	35: NW_088_d	36=28: NW_100_d	
									grey g A/P4000 <i>Lab*N0=17.8, 1.3, 0.7</i> <i>Lab*W0=95.3, 0.3, -4.9</i> <i>Lab*N1=17.7, 1.0, 0.7</i> <i>Lab*W1=95.3, 0.6, -5.0</i>
37: NW_000_d	38: NW_013_d	39: NW_025_d	40: NW_038_d	41: NW_050_d	42: NW_063_d	43: NW_075_d	44: NW_088_d	45=37: NW_100_d	
									metameric m P4000 <i>Lab*N1=17.7, 1.0, 0.7</i> <i>Lab*W1=95.3, 0.6, -5.0</i> <i>Lab*N=23.7, -5.0, -8.0</i> <i>Lab*W=95.5, 0.6, -5.1</i>
46: NW_000_d	47: NW_013_d	48: NW_025_d	49: NW_038_d	50: NW_050_d	51: NW_063_d	52: NW_075_d	53: NW_088_d	54: NW_100_d	

2-003130-L0 PS370-70



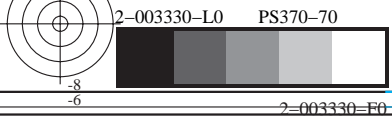
vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS TUB material: code=rh4ta
aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)



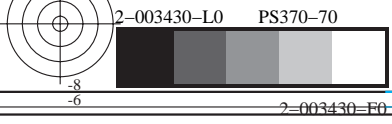
vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS TUB material: code=rh4ta
aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)

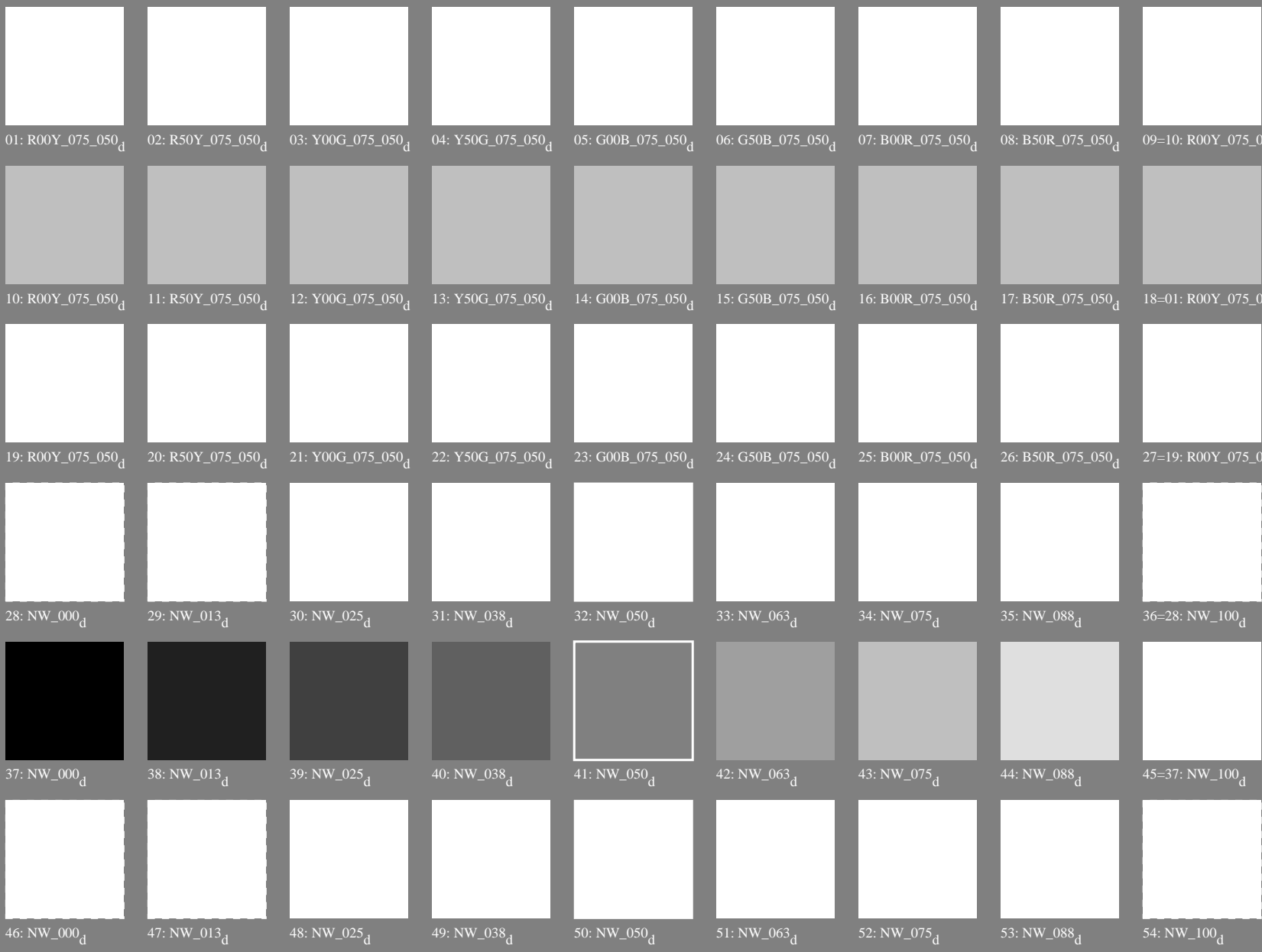


vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS TUB material: code=rh4ta
aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)



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



Series:
metameric
m
A

central
z
A/P4000

metameric
m
P4000

metameric
m
A
*Lab*N0=17.8, 1.3, 0.7*
*Lab*W0=95.3, 0.3, -4.9*
*Lab*N=23.1, -3.5, -9.1*
*Lab*W=95.4, 0.3, -5.0*

grey
g
A/P4000
*Lab*N0=17.8, 1.3, 0.7*
*Lab*W0=95.3, 0.3, -4.9*
*Lab*N1=17.7, 1.0, 0.7*
*Lab*W1=95.3, 0.6, -5.0*

metameric
m
P4000
*Lab*N1=17.7, 1.0, 0.7*
*Lab*W1=95.3, 0.6, -5.0*
*Lab*N=23.7, -5.0, -8.0*
*Lab*W=95.5, 0.6, -5.1*

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS
aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)
TUB material: code=rh4ta

n/j	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md		
0/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9	0.0 0.0	55.7 66.7 55.8	87.0 39.9	
1/657	R13Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	58.5 60.8 58.7	84.6 44.0	1.0 0.125 0.0	58.7 60.4 58.9	84.4 44.2	0.5 0.6	58.5 60.8 58.7	84.6 44.0	
2/666	R25Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	62.0 53.6 62.0	82.0 49.1	1.0 0.25 0.0	62.5 52.5 62.5	81.6 49.9	1.2 4.2	62.0 53.6 62.0	82.0 49.1	
3/675	R38Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	66.6 44.2 66.7	80.0 56.4	1.0 0.375 0.0	66.9 43.6 67.0	79.9 56.9	0.7 5.1	66.0 44.2 66.7	80.0 56.4	
4/684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2	0.0 5.9	71.8 34.6 71.7	79.6 64.2	
5/693	R63Y_100_100a	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	77.7 23.8 77.9	81.5 72.9	1.0 0.625 0.0	77.3 24.4 77.5	81.3 72.5	0.7 6.8	77.7 23.8 77.9	81.5 72.9	
6/702	R75Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	82.9 15.1 83.1	84.4 79.6	1.0 0.75 0.0	82.3 16.1 82.4	84.0 78.9	1.2 7.7	82.9 15.1 83.1	84.4 79.6	
7/711	R88Y_100_100a	1.0 0.875 0.0	1.0 1.0 0.5	83	1.0 0.883 0.0	87.1 8.3 87.4	87.8 84.5	1.0 0.875 0.0	86.8 8.7 87.1	87.5 84.2	0.5 8.3	87.1 8.3 87.4	87.8 84.5	
8/720	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3	0.0 8.9	90.5 2.6 91.3	91.4 88.3	
9/639	Y13G_100_100a	0.875 1.0 0.0	1.0 1.0 0.5	97	0.883 1.0 0.0	87.6 -1.2 84.7	84.7 90.8	0.875 1.0 0.0	87.4 -1.5 84.2	84.2 91.0	0.5 9.6	87.6 -1.2 84.7	84.7 90.8	
10/558	Y25G_100_100a	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	84.5 -4.5 78.8	79.0 93.3	0.75 1.0 0.0	84.1 -5.0 78.0	78.1 93.6	1.0 10.2	84.5 -4.5 78.8	79.0 93.3	
11/477	Y38G_100_100a	0.625 1.0 0.0	1.0 1.0 0.5	112	0.633 1.0 0.0	77.9 -10.1 70.6	71.4 98.1	0.625 1.0 0.0	77.5 -10.5 70.1	70.9 98.5	0.7 11.1	77.9 -10.1 70.6	71.4 98.1	
12/396	Y50G_100_100a	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	72.4 -16.7 58.9	61.3 105.8	0.5 1.0 0.0	72.4 -16.7 58.9	61.3 105.8	0.0 11.9	0.5 1.0 0.0	72.4 -16.7 58.9	61.3 105.8
13/315	Y63G_100_100a	0.375 1.0 0.0	1.0 1.0 0.5	128	0.366 1.0 0.0	67.1 -23.4 49.3	54.6 115.4	0.375 1.0 0.0	67.8 -22.5 50.0	54.8 114.2	1.3 12.8	0.366 1.0 0.0	67.1 -23.4 49.3	54.6 115.4
14/234	Y75G_100_100a	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	57.9 -35.3 36.5	134.0	0.25 1.0 0.0	58.5 -34.0 37.7	50.8 132.0	1.8 13.7	0.233 1.0 0.0	57.9 -35.3 36.5	50.8 134.0
15/153	Y88G_100_100a	0.125 1.0 0.0	1.0 1.0 0.5	143	0.116 1.0 0.0	53.7 -43.7 27.1	51.4 148.1	0.125 1.0 0.0	54.1 -42.4 27.6	50.7 146.9	1.4 14.3	0.116 1.0 0.0	53.7 -43.7 27.1	51.4 148.1
16/72	G00C_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	47.2 -60.1 14.6	61.9 166.3	0.0 1.0 0.0	47.2 -60.1 14.6	61.9 166.3	0.0 14.9	0.0 1.0 0.0	47.2 -60.1 14.6	61.9 166.3
17/73	G13C_100_100a	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.116	47.5 -60.5 6.9	60.9 173.4	0.0 1.0 0.125	47.6 -60.5 6.3	60.8 173.9	0.5 15.6	0.0 1.0 0.116	47.5 -60.5 6.9	60.9 173.4
18/74	G25C_100_100a	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.233	48.0 -59.7 -1.4	59.7 181.4	0.0 1.0 0.25	48.1 -59.5 -2.6	59.6 182.5	1.2 16.2	0.0 1.0 0.233	48.0 -59.7 -1.4	59.7 181.4
19/75	G38C_100_100a	0.0 1.0 0.375	1.0 1.0 0.5	172	0.0 1.0 0.366	48.4 -59.2 -12.7	60.6 192.1	0.0 1.0 0.375	48.5 -59.1 -13.4	60.6 192.8	0.7 17.1	0.0 1.0 0.366	48.4 -59.2 -12.7	60.6 192.1
20/76	G50C_100_100a	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	48.8 -58.7 -24.9	63.8 203.0	0.0 1.0 0.5	48.8 -58.7 -24.9	63.8 203.0	0.0 180	0.0 1.0 0.5	48.8 -58.7 -24.9	63.8 203.0
21/77	G63C_100_100a	0.0 1.0 0.625	1.0 1.0 0.5	188	0.0 1.0 0.633	49.3 -57.7 -35.4	67.7 211.5	0.0 1.0 0.625	49.3 -57.7 -34.8	67.4 211.0	0.6 188	0.0 1.0 0.633	49.3 -57.7 -35.4	67.7 211.5
22/78	G75C_100_100a	0.0 1.0 0.75	1.0 1.0 0.5	196	0.0 1.0 0.766	49.9 -56.3 -45.0	72.1 218.6	0.0 1.0 0.75	49.8 -56.4 -44.1	71.6 218.0	0.8 197	0.0 1.0 0.766	49.9 -56.3 -45.0	72.1 218.6
23/79	G88C_100_100a	0.0 1.0 0.875	1.0 1.0 0.5	203	0.0 1.0 0.883	50.3 -55.3 -51.2	75.4 222.8	0.0 1.0 0.875	50.3 -55.3 -50.8	75.1 222.5	0.4 203	0.0 1.0 0.883	50.3 -55.3 -51.2	75.4 222.8
24/80	C00B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	50.7 -53.9 -57.6	78.9 226.8	0.0 1.0 1.0	50.7 -53.9 -57.6	78.9 226.8	0.0 210	0.0 1.0 1.0	50.7 -53.9 -57.6	78.9 226.8
25/71	C13B_100_100a	0.0 0.875 1.0	1.0 1.0 0.5	217	0.0 0.883 1.0	48.2 -49.3 -57.0	75.4 229.1	0.0 0.875 1.0	48.0 -49.0 -56.9	75.2 229.2	0.3 216	0.0 0.883 1.0	48.2 -49.3 -57.0	75.4 229.1
26/62	C25B_100_100a	0.0 0.75 1.0	1.0 1.0 0.5	224	0.0 0.766 1.0	45.4 -43.6 -56.2	71.2 232.1	0.0 0.75 1.0	45.0 -42.8 -56.1	70.6 232.6	0.9 222	0.0 0.766 1.0	45.4 -43.6 -56.2	71.2 232.1
27/53	C38B_100_100a	0.0 0.625 1.0	1.0 1.0 0.5	232	0.0 0.633 1.0	41.7 -36.6 -55.2	66.3 236.4	0.0 0.625 1.0	41.5 -36.1 -55.1	65.9 236.7	0.4 231	0.0 0.633 1.0	41.7 -36.6 -55.2	66.3 236.4
28/44	C50B_100_100a	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	37.2 -27.1 -54.1	60.6 243.3	0.0 0.5 1.0	37.2 -27.1 -54.1	60.6 243.3	0.0 240	0.0 0.5 1.0	37.2 -27.1 -54.1	60.6 243.3
29/35	C63B_100_100a	0.0 0.375 1.0	1.0 1.0 0.5	248	0.0 0.366 1.0	32.8 -18.1 -53.0	56.0 251.1	0.0 0.375 1.0	33.1 -18.7 -53.0	56.2 250.5	0.6 248	0.0 0.366 1.0	32.8 -18.1 -53.0	56.0 251.1
30/26	C75B_100_100a	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.233 1.0	28.7 -8.5 -51.9	52.6 260.6	0.0 0.25 1.0	29.2 -9.6 -51.9	52.8 259.4	1.2 257	0.0 0.233 1.0	28.7 -8.5 -51.9	52.6 260.6
31/17	C88B_100_100a	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.116 1.0	25.0 -1.0 -51.3	51.3 268.8	0.0 0.125 1.0	25.2 -1.5 -51.3	51.3 268.3	0.4 263	0.0 0.116 1.0	25.0 -1.0 -51.3	51.3 268.8
32/8	B00M_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5	0.0 270	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
33/89	B13M_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	28.3 19.0 -42.4	46.4 294.1	0.125 0.0 1.0	28.7 19.8 -41.6	46.1 295.4	1.1 276	0.116 0.0 1.0	28.3 19.0 -42.4	46.4 294.1
34/170	B25M_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	31.6 25.8 -37.0	45.1 304.8	0.25 0.0 1.0	32.1 26.6 -36.2	45.0 306.3	1.2 282	0.233 0.0 1.0	31.6 25.8 -37.0	45.1 304.8
35/251	B38M_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	36.5 39.5 -24.6	46.6 328.0	0.375 0.0 1.0	36.8 40.3 -23.6	46.7 329.6	1.3 291	0.366 0.0 1.0	36.5 39.5 -24.6	46.6 328.0
36/332	B50M_100_100a	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 349.9	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 349.9	0.0 300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 349.9
37/413	B63M_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	46.4 54.0 -9.5	54.8 349.9	0.625 0.0 1.0	46.1 53.4 -10.1	54.4 349.2	0.8 308	0.633 0.0 1.0	46.4 54.0 -9.5	54.8 349.9
38/494	B75M_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	49.6 61.0 -0.6	61.0 359.4	0.75 0.0 1.0	49.6 61.0 -0.6	61.0 359.4	0.0 317	0.766 0.0 1.0	49.6 61.0 -0.6	61.0 359.4
39/575	B88M_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	50.1 61.5 -0.1	61.5 359.8	0.875 0.0 1.0	49.6 61.0 -0.6	61.0 359.4	0.8 323	0.883 0.0 1.0	50.1 61.5 -0.1	61.5 359.8
40/656	M00R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6	0.0 330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
41/655	M13R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	56.1 68.3 11.8	69.3 9.8	1.0 0.0 0.875	56.1 68.3 12.1	69.3 370.0	0.2 336	1.0 0.0 0.883	56.1 68.3 11.8	69.3 9.8
42/654	M25R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	56.0 68.1 16.1	70.0 13.3	1.0 0.0 0.75	56.0 68.0 16.7	70.1 373.8	0.6 342	1.0 0.0 0.766	56.0 68.1 16.1	70.0 13.3
43/653	M38R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	56.0 67.8 22.9	71.5 18.6	1.0 0.0 0.625	56.0 67.7 23.3	71.6 379.0	0.4 351	1.0 0.0 0.633	56.0 67.8 22.9	71.5 18.6
44/652	M50R_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0	1.0 0.0 0.5	55.9 67.5 30.1	73.9 384.0	0.0 360	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0
45/651	M63R_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	56.0 67.1 38.0	77.2 29.5	1.0 0.0 0.375	56.0 67.1 37.6	77.0 389.2	0.4 368	1.0 0.0 0.366	56.0 67.1 38.0	77.2 29.5
46/650	M75R_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	56.0 66.9 45.0	80.7 33.9	1.0 0.0 0.25	56.0 66.9 44.3	80.2 393.5	0.7 377	1.0 0.0 0.233	56.0 66.9 45.0	80.7 33.9
47/649	M88R_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	55.8 66.8 50.6	83.8 37.1	1.0 0.0 0.125	55.8 66.8 50.2	83.6 396.9	0.3 383	1.0 0.0 0.116	55.8 66.8 50.6	83.8 37.1
48/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9	0.0 389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
49/0	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	17.8 0.0 0.0	0.0 0.0	0.0 0.0 0.0	58.7 60.4 58.9	84.4 404.2	93.8 360	1.0 1.0 1.0	95.3 0.0 0.0	0.0 0.0 0.0
50/91	NW_013a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	27.5 0.0 0.0	0.0 0.0	0.125 0.125 0.125	17.8 0.0 0.0	0.0 9.6 360	0.0 0.0	1.0 1.0 1.0	95.3 0.0 0.0	0.0 0.0 0.0
51/182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	37.1 0.0 0.0	0.0 0.0	0.25 0.25 0.25	28.0 -0.5 -0.5	0.7 224.9	9.1 360	1.0 1.0 1.0		

n/j	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md
0/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9	0.0 0.0	389
1/666	R25Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	62.0 53.6 62.0	82.0 49.1	1.0 0.25 0.0	62.5 52.5 62.5	81.6 49.9	1.2 42	1.0 0.233 0.0
2/684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2	0.0 59	1.0 0.5 0.0
3/702	R75Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	82.9 15.1 83.1	84.4 79.6	1.0 0.75 0.0	82.3 16.1 82.4	84.0 78.9	1.2 77	1.0 0.766 0.0
4/720	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3	0.0 89	1.0 1.0 0.0
5/558	Y25G_100_100a	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	84.5 -4.5 78.8	79.0 93.3	0.75 1.0 0.0	84.1 -5.0 78.0	78.1 93.6	1.0 102	0.766 1.0 0.0
6/396	Y50G_100_100a	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	72.4 -16.7 58.9	61.3 105.8	0.5 1.0 0.0	72.4 -16.7 58.9	61.3 105.8	0.0 119	0.5 1.0 0.0
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 -35.3 36.5	50.8 134.0	0.25 1.0 0.0	58.5 -34.0 37.7	50.8 132.0	1.8 137	0.233 1.0 0.0
8/72	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	47.2 -60.1 14.6	61.9 166.3	0.0 1.0 0.0	47.2 -60.1 14.6	61.9 166.3	0.0 149	0.0 1.0 0.0
9/72	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	47.2 -60.1 14.6	61.9 166.3	0.0 1.0 0.0	47.2 -60.1 14.6	61.9 166.3	0.0 149	0.0 1.0 0.0
10/76	G25B_100_100a	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	48.8 -58.7 -24.9	63.8 203.0	0.0 1.0 0.5	48.8 -58.7 -24.9	63.8 203.0	0.0 180	0.0 1.0 0.5
11/80	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	50.7 -53.9 -57.6	78.9 226.8	0.0 1.0 1.0	50.7 -53.9 -57.6	78.9 226.8	0.0 210	0.0 1.0 1.0
12/44	G75B_100_100a	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	37.2 -27.1 -54.1	60.6 243.3	0.0 0.5 1.0	37.2 -27.1 -54.1	60.6 243.3	0.0 240	0.0 0.5 1.0
13/8	B00M_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5	0.0 270	0.0 0.0 1.0
14/332	B25R_100_100a	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 340.9	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 340.9	0.0 300	0.5 0.0 1.0
15/656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6	0.0 330	1.0 0.0 1.0
16/652	B75R_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0	0.0 360	1.0 0.0 0.5
17/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9	0.0 389	1.0 0.0 0.0
18/688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	75.5 33.3 27.9	43.5 39.9	1.0 0.5 0.5	73.5 33.0 31.6	45.7 43.7	4.2 389	1.0 0.0 0.0
19/706	R50Y_100_050a	1.0 0.75 0.5	1.0 0.5 0.75	60	1.0 0.75 0.5	83.5 17.3 35.8	39.8 64.2	1.0 0.75 0.5	83.8 15.4 35.8	39.0 66.7	1.9 59	1.0 0.5 0.0
20/724	Y00G_100_050a	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	92.9 1.3 45.6	45.7 88.3	1.0 1.0 0.5	92.9 0.7 41.5	41.5 88.9	4.2 89	1.0 1.0 0.0
21/562	Y50G_100_050a	0.75 1.0 0.5	1.0 0.5 0.75	120	0.75 1.0 0.5	83.8 -8.3 29.4	30.6 105.8	0.75 1.0 0.5	85.6 -6.8 27.9	28.7 103.8	2.8 119	0.5 1.0 0.0
22/400	G00B_100_050a	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	71.2 -30.0 7.3	30.9 166.3	0.5 1.0 0.5	74.5 -17.6 13.8	22.4 141.8	14.4 149	0.0 1.0 0.0
23/404	G50B_100_050a	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	73.0 -26.9 -28.8	39.4 226.8	0.5 1.0 1.0	77.4 -16.8 -23.2	28.7 234.0	12.3 210	0.0 1.0 1.0
24/368	B00R_100_050a	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	58.9 2.4 -25.1	25.2 275.5	0.5 0.5 1.0	59.8 12.4 -22.1	25.4 299.2	10.4 270	0.0 1.0 1.0
25/692	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	75.6 34.2 3.9	34.5 6.6	1.0 0.5 1.0	75.0 32.3 -0.4	32.3 359.2	4.8 330	1.0 0.0 1.0
26/688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	75.5 33.3 27.9	43.5 39.9	1.0 0.5 0.5	73.5 33.0 31.6	45.7 43.7	4.2 389	1.0 0.0 0.0
27/506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	56.1 33.3 27.9	43.5 39.9	0.75 0.25 0.25	57.2 35.8 33.0	48.7 42.6	5.7 389	1.0 0.0 0.0
28/524	R50Y_075_050a	0.75 0.5 0.25	0.75 0.5 0.5	60	0.75 0.5 0.25	64.2 17.3 35.8	39.8 64.2	0.75 0.5 0.25	68.6 15.5 37.8	40.9 67.6	5.1 59	1.0 0.5 0.0
29/542	Y00G_075_050a	0.75 0.75 0.25	0.75 0.5 0.5	90	0.75 0.75 0.25	73.5 1.3 45.6	45.7 88.3	0.75 0.75 0.25	78.0 0.0 43.6	43.6 89.9	5.0 89	1.0 1.0 0.0
30/380	Y50G_075_050a	0.5 0.75 0.25	0.75 0.5 0.5	120	0.5 0.75 0.25	64.4 -8.3 29.4	30.6 105.8	0.5 0.75 0.25	68.9 -8.2 31.4	32.5 104.7	4.8 119	0.5 1.0 0.0
31/218	G00B_075_050a	0.25 0.75 0.25	0.75 0.5 0.5	150	0.25 0.75 0.25	51.9 -30.0 7.3	30.9 166.3	0.25 0.75 0.25	55.6 -22.2 14.4	26.5 146.8	11.2 149	0.0 1.0 0.0
32/222	G50B_075_050a	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.75	53.6 -26.9 -28.8	39.4 226.8	0.25 0.75 0.75	58.6 -20.9 -27.3	34.4 232.5	7.9 210	0.0 1.0 1.0
33/186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	39.6 2.4 -25.1	25.2 275.5	0.25 0.25 0.75	41.9 8.7 -25.4	26.8 288.9	6.7 270	0.0 0.0 1.0
34/510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	56.2 34.2 3.9	34.5 6.6	0.75 0.25 0.75	58.4 35.6 -0.2	35.6 359.6	4.9 330	1.0 0.0 1.0
35/506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	56.1 33.3 27.9	43.5 39.9	0.75 0.25 0.25	57.2 35.8 33.0	48.7 42.6	5.7 389	1.0 0.0 0.0
36/324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.0	36.8 33.3 27.9	43.5 39.9	0.5 0.0 0.0	38.8 38.4 31.7	49.8 39.5	6.6 389	1.0 0.0 0.0
37/342	R50Y_050_050a	0.5 0.25 0.0	0.5 0.5 0.25	60	0.5 0.25 0.0	44.8 17.3 35.8	39.8 64.2	0.5 0.25 0.0	50.3 15.5 40.5	43.4 69.0	7.4 59	1.0 0.5 0.0
38/360	Y00G_050_050a	0.5 0.5 0.0	0.5 0.5 0.25	90	0.5 0.5 0.0	54.1 1.3 45.6	45.7 88.3	0.5 0.5 0.0	59.7 -0.1 48.4	48.4 90.2	6.3 89	1.0 1.0 0.0
39/198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	45.1 -8.3 29.4	30.6 105.8	0.25 0.5 0.0	49.1 -10.8 32.9	34.6 108.2	5.8 119	0.5 1.0 0.0
40/36	G00B_050_050a	0.0 0.5 0.0	0.5 0.5 0.25	150	0.0 0.5 0.0	32.5 -30.0 7.3	30.9 166.3	0.0 0.5 0.0	37.5 -28.5 13.0	31.3 155.4	7.7 149	0.0 1.0 0.0
41/40	G50B_050_050a	0.0 0.5 0.5	0.5 0.5 0.25	210	0.0 0.5 0.5	34.2 -26.9 -28.8	39.4 226.8	0.0 0.5 0.5	39.9 -26.4 -31.0	40.7 229.4	6.1 210	0.0 1.0 1.0
42/4	B00R_050_050a	0.0 0.0 0.5	0.5 0.5 0.25	270	0.0 0.0 0.5	20.2 2.4 -25.1	25.2 275.5	0.0 0.0 0.5	21.8 9.2 -27.3	28.8 288.6	7.3 270	0.0 0.0 1.0
43/328	B50R_050_050a	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	36.9 34.2 3.9	34.5 6.6	0.5 0.0 0.5	39.2 40.1 1.2	40.1 1.7	6.8 330	1.0 0.0 1.0
44/324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.0	36.8 33.3 27.9	43.5 39.9	0.5 0.0 0.0	38.8 38.4 31.7	49.8 39.5	6.6 389	1.0 0.0 0.0
45/0	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	17.8 0.0 0.0	0.0 0.0	0.0 0.0 0.0	17.8 0.0 0.0	0.0 0.0	0.0 360	1.0 1.0 1.0
46/91	NW_013a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	27.5 0.0 0.0	0.0 0.0	0.125 0.125 0.125	28.0 -0.5 -0.5	0.7 224.9	0.9 360	1.0 1.0 1.0
47/182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	37.1 0.0 0.0	0.0 0.0	0.25 0.25 0.25	42.2 -0.9 -0.8	1.2 222.2	5.1 360	1.0 1.0 1.0
48/273	NW_038a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	46.8 0.0 0.0	0.0 0.0	0.375 0.375 0.375	54.9 -0.8 -0.7	1.1 220.4	8.1 360	1.0 1.0 1.0
49/364	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	56.5 0.0 0.0	0.0 0.0	0.5 0.5 0.5	63.2 -0.8 -0.7	1.1 220.8	6.8 360	1.0 1.0 1.0
50/455	NW_063a	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	66.2 0.0 0.0	0.0 0.0	0.625 0.625 0.625	72.0 -0.6 -0.5	0.8 220.9	5.8 360	1.0 1.0 1.0
51/546	NW_075a	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	75.9 0.0 0.0	0.0 0.0	0.75 0.75 0.75	80.6 -0.5 -0.5	0.7 222.2	4.8 360	1.0 1.0 1.0
52/637	NW_088a	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	85.6 0.0 0.0	0.0 0.0	0.875 0.875 0.875	89.5 -0.2 -0.2	0.3 216.6	3.9 360	1.0 1.0 1.0
53/728	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.3 0.0 0.0	0.0 0.0	1.0 1.0 1.0	95.3 0.0 0.0	0.0 0.0	0.0 360	1.0 1.0 1.0

delta E* = 4.0

gráfico TUB-PS37; reproducción en color
 colores y diferencia en color, ΔE^* , 3D=0, de=0, cmy0

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

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS
 aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)
 TUB material: code=rh4ta

Table with 8 columns of color data (HIC, rgb, icl, hsi, LabCh, rgb, LabCh, DE, hsi, rgb, LabCh) and 80 rows of color patches (0-79). Includes a 'delta E*' = 3.8' label at the bottom right of the table area.

gráfico TUB-PS37; reproducción en color colores y diferencia en color, ΔE*, 3D=0, de=0, cmy0

entrada: rgb/cmyk -> rgb salida: transfiera a cmy0d

vea archivos semejantes: http://130.149.60.45/~farbmetrik/PS37/PS37.HTM información técnica: http://www.ps.bam.de o http://130.149.60.45/~farbmetrik

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS TUB material: code=rh4ta aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)



vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS TUB material: code=rh4ta
 aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	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	22.5 8.3 6.9	10.8 39.9	0.125 0.0 0.0	23.6 7.8 7.5	10.9 43.7 1.2	389	1.0 0.0 0.0	55.7 66.7 55.8 87.0 39.9
82	B50R_012_012a	0.125 0.0 0.125	0.125 0.125 0.062	330	0.125 0.0 0.125	22.5 8.5 0.9	8.6 6.6	0.125 0.0 0.125	22.9 8.5 -0.4	8.5 35.7 1.4	330	1.0 0.0 1.0	56.0 68.5 7.9 69.0 6.6
83	B25R_025_025a	0.125 0.0 0.25	0.25 0.25 0.125	300	0.125 0.0 0.25	23.8 11.9 -4.1	12.6 340.9	0.125 0.0 0.25	27.5 13.7 -6.2	15.1 335.5 4.5	300	0.5 0.0 1.0	42.1 47.7 -16.4 50.5 340.9
84	B15R_037_037a	0.125 0.0 0.375	0.375 0.375 0.187	289	0.118 0.0 0.375	24.1 12.9 -11.3	17.2 318.7	0.125 0.0 0.375	28.5 16.5 -12.8	20.9 322.0 5.9	288	0.316 0.0 1.0	44.6 34.5 -30.2 45.9 318.7
85	B11R_050_050a	0.125 0.0 0.5	0.5 0.5 0.25	284	0.116 0.0 0.5	24.7 12.9 -18.5	22.5 304.8	0.125 0.0 0.5	27.5 19.2 -19.4	27.3 314.5 6.9	282	0.233 0.0 1.0	31.6 25.8 -37.0 45.1 304.8
86	B09R_062_062a	0.125 0.0 0.625	0.625 0.625 0.312	281	0.114 0.0 0.625	25.6 14.4 -24.5	28.5 300.5	0.125 0.0 0.625	27.7 19.8 -26.2	32.9 307.0 5.9	279	0.183 0.0 1.0	30.3 23.1 -39.3 45.6 300.5
87	B07R_075_075a	0.125 0.0 0.75	0.75 0.75 0.375	279	0.112 0.0 0.75	26.5 16.0 -30.5	34.4 297.6	0.125 0.0 0.75	28.0 20.3 -31.9	37.8 302.5 4.8	278	0.15 0.0 1.0	29.4 21.3 -40.7 45.9 297.6
88	B06R_087_087a	0.125 0.0 0.875	0.875 0.875 0.437	278	0.110 0.0 0.875	27.5 17.8 -36.1	40.3 296.2	0.125 0.0 0.875	28.8 21.1 -37.1	42.7 299.6 3.7	277	0.133 0.0 1.0	28.9 20.3 -41.3 46.1 296.2
89	B05R_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	28.3 19.0 -42.4	46.4 294.1	0.125 0.0 1.0	28.7 19.8 -41.6	46.1 295.4 1.1	276	0.116 0.0 1.0	28.3 19.0 -42.4 46.4 294.1
90	Y00G_012_012a	0.125 0.125 0.0	0.125 0.125 0.062	90	0.125 0.125 0.0	26.9 0.3 11.4	11.4 88.3	0.125 0.125 0.0	27.9 -0.8 9.4	9.5 95.1 2.5	89	1.0 1.0 0.0	90.5 2.6 91.3 91.4 88.3
91	NW_012a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	27.5 0.0 0.0	0.0 0.0	0.125 0.125 0.125	28.0 -0.5 -0.5	0.7 224.9 0.9	360	1.0 1.0 1.0	95.3 0.0 0.0 0.0 0.0
92	BO0R_025_012a	0.125 0.125 0.25	0.25 0.125 0.187	270	0.124 0.124 0.25	28.1 0.6 -6.2	6.3 275.5	0.125 0.125 0.25	31.7 1.9 -8.7	8.9 282.5 4.6	270	0.0 0.0 1.0	22.6 4.8 -50.3 50.5 275.5
93	BO0R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.124 0.375	28.7 1.2 -12.5	12.6 275.5	0.125 0.125 0.375	33.5 4.6 -14.4	15.1 287.6 6.2	270	0.0 0.0 1.0	22.6 4.8 -50.3 50.5 275.5
94	BO0R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.124 0.5	29.3 1.8 -18.8	18.9 275.5	0.125 0.125 0.5	32.8 7.6 -20.4	21.8 290.4 6.9	270	0.0 0.0 1.0	22.6 4.8 -50.3 50.5 275.5
95	BO0R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	29.9 2.4 -25.1	25.2 275.5	0.125 0.125 0.625	33.0 9.1 -26.3	27.8 289.2 7.5	270	0.0 0.0 1.0	22.6 4.8 -50.3 50.5 275.5
96	BO0R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	30.5 3.0 -31.4	31.6 275.5	0.125 0.125 0.75	32.4 10.7 -32.5	34.2 288.2 7.9	270	0.0 0.0 1.0	22.6 4.8 -50.3 50.5 275.5
97	BO0R_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	31.1 3.6 -37.7	37.9 275.5	0.125 0.125 0.875	32.8 12.5 -37.8	39.8 288.3 9.0	270	0.0 0.0 1.0	22.6 4.8 -50.3 50.5 275.5
98	BO0R_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	31.7 4.2 -44.0	44.2 275.5	0.125 0.125 1.0	32.1 12.3 -42.7	44.4 286.1 8.1	270	0.0 0.0 1.0	22.6 4.8 -50.3 50.5 275.5
99	Y50G_025_012a	0.125 0.25 0.0	0.25 0.25 0.125	150	0.125 0.25 0.0	31.4 -4.1	14.7 15.3 105.8	0.125 0.25 0.0	36.4 -6.0	16.8 17.8 109.7 5.7	119	0.5 1.0 0.0	72.4 -16.7 58.9 61.3 105.8
100	GO0B_025_012a	0.125 0.25 0.125	0.25 0.125 0.187	120	0.124 0.25 0.124	31.1 -7.5	1.8 7.7 166.3	0.125 0.25 0.125	34.9 -7.7	3.5 8.5 155.4 4.1	149	0.0 1.0 0.0	47.2 -60.1 14.6 61.9 166.3
101	G50B_025_012a	0.125 0.25 0.25	0.25 0.125 0.187	210	0.124 0.25 0.25	31.6 -6.7	-7.2 9.8 226.8	0.125 0.25 0.25	35.6 -7.3	-9.3 11.9 231.8 4.6	210	0.0 1.0 0.0	50.7 -53.9 -57.6 78.9 226.8
102	G75B_037_025a	0.125 0.25 0.375	0.375 0.25 0.25	240	0.124 0.25 0.375	32.3 -6.7	-13.5 15.1 243.3	0.125 0.25 0.375	38.2 -5.2	-14.7 15.7 250.3 6.1	240	0.0 0.5 1.0	37.2 -27.1 -54.1 60.6 243.3
103	G84B_050_037a	0.125 0.25 0.5	0.5 0.375 0.312	251	0.124 0.243 0.5	32.5 -5.4	-19.7 20.5 254.7	0.125 0.25 0.5	38.2 -2.8	-20.5 20.7 262.0 6.2	251	0.0 0.316 1.0	31.3 -14.4 -52.7 54.6 254.7
104	G88B_062_050a	0.125 0.25 0.625	0.625 0.5 0.375	256	0.125 0.241 0.625	32.9 -4.2	-25.9 26.3 260.6	0.125 0.25 0.625	38.0 -0.8	-26.5 26.6 268.2 6.1	257	0.0 0.233 1.0	28.7 -8.5 -51.9 52.6 260.6
105	G90B_075_062a	0.125 0.25 0.75	0.75 0.625 0.437	259	0.125 0.239 0.75	33.3 -3.2	-32.3 32.5 264.1	0.125 0.25 0.75	36.7 1.8	-32.8 32.9 273.2 6.2	260	0.0 0.183 1.0	27.1 -5.2 -51.8 52.0 264.1
106	G92B_087_075a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.237 0.875	33.6 -2.3	-38.6 38.7 266.5	0.125 0.25 0.875	36.5 3.5	-38.7 38.8 275.2 6.5	262	0.0 0.15 1.0	25.0 -3.1 -51.5 51.6 266.5
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.241 1.0	34.0 -1.7	-45.0 45.0 267.7	0.125 0.25 1.0	35.9 4.1	-43.0 43.2 275.5 6.4	262	0.0 0.133 1.0	25.5 -2.0 -51.4 51.4 267.7
108	Y68G_037_037a	0.125 0.375 0.0	0.375 0.375 0.187	131	0.118 0.375 0.0	34.9 -10.7	16.7 19.8 122.5	0.125 0.375 0.0	40.0 -12.3	20.2 23.7 121.3 6.4	131	0.316 1.0 0.0	63.4 -28.5 44.6 53.0 122.5
109	GO0B_037_025a	0.125 0.375 0.125	0.375 0.25 0.25	150	0.124 0.375 0.124	34.8 -15.0	3.6 15.4 166.3	0.125 0.375 0.125	39.8 -11.3	7.7 15.2 149.2 6.7	149	0.0 1.0 0.0	47.2 -60.1 14.6 61.9 166.3
110	G25B_037_025a	0.125 0.375 0.25	0.375 0.25 0.25	180	0.124 0.375 0.25	35.2 -14.6	-6.2 15.9 203.0	0.125 0.375 0.25	40.8 -12.8	-5.6 14.0 203.5 5.8	180	0.0 1.0 0.5	48.8 -58.7 -24.9 63.8 203.0
111	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	35.7 -13.4	-14.4 19.7 226.8	0.125 0.375 0.375	41.7 -12.3	-15.7 20.0 231.7 6.2	210	0.0 1.0 1.0	50.7 -53.9 -57.6 78.9 226.8
112	G65B_050_037a	0.125 0.375 0.5	0.5 0.375 0.312	229	0.124 0.381 0.5	37.0 -14.7	-20.8 25.5 234.8	0.125 0.375 0.5	42.6 -12.0	-21.6 24.7 241.0 6.2	228	0.0 0.683 1.0	43.1 -39.2 -55.6 68.1 234.8
113	G75B_062_050a	0.125 0.375 0.625	0.625 0.5 0.375	240	0.125 0.375 0.625	37.2 -13.5	-20.7 30.3 243.3	0.125 0.375 0.625	42.9 -10.3	-27.5 29.3 249.4 6.5	240	0.0 0.5 1.0	37.2 -27.1 -54.1 60.6 243.3
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.364 0.75	37.2 -12.0	-33.2 35.3 250.0	0.125 0.375 0.75	41.0 -7.2	-33.6 34.3 257.7 6.1	247	0.0 0.383 1.0	33.3 -19.2 -53.1 56.5 250.0
115	G84B_087_075a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.362 0.875	37.6 -10.8	-39.5 41.0 254.7	0.125 0.375 0.875	41.7 -5.9	-39.0 39.5 261.3 6.3	251	0.0 0.316 1.0	31.3 -14.4 -52.7 54.6 254.7
116	G86B_100_087a	0.125 0.375 1.0	1.0 0.875 0.562	254	0.125 0.358 1.0	37.9 -9.4	-45.6 46.6 258.2	0.125 0.375 1.0	39.3 -3.4	-43.8 44.0 265.5 6.4	255	0.0 0.266 1.0	29.7 -10.8 -52.2 53.3 258.2
117	Y76G_050_050a	0.125 0.5 0.0	0.5 0.5 0.25	136	0.116 0.5 0.0	37.8 -17.6	18.2 25.4 134.0	0.125 0.5 0.0	43.0 -18.3	22.4 28.9 129.1 6.7	137	0.233 1.0 0.0	57.9 -35.3 36.5 50.8 134.0
118	GO0B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.124	38.5 -22.5	5.4 23.2 166.3	0.125 0.5 0.125	43.3 -18.7	11.1 21.8 149.2 8.3	149	0.0 1.0 0.0	47.2 -60.1 14.6 61.9 166.3
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.243	38.9 -22.3	-3.1 22.5 188.0	0.125 0.5 0.25	44.0 -18.3	-1.6 18.4 184.9 6.6	168	0.0 1.0 0.316	48.3 -59.6 -8.4 60.1 188.0
120	G34B_050_037a	0.125 0.5 0.375	0.5 0.375 0.312	191	0.124 0.5 0.381	39.4 -21.5	-14.6 26.0 214.3	0.125 0.5 0.375	45.3 -18.1	-13.7 22.7 217.2 6.9	191	0.0 1.0 0.683	49.5 -57.3 -39.1 69.4 214.3
121	G50B_050_037a	0.125 0.5 0.5	0.5 0.375 0.312	210	0.124 0.5 0.5	39.8 -20.2	-21.6 29.6 226.8	0.125 0.5 0.5	45.2 -17.7	-22.3 28.5 231.5 5.9	210	0.0 1.0 1.0	50.7 -53.9 -57.6 78.9 226.8
122	G61B_062_050a	0.125 0.5 0.625	0.625 0.5 0.375	224	0.125 0.508 0.625	41.3 -21.8	-28.1 35.6 232.1	0.125 0.5 0.625	46.8 -17.9	-28.2 33.4 237.5 6.7	222	0.0 0.766 1.0	45.4 -43.6 -56.2 71.2 232.1
123	G69B_075_062a	0.125 0.5 0.75	0.75 0.625 0.437	233	0.125 0.51 0.75	42.1 -22.2	-34.4 41.0 237.1	0.125 0.5 0.75	45.7 -17.2	-34.3 38.4 233.3 6.1	232	0.0 0.616 1.0	41.2 -35.5 -65.1 65.6 237.1
124	G75B_087_075a	0.125 0.5 0.875	0.875 0.75 0.5	240	0.125 0.5 0.875	42.0 -20.3	-46.6 45.4 243.3	0.125 0.5 0.875	46.0 -15.4	-40.2 43.0 248.9 6.2	240	0.0 0.5 1.0	37.2 -27.1 -54.1 60.6 243.3
125	G79B_100_087a	0.125 0.5 1.0	1.0 0.875 0.562	245	0.125 0.489 1.0	42.0 -18.7	-46.8 50.5 248.1	0.125 0.5 1.0	43.5 -11.8	-44.7 46.2 255.1 7.3	245	0.0 0.416 1.0	34.4 -21.4 -53.5 57.7 248.1
126	Y81G_062_062a	0.125 0.625 0.0	0.625 0.625 0.312	139	0.114 0.625 0.0	41.8 -24.3	20.4 31.7 139.9	0.125 0.625 0.0	46.4 -24.5	24.5 34.7 135.0 6.2	140	0.183 1.0 0.0	56.1 -38.8 32.6 50.7 139.9
127	GO0B_062_050a	0.125 0.625 0.125	0.625 0.5 0.375	150	0.125 0.625 0.125	42.2 -30.0	7.3 30.9 166.3	0.125 0.625 0.125	46.8 -24.5	13.8 28.2 150.5 9.7	149	0.0 1.0 0.0	47.2 -60.1 14.6 61.9 166.3
128	G11B_062_050a	0.125 0.625 0.25	0.625 0.5 0.375	164	0.125 0.625 0.241	42.6 -29.8	-0.7 29.8 181.4	0.125 0.625 0.25	47.6 -24.5	2.1 24.6 175.5 7.8	162	0.0 1.0 0.233	48.0 -59.7 -1.4 59.7 181.4
129	G25B_062_050a	0.125 0.625 0.375	0.625 0.5 0.375	180	0.125 0.625 0.375	42.9 -29.3	-12.4 31.9 203.0	0.125 0.625 0.37					

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37LONA.TXT /.PS
 aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)
 TUB material: code=rh4ta

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	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	27.3 16.6 13.9	21.7 39.9	0.25 0.0 0.0	29.6 18.2 17.4	25.2 43.7 4.4	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
163	R00Y_025_025a	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.125	27.3 16.6 7.5	18.4 24.0	0.25 0.0 0.125	29.6 18.7 7.1	20.0 20.8 2.9	360	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0
164	B50R_025_025a	0.25 0.0 0.25	0.25 0.25 0.125	330	0.25 0.0 0.25	27.3 17.1 19.1	17.2 6.6	0.25 0.0 0.25	30.0 19.5 -0.8	19.5 35.7 4.5	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
165	B34R_037_037a	0.25 0.0 0.375	0.25 0.375 0.187	311	0.256 0.0 0.375	29.0 21.4 -2.2	21.5 35.3	0.25 0.0 0.375	32.3 24.0 -5.2	24.6 34.7 5.1	311	0.683 0.0 1.0	47.8 57.1 -6.0	57.5 35.9
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 23.8 -8.2	25.2 340.9	0.25 0.0 0.5	31.7 26.2 -11.7	28.7 33.9 4.5	300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 340.9
167	B19R_062_062a	0.25 0.0 0.625	0.625 0.625 0.312	293	0.239 0.0 0.625	29.9 25.5 -14.5	29.3 330.3	0.25 0.0 0.625	31.9 27.6 -18.2	33.0 32.6 4.7	292	0.383 0.0 1.0	37.1 40.8 -23.2	47.0 330.3
168	B15R_075_075a	0.25 0.0 0.75	0.75 0.75 0.375	289	0.237 0.0 0.75	30.4 25.9 -22.7	34.4 318.7	0.25 0.0 0.75	32.1 28.0 -24.6	37.2 31.8 3.2	288	0.316 0.0 1.0	34.6 34.5 -30.2	45.9 318.7
169	B13R_087_087a	0.25 0.0 0.875	0.875 0.875 0.437	286	0.233 0.0 0.875	30.8 25.1 -30.5	39.5 309.4	0.25 0.0 0.875	32.5 28.2 -30.8	41.8 31.4 3.5	284	0.266 0.0 1.0	32.7 28.7 -34.9	45.2 309.4
170	B11R_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	31.6 25.8 -37.0	45.1 304.8	0.25 0.0 1.0	32.1 26.6 -36.2	45.0 30.6 1.2	282	0.233 0.0 1.0	31.6 25.8 -37.0	45.1 304.8
171	R50Y_025_025a	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.125 0.0	31.3 8.6 17.9	19.9 64.2	0.25 0.125 0.0	36.2 7.1 21.1	22.3 7.1 6.1	59	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2
172	R00Y_025_012a	0.25 0.125 0.125	0.25 0.125 0.187	390	0.25 0.124 0.124	32.2 8.3 6.9	10.8 39.9	0.25 0.125 0.125	35.2 9.2 10.4	13.9 48.5 4.6	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
173	B50R_025_012a	0.25 0.125 0.25	0.25 0.125 0.187	330	0.25 0.124 0.25	32.2 8.5 0.9	8.6 6.6	0.25 0.125 0.25	35.4 9.6 -1.4	9.7 35.1 4.1	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
174	B25R_037_025a	0.25 0.125 0.375	0.375 0.25 0.25	300	0.25 0.124 0.375	33.5 11.9 -4.1	12.6 340.9	0.25 0.125 0.375	38.2 12.1 -6.9	13.9 33.0 5.4	300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 340.9
175	B15R_050_037a	0.25 0.125 0.5	0.5 0.375 0.312	289	0.243 0.124 0.5	33.8 12.9 -11.3	17.2 318.7	0.25 0.125 0.5	36.6 14.3 -13.3	19.6 31.6 3.7	288	0.316 0.0 1.0	34.6 34.5 -30.2	45.9 318.7
176	B11R_062_050a	0.25 0.125 0.625	0.625 0.5 0.375	284	0.241 0.125 0.625	34.4 12.9 -18.5	22.5 304.8	0.25 0.125 0.625	37.2 16.0 -19.3	25.1 30.6 4.2	282	0.233 0.0 1.0	31.6 25.8 -37.0	45.1 304.8
177	B09R_075_062a	0.25 0.125 0.75	0.75 0.625 0.437	281	0.239 0.125 0.75	35.3 14.4 -24.5	28.5 300.5	0.25 0.125 0.75	36.4 17.7 -25.2	30.8 30.5 3.4	279	0.183 0.0 1.0	30.3 23.1 -39.3	45.6 300.5
178	B07R_087_057a	0.25 0.125 0.875	0.875 0.75 0.5	279	0.237 0.125 0.875	36.2 16.0 -30.5	34.4 297.6	0.25 0.125 0.875	36.5 19.0 -31.9	37.1 30.8 3.3	278	0.15 0.0 1.0	29.4 21.3 -40.7	45.9 297.6
179	B06R_100_087a	0.25 0.125 1.0	1.0 0.875 0.562	278	0.241 0.125 1.0	37.2 17.8 -36.1	40.3 296.2	0.25 0.125 1.0	35.4 19.0 -37.5	42.1 29.6 2.6	277	0.133 0.0 1.0	28.9 20.3 -41.3	46.1 296.2
180	Y00G_025_025a	0.25 0.25 0.0	0.25 0.25 0.125	90	0.25 0.25 0.0	36.0 0.6 22.8	22.8 88.3	0.25 0.25 0.0	40.3 -0.9 23.9	24.0 92.2 4.7	89	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3
181	Y00G_025_012a	0.25 0.25 0.125	0.25 0.125 0.187	90	0.25 0.25 0.124	36.5 0.3 11.4	11.4 88.3	0.25 0.25 0.125	40.9 -0.9 12.6	12.6 94.4 4.7	89	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3
182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	37.1 0.0 0.0	0.0 0.0	0.25 0.25 0.25	42.2 -0.9 -0.8	1.2 222.2 5.1	360	1.0 0.0 1.0	95.3 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	37.7 0.6 -6.2	6.3 275.5	0.25 0.25 0.375	43.2 1.5 -7.8	7.9 280.9 5.7	270	1.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
184	B00R_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.249 0.5	38.3 1.2 -12.5	12.6 275.5	0.25 0.25 0.5	42.5 3.6 -14.0	14.5 284.7 5.0	270	1.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
185	B00R_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	39.0 1.8 -18.8	18.9 275.5	0.25 0.25 0.625	42.9 6.1 -19.3	20.2 287.7 5.9	270	1.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	39.6 2.4 -25.1	25.2 275.5	0.25 0.25 0.75	41.9 8.7 -25.4	26.8 288.9 6.7	270	1.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
187	B00R_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	40.2 3.0 -31.4	31.6 275.5	0.25 0.25 0.875	41.0 10.5 -32.1	33.8 288.2 7.5	270	1.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
188	B00R_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	40.8 3.6 -37.7	37.9 275.5	0.25 0.25 1.0	39.8 11.1 -37.3	38.9 286.5 7.5	270	1.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
189	Y31G_037_037a	0.25 0.375 0.0	0.375 0.375 0.187	109	0.256 0.375 0.0	41.3 -3.0 27.7	27.8 96.2	0.25 0.375 0.0	46.3 -5.1 29.9	30.3 99.8 5.8	108	0.683 1.0 0.0	80.6 -8.1 73.8	74.3 96.2
190	Y50G_037_025a	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	41.1 -4.1 14.7	15.3 105.8	0.25 0.375 0.125	46.3 -5.9 17.0	18.0 109.3 5.9	119	0.5 1.0 0.0	72.4 -16.7 58.9	61.3 105.8
191	G00B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	40.8 -7.5 1.8	7.7 166.3	0.25 0.375 0.25	46.6 -6.4 3.5	7.3 151.1 6.0	149	0.0 1.0 0.0	47.2 -60.1 14.6	61.9 166.3
192	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	41.3 -6.7 -7.2	9.8 226.8	0.25 0.375 0.375	47.9 -6.2 -8.0	10.2 232.1 6.7	210	0.0 1.0 1.0	50.7 -53.9 -57.6	78.9 226.8
193	G75B_050_025a	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.375 0.5	42.0 -6.7 -13.5	15.1 243.3	0.25 0.375 0.5	47.8 -4.9 -14.3	15.1 251.0 6.1	240	0.0 0.5 1.0	37.2 -27.1 -54.1	60.6 243.3
194	G84B_062_037a	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.368 0.625	42.2 -5.4 -19.7	20.5 254.7	0.25 0.375 0.625	47.5 -2.7 -20.2	20.4 262.3 5.9	251	0.0 0.316 1.0	31.3 -14.4 -52.7	54.6 254.7
195	G88B_075_050a	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.366 0.75	42.6 -4.2 -25.9	26.3 260.6	0.25 0.375 0.75	46.4 0.1 -26.1	26.1 270.3 5.8	257	0.0 0.233 1.0	28.7 -8.5 -51.9	52.6 260.6
196	G90B_087_062a	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.364 0.875	43.0 -3.2 -32.3	32.5 264.1	0.25 0.375 0.875	45.9 1.6 -32.3	32.4 272.9 5.7	260	0.0 0.183 1.0	27.1 -5.2 -51.8	52.0 264.1
197	G92B_100_075a	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.362 1.0	43.3 -2.3 -38.6	38.7 266.5	0.25 0.375 1.0	43.1 3.7 -38.2	38.4 275.6 6.1	262	0.0 0.15 1.0	26.0 -3.1 -51.5	51.6 266.5
198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.25 0.125	120	0.25 0.5 0.0	45.1 -8.3 29.4	30.6 105.8	0.25 0.5 0.0	49.1 -10.8 32.9	34.6 108.2 5.8	119	0.5 1.0 0.0	72.4 -16.7 58.9	61.3 105.8
199	Y68G_050_037a	0.25 0.5 0.125	0.5 0.375 0.312	131	0.243 0.5 0.124	44.6 -10.7 16.7	19.8 122.5	0.25 0.5 0.125	48.6 -11.8 19.9	23.1 120.8 5.2	131	0.316 1.0 0.0	63.4 -28.5 44.6	53.0 122.5
200	G00B_050_025a	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.249	44.5 -15.0 3.6	15.4 166.3	0.25 0.5 0.25	49.7 -11.8 7.4	13.9 147.9 7.1	149	0.0 1.0 0.0	47.2 -60.1 14.6	61.9 166.3
201	G25B_050_025a	0.25 0.5 0.375	0.5 0.25 0.375	180	0.249 0.5 0.375	44.9 -14.6 -6.2	15.9 203.0	0.25 0.5 0.375	50.8 -11.6 -4.7	12.5 202.1 6.8	180	0.0 1.0 0.5	48.8 -58.7 -24.9	63.8 203.0
202	G50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	45.4 -13.4 -14.4	19.7 226.8	0.25 0.5 0.5	51.5 -11.3 -14.8	18.6 232.4 6.4	210	0.0 1.0 1.0	50.7 -53.9 -57.6	78.9 226.8
203	G65B_062_037a	0.25 0.5 0.625	0.625 0.375 0.437	229	0.25 0.506 0.625	46.7 -14.7 -20.8	25.5 234.8	0.25 0.5 0.625	52.4 -10.5 -20.5	23.0 242.9 7.1	228	0.0 0.683 1.0	43.1 -39.2 -55.6	68.1 234.8
204	G75B_075_050a	0.25 0.5 0.75	0.75 0.5 0.5	240	0.25 0.5 0.75	46.9 -13.5 -27.0	30.3 243.3	0.25 0.5 0.75	51.4 -8.5 -26.5	27.9 252.1 6.7	240	0.0 0.5 1.0	37.2 -27.1 -54.1	60.6 243.3
205	G80B_087_062a	0.25 0.5 0.875	0.875 0.625 0.562	247	0.25 0.489 0.875	46.9 -12.0 -33.2	35.3 250.0	0.25 0.5 0.875	50.3 -6.5 -33.1	33.7 258.8 6.4	247	0.0 0.383 1.0	33.3 -19.2 -53.1	56.5 250.0
206	G84B_100_075a	0.25 0.5 1.0	1.0 0.75 0.625	251	0.25 0.487 1.0	47.3 -10.8 -39.5	41.0 254.7	0.25 0.5 1.0	47.6 -3.4 -38.4	38.5 264.8 7.4	251	0.0 0.316 1.0	31.3 -14.4 -52.7	54.6 254.7
207	Y61G_062_062a	0.25 0.625 0.0	0.625 0.625 0.312	127	0.239 0.625 0.0	49.2 -13.8 31.6	34.5 113.7	0.25 0.625 0.0	52.2 -16.5 35.0	38.7 115.3 5.2	127	0.383 1.0 0.0	68.1 -22.2 50.6	55.3 113.7
208	Y76G_062_050a	0.25 0.625 0.125	0.625 0.5 0.375	136	0.241 0.625 0.125	47.5 -17.6 18.2	25.4 134.0	0.25 0.625 0.125	51.9 -17.1 23.4	29.0 126.1 6.8	137	0.233 1.0 0.0	57.9 -35.3 36.5	50.8 134.0
209	G00B_062_037a	0.25 0.625 0.25	0.625 0.375 0.437	150	0.25 0.625 0.25	48.2 -22.5 5.4	23.2 166.3	0.25 0.625 0.25	53.1 -17.0 11.0	20.2 147.0 9.2	149	0.0 1.0 0.0	47.2 -60.1 14.6	61.9 166.3
210	G15B_062_037a	0.25 0.625 0.375	0.625 0.375 0.437	169	0.25 0.625 0.368	48.6 -2								

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS
 aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)
 TUB material: code=rha4ta

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	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	32.0 25.0 20.9	32.6 39.9	0.375 0.0 0.0	33.9 29.1 25.4	38.7 41.1 6.4	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
244	R18Y_037_037a	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.118	32.1 25.1 15.2	29.4 31.2	0.375 0.0 0.125	34.4 29.3 16.7	33.8 29.7 4.9	371	1.0 0.0 0.316	56.0 67.1 40.7	78.5 31.2
245	B65R_037_037a	0.375 0.0 0.25	0.375 0.375 0.187	349	0.375 0.0 0.256	32.1 25.4 7.6	26.6 16.6	0.375 0.0 0.25	34.3 30.1 7.2	31.0 13.4 5.1	348	1.0 0.0 0.683	56.0 67.9 20.2	70.9 16.6
246	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	32.1 25.7 2.9	25.8 6.6	0.375 0.0 0.375	34.4 30.5 0.5	30.5 1.0 5.8	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
247	B38R_050_050a	0.375 0.0 0.5	0.5 0.5 0.25	316	0.383 0.0 0.5	33.7 30.5 -0.3	30.5 359.4	0.375 0.0 0.5	35.3 35.0 -3.0	35.2 355.0 5.5	317	0.766 0.0 1.0	49.6 61.0 -0.6	61.0 359.4
248	B30R_062_062a	0.375 0.0 0.625	0.625 0.625 0.312	307	0.385 0.0 0.625	35.3 33.2 -6.6	33.8 348.7	0.375 0.0 0.625	37.0 38.2 -7.7	38.9 348.5 5.3	307	1.0 0.616 0.0	45.9 53.1 -10.6	54.1 348.7
249	B25R_075_075a	0.375 0.0 0.75	0.75 0.75 0.375	300	0.375 0.0 0.75	36.0 35.8 -12.3	37.9 340.9	0.375 0.0 0.75	36.7 39.1 -14.0	41.5 340.2 3.7	300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 340.9
250	B20R_087_087a	0.375 0.0 0.875	0.875 0.875 0.437	295	0.364 0.0 0.875	35.9 37.5 -18.8	42.0 333.4	0.375 0.0 0.875	36.9 40.4 -19.5	44.9 334.1 3.1	294	0.416 0.0 1.0	38.5 42.9 -21.4	48.0 333.4
251	B18R_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	36.5 39.5 -24.6	46.6 328.0	0.375 0.0 1.0	36.8 40.3 -23.6	46.7 329.6 1.3	291	1.0 0.366 0.0	1.0 36.5 39.5	-24.6 46.6 328.0
252	R31Y_037_037a	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.118 0.0	35.4 17.9 24.4	30.2 53.6	0.375 0.125 0.0	39.6 17.8 28.5	33.6 58.0 5.9	48	1.0 0.316 0.0	64.9 47.8 65.0	80.7 53.6
253	R00Y_037_025a	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.124	36.9 16.6 13.9	21.7 39.9	0.375 0.125 0.125	39.6 18.7 18.4	26.3 44.5 5.6	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
254	R00Y_037_025a	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.25	37.0 16.8 7.5	18.4 24.0	0.375 0.125 0.25	39.9 19.0 7.3	20.4 20.9 3.6	360	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0
255	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.124 0.375	37.0 17.1 1.9	17.2 6.6	0.375 0.125 0.375	40.5 19.3 -1.2	19.4 356.3 5.2	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
256	B34R_050_037a	0.375 0.125 0.5	0.5 0.375 0.312	311	0.381 0.124 0.5	38.7 21.4 -2.2	21.5 353.9	0.375 0.125 0.5	40.9 23.9 -5.1	24.5 347.8 4.4	311	1.0 0.683 0.0	1.0 47.8 57.1	-6.0 57.5 353.9
257	B25R_062_050a	0.375 0.125 0.625	0.625 0.5 0.375	300	0.375 0.125 0.625	39.6 23.8 -8.2	25.2 340.9	0.375 0.125 0.625	41.7 26.3 -10.1	28.1 338.9 3.7	300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 340.9
258	B19R_075_062a	0.375 0.125 0.75	0.75 0.625 0.437	293	0.364 0.125 0.75	39.5 25.5 -14.5	29.3 330.3	0.375 0.125 0.75	41.3 29.0 -15.9	33.1 313.3 4.1	292	0.383 0.0 1.0	37.1 40.8 -23.2	47.0 330.3
259	B15R_087_050a	0.375 0.125 0.875	0.875 0.75 0.5	289	0.362 0.125 0.875	40.1 25.9 -20.7	34.4 318.7	0.375 0.125 0.875	41.2 31.9 -20.9	38.2 326.8 6.4	288	0.316 0.1 0.0	34.6 34.5 -30.2	45.9 318.7
260	B13R_100_087a	0.375 0.125 1.0	1.0 0.875 0.562	286	0.358 0.125 1.0	40.5 25.1 -32.5	39.5 309.4	0.375 0.125 1.0	40.4 32.6 -25.7	41.5 321.6 8.8	284	0.266 0.0 1.0	32.7 28.7 -34.9	45.2 309.4
261	R68Y_037_037a	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.256 0.0	41.0 7.7 29.9	30.9 75.5	0.375 0.25 0.0	47.3 7.0 33.7	34.5 78.2 7.4	71	1.0 0.683 0.0	79.7 20.6 79.9	82.5 75.5
262	R50Y_037_025a	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.25 0.124	41.0 8.6 17.9	19.9 64.2	0.375 0.25 0.125	47.4 8.2 21.6	23.1 69.1 7.4	59	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2
263	R00Y_037_012a	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.249	41.9 8.3 6.9	10.8 39.9	0.375 0.25 0.25	47.9 8.8 9.5	12.9 47.1 6.5	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
264	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.249 0.375	41.9 8.5 0.9	8.6 6.6	0.375 0.25 0.375	48.3 9.0 -1.4	9.1 350.7 6.8	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
265	B25R_050_025a	0.375 0.25 0.5	0.5 0.25 0.375	300	0.375 0.249 0.5	43.2 11.9 -4.1	12.6 340.9	0.375 0.25 0.5	48.3 12.5 -6.4	14.1 332.8 5.6	300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 340.9
266	B15R_062_037a	0.375 0.25 0.625	0.625 0.375 0.437	289	0.368 0.25 0.625	43.4 12.9 -11.3	17.2 318.7	0.375 0.25 0.625	48.2 15.3 -11.9	19.4 322.1 5.3	288	0.316 0.0 1.0	34.6 34.5 -30.2	45.9 318.7
267	B11R_075_050a	0.375 0.25 0.75	0.75 0.5 0.5	284	0.366 0.25 0.75	44.1 12.9 -18.5	22.5 304.8	0.375 0.25 0.75	47.4 18.7 -17.0	25.3 317.6 6.8	282	0.233 0.0 1.0	31.6 25.8 -37.0	45.1 304.8
268	B09R_087_062a	0.375 0.25 0.875	0.875 0.625 0.562	281	0.364 0.25 0.875	45.0 14.4 -24.5	28.5 305.5	0.375 0.25 0.875	47.3 22.3 -21.8	31.2 315.5 8.5	279	0.183 0.0 1.0	30.2 23.1 -39.3	45.6 305.5
269	B07R_100_075a	0.375 0.25 1.0	1.0 0.75 0.625	279	0.362 0.25 1.0	45.9 16.0 -30.5	34.4 297.6	0.375 0.25 1.0	45.6 23.5 -26.8	35.4 311.6 8.5	278	0.15 0.0 1.0	29.4 21.3 -40.7	45.9 297.6
270	Y00G_037_037a	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.375 0.0	45.0 1.0 34.2	34.2 88.3	0.375 0.375 0.0	52.1 -0.4 37.1	37.1 90.7 7.8	89	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3
271	Y00G_037_025a	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.375 0.124	45.6 0.6 22.8	88.3	0.375 0.375 0.125	52.8 -0.7 24.3	24.3 91.6 7.4	89	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3
272	Y00G_037_012a	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.375 0.249	46.2 0.3 11.4	11.4 88.3	0.375 0.375 0.25	53.4 -0.8 11.1	11.1 94.5 7.2	89	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3
273	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	46.8 0.0 0.0	0.0 0.0	0.0 0.0 0.0	54.9 -0.8 -0.7	1.1 220.4 8.1	360	1.0 1.0 1.0	95.3 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	47.4 0.6 -6.2	6.3 275.5	0.375 0.375 0.5	54.1 2.8 -6.6	7.2 293.1 7.0	270	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
275	B00R_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	48.0 1.2 -12.5	12.6 275.5	0.375 0.375 0.625	53.9 5.6 -12.1	13.3 295.0 7.3	270	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
276	B00R_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	48.6 1.8 -18.8	18.9 275.5	0.375 0.375 0.75	52.9 9.3 -17.4	19.7 298.1 8.7	270	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
277	B00R_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	49.2 2.4 -25.1	25.2 275.5	0.375 0.375 0.875	52.7 12.6 -22.2	25.6 299.6 11.1	270	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
278	B00R_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	49.8 3.0 -31.4	31.6 275.5	0.375 0.375 1.0	50.5 15.6 -26.9	31.1 300.1 13.3	270	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5
279	Y23G_050_050a	0.375 0.5 0.0	0.5 0.5 0.25	104	0.383 0.5 0.0	51.2 -2.2 39.4	39.5 93.3	0.375 0.5 0.0	57.0 -3.9 43.7	43.9 95.2 7.4	102	0.766 1.0 0.0	82.5 -4.5 78.8	79.0 93.3
280	Y31G_050_037a	0.375 0.5 0.125	0.5 0.375 0.312	109	0.381 0.5 0.124	51.0 -3.0 27.7	27.8 96.2	0.375 0.5 0.125	57.0 -4.5 30.2	30.6 98.5 6.6	108	0.683 1.0 0.0	80.6 -8.1 73.8	74.3 96.2
281	Y50G_050_025a	0.375 0.5 0.25	0.5 0.25 0.375	120	0.375 0.5 0.249	50.8 -4.1 14.7	15.3 105.8	0.375 0.5 0.25	57.4 -5.1 15.8	16.6 108.0 6.8	119	0.5 1.0 0.0	72.4 -16.7 58.9	61.3 105.8
282	G00B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	50.5 -7.5 1.8	7.7 166.3	0.375 0.5 0.375	58.3 -5.2 3.7	6.4 144.4 8.3	149	0.0 1.0 1.0	47.2 -60.1 14.6	61.9 166.3
283	G50B_050_012a	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.5	50.9 -6.7 -7.2	9.8 226.8	0.375 0.5 0.5	58.7 -5.1 -6.6	8.4 232.1 7.8	210	0.0 1.0 1.0	50.7 -53.9 -57.6	78.9 226.8
284	G75B_062_025a	0.375 0.5 0.625	0.625 0.25 0.5	240	0.375 0.5 0.625	51.7 -6.7 -13.5	15.1 243.3	0.375 0.5 0.625	59.2 -2.9 -12.5	12.8 256.8 8.5	240	0.0 0.5 1.0	37.2 -27.1 -54.1	60.6 243.3
285	G84B_075_037a	0.375 0.5 0.75	0.75 0.375 0.562	251	0.375 0.493 0.75	51.9 -5.4 -19.7	20.5 254.7	0.375 0.5 0.75	58.3 0.4 -17.8	17.8 271.5 8.8	251	0.0 0.316 1.0	31.3 -14.4 -52.7	54.6 254.7
286	G88B_087_050a	0.375 0.5 0.875	0.875 0.5 0.625	256	0.375 0.491 0.875	52.3 -4.2 -25.9	26.3 260.6	0.375 0.5 0.875	58.5 3.7 -22.7	23.0 279.3 10.5	257	0.0 0.233 1.0	28.7 -8.5 -51.9	52.6 260.6
287	G90B_100_062a	0.375 0.5 1.0	1.0 0.625 0.687	259	0.375 0.489 1.0	52.7 -3.2 -32.3	32.5 264.1	0.375 0.5 1.0	55.0 8.4 -27.4	28.7 281.1 12.9	260	0.0 0.183 1.0	27.1 -5.2 -51.8	52.0 264.1
288	Y38G_062_062a	0.375 0.625 0.0	0.625 0.625 0.312	113	0.385 0.625 0.0	54.9 -6.8 43.3	43.9 99.0	0.375 0.625 0.0	60.7 -7.8 47.0	47.6 99.4 6.9	112	0.616 1.0 0.0	77.1 -11.0 69.4	70.3 99.0
289	Y50G_062_050a	0.375 0.625 0.125	0.625 0.5 0.375	120	0.375 0.625 0.125	54.7 -8.3 29.4	30.6 105.8	0.375 0.625 0.125	60.5 -8.8 33.6	34.7 104.7 7.1	119	0.5 1.0 0.0	72.4 -16.7 58.9	61.3 105.8
290	Y68G_062_037a	0.375 0.625 0.25	0.625 0.375 0.437	131	0.368 0.625 0.25	54.3 -10.7 16.7	19.8 122.5	0.375 0.625 0.25	60.5 -9.6 19.3	21.6 116.5 6.8	131	0.316 1.0 0.0	63.4 -28.5 44.6	53.0 12

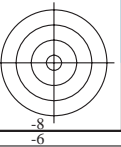
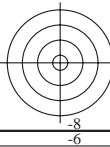
Table with columns for various color channels (n, HIC*Fa, rgb*Fa, icf*Fa, hsi*Fa, rgb*Fa, LabCh*Fa, DE*Fa, hsiMd, rgb*Ma, LabCh*Ma) and rows of numerical data for color calibration.

gráfico TUB-PS37; reproducción en color colores y diferencia en color, ΔE*, 3D=0, de=0, cmy0

entrada: rgb/cmyk -> rgb salida: transferia a cmy0d

vea archivos semejantes: http://130.149.60.45/~farbmetrik/PS37/PS37.HTM información técnica: http://www.ps.bam.de o http://130.149.60.45/~farbmetrik

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS TUB material: code=rh4ta aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)



n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	rgb*Md	LabCh*Md		
405	R00Y_062_062a	0.625 0.0 0.0	0.625 0.625 0.312	390	0.625 0.0 0.0	41.5 41.7 34.9	39.9	0.625 0.0 0.0	43.1 45.8 38.0	59.5 39.6 5.4	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
406	R31Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.114	41.6 41.8 29.6	51.2 35.3	0.625 0.0 0.125	43.0 45.9 31.3	55.6 34.3 4.6	380	1.0 0.0 0.183	55.9 66.9 47.4	82.0 35.3
407	R11Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.239	41.6 42.0 23.1	48.0 28.8	0.625 0.0 0.25	43.2 46.5 23.2	52.0 26.5 4.7	367	1.0 0.0 0.383	56.0 67.2 37.1	76.8 28.8
408	B69R_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.385	41.7 42.3 14.8	44.8 19.3	0.625 0.0 0.375	43.1 46.8 14.5	49.1 17.2 4.7	352	1.0 0.0 0.616	56.0 67.2 23.8	71.8 19.3
409	B59R_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.625 0.0 0.51	41.7 42.6 8.9	43.5 11.8	0.625 0.0 0.5	43.4 47.5 7.4	48.0 8.8 5.3	339	1.0 0.0 0.816	56.0 68.2 14.2	69.7 11.8
410	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	41.6 42.8 4.9	43.1 6.6	0.625 0.0 0.625	43.7 47.7 2.1	47.7 2.6 5.9	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
411	B42R_075_075a	0.625 0.0 0.75	0.75 0.75 0.375	321	0.637 0.0 0.75	41.7 45.7 -0.4	45.7 359.4	0.625 0.0 0.75	45.4 51.0 -1.0	51.0 358.8 6.4	322	0.85 0.0 1.0	49.6 61.0 -0.6	61.0 359.4
412	B36R_087_087a	0.625 0.0 0.875	0.875 0.875 0.437	314	0.641 0.0 0.875	45.3 52.5 -1.7	52.6 358.0	0.625 0.0 0.875	46.7 54.0 -4.6	54.2 355.1 3.5	315	0.733 0.0 1.0	49.2 60.0 -2.0	61.0 358.0
413	B31R_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	46.4 54.0 -9.5	54.8 349.9	0.625 0.0 1.0	46.1 53.4 -10.1	54.4 349.2 0.8	308	0.633 0.0 1.0	46.4 54.0 -9.5	54.8 349.9
414	R18Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	41	0.625 0.114 0.0	44.5 35.4 37.9	51.9 46.9	0.625 0.125 0.0	47.8 36.2 42.0	55.5 49.1 5.3	39	1.0 0.183 0.0	60.5 56.7 60.7	83.1 46.9
415	R00Y_062_050a	0.625 0.125 0.125	0.625 0.5 0.375	390	0.625 0.125 0.125	46.4 33.3 27.9	43.5 39.9	0.625 0.125 0.125	48.3 36.3 33.9	49.7 43.0 6.9	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
416	R26Y_062_050a	0.625 0.125 0.25	0.625 0.5 0.375	376	0.625 0.125 0.241	46.5 33.4 22.5	40.3 33.9	0.625 0.125 0.25	48.2 36.4 24.5	43.9 34.0 3.9	377	1.0 0.0 0.233	56.0 66.9 45.0	80.7 33.9
417	R00Y_062_050a	0.625 0.125 0.375	0.625 0.5 0.375	360	0.625 0.125 0.375	46.5 33.7 15.0	36.9 24.0	0.625 0.125 0.375	48.7 36.2 14.6	39.0 21.9 3.3	360	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0
418	B61R_062_050a	0.625 0.125 0.5	0.625 0.5 0.375	344	0.625 0.125 0.508	46.6 34.0 8.0	35.0 13.3	0.625 0.125 0.5	49.1 36.6 6.0	37.1 9.3 4.1	342	1.0 0.0 0.766	56.0 68.1 16.1	70.0 13.3
419	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	46.5 34.2 3.9	34.5 6.6	0.625 0.125 0.625	49.3 36.6 -0.2	36.6 359.6 5.5	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
420	B40R_075_062a	0.625 0.125 0.75	0.75 0.625 0.437	319	0.635 0.125 0.75	47.4 38.1 -0.3	38.1 359.4	0.625 0.125 0.75	50.4 40.2 -3.4	40.4 355.0 4.8	320	0.816 0.0 1.0	49.6 61.0 -0.6	61.0 359.4
421	B34R_087_075a	0.625 0.125 0.875	0.875 0.75 0.5	311	0.637 0.125 0.875	50.0 42.8 -4.5	43.1 353.9	0.625 0.125 0.875	51.2 44.4 -7.3	45.0 350.6 3.4	311	0.683 0.0 1.0	47.8 57.1 -6.0	57.5 353.9
422	B29R_100_087a	0.625 0.125 1.0	1.0 0.875 0.562	305	0.635 0.125 1.0	51.1 45.2 -10.8	46.5 346.4	0.625 0.125 1.0	50.2 45.4 -12.2	47.1 344.8 1.6	305	0.583 0.0 1.0	44.8 51.6 -12.4	53.1 346.4
423	R38Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	53	0.625 0.239 0.0	48.7 26.8 42.1	49.9 57.4	0.625 0.25 0.0	53.4 25.7 46.8	53.4 61.1 6.7	52	1.0 0.383 0.0	67.2 43.0 67.3	79.9 57.4
424	R23Y_062_050a	0.625 0.25 0.125	0.625 0.5 0.375	44	0.625 0.241 0.125	49.6 26.8 31.0	41.0 49.1	0.625 0.25 0.125	53.3 26.0 36.6	44.9 54.6 6.8	42	1.0 0.233 0.0	62.0 53.6 62.0	82.0 49.1
425	R00Y_062_037a	0.625 0.25 0.25	0.625 0.375 0.437	390	0.625 0.25 0.25	51.4 25.0 20.9	32.6 39.9	0.625 0.25 0.25	53.8 25.6 25.8	36.4 45.1 5.5	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
426	R18Y_062_037a	0.625 0.25 0.375	0.625 0.375 0.437	371	0.625 0.25 0.368	51.5 25.1 15.2	29.4 31.2	0.625 0.25 0.375	54.4 25.5 16.1	30.1 32.2 3.0	371	1.0 0.0 0.316	56.0 67.1 40.7	78.5 31.2
427	B65R_062_037a	0.625 0.25 0.5	0.625 0.375 0.437	349	0.625 0.25 0.506	51.5 25.4 7.6	26.6 16.6	0.625 0.25 0.5	54.7 25.7 5.9	26.4 13.0 3.6	348	1.0 0.0 0.683	56.0 67.9 20.2	70.9 16.6
428	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	51.5 25.7 2.9	25.8 6.6	0.625 0.25 0.625	55.4 25.8 -1.2	25.8 357.1 5.8	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
429	B38R_075_050a	0.625 0.25 0.75	0.75 0.5 0.5	316	0.633 0.25 0.75	53.1 30.5 -0.3	30.5 359.4	0.625 0.25 0.75	55.7 29.9 -4.9	30.3 350.6 5.3	317	0.766 0.0 1.0	49.6 61.0 -0.6	61.0 359.4
430	B30R_087_062a	0.625 0.25 0.875	0.875 0.625 0.562	307	0.635 0.25 0.875	54.7 33.2 -6.6	33.8 348.7	0.625 0.25 0.875	56.2 33.6 -9.3	34.9 344.5 3.0	307	0.616 0.0 1.0	45.9 53.1 -10.6	54.1 348.7
431	B25R_100_075a	0.625 0.25 1.0	1.0 0.75 0.625	300	0.625 0.25 1.0	55.4 35.8 -12.3	37.9 340.9	0.625 0.25 1.0	54.7 24.9 -14.5	37.8 337.4 2.4	300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 340.9
432	R61Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	67	0.625 0.385 0.0	54.8 15.7 48.2	50.7 71.9	0.625 0.375 0.0	59.5 14.6 51.9	54.0 74.2 6.0	67	1.0 0.616 0.0	77.0 25.1 77.2	81.2 71.9
433	R50Y_062_050a	0.625 0.375 0.125	0.625 0.5 0.375	60	0.625 0.375 0.125	54.5 17.3 35.8	39.8 64.2	0.625 0.375 0.125	59.3 15.4 40.3	43.2 69.1 6.8	59	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2
434	R31Y_062_037a	0.625 0.375 0.25	0.625 0.375 0.437	49	0.625 0.368 0.25	54.8 17.9 24.4	30.2 53.6	0.625 0.375 0.25	59.5 15.8 28.4	32.6 60.8 6.5 4.8	49	1.0 0.316 0.0	64.9 47.8 65.0	80.7 53.6
435	R00Y_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	56.3 16.6 19.9	24.9 21.7	0.625 0.375 0.375	60.2 15.4 17.5	23.3 48.5 5.3	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
436	R00Y_062_025a	0.625 0.375 0.5	0.625 0.25 0.5	360	0.625 0.375 0.5	56.3 16.8 7.5	18.4 24.0	0.625 0.375 0.5	60.8 15.3 7.1	16.9 24.8 4.7	360	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0
437	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	56.4 17.1 1.9	17.2 6.6	0.625 0.375 0.625	61.5 15.4 -1.6	15.5 354.0 6.5	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
438	B34R_075_037a	0.625 0.375 0.75	0.75 0.375 0.562	311	0.631 0.375 0.75	58.1 21.4 -2.2	21.5 353.9	0.625 0.375 0.75	61.2 19.6 -6.0	20.5 342.9 5.2	311	0.683 0.0 1.0	47.8 57.1 -6.0	57.5 353.9
439	B25R_087_050a	0.625 0.375 0.875	0.875 0.5 0.625	300	0.625 0.375 0.875	59.0 23.8 -2.2	25.2 340.9	0.625 0.375 0.875	61.5 22.8 -10.7	25.2 334.6 3.7	300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 340.9
440	B19R_100_062a	0.625 0.375 1.0	1.0 0.625 0.687	293	0.614 0.375 1.0	58.9 25.5 -14.5	29.3 330.3	0.625 0.375 1.0	58.6 25.2 -16.6	30.1 326.6 2.1	292	0.383 0.0 1.0	37.1 40.8 -23.2	47.0 330.3
441	R81Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	79	0.625 0.51 0.0	59.6 7.6 53.1	53.6 81.7	0.625 0.5 0.0	64.4 6.4 56.2	56.6 83.4 5.8 8.0	80	1.0 0.816 0.0	84.7 12.2 85.0	85.9 81.7
442	R76Y_062_050a	0.625 0.5 0.125	0.625 0.5 0.375	76	0.625 0.508 0.125	60.0 7.5 41.5	42.2 79.6	0.625 0.5 0.125	64.8 6.5 44.2	44.7 81.5 5.5 7.7	77	1.0 0.766 0.0	82.9 15.1 83.1	84.4 79.6
443	R68Y_062_037a	0.625 0.5 0.25	0.625 0.375 0.437	71	0.625 0.506 0.25	60.4 7.7 29.9	30.9 75.5	0.625 0.5 0.25	65.3 6.5 31.2	31.9 78.1 5.2 7.1	71	1.0 0.683 0.0	79.7 20.6 79.9	82.5 75.5
444	R50Y_062_025a	0.625 0.5 0.375	0.625 0.25 0.5	60	0.625 0.5 0.375	60.3 8.6 17.9	19.9 64.2	0.625 0.5 0.375	65.9 6.7 19.2	20.3 70.7 6.0 5.9	60	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2
445	R00Y_062_012a	0.625 0.5 0.5	0.625 0.125 0.562	390	0.625 0.5 0.5	61.3 8.3 6.9	10.8 39.9	0.625 0.5 0.5	66.6 6.8 8.0	10.5 49.6 5.6 38.9	10	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
446	B50R_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	300	0.625 0.5 0.625	61.3 8.5 0.9	8.6 6.6	0.625 0.5 0.625	66.8 6.8 -1.4	7.0 348.0 6.3	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
447	B25R_075_025a	0.625 0.5 0.75	0.75 0.25 0.625	330	0.625 0.5 0.75	62.6 11.9 -4.1	12.6 340.9	0.625 0.5 0.75	67.0 10.1 -6.3	12.0 327.8 5.2	300	0.5 0.0 1.0	42.1 47.7 -16.4	50.5 340.9
448	B15R_087_037a	0.625 0.5 0.875	0.875 0.375 0.687	289	0.618 0.5 0.875	62.8 12.9 -11.3	17.2 318.7	0.625 0.5 0.875	66.6 13.1 -11.6	17.6 318.4 3.8	288	0.316 0.0 1.0	34.6 34.5 -30.2	45.9 318.7
449	B11R_100_050a	0.625 0.5 1.0	1.0 0.5 0.75	284	0.616 0.5 1.0	63.4 12.9 -18.5	22.5 304.8	0.625 0.5 1.0	63.5 17.3 -17.0	24.3 315.5 4.7	282	0.233 0.0 1.0	31.6 25.8 -37.0	45.1 304.8
450	Y00G_062_062a	0.625 0.625 0.0	0.625 0.625 0.312	90	0.625 0.625 0.0	63.2 1.6 57.1	57.1 88.3	0.625 0.625 0.0	68.1 0.4 60.4	60.4 89.5 6.0 8.9	10	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3
451	Y00G_062_050a	0.625 0.625 0.125	0.625 0.5 0.375	90	0.625 0.625 0.125	63.8 1.3 45.6	45.7 88.3	0.625 0.625 0.125	68.9 0.1 47.8	47.8 89.8 5.6 8.9	10	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3
452	Y00G_062_037a	0.625 0.625 0.25	0.625 0.375 0.437	90	0.625 0.625 0.25	64.4 1.0 34.2	34.2 88.3	0.625 0.625 0.25	69.7 -0.2 34.0	34.0 90.3 5.4 8.9	10	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3
453	Y00G_062_025a	0.625 0.625 0.375	0.625 0.25 0.5	90	0.625 0.625 0.375	65.0 0.6 22.8	22.8 88.3	0.625 0.625 0.375	70.2 -0.4 21.0	21.0 91.3 5.6 8.9	10	1.0 1.0 0.0	90.5 2.6 91.3	91.4 88.3

Table with columns for various color channels (n, HIC*Fa, rgb*Fa, icf*Fa, hsi*Fa, rgb**Fa, LabCh*Fa, DE**Fa, hsiMd, rgb**Md, LabCh**Md) and rows of numerical data representing color calibration parameters.

gráfico TUB-PS37; reproducción en color colores y diferencia en color, ΔE*, 3D=0, de=0, cmy0

entrada: rgb/cmyk -> rgbd salida: transfiera a cmy0d

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Ma	rgb*Ma	LabCh*Ma		
567	R00Y_087_087a	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.0	51.0 58.4 48.8	76.1 39.9	0.875 0.0 0.0	52.2 61.7 50.0	79.4 39.0 3.7	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
568	R36Y_087_087a	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.116	51.1 58.5 43.6	72.9 36.7	0.875 0.0 0.125	52.3 61.9 44.5	76.2 35.7 3.7	382	1.0 0.0 0.133	55.8 66.8 49.8	83.4 36.7
569	R23Y_087_087a	0.875 0.0 0.25	0.875 0.875 0.437	374	0.875 0.0 0.233	51.2 58.6 37.9	69.8 32.9	0.875 0.0 0.25	52.5 61.9 38.4	72.9 31.8 3.6	375	1.0 0.0 0.266	56.0 67.0 43.4	79.8 32.9
570	R08Y_087_087a	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.364	51.2 58.9 30.7	66.5 27.5	0.875 0.0 0.375	52.5 62.3 30.5	69.4 26.1 3.5	365	1.0 0.0 0.416	55.9 67.4 35.0	76.0 27.5
571	B70R_087_087a	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.51	51.2 59.3 22.3	63.3 20.6	0.875 0.0 0.5	52.5 62.7 22.5	66.7 19.7 3.7	354	1.0 0.0 0.583	56.0 67.7 25.5	72.4 20.6
572	B63R_087_087a	0.875 0.0 0.625	0.875 0.875 0.437	346	0.875 0.0 0.641	51.2 59.5 15.4	61.5 14.5	0.875 0.0 0.625	52.6 62.9 15.7	64.8 14.0 3.6	344	1.0 0.0 0.733	56.0 68.0 17.6	70.3 14.5
573	B56R_087_087a	0.875 0.0 0.75	0.875 0.875 0.437	338	0.875 0.0 0.758	51.3 59.7 10.8	60.7 10.3	0.875 0.0 0.75	52.5 63.2 10.6	64.1 9.5 3.6	337	1.0 0.0 0.866	56.1 68.3 12.4	69.4 10.3
574	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	51.2 59.9 6.9	60.3 6.6	0.875 0.0 0.875	52.6 63.6 6.2	63.9 5.6 4.0	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
575	B44R_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	50.1 61.5 -0.1	61.5 35.8	0.875 0.0 1.0	53.1 64.8 3.5	64.9 3.1 5.7	323	0.883 0.0 1.0	50.1 61.5 -0.1	61.5 35.8
576	R13Y_087_087a	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.116 0.0	53.8 52.4 51.8	73.7 44.6	0.875 0.125 0.0	56.2 53.8 52.6	75.3 44.3 2.8	37	1.0 0.133 0.0	58.9 59.9 59.2	84.2 44.6
577	R00Y_087_075a	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.125	55.9 50.0 41.8	65.2 39.9	0.875 0.125 0.125	56.2 53.6 46.2	70.8 40.7 5.6	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
578	R35Y_087_075a	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.237	56.0 50.1 36.7	62.2 36.2	0.875 0.125 0.25	56.4 53.6 39.4	66.6 36.3 4.4	382	1.0 0.0 0.15	55.8 66.9 49.0	82.9 36.2
579	R18Y_087_075a	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.362	56.1 50.3 30.5	58.8 31.2	0.875 0.125 0.375	56.7 53.6 30.5	61.7 29.6 3.3	371	1.0 0.0 0.316	56.0 67.1 40.7	78.5 31.2
580	R00Y_087_075a	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.5	56.0 50.6 22.5	55.4 24.0	0.875 0.125 0.5	56.7 54.0 21.7	58.2 21.8 3.4	360	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0
581	B65R_087_075a	0.875 0.125 0.625	0.875 0.75 0.5	349	0.875 0.125 0.637	56.1 50.9 15.2	53.2 16.6	0.875 0.125 0.625	57.1 53.9 13.9	55.7 14.5 3.3	348	1.0 0.0 0.683	56.0 67.9 20.2	70.9 16.6
582	B57R_087_075a	0.875 0.125 0.75	0.875 0.75 0.5	339	0.875 0.125 0.762	56.2 51.2 9.7	52.1 10.8	0.875 0.125 0.75	57.0 54.2 7.7	54.8 8.1 3.7	337	1.0 0.0 0.85	56.1 68.2 13.0	69.5 10.8
583	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	56.1 51.4 5.9	51.7 6.6	0.875 0.125 0.875	57.3 54.2 3.0	54.3 3.2 4.2	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
584	B43R_100_087a	0.875 0.125 1.0	1.0 0.875 0.562	322	0.883 0.125 1.0	55.3 53.3 -0.5	53.3 35.9	0.875 0.125 1.0	57.1 56.4 0.1	56.4 0.1 3.5	322	0.866 0.0 1.0	49.6 61.0 -0.6	61.0 35.9
585	R26Y_087_087a	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.233 0.0	57.4 44.9 55.2	71.2 50.8	0.875 0.25 0.0	60.2 44.8 57.7	73.1 52.1 3.7	44	1.0 0.266 0.0	63.1 51.3 63.1	81.4 50.8
586	R15Y_087_075a	0.875 0.25 0.125	0.875 0.75 0.5	39	0.875 0.237 0.125	58.7 44.1 44.8	62.9 45.4	0.875 0.25 0.125	60.5 44.8 49.0	66.4 47.5 4.6	37	1.0 0.15 0.0	59.4 58.8 59.7	83.8 45.4
587	R00Y_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	60.9 41.7 34.9	54.3 39.9	0.875 0.25 0.25	61.3 43.5 40.5	59.5 42.9 5.9	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
588	R31Y_087_062a	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.364	61.0 41.8 29.6	51.2 35.3	0.875 0.25 0.375	61.6 43.6 31.6	53.9 35.9 2.7	380	1.0 0.0 0.183	55.9 66.9 47.4	82.0 35.3
589	R11Y_087_062a	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.489	61.0 42.0 23.1	48.0 28.8	0.875 0.25 0.5	61.8 43.6 22.1	48.9 26.9 2.0	367	1.0 0.0 0.383	56.0 67.2 37.1	76.8 28.8
590	B69R_087_062a	0.875 0.25 0.625	0.875 0.625 0.562	353	0.875 0.25 0.635	61.0 42.3 14.8	44.8 19.1	0.875 0.25 0.625	62.3 43.5 13.1	45.5 16.7 2.4	352	1.0 0.0 0.616	56.0 67.7 23.8	71.8 19.3
591	B59R_087_062a	0.875 0.25 0.75	0.875 0.625 0.562	341	0.875 0.25 0.76	61.1 42.6 8.9	43.5 11.8	0.875 0.25 0.75	62.3 43.7 6.3	44.2 8.2 3.1	339	1.0 0.0 0.816	56.0 68.2 14.2	69.7 11.8
592	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	61.0 42.8 4.9	42.1 6.6	0.875 0.25 0.875	62.7 43.6 0.7	43.7 0.9 4.6	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
593	B42R_100_075a	0.875 0.25 1.0	1.0 0.75 0.625	321	0.887 0.25 1.0	61.0 45.7 -0.4	45.7 35.9	0.875 0.25 1.0	62.5 46.4 -1.8	46.4 35.7 2.1	322	0.85 0.0 1.0	49.6 61.0 -0.6	61.0 35.9
594	R41Y_087_087a	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.364 0.0	62.2 35.5 60.1	69.8 59.3	0.875 0.375 0.0	65.5 34.6 62.5	71.5 60.9 4.1	54	1.0 0.416 0.0	68.5 40.6 68.7	79.8 59.3
595	R31Y_087_075a	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.362 0.125	62.8 35.8 48.8	60.5 53.6	0.875 0.375 0.125	65.7 34.6 52.5	62.9 56.6 4.9	48	1.0 0.316 0.0	64.9 47.8 65.0	80.7 53.6
596	R18Y_087_062a	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.364 0.25	63.8 35.4 37.9	51.9 46.9	0.875 0.375 0.25	66.2 34.1 42.9	54.8 51.4 5.6	39	1.0 0.183 0.0	60.5 56.7 60.7	83.1 46.9
597	R00Y_087_050a	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	65.8 33.3 27.9	43.5 39.9	0.875 0.375 0.375	66.9 33.2 32.3	46.3 44.1 4.5	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
598	R26Y_087_050a	0.875 0.375 0.5	0.875 0.5 0.625	376	0.875 0.375 0.491	65.9 33.4 22.5	40.3 33.9	0.875 0.375 0.5	67.2 33.4 22.9	40.5 34.5 1.3	377	1.0 0.0 0.233	56.0 66.9 45.0	80.7 33.9
599	R00Y_087_050a	0.875 0.375 0.625	0.875 0.5 0.625	360	0.875 0.375 0.625	65.9 33.7 15.0	36.9 24.0	0.875 0.375 0.625	67.9 33.0 13.6	35.7 22.4 2.6	360	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0
600	B61R_087_050a	0.875 0.375 0.75	0.875 0.5 0.625	344	0.875 0.375 0.758	66.0 34.0 8.0	35.0 13.3	0.875 0.375 0.75	68.3 32.9 5.7	33.4 9.8 3.4	342	1.0 0.0 0.766	56.0 68.1 16.1	70.0 13.3
601	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	65.9 34.2 3.9	34.5 6.6	0.875 0.375 0.875	68.7 32.6 -0.4	32.7 35.9 5.4	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
602	B40R_100_062a	0.875 0.375 1.0	1.0 0.625 0.687	319	0.885 0.375 1.0	66.8 38.1 -0.3	38.1 35.9	0.875 0.375 1.0	67.2 37.0 -3.6	37.1 35.4 3.4	320	0.816 0.0 1.0	49.6 61.0 -0.6	61.0 35.9
603	R58Y_087_087a	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.51 0.0	68.3 24.4 66.3	70.6 69.7	0.875 0.5 0.0	71.6 23.5 68.3	72.3 71.0 4.0	65	1.0 0.583 0.0	75.5 27.9 75.8	80.7 69.7
604	R50Y_087_075a	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.5 0.125	68.0 25.9 53.8	59.7 64.2	0.875 0.5 0.125	71.6 24.0 56.9	61.7 67.1 5.1	59	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2
605	R38Y_087_062a	0.875 0.5 0.25	0.875 0.625 0.562	53	0.875 0.489 0.25	68.1 26.8 42.1	49.9 57.4	0.875 0.5 0.25	71.8 24.4 60.0	62.1 62.0 5.9	52	1.0 0.383 0.0	67.2 43.0 67.3	79.9 57.4
606	R23Y_087_050a	0.875 0.5 0.375	0.875 0.5 0.625	44	0.875 0.491 0.375	68.9 26.8 31.0	41.0 49.1	0.875 0.5 0.375	72.5 23.7 34.5	41.8 55.4 5.8	42	1.0 0.233 0.0	62.0 53.6 62.0	82.0 49.1
607	R00Y_087_037a	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.5	70.8 25.0 20.9	32.6 39.9	0.875 0.5 0.5	73.2 23.0 23.9	33.2 46.1 4.3	389	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9
608	R18Y_087_037a	0.875 0.5 0.625	0.875 0.375 0.687	371	0.875 0.5 0.618	70.8 25.1 15.2	29.4 31.2	0.875 0.5 0.625	73.7 22.8 14.8	27.2 32.9 3.7	371	1.0 0.0 0.316	56.0 67.1 40.7	78.5 31.2
609	B63R_087_037a	0.875 0.5 0.75	0.875 0.375 0.687	349	0.875 0.5 0.756	70.9 25.4 7.6	29.6 16.6	0.875 0.5 0.75	74.4 22.4 6.0	23.2 14.9 4.9	348	1.0 0.0 0.683	56.0 67.9 20.2	70.9 16.6
610	B50R_087_037a	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	70.8 25.7 2.9	25.8 6.6	0.875 0.5 0.875	74.8 22.4 -1.0	22.5 35.7 6.5	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6
611	B38R_100_050a	0.875 0.5 1.0	1.0 0.5 0.75	316	0.883 0.5 1.0	72.5 30.5 -0.3	30.5 35.9	0.875 0.5 1.0	72.1 28.3 -4.6	28.7 35.0 4.8	317	0.766 0.0 1.0	49.6 61.0 -0.6	61.0 35.9
612	R73Y_087_087a	0.875 0.625 0.0	0.875 0.875 0.437	74	0.875 0.641 0.0	73.7 15.1 71.6	73.2 78.0	0.875 0.625 0.0	76.5 15.2 72.9	74.4 78.1 3.0	75	1.0 0.733 0.0	81.7 17.2 81.8	83.6 78.0
613	R68Y_087_075a	0.875 0.625 0.125	0.875 0.75 0.5	71	0.875 0.637 0.125	73.9 15.4 59.9	61.9 75.5	0.875 0.625 0.125	76.8 15.1 60.7	62.5 75.9 3.0	71	1.0 0.683 0.0	79.7 20.6 79.9	82.5 75.5
614	R61Y_087_062a	0.875 0.625 0.25	0.875 0.625 0.562	67	0.875 0.635 0.25	74.1 15.7 48.2	50.7 71.9	0.875 0.625 0.25	77.2 15.0 49.3	51.5 73.0 3.3	67	1.0 0.616 0.0	77.0 25.1 77.2	81.2 71.9
615	R50Y_087_050a	0.875 0.625 0.375	0.875 0.5 0.625	60	0.875 0.625 0.375	73.8 17.3 35.8	39.8 64.2	0.875 0.625 0.375	77.5 15.2 36.7	39.8 67.5 4.2	59	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2
616	R31Y_087_037a	0.875 0.62												

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md
648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9	1.0 0.0 0.0	55.7 66.7 55.8	87.0 39.9	0.0 0.0	55.7 66.7 55.8
649	R38Y_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	55.8 66.8 50.6	83.8 37.1	1.0 0.0 0.125	55.8 66.8 50.6	83.8 37.1	0.0 0.116	55.8 66.8 50.6
650	R26Y_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	56.0 66.9 45.0	80.7 33.9	1.0 0.0 0.25	56.0 66.9 45.0	80.7 33.9	0.0 0.233	56.0 66.9 45.0
651	R13Y_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	56.0 67.1 38.0	77.2 29.5	1.0 0.0 0.375	56.0 67.1 38.0	77.2 29.5	0.0 0.366	56.0 67.1 38.0
652	R00Y_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0	1.0 0.0 0.5	55.9 67.5 30.1	73.9 24.0	0.0 0.5	55.9 67.5 30.1
653	B68R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	56.0 67.8 22.9	71.5 18.6	1.0 0.0 0.625	56.0 67.8 22.9	71.5 18.6	0.0 0.633	56.0 67.8 22.9
654	B61R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	56.0 68.1 16.1	70.0 13.3	1.0 0.0 0.75	56.0 68.0 16.7	70.1 13.8 0.6	0.0 0.766	56.0 68.1 16.1
655	B55R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	56.1 68.3 11.8	69.3 9.8	1.0 0.0 0.875	56.1 68.3 12.1	69.3 10.0 0.2	0.0 0.883	56.1 68.3 11.8
656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6 0.0	0.0 1.0	56.0 68.5 7.9
657	R11Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	58.5 60.8 58.7	84.6 44.0	1.0 0.125 0.0	58.7 60.4 58.9	84.4 44.2 0.5	0.0 0.116	58.5 60.8 58.7
658	R00Y_100_087a	1.0 0.125 0.125	1.0 0.875 0.562	390	1.0 0.125 0.125	60.7 58.4 48.8	76.1 39.9	1.0 0.125 0.125	59.3 59.8 52.3	79.5 41.2 4.0	0.0 0.125	55.7 66.7 55.8
659	R36Y_100_087a	1.0 0.125 0.25	1.0 0.875 0.562	382	1.0 0.125 0.241	60.7 58.5 43.6	72.9 36.7	1.0 0.125 0.25	59.5 59.5 45.2	74.8 37.2 2.3	0.0 0.133	55.8 66.8 49.8
660	R23Y_100_087a	1.0 0.125 0.375	1.0 0.875 0.562	374	1.0 0.125 0.358	60.9 58.6 37.9	69.9 32.9	1.0 0.125 0.375	59.6 59.6 38.5	71.0 32.8 1.7	0.0 0.266	56.0 67.0 43.4
661	R08Y_100_087a	1.0 0.125 0.5	1.0 0.875 0.562	365	1.0 0.125 0.489	60.8 58.9 30.7	66.5 27.5	1.0 0.125 0.5	59.6 60.0 30.0	67.1 26.5 1.7	0.0 0.416	55.9 67.4 35.0
662	B70R_100_087a	1.0 0.125 0.625	1.0 0.875 0.562	355	1.0 0.125 0.635	60.9 59.3 22.3	63.3 20.6	1.0 0.125 0.625	60.0 59.8 21.4	63.5 19.7 1.3	0.0 0.583	56.0 67.7 25.5
663	B63R_100_087a	1.0 0.125 0.75	1.0 0.875 0.562	346	1.0 0.125 0.766	60.9 59.5 15.4	61.5 14.5	1.0 0.125 0.75	59.9 60.1 14.5	61.8 13.5 1.4	0.0 0.733	56.0 68.0 17.6
664	B56R_100_087a	1.0 0.125 0.875	1.0 0.875 0.562	338	1.0 0.125 0.883	61.0 59.7 10.8	60.7 10.3	1.0 0.125 0.875	60.0 60.2 9.4	60.9 8.8 1.7	0.0 0.866	56.1 68.3 12.4
665	B50R_100_087a	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	60.9 59.9 6.9	60.3 6.6	1.0 0.125 1.0	60.3 60.0 4.4	60.2 4.2 2.6	0.0 1.0	56.0 68.5 7.9
666	R23Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	62.0 53.6 62.0	82.0 49.1	1.0 0.25 0.0	62.5 52.5 62.5	81.6 49.9 1.2	0.0 0.233	62.0 53.6 62.0
667	R13Y_100_087a	1.0 0.25 0.125	1.0 0.875 0.562	38	1.0 0.241 0.125	63.5 52.4 51.8	73.7 44.6	1.0 0.25 0.125	63.2 51.5 54.5	75.5 46.7 3.2	0.0 0.133	60.9 59.9 59.2
668	R00Y_100_075a	1.0 0.25 0.25	1.0 0.75 0.625	390	1.0 0.25 0.25	65.6 50.0 41.8	65.2 39.9	1.0 0.25 0.25	63.8 50.5 46.8	68.9 42.8 5.2	0.0 0.0	55.7 66.7 55.8
669	R35Y_100_075a	1.0 0.25 0.375	1.0 0.75 0.625	381	1.0 0.25 0.362	65.7 50.1 36.7	62.2 36.2	1.0 0.25 0.375	64.1 50.4 39.4	63.9 38.0 3.0	0.0 0.15	55.8 66.9 49.0
670	R18Y_100_075a	1.0 0.25 0.5	1.0 0.75 0.625	371	1.0 0.25 0.487	65.8 50.3 30.5	58.8 31.2	1.0 0.25 0.5	64.2 50.4 30.8	59.1 31.3 1.6	0.0 0.316	56.0 67.1 40.7
671	R00Y_100_075a	1.0 0.25 0.625	1.0 0.75 0.625	360	1.0 0.25 0.625	65.7 50.6 22.5	55.4 24.0	1.0 0.25 0.625	64.8 49.9 21.5	54.4 23.3 1.5	0.0 0.5	55.9 67.5 30.1
672	B65R_100_075a	1.0 0.25 0.75	1.0 0.75 0.625	349	1.0 0.25 0.762	65.8 50.9 15.2	53.2 16.6	1.0 0.25 0.75	64.8 50.2 13.5	52.0 15.1 2.0	0.0 0.683	56.0 67.9 20.2
673	B57R_100_075a	1.0 0.25 0.875	1.0 0.75 0.625	339	1.0 0.25 0.887	65.9 51.2 9.7	52.1 10.8	1.0 0.25 0.875	65.3 50.7 7.3	50.5 8.2 2.7	0.0 0.85	56.1 68.2 13.0
674	B50R_100_075a	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	65.8 51.4 5.9	51.7 6.6	1.0 0.25 1.0	65.2 50.2 2.2	50.3 2.5 3.9	0.0 1.0	56.0 68.5 7.9
675	R36Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	66.6 44.2 66.7	80.0 56.4	1.0 0.375 0.0	66.9 43.6 67.0	79.9 56.9 0.7	0.0 0.366	66.6 44.2 66.7
676	R26Y_100_087a	1.0 0.375 0.125	1.0 0.875 0.562	46	1.0 0.358 0.125	67.1 44.9 55.2	71.2 50.8	1.0 0.375 0.125	67.0 43.9 57.6	72.4 52.7 2.5	0.0 0.4	66.7 43.1 51.3
677	R15Y_100_075a	1.0 0.375 0.25	1.0 0.75 0.625	39	1.0 0.362 0.25	68.4 44.1 44.8	62.9 45.4	1.0 0.375 0.25	67.6 43.1 48.7	65.1 48.5 4.1	0.0 0.15	59.4 58.8 59.7
678	R00Y_100_062a	1.0 0.375 0.375	1.0 0.625 0.687	390	1.0 0.375 0.375	70.6 41.7 34.9	54.3 39.9	1.0 0.375 0.375	68.6 41.4 39.9	57.5 43.9 5.4	0.0 0.0	55.7 66.7 55.8
679	R31Y_100_062a	1.0 0.375 0.5	1.0 0.625 0.687	379	1.0 0.375 0.489	70.6 41.8 29.6	51.2 35.3	1.0 0.375 0.5	68.8 41.5 31.2	51.9 36.9 2.4	0.0 0.183	55.9 66.9 47.4
680	R11Y_100_062a	1.0 0.375 0.625	1.0 0.625 0.687	367	1.0 0.375 0.614	70.7 42.0 23.1	48.0 28.8	1.0 0.375 0.625	69.4 40.8 22.3	46.5 27.7 1.9	0.0 0.367	56.0 67.2 37.1
681	B69R_100_062a	1.0 0.375 0.75	1.0 0.625 0.687	353	1.0 0.375 0.76	70.7 42.3 14.8	44.8 19.3	1.0 0.375 0.75	69.8 40.9 13.1	42.9 17.8 2.4	0.0 0.616	56.0 67.7 23.8
682	B59R_100_062a	1.0 0.375 0.875	1.0 0.625 0.687	341	1.0 0.375 0.885	70.7 42.6 8.9	43.5 11.8	1.0 0.375 0.875	70.2 40.6 6.1	41.1 8.6 3.4	0.0 0.816	56.0 68.2 14.2
683	B50R_100_062a	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	70.7 42.8 4.9	43.1 6.6	1.0 0.375 1.0	70.2 40.9 0.6	40.9 0.9 4.7	0.0 390	56.0 68.5 7.9
684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2	1.0 0.5 0.0	71.8 34.6 71.7	79.6 64.2 0.0	0.0 0.5	71.8 34.6 71.7
685	R41Y_100_087a	1.0 0.5 0.125	1.0 0.875 0.562	55	1.0 0.489 0.125	71.9 35.5 60.1	69.8 59.3	1.0 0.5 0.125	71.6 35.1 61.1	70.5 60.1 1.1	0.0 0.416	68.5 40.6 68.7
686	R31Y_100_075a	1.0 0.5 0.25	1.0 0.75 0.625	49	1.0 0.487 0.25	72.5 35.8 48.8	60.5 53.6	1.0 0.5 0.25	72.1 34.6 51.3	61.9 55.9 2.8	0.0 0.316	64.9 47.8 65.0
687	R18Y_100_062a	1.0 0.5 0.375	1.0 0.625 0.687	41	1.0 0.489 0.375	73.5 35.4 37.9	51.9 46.9	1.0 0.5 0.375	72.6 34.2 42.0	54.2 50.8 4.3	0.0 0.183	60.5 56.7 60.7
688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	75.5 33.3 27.9	43.5 39.9	1.0 0.5 0.5	73.5 33.0 31.6	45.7 43.7 4.2	0.0 0.0	55.7 66.7 55.8
689	R26Y_100_050a	1.0 0.5 0.625	1.0 0.5 0.75	376	1.0 0.5 0.616	75.6 33.4 22.5	40.3 33.9	1.0 0.5 0.625	74.1 32.5 22.3	39.5 34.4 1.7	0.0 0.233	56.0 66.9 45.0
690	R50Y_100_050a	1.0 0.5 0.75	1.0 0.5 0.75	360	1.0 0.5 0.75	75.6 33.7 15.0	36.9 24.0	1.0 0.5 0.75	74.5 31.3 13.8	35.1 23.1 2.1	0.0 0.5	55.9 67.5 30.1
691	B61R_100_050a	1.0 0.5 0.875	1.0 0.5 0.75	344	1.0 0.5 0.883	75.6 34.0 8.0	35.0 13.3	1.0 0.5 0.875	74.8 32.4 6.0	33.0 10.5 2.7	0.0 0.766	56.0 68.1 16.1
692	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	75.6 34.2 3.9	34.5 6.6	1.0 0.5 1.0	75.0 32.3 -0.4	32.3 35.9 2.4	0.0 1.0	56.0 68.5 7.9
693	R63Y_100_100a	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	77.7 23.8 77.9	81.5 72.9	1.0 0.625 0.0	77.3 24.4 77.5	81.3 72.5 0.7	0.0 0.633	77.7 23.8 77.9
694	R58Y_100_087a	1.0 0.625 0.125	1.0 0.875 0.562	65	1.0 0.635 0.125	78.0 24.4 66.3	70.6 69.7	1.0 0.625 0.125	77.9 23.7 66.7	70.8 70.3 0.8	0.0 0.583	75.5 23.9 75.8
695	R50Y_100_075a	1.0 0.625 0.25	1.0 0.75 0.625	60	1.0 0.625 0.25	77.7 25.9 53.8	59.7 64.2	1.0 0.625 0.25	78.0 23.9 56.1	60.9 66.9 3.1	0.0 0.5	71.8 34.6 71.7
696	R38Y_100_062a	1.0 0.625 0.375	1.0 0.625 0.687	53	1.0 0.614 0.375	77.7 26.8 42.1	49.9 57.4	1.0 0.625 0.375	78.2 24.0 44.9	51.0 61.8 4.0	0.0 0.383	67.2 43.0 67.3
697	R23Y_100_050a	1.0 0.625 0.5	1.0 0.5 0.75	44	1.0 0.616 0.5	78.6 26.8 31.0	41.0 49.1	1.0 0.625 0.5	78.8 23.6 33.9	41.3 55.0 4.2	0.0 0.233	62.0 53.6 62.0
698	R00Y_100_037a	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	80.4 25.0 20.9	32.6 39.9	1.0 0.625 0.625	79.8 22.6 23.8	32.8 46.4 3.8	0.0 0.0	55.7 66.7 55.8
699	R18Y_100_037a	1.0 0.625 0.75	1.0 0.375 0.812	371	1.0 0.625 0.743	80.5 25.1 15.2	29.4 31.2	1.0 0.625 0.75	80.1 22.6 15.0	27.1 33.6 2.6	0.0 0.316	56.0 67.1 40.7
700	B65R_100_037a	1.0 0.625 0.875	1.0 0.375 0.812	349	1.0 0.625 0.881	80.5 25.4 7.6	26.6 16.6	1.0 0.625 0.875	80.8 22.1 6.1	22.9 15.6 3.6	0.0 0.683	56.0 67.9 20.2
701	B50R_100_037a	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	80.5 25.7 2.9	25.8 6.6	1.0 0.625 1.0	81.1 22.1 -0.8	22.2 35.7 8.2	0.0 1.0	56.0 68.5 7.9
702	R76Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	82.9 15.1 83.1	84.4 79.6	1.0 0.75 0.0	82.3 16.1 82.4	84.0 78.9 1.2	0.0 0.766	82.9 15.1 83.1
703	R63Y_100_087a	1.0 0.75 0.125	1.0 0.875 0.562	74	1.0 0.766 0.125	83.4 15.1 71.6	73.2 78.0	1.0 0.75 0.125	82.8 15.6 71.1	72.8 77.6 0.9	0.0 0.733	81.7 17.2 81.8
704	R53Y_100_075a	1.0 0.75 0.25	1.0 0.75 0.625	71	1.0 0.762 0.25	83.6 15.4 59.9	61.9 75.5	1.0 0.75 0.25	83.2 15.2 59.6	61.5 75.6 0.5	0.0 0.683	79.7 20.6 79

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md
729	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.3 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.2 0.0 0.0	0.1 96.7 0.1	360
730	G50B_100_012a	0.875 1.0 1.0	1.0 1.0 1.0	1.0 0.125 0.937	210	0.875 1.0 1.0	89.7 -6.7 -7.2	9.8 226.8	0.875 1.0 1.0	91.3 -3.8 -5.2	6.5 233.1 3.8	210
731	G50B_100_025a	0.75 1.0 1.0	1.0 1.0 1.0	1.0 0.25 0.875	210	0.75 1.0 1.0	84.1 -13.4 -14.4	19.7 226.8	0.75 1.0 1.0	86.8 -7.7 -10.8	13.3 234.4 7.3	210
732	G50B_100_037a	0.625 1.0 1.0	1.0 1.0 1.0	1.0 0.375 0.812	210	0.625 1.0 1.0	78.5 -20.2 -21.6	29.6 226.8	0.625 1.0 1.0	82.1 -12.1 -16.8	20.8 234.3 10.0	210
733	G50B_100_050a	0.5 1.0 1.0	1.0 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	73.0 -26.9 -28.8	39.4 226.8	0.5 1.0 1.0	76.0 -17.9 -24.6	30.5 233.9 10.3	210
734	G50B_100_062a	0.375 1.0 1.0	1.0 1.0 1.0	1.0 0.625 0.687	210	0.375 1.0 1.0	67.4 -33.7 -36.0	49.3 226.8	0.375 1.0 1.0	70.3 -23.8 -31.8	39.7 233.1 11.1	210
735	G50B_100_075a	0.25 1.0 1.0	1.0 1.0 1.0	1.0 0.75 0.625	210	0.25 1.0 1.0	61.8 -40.4 -43.2	59.2 226.8	0.25 1.0 1.0	63.7 -31.6 -40.3	51.3 231.8 9.4	210
736	G50B_100_087a	0.125 1.0 1.0	1.0 1.0 1.0	1.0 0.875 0.562	210	0.125 1.0 1.0	56.3 -47.2 -50.4	69.1 226.8	0.125 1.0 1.0	57.2 -40.1 -48.4	62.9 230.3 7.4	210
737	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 1.0	1.0 0.5 210	210	0.0 1.0 1.0	50.7 -53.9 -57.6	78.9 226.8	0.0 1.0 1.0	49.1 -54.4 -58.7	80.1 227.1 2.0	210
738	ROOY_100_012a	1.0 0.875 0.875	1.0 1.0 1.0	1.0 0.125 0.937	390	1.0 0.875 0.875	90.3 8.3 6.9	10.8 39.9	1.0 0.875 0.875	90.5 6.5 8.5	10.8 52.6 2.4	389
739	NW_087a	0.875 0.875 0.875	0.875 0.0 0.875	0.875 0.875 360	360	0.875 0.875 0.875	85.6 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	89.2 -0.2 0.0	0.2 199.2 3.6	360
740	G50B_087_012a	0.75 0.875 0.875	0.875 0.125 0.812	210	210	0.75 0.875 0.875	80.0 -6.7 -7.2	9.8 226.8	0.75 0.875 0.875	85.1 -4.1 -5.5	6.9 232.7 5.9	210
741	G50B_087_025a	0.625 0.875 0.875	0.875 0.25 0.75	210	210	0.625 0.875 0.875	74.4 -13.4 -14.4	19.7 226.8	0.625 0.875 0.875	80.3 -8.2 -11.2	13.9 233.5 8.5	210
742	G50B_087_037a	0.5 0.875 0.875	0.875 0.375 0.687	210	210	0.5 0.875 0.875	68.9 -20.2 -21.6	29.6 226.8	0.5 0.875 0.875	74.3 -13.8 -18.8	23.4 233.8 8.8	210
743	G50B_087_050a	0.375 0.875 0.875	0.875 0.5 0.625	210	210	0.375 0.875 0.875	63.3 -26.9 -28.8	39.4 226.8	0.375 0.875 0.875	68.5 -19.6 -26.0	32.5 233.0 9.4	210
744	G50B_087_062a	0.25 0.875 0.875	0.875 0.625 0.562	210	210	0.25 0.875 0.875	57.7 -33.7 -36.0	49.3 226.8	0.25 0.875 0.875	61.7 -27.0 -34.7	44.0 232.0 7.8	210
745	G50B_087_075a	0.125 0.875 0.875	0.875 0.75 0.5 210	210	210	0.125 0.875 0.875	52.2 -40.4 -43.2	59.2 226.8	0.125 0.875 0.875	55.3 -35.3 -42.8	55.5 230.4 6.0	210
746	G50B_087_087a	0.0 0.875 0.875	0.875 0.875 0.437	210	210	0.0 0.875 0.875	46.6 -47.2 -50.4	69.1 226.8	0.0 0.875 0.875	47.7 -48.5 -52.7	71.7 227.3 2.9	210
747	ROOY_100_025a	1.0 0.75 0.75	1.0 0.25 0.875	390	390	1.0 0.75 0.75	85.4 16.6 13.9	21.7 39.9	1.0 0.75 0.75	84.2 15.4 17.1	23.0 48.1 3.6	389
748	ROOY_087_012a	0.875 0.75 0.75	0.75 0.125 0.812	390	390	0.875 0.75 0.75	80.6 8.3 6.9	10.8 39.9	0.875 0.75 0.75	84.1 6.6 8.8	11.0 53.1 4.3	389
749	NW_075a	0.75 0.75 0.75	0.75 0.0 0.75	360	360	0.75 0.75 0.75	75.9 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	80.4 -4.4 -0.3	0.5 220.1 4.5	360
750	G50B_075_012a	0.625 0.75 0.75	0.75 0.125 0.687	210	210	0.625 0.75 0.75	70.3 -6.7 -7.2	9.8 226.8	0.625 0.75 0.75	76.5 -4.4 -5.7	7.2 232.0 6.7	210
751	G50B_075_025a	0.5 0.75 0.75	0.75 0.25 0.625	210	210	0.5 0.75 0.75	64.7 -13.4 -14.4	19.7 226.8	0.5 0.75 0.75	71.2 -9.0 -12.1	15.1 233.3 8.1	210
752	G50B_075_037a	0.375 0.75 0.75	0.75 0.375 0.562	210	210	0.375 0.75 0.75	59.2 -20.2 -21.6	29.6 226.8	0.375 0.75 0.75	65.1 -14.7 -19.6	24.6 233.0 8.2	210
753	G50B_075_050a	0.25 0.75 0.75	0.75 0.5 0.5 210	210	210	0.25 0.75 0.75	53.6 -26.9 -28.8	39.4 226.8	0.25 0.75 0.75	58.9 -21.3 -27.6	34.8 232.3 7.8	210
754	G50B_075_062a	0.125 0.75 0.75	0.75 0.625 0.437	210	210	0.125 0.75 0.75	48.0 -33.7 -36.0	49.3 226.8	0.125 0.75 0.75	51.7 -29.7 -36.4	47.0 231.7 5.4	210
755	G50B_075_075a	0.0 0.75 0.75	0.75 0.75 0.375	210	210	0.0 0.75 0.75	42.5 -40.4 -43.2	59.2 226.8	0.0 0.75 0.75	44.8 -41.2 -45.7	61.5 227.9 3.4	210
756	ROOY_100_037a	1.0 0.625 0.625	1.0 0.375 0.812	390	390	1.0 0.625 0.625	80.4 25.0 20.9	32.6 39.9	1.0 0.625 0.625	79.0 23.4 25.6	34.7 47.5 5.1	389
757	ROOY_087_025a	0.875 0.625 0.625	0.875 0.25 0.75	390	390	0.875 0.625 0.625	75.7 16.6 13.9	21.7 39.9	0.875 0.625 0.625	77.4 16.2 17.8	24.1 47.7 4.3	389
758	ROOY_075_012a	0.75 0.625 0.625	0.75 0.125 0.687	390	390	0.75 0.625 0.625	70.9 8.3 6.9	10.8 39.9	0.75 0.625 0.625	75.3 6.9 8.7	11.1 51.3 4.9	389
759	NW_062a	0.625 0.625 0.625	0.625 0.0 0.625	360	360	0.625 0.625 0.625	66.2 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	72.8 -0.5 -0.4	0.7 214.6 6.6	360
760	G50B_062_012a	0.5 0.625 0.625	0.625 0.125 0.562	210	210	0.5 0.625 0.625	60.6 -6.7 -7.2	9.8 226.8	0.5 0.625 0.625	67.7 -5.0 -6.4	8.1 231.9 7.2	210
761	G50B_062_025a	0.375 0.625 0.625	0.625 0.25 0.5 210	210	210	0.375 0.625 0.625	55.1 -13.4 -14.4	19.7 226.8	0.375 0.625 0.625	62.2 -10.0 -13.3	16.7 232.9 7.9	210
762	G50B_062_037a	0.25 0.625 0.625	0.625 0.375 0.437	210	210	0.25 0.625 0.625	49.5 -20.2 -21.6	29.6 226.8	0.25 0.625 0.625	55.9 -16.4 -21.3	26.9 232.3 7.4	210
763	G50B_062_050a	0.125 0.625 0.625	0.625 0.5 0.375	210	210	0.125 0.625 0.625	43.9 -26.9 -28.8	39.4 226.8	0.125 0.625 0.625	48.8 -24.5 -30.2	38.9 230.9 5.6	210
764	G50B_062_062a	0.0 0.625 0.625	0.625 0.625 0.312	210	210	0.0 0.625 0.625	38.4 -33.7 -36.0	49.3 226.8	0.0 0.625 0.625	41.5 -35.1 -39.6	52.9 228.4 5.0	210
765	ROOY_100_050a	1.0 0.5 0.5 1.0 0.5 0.5 390	1.0 0.5 0.5 390	390	390	1.0 0.5 0.5 390	75.5 33.3 27.9	43.5 39.9	1.0 0.5 0.5 72.0	34.9 33.2	48.2 43.6 6.5	389
766	ROOY_087_037a	0.875 0.5 0.5 0.875 0.375 0.687	0.875 0.375 0.687	390	390	0.875 0.5 0.5 70.8	25.0 20.9	32.6 39.9	0.875 0.5 0.5 71.8	25.0 26.1	36.1 46.3 5.3	389
767	ROOY_075_025a	0.75 0.5 0.5 0.75 0.25 0.625	0.75 0.25 0.625	390	390	0.75 0.5 0.5 66.0	16.6 13.9	21.7 39.9	0.75 0.5 0.5 68.8	16.5 17.5	24.0 46.7 4.5	389
768	ROOY_062_012a	0.625 0.5 0.5 0.625 0.125 0.562	0.625 0.125 0.562	390	390	0.625 0.5 0.5 61.3	8.3 6.9	10.8 39.9	0.625 0.5 0.5 66.9	7.6 8.5	11.5 48.3 5.8	389
769	NW_050a	0.5 0.5 0.5 0.5 0.0 0.5 360	0.5 0.5 0.5 360	360	360	0.5 0.5 0.5 65.6	0.0 0.0	0.0 0.0	0.5 0.5 0.5 64.4	-0.6 -0.4	0.8 215.7 7.9	360
770	G50B_050_012a	0.375 0.5 0.5 0.5 0.125 0.437	0.375 0.125 0.437	210	210	0.375 0.5 0.5 50.9	-6.7 -7.2	9.8 226.8	0.375 0.5 0.5 59.1	-5.4 -6.8	8.7 231.6 8.2	210
771	G50B_050_025a	0.25 0.5 0.5 0.5 0.25 0.375	0.25 0.25 0.375	210	210	0.25 0.5 0.5 45.4	-13.4 -14.4	19.7 226.8	0.25 0.5 0.5 52.5	-11.7 -15.1	19.1 232.2 7.4	210
772	G50B_050_037a	0.125 0.5 0.5 0.5 0.375 0.312	0.125 0.375 0.312	210	210	0.125 0.5 0.5 39.8	-20.2 -21.6	29.6 226.8	0.125 0.5 0.5 45.7	-18.7 -23.5	30.1 231.4 6.3	210
773	G50B_050_050a	0.0 0.5 0.5 0.5 0.5 0.25 210	0.0 0.5 0.25 210	210	210	0.0 0.5 0.5 34.2	-26.9 -28.8	39.4 226.8	0.0 0.5 0.5 38.7	-28.3 -32.7	43.3 229.1 6.0	210
774	ROOY_100_062a	1.0 0.375 0.375 1.0 0.625 0.687	1.0 0.625 0.687	390	390	1.0 0.375 0.375 70.6	41.7 34.9	54.3 39.9	1.0 0.375 0.375 66.3	44.7 41.6	61.1 42.9 8.5	389
775	ROOY_087_050a	0.875 0.375 0.375 0.875 0.5 0.625	0.875 0.5 0.625	390	390	0.875 0.375 0.375 65.8	33.3 27.9	43.5 39.9	0.875 0.375 0.375 65.1	36.5 34.4	50.2 43.3 7.2	389
776	ROOY_075_037a	0.75 0.375 0.375 0.75 0.375 0.562	0.75 0.375 0.562	390	390	0.75 0.375 0.375 61.1	25.0 20.9	32.6 39.9	0.75 0.375 0.375 62.0	27.7 26.4	38.3 43.6 6.1	389
777	ROOY_062_025a	0.625 0.375 0.375 0.625 0.25 0.5 390	0.625 0.25 0.5 390	390	390	0.625 0.375 0.375 56.3	16.6 13.9	21.7 39.9	0.625 0.375 0.375 60.3	17.5 18.1	25.2 46.0 5.8	389
778	ROOY_050_012a	0.5 0.375 0.375 0.5 0.125 0.437	0.5 0.125 0.437	390	390	0.5 0.375 0.375 51.6	8.3 6.9	10.8 39.9	0.5 0.375 0.375 57.7	8.8 9.2	12.7 46.5 6.5	389
779	NW_037a	0.375 0.375 0.375 0.375 0.0 0.375 360	0.375 0.375 360	360	360	0.375 0.375 0.375 46.8	0.0 0.0	0.0 0.0	0.375 0.375 0.375 55.7	-0.7 -0.5	0.9 215.9 8.9	360
780	G50B_037_012a	0.25 0.375 0.375 0.375 0.125 0.312	0.25 0.125 0.312	210	210	0.25 0.375 0.375 41.3	-6.7 -7.2	9.8 226.8	0.25 0.375 0.375 50.1	-6.3 -8.1	10.3 231.8 8.8	210
781	G50B_037_025a	0.125 0.375 0.375 0.375 0.25 0.25 210	0.125 0.25 0.25 210	210	210	0.125 0.375 0.375 35.7	-13.4 -14.4	19.7 226.8	0.125 0.375 0.375 42.9	-13.4 -16.9	21.6 231.5 7.6	210
782	G50B_037_037a	0.0 0.375 0.375 0.375 0.375 0.187	0.0 0.375 0.187 210	210	210	0.0 0.375 0.375 30.1	-20.2 -21.6	29.6 226.8	0.0 0.375 0.375 36.1	-21.9 -26.1	34.1 229.9	

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS TUB material: code=rh4ta
 aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	rgb*Md	LabCh*Md	
810	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	0.0 0.0 1.0	360	1.0 1.0 1.0	95.3 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.3 0.0 0.0	360	1.0 1.0 1.0	95.3 0.0 0.0
811	BOOR_100_012a	0.875 0.875 1.0	1.0 0.125 0.937	270	0.875 0.875 1.0	86.2 0.6 0.6	-6.2 6.3 275.5	0.875 0.875 1.0	87.1 2.6 -5.9	6.5 293.9 2.2	270	0.0 0.0 1.0	22.6 4.8 -50.3
812	BOOR_100_025a	0.75 0.75 1.0	1.0 0.25 0.875	270	0.75 0.75 1.0	77.1 1.2 -12.5	12.6 275.5	0.75 0.75 1.0	77.9 6.6 -11.1	12.9 300.9 5.6	270	0.0 0.0 1.0	22.6 4.8 -50.3
813	BOOR_100_037a	0.625 0.625 1.0	1.0 0.375 0.812	270	0.625 0.625 1.0	68.0 1.8 -18.8	18.9 275.5	0.625 0.625 1.0	69.0 9.0 -16.7	19.0 298.4 7.6	270	0.0 0.0 1.0	22.6 4.8 -50.3
814	BOOR_100_050a	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	58.9 2.4 -25.1	25.2 275.5	0.5 0.5 1.0	57.6 13.1 -23.0	26.5 297.7 11.0	270	0.0 0.0 1.0	22.6 4.8 -50.3
815	BOOR_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	49.8 3.0 -31.4	31.6 275.5	0.375 0.375 1.0	47.9 14.9 -29.2	32.8 299.0 12.2	270	0.0 0.0 1.0	22.6 4.8 -50.3
816	BOOR_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	40.8 3.6 -37.7	37.9 275.5	0.25 0.25 1.0	39.1 15.1 -35.1	38.3 293.2 11.8	270	0.0 0.0 1.0	22.6 4.8 -50.3
817	BOOR_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	31.7 4.2 -44.0	44.2 275.5	0.125 0.125 1.0	30.0 15.3 -41.1	43.9 290.4 11.5	270	0.0 0.0 1.0	22.6 4.8 -50.3
818	BOOR_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	22.6 4.8 -50.3	50.5 275.5	0.0 0.0 1.0	22.3 7.3 -49.0	49.5 278.5 2.8	270	0.0 0.0 1.0	22.6 4.8 -50.3
819	YOOG_100_012a	1.0 1.0 0.875	1.0 0.125 0.937	90	1.0 1.0 0.875	94.7 0.3 11.4	11.4 88.3	1.0 1.0 0.875	94.7 0.0 9.8	9.8 89.8 1.6	89	1.0 1.0 1.0	90.5 2.6 91.3
820	NW_087a	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	85.6 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	89.1 -0.1 -0.1	0.2 214.6 3.5	360	1.0 1.0 1.0	95.3 0.0 0.0
821	BOOR_087_012a	0.75 0.75 0.875	0.875 0.125 0.812	270	0.75 0.75 0.875	76.5 0.6 -6.2	6.3 275.5	0.75 0.75 0.875	81.1 2.4 -5.9	6.4 292.1 4.9	270	0.0 0.0 1.0	22.6 4.8 -50.3
822	BOOR_087_025a	0.625 0.625 0.875	0.875 0.25 0.75	270	0.625 0.625 0.875	67.4 1.2 -12.5	12.6 275.5	0.625 0.625 0.875	71.1 6.8 -11.3	13.2 301.0 6.8	270	0.0 0.0 1.0	22.6 4.8 -50.3
823	BOOR_087_037a	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.5 0.875	58.3 1.8 -18.8	18.9 275.5	0.5 0.5 0.875	60.7 8.3 -18.3	20.1 294.5 6.9	270	0.0 0.0 1.0	22.6 4.8 -50.3
824	BOOR_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	49.2 2.4 -25.1	25.2 275.5	0.375 0.375 0.875	50.4 11.8 -24.0	26.7 296.2 9.5	270	0.0 0.0 1.0	22.6 4.8 -50.3
825	BOOR_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	40.2 3.0 -31.4	31.6 275.5	0.25 0.25 0.875	40.1 13.5 -30.8	33.6 293.7 10.5	270	0.0 0.0 1.0	22.6 4.8 -50.3
826	BOOR_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	31.1 3.6 -37.7	37.9 275.5	0.125 0.125 0.875	30.1 14.8 -37.2	40.1 291.6 11.1	270	0.0 0.0 1.0	22.6 4.8 -50.3
827	BOOR_087_087a	0.0 0.0 0.875	0.875 0.875 0.437	270	0.0 0.0 0.875	22.0 4.2 -44.0	44.2 275.5	0.0 0.0 0.875	22.3 8.7 -44.6	45.4 281.1 4.5	270	0.0 0.0 1.0	22.6 4.8 -50.3
828	YOOG_100_025a	1.0 1.0 0.75	1.0 0.25 0.875	90	1.0 1.0 0.75	94.1 0.6 22.8	22.8 88.3	1.0 1.0 0.75	93.9 0.2 20.1	20.1 89.2 2.7	89	1.0 1.0 1.0	90.5 2.6 91.3
829	YOOG_087_012a	0.875 0.875 0.75	0.875 0.125 0.812	90	0.875 0.875 0.75	85.0 0.3 11.4	11.4 88.3	0.875 0.875 0.75	88.5 -0.1 10.0	10.0 90.8 3.8	89	1.0 1.0 1.0	90.5 2.6 91.3
830	NW_075a	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	75.9 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	80.5 -0.4 -0.2	0.5 216.0 4.6	360	1.0 1.0 1.0	95.3 0.0 0.0
831	BOOR_075_012a	0.625 0.625 0.75	0.75 0.125 0.687	270	0.625 0.625 0.75	66.8 0.6 -6.2	6.3 275.5	0.625 0.625 0.75	72.2 2.4 -6.2	6.7 291.5 5.7	270	0.0 0.0 1.0	22.6 4.8 -50.3
832	BOOR_075_025a	0.5 0.5 0.75	0.75 0.25 0.625	270	0.5 0.5 0.75	57.7 1.2 -12.5	12.6 275.5	0.5 0.5 0.75	61.8 6.1 -11.9	13.3 297.1 6.3	270	0.0 0.0 1.0	22.6 4.8 -50.3
833	BOOR_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	48.6 1.8 -18.8	18.9 275.5	0.375 0.375 0.75	51.1 9.0 -18.4	20.5 296.0 7.6	270	0.0 0.0 1.0	22.6 4.8 -50.3
834	BOOR_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	39.6 2.4 -25.1	25.2 275.5	0.25 0.25 0.75	40.7 11.5 -24.7	27.5 295.1 9.2	270	0.0 0.0 1.0	22.6 4.8 -50.3
835	BOOR_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	30.5 3.0 -31.4	31.6 275.5	0.125 0.125 0.75	30.2 13.2 -31.5	34.2 292.7 10.1	270	0.0 0.0 1.0	22.6 4.8 -50.3
836	BOOR_075_075a	0.0 0.0 0.75	0.75 0.75 0.375	270	0.0 0.0 0.75	21.4 3.6 -37.7	37.9 275.5	0.0 0.0 0.75	21.7 10.5 -38.4	39.8 285.3 6.9	270	0.0 0.0 1.0	22.6 4.8 -50.3
837	YOOG_100_037a	1.0 1.0 0.625	1.0 0.375 0.812	90	1.0 1.0 0.625	93.5 1.0 34.2	34.2 88.3	1.0 1.0 0.625	93.3 0.6 31.6	31.6 88.8 2.6	89	1.0 1.0 1.0	90.5 2.6 91.3
838	YOOG_087_025a	0.875 0.875 0.625	0.875 0.25 0.75	90	0.875 0.875 0.625	84.4 0.6 22.8	22.8 88.3	0.875 0.875 0.625	87.9 0.1 21.1	21.1 89.7 3.9	89	1.0 1.0 1.0	90.5 2.6 91.3
839	YOOG_075_012a	0.75 0.75 0.625	0.75 0.125 0.687	90	0.75 0.75 0.625	75.3 0.3 11.4	11.4 88.3	0.75 0.75 0.625	80.1 -0.3 10.0	10.0 92.1 5.0	89	1.0 1.0 1.0	90.5 2.6 91.3
840	NW_062a	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	66.2 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	73.0 -0.5 -0.4	0.7 215.4 6.8	360	1.0 1.0 1.0	95.3 0.0 0.0
841	BOOR_062_012a	0.5 0.5 0.625	0.625 0.125 0.562	270	0.5 0.5 0.625	57.1 0.6 -6.2	6.3 275.5	0.5 0.5 0.625	63.3 2.4 -6.7	7.1 290.3 6.4	270	0.0 0.0 1.0	22.6 4.8 -50.3
842	BOOR_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	48.0 1.2 -12.5	12.6 275.5	0.375 0.375 0.625	53.0 5.5 -12.6	13.8 293.6 6.6	270	0.0 0.0 1.0	22.6 4.8 -50.3
843	BOOR_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	39.0 1.8 -18.8	18.9 275.5	0.25 0.25 0.625	42.2 8.6 -19.0	20.9 294.4 7.5	270	0.0 0.0 1.0	22.6 4.8 -50.3
844	BOOR_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	29.9 2.4 -25.1	25.2 275.5	0.125 0.125 0.625	30.9 11.2 -26.1	28.4 293.1 8.8	270	0.0 0.0 1.0	22.6 4.8 -50.3
845	BOOR_062_062a	0.0 0.0 0.625	0.625 0.625 0.312	270	0.0 0.0 0.625	20.8 3.0 -31.4	31.6 275.5	0.0 0.0 0.625	21.2 10.3 -33.5	35.1 287.1 7.6	270	0.0 0.0 1.0	22.6 4.8 -50.3
846	YOOG_100_050a	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	92.9 1.3 45.6	45.7 88.3	1.0 1.0 0.5	92.6 1.0 43.5	43.5 88.6 2.2	89	1.0 1.0 1.0	90.5 2.6 91.3
847	YOOG_087_037a	0.875 0.875 0.5	0.875 0.375 0.687	90	0.875 0.875 0.5	83.8 1.0 34.2	34.2 88.3	0.875 0.875 0.5	87.2 0.4 32.5	32.5 89.2 3.8	89	1.0 1.0 1.0	90.5 2.6 91.3
848	YOOG_075_025a	0.75 0.75 0.5	0.75 0.25 0.625	90	0.75 0.75 0.5	74.7 0.6 22.8	22.8 88.3	0.75 0.75 0.5	79.4 -0.1 21.4	21.4 90.2 5.0	89	1.0 1.0 1.0	90.5 2.6 91.3
849	YOOG_062_012a	0.625 0.625 0.5	0.625 0.125 0.562	90	0.625 0.625 0.5	65.6 0.3 11.4	11.4 88.3	0.625 0.625 0.5	72.5 -0.5 10.2	10.2 92.9 7.0	89	1.0 1.0 1.0	90.5 2.6 91.3
850	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	56.5 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	64.3 -0.6 -0.5	0.8 216.4 7.8	360	1.0 1.0 1.0	95.3 0.0 0.0
851	BOOR_050_012a	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.375 0.5	47.4 0.6 -6.2	6.3 275.5	0.375 0.375 0.5	53.9 3.1 -7.0	7.6 293.8 6.9	270	0.0 0.0 1.0	22.6 4.8 -50.3
852	BOOR_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.25 0.25 0.5	38.3 1.2 -12.5	12.6 275.5	0.25 0.25 0.5	43.0 5.6 -13.9	15.0 292.0 6.5	270	0.0 0.0 1.0	22.6 4.8 -50.3
853	BOOR_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.125 0.125 0.5	29.3 1.8 -18.8	18.9 275.5	0.125 0.125 0.5	31.3 9.4 -20.6	22.6 294.5 8.0	270	0.0 0.0 1.0	22.6 4.8 -50.3
854	BOOR_050_050a	0.0 0.0 0.5	0.5 0.5 0.25	270	0.0 0.0 0.5	20.2 2.4 -25.1	25.2 275.5	0.0 0.0 0.5	21.3 10.5 -27.2	29.1 291.1 8.3	270	0.0 0.0 1.0	22.6 4.8 -50.3
855	YOOG_100_062a	1.0 1.0 0.375	1.0 0.625 0.687	90	1.0 1.0 0.375	92.3 1.6 57.1	57.1 88.3	1.0 1.0 0.375	91.9 1.5 56.0	56.1 88.3 1.1	89	1.0 1.0 1.0	90.5 2.6 91.3
856	YOOG_087_050a	0.875 0.875 0.375	0.875 0.5 0.625	90	0.875 0.875 0.375	83.2 1.3 45.6	45.7 88.3	0.875 0.875 0.375	86.6 0.8 45.2	45.2 88.8 3.5	89	1.0 1.0 1.0	90.5 2.6 91.3
857	YOOG_075_037a	0.75 0.75 0.375	0.75 0.375 0.562	90	0.75 0.75 0.375	74.1 1.0 34.2	34.2 88.3	0.75 0.75 0.375	78.9 0.1 33.6	33.6 89.6 4.8	89	1.0 1.0 1.0	90.5 2.6 91.3
858	YOOG_062_025a	0.625 0.625 0.375	0.625 0.25 0.5	90	0.625 0.625 0.375	65.0 0.6 22.8	22.8 88.3	0.625 0.625 0.375	71.8 -0.3 22.3	22.3 90.7 6.8	89	1.0 1.0 1.0	90.5 2.6 91.3
859	YOOG_050_012a	0.5 0.5 0.375	0.5 0.125 0.437	90	0.5 0.5 0.375	55.9 0.3 11.4	11.4 88.3	0.5 0.5 0.375	63.9 -0.6 10.9	10.9 93.4 8.1	89	1.0 1.0 1.0	90.5 2.6 91.3
860	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	46.8 0.0 0.0	0.0 0.0 0.0	0.375 0.375 0.375	56.3 -0.7 -0.6	0.9 218.7 9.4	360	1.0 1.0 1.0	95.3 0.0 0.0
861	BOOR_037_012a	0.25 0.25 0.375	0.375 0.125 0.312	270	0.25 0.25 0.375	37.7 0.6 -6.2	6.3 275.5	0.25 0.25 0.375	44.5 3.3 -7.7	8.4 293.4 7.4	270	0.0 0.0 1.0	22.6 4.8 -50.3
862	BOOR_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.125 0.125 0.375	28.7 1.2 -12.5							

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS
 aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)
 TUB material: code=rh4ta

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md
891	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.3 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.2 0.0 0.0	0.0 0.0	333.8 0.0	360
892	B50R_100_012a	1.0 0.875 1.0	1.0 0.125 0.937	330	1.0 0.875 1.0	90.3 8.5 0.9	8.6 6.6	1.0 0.875 1.0	91.1 6.5 -0.7	6.5	353.0 2.8	330
893	B50R_100_025a	1.0 0.75 1.0	1.0 0.25 0.875	330	1.0 0.75 1.0	85.4 17.1 1.9	17.2 6.6	1.0 0.75 1.0	85.8 14.6 -0.9	14.7	356.1 3.8	330
894	B50R_100_037a	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	80.5 25.7 2.9	25.8 6.6	1.0 0.625 1.0	80.9 22.4 -0.8	22.4	357.8 5.0	330
895	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	75.6 34.2 3.9	34.5 6.6	1.0 0.5 1.0	74.2 33.6 -0.1	33.6	359.6 4.4	330
896	B50R_100_062a	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	70.7 42.8 4.9	43.1 6.6	1.0 0.375 1.0	68.7 43.1 1.2	43.1	1.6 4.2	330
897	B50R_100_075a	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	65.8 51.4 5.9	51.7 6.6	1.0 0.25 1.0	63.7 52.6 3.6	52.7	3.9 3.4	330
898	B50R_100_087a	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	60.9 59.9 6.9	60.3 6.6	1.0 0.125 1.0	58.4 62.6 6.9	63.0	6.2 3.6	330
899	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	56.0 68.5 7.9	69.0 6.6	1.0 0.0 1.0	54.6 69.7 11.1	70.6	9.1 3.6	330
900	GO0B_100_012a	0.875 1.0 0.875	1.0 0.125 0.937	150	0.875 1.0 0.875	89.2 -7.5 1.8	7.7 166.3	0.875 1.0 0.875	90.7 -3.8 4.4	5.9	130.8 4.7	149
901	NW_087a	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	85.6 0.0 0.0	0.0 0.0	0.875 0.875 0.875	89.2 -0.1 -0.1	0.2	216.1 3.6	360
902	B50R_087_012a	0.875 0.75 0.875	0.875 0.125 0.812	330	0.875 0.75 0.875	80.7 8.5 0.9	8.6 6.6	0.875 0.75 0.875	85.2 6.3 -0.8	6.4	352.1 5.3	330
903	B50R_087_025a	0.875 0.625 0.875	0.875 0.25 0.75	330	0.875 0.625 0.875	75.7 17.1 1.9	17.2 6.6	0.875 0.625 0.875	79.2 15.4 -1.1	15.4	355.9 4.9	330
904	B50R_087_037a	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	70.8 25.7 2.9	25.8 6.6	0.875 0.5 0.875	74.0 23.8 -0.8	23.8	357.8 5.3	330
905	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	65.9 34.2 3.9	34.5 6.6	0.875 0.375 0.875	67.4 34.7 0.0	34.7	359.9 4.2	330
906	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.625	330	0.875 0.25 0.875	61.0 42.8 4.9	43.1 6.6	0.875 0.25 0.875	61.1 46.0 1.9	46.0	2.3 4.4	330
907	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	56.1 51.4 5.9	51.7 6.6	0.875 0.125 0.875	55.7 56.2 4.9	56.4	5.0 4.9	330
908	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	51.2 59.9 6.9	60.3 6.6	0.875 0.0 0.875	51.4 64.6 9.0	65.2	7.9 5.0	330
909	GO0B_100_025a	0.75 1.0 0.75	1.0 0.25 0.875	150	0.75 1.0 0.75	83.2 -15.0 3.6	15.4 166.3	0.75 1.0 0.75	85.8 -7.6 8.6	11.5	131.3 9.2	149
910	GO0B_087_012a	0.75 0.875 0.75	0.875 0.125 0.812	150	0.75 0.875 0.75	79.6 -7.5 1.8	7.7 166.3	0.75 0.875 0.75	84.5 -4.2 4.4	6.1	133.9 6.4	149
911	NW_075a	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	75.9 0.0 0.0	0.0 0.0	0.75 0.75 0.75	81.1 -0.4 -0.2	0.4	212.9 5.2	360
912	B50R_075_012a	0.75 0.625 0.75	0.75 0.125 0.687	330	0.75 0.625 0.75	71.0 8.5 0.9	8.6 6.6	0.75 0.625 0.75	76.2 6.7 -1.1	6.8	350.4 5.9	330
913	B50R_075_025a	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	66.1 17.1 1.9	17.2 6.6	0.75 0.5 0.75	70.3 15.8 -1.3	15.9	355.2 5.5	330
914	B50R_075_037a	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	61.1 25.7 2.9	25.8 6.6	0.75 0.375 0.75	64.1 26.0 -0.9	26.1	357.9 4.9	330
915	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	56.2 34.2 3.9	34.5 6.6	0.75 0.25 0.75	57.5 37.2 0.5	37.2	0.7 4.7	330
916	B50R_075_062a	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	51.3 42.8 4.9	43.1 6.6	0.75 0.125 0.75	51.6 47.7 3.0	47.8	3.7 5.2	330
917	B50R_075_075a	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	46.4 51.4 5.9	51.7 6.6	0.75 0.0 0.75	47.0 57.1 6.5	57.4	6.5 5.7	330
918	GO0B_100_037a	0.625 1.0 0.625	1.0 0.375 0.812	150	0.625 1.0 0.625	77.2 -22.5 5.4	23.2 166.3	0.625 1.0 0.625	79.8 -12.2 12.6	17.6	134.2 12.7	149
919	GO0B_087_025a	0.625 0.875 0.625	0.875 0.25 0.75	150	0.625 0.875 0.625	73.6 -15.0 3.6	15.4 166.3	0.625 0.875 0.625	79.0 -8.3 8.7	12.1	133.7 10.0	149
920	GO0B_075_012a	0.625 0.75 0.625	0.75 0.125 0.687	150	0.625 0.75 0.625	69.9 -7.5 1.8	7.7 166.3	0.625 0.75 0.625	76.0 -4.5 4.2	6.2	136.7 7.2	149
921	NW_062a	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	66.2 0.0 0.0	0.0 0.0	0.625 0.625 0.625	73.2 -0.5 -0.4	0.7	218.4 7.0	360
922	B50R_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	61.3 8.5 0.9	8.6 6.6	0.625 0.5 0.625	67.7 7.4 -1.3	7.5	350.0 6.9	330
923	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	56.4 17.1 1.9	17.2 6.6	0.625 0.375 0.625	62.2 16.4 -1.3	16.5	355.4 6.7	330
924	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	51.5 25.7 2.9	25.8 6.6	0.625 0.25 0.625	55.1 28.0 -0.6	28.0	358.7 5.6	330
925	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	46.5 34.2 3.9	34.5 6.6	0.625 0.125 0.625	48.6 39.4 1.3	39.4	1.9 6.1	330
926	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	41.6 42.8 4.9	43.1 6.6	0.625 0.0 0.625	43.2 49.8 4.6	50.0	5.2 7.1	330
927	GO0B_100_050a	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	71.2 -30.0 7.3	30.9 166.3	0.5 1.0 0.5	72.9 -18.3 15.2	23.8	140.3 14.2	149
928	GO0B_087_037a	0.5 0.875 0.5	0.875 0.375 0.687	150	0.5 0.875 0.5	67.5 -22.5 5.4	23.2 166.3	0.5 0.875 0.5	72.1 -14.2 11.3	18.2	141.5 11.1	149
929	GO0B_075_025a	0.5 0.75 0.5	0.75 0.25 0.625	150	0.5 0.75 0.5	63.9 -15.0 3.6	15.4 166.3	0.5 0.75 0.5	70.1 -9.1 8.2	12.3	137.8 9.7	149
930	GO0B_062_012a	0.5 0.625 0.5	0.625 0.125 0.562	150	0.5 0.625 0.5	60.2 -7.5 1.8	7.7 166.3	0.5 0.625 0.5	67.7 -5.0 3.8	6.3	142.3 8.1	149
931	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	56.5 0.0 0.0	0.0 0.0	0.5 0.5 0.5	65.4 -0.6 -0.5	0.8	217.3 8.9	360
932	B50R_050_012a	0.5 0.375 0.5	0.5 0.125 0.437	330	0.5 0.375 0.5	51.6 8.5 0.9	8.6 6.6	0.5 0.375 0.5	58.8 8.6 -1.3	8.7	350.9 7.5	330
933	B50R_050_025a	0.5 0.25 0.5	0.5 0.25 0.375	330	0.5 0.25 0.5	46.7 17.1 1.9	17.2 6.6	0.5 0.25 0.5	52.4 19.3 -1.2	19.4	356.2 6.9	330
934	B50R_050_037a	0.5 0.125 0.5	0.5 0.375 0.312	330	0.5 0.125 0.5	41.8 25.7 2.9	25.8 6.6	0.5 0.125 0.5	44.7 31.4 0.1	31.4	0.2 7.0	330
935	B50R_050_050a	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	36.9 34.2 3.9	34.5 6.6	0.5 0.0 0.5	39.2 42.3 3.0	42.4	4.1 8.4	330
936	GO0B_100_062a	0.375 1.0 0.375	1.0 0.625 0.687	150	0.375 1.0 0.375	65.2 -37.6 9.1	38.7 166.3	0.375 1.0 0.375	66.3 -25.1 17.5	30.6	145.1 15.0	149
937	GO0B_087_050a	0.375 0.875 0.375	0.875 0.5 0.625	150	0.375 0.875 0.375	61.5 -30.0 7.3	30.9 166.3	0.375 0.875 0.375	65.9 -20.4 14.7	25.2	144.2 12.9	149
938	GO0B_075_037a	0.375 0.75 0.375	0.75 0.375 0.562	150	0.375 0.75 0.375	57.9 -22.5 5.4	23.2 166.3	0.375 0.75 0.375	63.5 -15.4 11.3	19.2	143.6 10.8	149
939	GO0B_062_025a	0.375 0.625 0.375	0.625 0.25 0.5	150	0.375 0.625 0.375	54.2 -15.0 3.6	15.4 166.3	0.375 0.625 0.375	61.7 -10.3 8.1	13.1	141.9 9.9	149
940	GO0B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	50.5 -7.5 1.8	7.7 166.3	0.375 0.5 0.375	59.5 -5.5 4.1	6.9	143.3 9.4	149
941	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	46.8 0.0 0.0	0.0 0.0	0.375 0.375 0.375	57.4 -0.7 -0.5	0.9	218.5 10.6	360
942	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.25 0.375	41.9 8.5 0.9	8.6 6.6	0.375 0.25 0.375	49.8 10.0 -1.3	10.1	352.1 8.4	330
943	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.125 0.375	37.0 17.1 1.9	17.2 6.6	0.375 0.125 0.375	42.4 21.7 -0.7	21.7	358.1 7.6	330
944	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	32.1 25.7 2.9	25.8 6.6	0.375 0.0 0.375	35.9 33.7 1.6	33.7	2.8 8.9	330
945	GO0B_100_075a	0.25 1.0 0.25	1.0 0.75 0.625	150	0.25 1.0 0.25	59.2 -45.1 10.9	46.4 166.3	0.25 1.0 0.25	59.2 -33.7 17.5	38.0	152.5 13.0	149
946	GO0B_087_062a	0.25 0.875 0.25	0.875 0.625 0.562	150	0.25 0.875 0.25	55.5 -37.6 9.1	38.7 166.3	0.25 0.875 0.25	58.4 -28.5 16.8	33.1	149.4 12.2	149
947	GO0B_075_050a	0.25 0.75 0.25	0.75 0.5 0.5	150	0.25 0.75 0.25	51.9 -30.0 7.3	30.9 166.3	0.25 0.75 0.25	56.6 -22.3 14.4	26.6	147.0 11.5	149
948	GO0B_062_037a	0.25 0.625 0.25	0.625 0.375 0.437	150	0.25 0.625 0.25	48.2 -22.5 5.4	23.2 166.3	0.25 0.625 0.25	54.5 -17.4 11.3	20.8	146.8 10.0	149
949	GO0B_050_025a	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.249	44.5 -15.0 3.6	15.4 166.3	0.25 0.5 0.25	52.2 -12.2 7.8	14.5	147.2 9.2	149
950	GO0B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	40.8 -7.5 1.8	7.7 166.3	0.25 0.375 0.25	50.4 -6.5 4.2	7.8	146.7 9.9	149
951	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	37.1 0.0 0.0	0.0					

vea archivos semejantes: <http://130.149.60.45/~farbmetrik/PS37/PS37.HTM>
 información técnica: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB matrícula: 20130201-PS37/PS37L0NA.TXT /.PS TUB material: code=rh4ta
 aplicación para la medida salida en la impresión offset, separación cmy0 (CMYK)

n	HIC*Fd	rgb_Fd	icf_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md
1053	NW_086a	0.866 0.866 0.866	0.866 0.0 0.866	360	0.866 0.866 0.866	84.9 0.0 0.0	0.866 0.866 0.866	89.2 -0.2 0.0	199.7 4.3 360	1.0 1.0 1.0	95.3 0.0 0.0	
1054	NW_093a	0.933 0.933 0.933	0.933 0.0 0.933	360	0.933 0.933 0.933	90.1 0.0 0.0	0.933 0.933 0.933	92.0 -0.1 0.0	181.7 1.9 360	1.0 1.0 1.0	95.3 0.0 0.0	
1055	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.3 0.0 0.0	1.0 1.0 1.0	95.3 0.0 0.0	115.3 0.0 360	1.0 1.0 1.0	95.3 0.0 0.0	
1056	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	17.8 0.0 0.0	0.0 0.0 0.0	18.8 0.0 0.1	82.2 1.0 360	1.0 1.0 1.0	95.3 0.0 0.0	
1057	NW_006a	0.066 0.066 0.066	0.066 0.0 0.066	360	0.066 0.066 0.066	22.9 0.0 0.0	0.066 0.066 0.066	22.4 -0.1 0.0	158.0 0.5 360	1.0 1.0 1.0	95.3 0.0 0.0	
1058	NW_013a	0.133 0.133 0.133	0.133 0.0 0.133	360	0.133 0.133 0.133	28.1 0.0 0.0	0.133 0.133 0.133	30.4 -0.5 -0.6	226.4 2.4 360	1.0 1.0 1.0	95.3 0.0 0.0	
1059	NW_020a	0.2 0.2 0.2	0.2 0.0 0.2	360	0.2 0.2 0.2	33.3 0.0 0.0	0.2 0.2 0.2	38.9 -0.8 1.2	226.6 5.7 360	1.0 1.0 1.0	95.3 0.0 0.0	
1060	NW_026a	0.266 0.266 0.266	0.266 0.0 0.266	360	0.266 0.266 0.266	38.4 0.0 0.0	0.266 0.266 0.266	45.5 -0.8 -0.8	223.4 7.2 360	1.0 1.0 1.0	95.3 0.0 0.0	
1061	NW_033a	0.333 0.333 0.333	0.333 0.0 0.333	360	0.333 0.333 0.333	43.6 0.0 0.0	0.333 0.333 0.333	51.8 -0.8 -0.7	220.3 8.3 360	1.0 1.0 1.0	95.3 0.0 0.0	
1062	NW_040a	0.4 0.4 0.4	0.4 0.0 0.4	360	0.4 0.4 0.4	48.8 0.0 0.0	0.4 0.4 0.4	57.2 -0.8 -0.7	219.8 8.5 360	1.0 1.0 1.0	95.3 0.0 0.0	
1063	NW_046a	0.466 0.466 0.466	0.466 0.0 0.466	360	0.466 0.466 0.466	53.9 0.0 0.0	0.466 0.466 0.466	61.6 -0.8 -0.6	220.1 7.8 360	1.0 1.0 1.0	95.3 0.0 0.0	
1064	NW_053a	0.533 0.533 0.533	0.533 0.0 0.533	360	0.533 0.533 0.533	59.1 0.0 0.0	0.533 0.533 0.533	66.8 -0.7 -0.6	219.8 7.8 360	1.0 1.0 1.0	95.3 0.0 0.0	
1065	NW_060a	0.6 0.6 0.6	0.6 0.0 0.6	360	0.6 0.6 0.6	64.3 0.0 0.0	0.6 0.6 0.6	71.9 -0.6 -0.5	217.4 7.7 360	1.0 1.0 1.0	95.3 0.0 0.0	
1066	NW_066a	0.666 0.666 0.666	0.666 0.0 0.666	360	0.666 0.666 0.666	69.4 0.0 0.0	0.666 0.666 0.666	76.6 -0.5 -0.4	218.0 7.2 360	1.0 1.0 1.0	95.3 0.0 0.0	
1067	NW_073a	0.734 0.734 0.734	0.734 0.0 0.734	360	0.734 0.734 0.734	74.6 0.0 0.0	0.734 0.734 0.734	80.7 -0.4 -0.3	213.1 6.0 360	1.0 1.0 1.0	95.3 0.0 0.0	
1068	NW_080a	0.8 0.8 0.8	0.8 0.0 0.8	360	0.8 0.8 0.8	79.8 0.0 0.0	0.8 0.8 0.8	84.6 -0.3 -0.2	208.6 4.8 360	1.0 1.0 1.0	95.3 0.0 0.0	
1069	NW_086a	0.866 0.866 0.866	0.866 0.0 0.866	360	0.866 0.866 0.866	84.9 0.0 0.0	0.866 0.866 0.866	89.1 -0.2 -0.1	209.9 4.2 360	1.0 1.0 1.0	95.3 0.0 0.0	
1070	NW_093a	0.933 0.933 0.933	0.933 0.0 0.933	360	0.933 0.933 0.933	90.1 0.0 0.0	0.933 0.933 0.933	92.1 0.0 0.0	141.9 2.0 360	1.0 1.0 1.0	95.3 0.0 0.0	
1071	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.3 0.0 0.0	1.0 1.0 1.0	95.3 0.0 0.0	104.4 0.0 360	1.0 1.0 1.0	95.3 0.0 0.0	
1072	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	17.8 0.0 0.0	0.0 0.0 0.0	20.1 0.3 0.5	61.6 2.4 360	1.0 1.0 1.0	95.3 0.0 0.0	
1073	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.3 0.0 0.0	1.0 1.0 1.0	95.4 0.0 -0.1	260.6 0.1 360	1.0 1.0 1.0	95.3 0.0 0.0	
1074	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	55.7 66.7 55.8	1.0 0.0 0.0	53.6 68.7 56.5	89.0 39.4 3.0	389	1.0 0.0 0.0	55.7 66.7 55.8
1075	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	50.7 -53.9 -57.6	0.0 1.0 1.0	48.2 -54.9 -59.7	81.1 227.4 3.4	210	0.0 1.0 1.0	50.7 -53.9 -57.6
1076	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	90.5 2.6 91.3	1.0 1.0 0.0	89.7 3.5 91.8	91.9 87.7 1.2	89	1.0 1.0 0.0	90.5 2.6 91.3
1077	B00R_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	22.6 4.8 -50.3	0.0 0.0 1.0	20.6 7.4 -48.2	48.7 278.7 3.8	270	0.0 0.0 1.0	22.6 4.8 -50.3
1078	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	47.2 -60.1 14.6	0.0 1.0 0.0	43.5 -62.7 11.4	63.7 169.6 5.4	149	0.0 1.0 0.0	47.2 -60.1 14.6
1079	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	56.0 68.5 7.9	1.0 0.0 1.0	53.4 70.9 14.5	72.4 11.6 7.4	330	1.0 0.0 1.0	56.0 68.5 7.9

delta E* = 4.2

2-0032130-F0

PS370-7N, 22/22-F

gráfico TUB-PS37; reproducción en color
 colores y diferencia en color, ΔE^* , 3D=0, de=0, cmy0

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

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