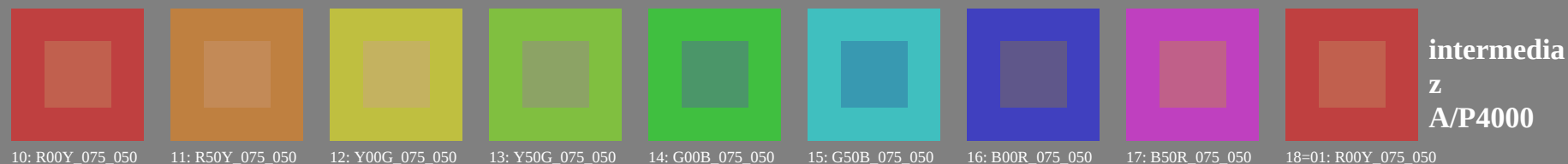
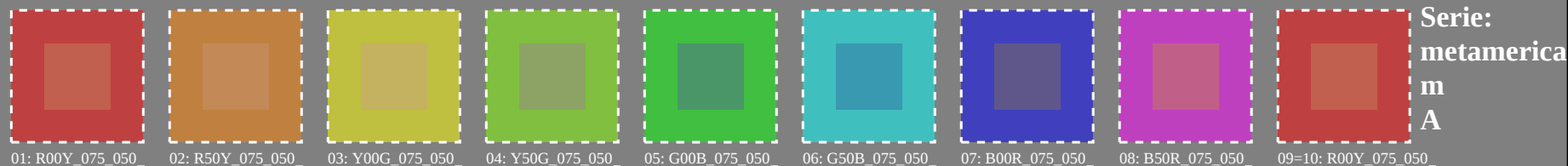
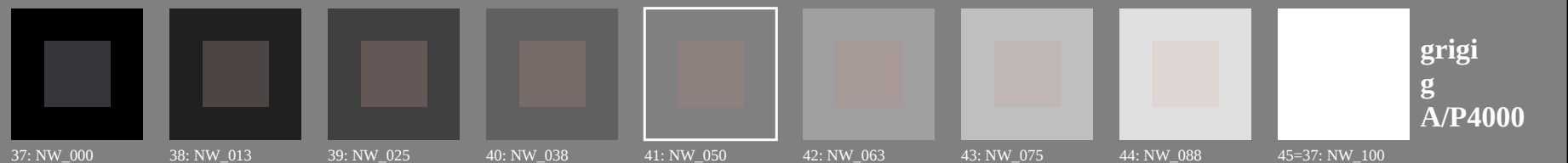
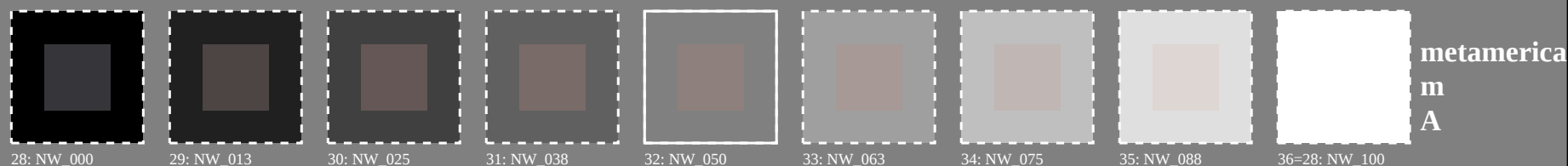


Tavola 3 di campioni di colore per la resa del colore: colori matamerici sotto illuminanti A e P4000; display standard (sRGB)



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

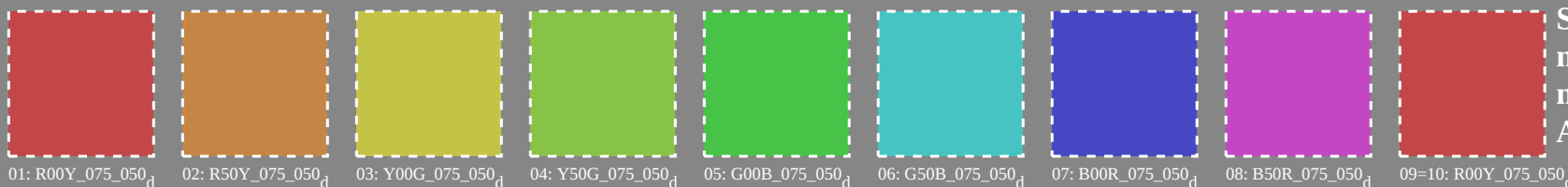
metamerica
m
P4000



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

metamerica
m
P4000

Tavola 3 di campioni di colore per la resa del colore: colori matamerici sotto illuminanti A e P4000; display standard (sRGB); rg



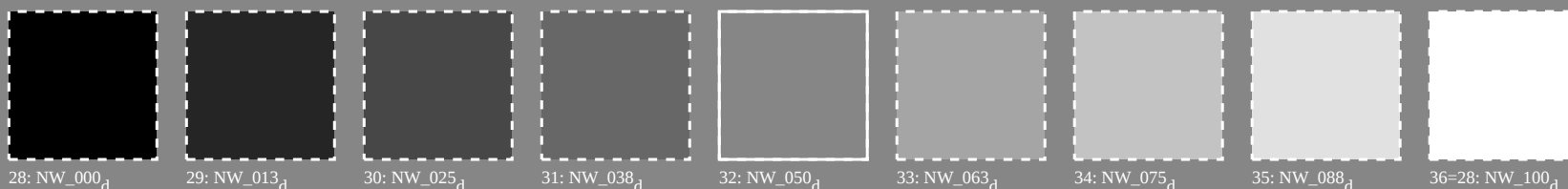
Serie:
metamerica
m
A



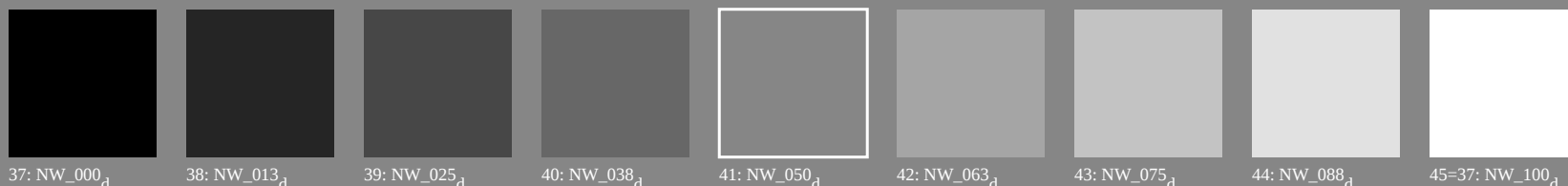
intermedia
z
A/P4000

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

metamerica
m
P4000



metamerica
m
A
Lab*N=0.0, 0.0, 0.0
Lab*W=95.4, 0.0, 0.0



grigi
g
A/P4000

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

metamerica
m
P4000

4-003130-L0 PI310-70

Grafico TUB-PI31; riproduzione del colore
54 colori; metamerica per A&P4000, 3D=0, de=0, sRGB

Input: *rgb/cmyk* -> *rgb_d*
Output: trasferire a *rgb_d*

vedi file simili: <http://farbe.li.tu-berlin.de/PI31/PI31.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n/j	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsi_Md	rgb*Md	LabCh*Md		
0/648	R00Y_100_100d	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5	100.4 39.9 0.0	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
1/657	R13Y_100_100d	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	51.4 74.1 64.9	98.5 41.2	1.0 0.125 0.0	51.5 73.9 64.9	98.3 41.3 0.2	36	1.0 0.116 0.0	51.4 74.1 64.9	98.5 41.2
2/666	R25Y_100_100d	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.7 67.6 65.8	94.4 44.2	1.0 0.25 0.0	54.0 66.7 65.9	93.8 44.6 1.0	42	1.0 0.233 0.0	53.7 67.6 65.8	94.4 44.2
3/675	R38Y_100_100d	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	57.9 56.2 67.9	88.1 50.3	1.0 0.375 0.0	58.2 55.4 67.9	87.7 50.7 0.7	51	1.0 0.366 0.0	57.9 56.2 67.9	88.1 50.3
4/684	R50Y_100_100d	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7 0.0	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
5/693	R63Y_100_100d	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	70.5 24.7 75.4	79.4 71.8	1.0 0.625 0.0	70.1 25.8 75.0	79.3 71.0 1.2	68	1.0 0.633 0.0	70.5 24.7 75.4	79.4 71.8
6/702	R75Y_100_100d	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	78.2 7.8 80.6	81.0 84.4	1.0 0.75 0.0	77.2 9.8 79.7	80.3 82.9 2.3	77	1.0 0.766 0.0	78.2 7.8 80.6	81.0 84.4
7/711	R88Y_100_100d	1.0 0.875 0.0	1.0 1.0 0.5	83	1.0 0.883 0.0	85.3 -6.7 85.5	85.8 94.4	1.0 0.875 0.0	84.8 -5.7 85.0	85.2 93.8 1.1	83	1.0 0.883 0.0	85.3 -6.7 85.5	85.8 94.4
8/720	Y00G_100_100d	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8	1.0 1.0 0.0	92.6 -20.6 90.7	93.0 102.8 0.0	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
9/639	Y13G_100_100d	0.875 1.0 0.0	1.0 1.0 0.5	97	0.883 1.0 0.0	90.5 -32.2 88.3	94.0 110.0	0.875 1.0 0.0	90.4 -33.0 88.1	94.1 110.5 0.8	96	0.883 1.0 0.0	90.5 -32.2 88.3	94.0 110.0
10/558	Y25G_100_100d	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	88.7 -43.3 86.2	96.5 116.6	0.75 1.0 0.0	88.5 -44.9 85.8	96.8 117.6 1.6	102	0.766 1.0 0.0	88.7 -43.3 86.2	96.5 116.6
11/477	Y38G_100_100d	0.625 1.0 0.0	1.0 1.0 0.5	112	0.633 1.0 0.0	87.0 -55.0 84.1	100.5 123.2	0.625 1.0 0.0	86.9 -55.7 83.9	100.7 123.6 0.7	111	0.633 1.0 0.0	87.0 -55.0 84.1	100.5 123.2
12/396	Y50G_100_100d	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3 0.0	119	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
13/315	Y63G_100_100d	0.375 1.0 0.0	1.0 1.0 0.5	128	0.366 1.0 0.0	84.7 -73.2 81.2	109.3 132.0	0.375 1.0 0.0	84.7 -72.8 81.2	109.1 131.8 0.3	128	0.366 1.0 0.0	84.7 -73.2 81.2	109.3 132.0
14/234	Y75G_100_100d	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	84.0 -78.7 80.4	112.5 134.3	0.25 1.0 0.0	84.1 -78.2 80.4	112.2 134.1 0.4	137	0.233 1.0 0.0	84.0 -78.7 80.4	112.5 134.3
15/153	Y88G_100_100d	0.125 1.0 0.0	1.0 1.0 0.5	143	0.116 1.0 0.0	83.7 -81.5 80.0	114.2 135.5	0.125 1.0 0.0	83.7 -81.4 80.0	114.2 135.5 0.1	143	0.116 1.0 0.0	83.7 -81.5 80.0	114.2 135.5
16/72	G00C_100_100d	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0 0.0	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
17/73	G13C_100_100d	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.116	83.6 -82.1 76.8	112.5 136.9	0.0 1.0 0.125	83.6 -82.1 76.5	112.3 137.0 0.2	156	0.0 1.0 0.116	83.6 -82.1 76.8	112.5 136.9
18/74	G25C_100_100d	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.233	83.7 -80.8 70.1	106.9 139.0	0.0 1.0 0.25	83.8 -80.5 69.1	106.1 139.3 1.0	162	0.0 1.0 0.233	83.7 -80.8 70.1	106.9 139.0
19/75	G38C_100_100d	0.0 1.0 0.375	1.0 1.0 0.5	172	0.0 1.0 0.366	84.0 -78.0 58.8	97.7 142.9	0.0 1.0 0.375	84.0 -77.7 58.1	97.1 143.2 0.7	171	0.0 1.0 0.366	84.0 -78.0 58.8	97.7 142.9
20/76	G50C_100_100d	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	84.3 -73.7 44.9	86.4 148.6	0.0 1.0 0.5	84.3 -73.7 44.9	86.3 148.6 0.0	180	0.0 1.0 0.5	84.3 -73.7 44.9	86.4 148.6
21/77	G63C_100_100d	0.0 1.0 0.625	1.0 1.0 0.5	188	0.0 1.0 0.633	84.8 -68.1 29.5	74.3 156.5	0.0 1.0 0.625	84.7 -68.5 30.6	75.0 155.9 1.1	188	0.0 1.0 0.633	84.8 -68.1 29.5	74.3 156.5
22/78	G75C_100_100d	0.0 1.0 0.75	1.0 1.0 0.5	196	0.0 1.0 0.766	85.4 -61.2 13.7	62.8 167.3	0.0 1.0 0.75	85.3 -62.0 15.8	64.0 165.6 2.3	197	0.0 1.0 0.766	85.4 -61.2 13.7	62.8 167.3
23/79	G88C_100_100d	0.0 1.0 0.875	1.0 1.0 0.5	203	0.0 1.0 0.883	86.1 -54.1 0.0	54.1 180.0	0.0 1.0 0.875	86.0 -54.5 1.0	54.5 178.8 1.1	203	0.0 1.0 0.883	86.1 -54.1 0.0	54.1 180.0
24/80	C00B_100_100d	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3 0.0	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
25/71	C13B_100_100d	0.0 0.875 1.0	1.0 1.0 0.5	217	0.0 0.883 1.0	78.5 -33.4 -26.3	42.5 218.2	0.0 0.875 1.0	77.9 -32.3 -27.0	42.1 219.8 1.3	216	0.0 0.883 1.0	78.5 -33.4 -26.3	42.5 218.2
26/62	C25B_100_100d	0.0 0.75 1.0	1.0 1.0 0.5	224	0.0 0.766 1.0	70.2 -19.5 -39.3	43.9 243.6	0.0 0.75 1.0	69.1 -17.0 -40.7	44.1 247.2 3.0	222	0.0 0.766 1.0	70.2 -19.5 -39.3	43.9 243.6
27/53	C38B_100_100d	0.0 0.625 1.0	1.0 1.0 0.5	232	0.0 0.633 1.0	60.9 -1.5 -53.9	53.9 268.3	0.0 0.625 1.0	60.3 -0.1 -54.6	54.6 269.8 1.7	231	0.0 0.633 1.0	60.9 -1.5 -53.9	53.9 268.3
28/44	C50B_100_100d	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0 0.0	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
29/35	C63B_100_100d	0.0 0.375 1.0	1.0 1.0 0.5	248	0.0 0.366 1.0	43.4 38.7 -82.0	90.7 295.3	0.0 0.375 1.0	43.8 37.6 -81.2	89.5 294.8 1.4	248	0.0 0.366 1.0	43.4 38.7 -82.0	90.7 295.3
30/26	C75B_100_100d	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6	0.0 0.25 1.0	37.1 55.9 -92.3	107.9 301.1 2.1	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6
31/17	C88B_100_100d	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.116 1.0	32.3 70.0 -100.3	122.3 304.9	0.0 0.125 1.0	32.4 69.6 -100.0	121.9 304.8 0.5	263	0.0 0.116 1.0	32.3 70.0 -100.3	122.3 304.9
32/8	B00M_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2 0.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
33/89	B13M_100_100d	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	30.9 76.2 -102.5	127.8 306.6	0.125 0.0 1.0	31.0 76.2 -102.5	127.7 306.6 0.0	276	0.116 0.0 1.0	30.9 76.2 -102.5	127.8 306.6
34/170	B25M_100_100d	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4	0.25 0.0 1.0	32.6 76.8 -99.8	125.9 307.5 0.4	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
35/251	B38M_100_100d	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	34.9 77.9 -95.7	123.0 309.1	0.375 0.0 1.0	35.1 77.9 -95.5	123.3 309.2 0.3	291	0.366 0.0 1.0	34.9 77.9 -95.7	123.0 309.1
36/332	B50M_100_100d	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.4 311.6	0.5 0.0 1.0	38.5 79.8 -89.7	120.1 311.6 0.0	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.4 311.6
37/413	B63M_100_100d	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	43.0 82.7 -82.2	116.6 315.1	0.625 0.0 1.0	42.7 82.5 -82.8	116.8 314.8 0.6	308	0.633 0.0 1.0	43.0 82.7 -82.2	116.6 315.1
38/494	B75M_100_100d	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4	0.75 0.0 1.0	47.2 85.8 -75.1	114.1 318.8 1.3	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4
39/575	B88M_100_100d	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	52.5 90.1 -66.3	111.9 323.6	0.875 0.0 1.0	52.1 89.8 -66.9	112.0 323.3 0.7	323	0.883 0.0 1.0	52.5 90.1 -66.3	111.9 323.6
40/656	M00R_100_100d	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	1.0 0.0 1.0	57.2 94.3 -58.4	111.0 328.2 0.0	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
41/655	M13R_100_100d	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	55.7 90.6 -44.8	101.1 333.6	1.0 0.0 0.875	55.6 90.3 -43.9	100.4 334.0 0.9	336	1.0 0.0 0.883	55.7 90.6 -44.8	101.1 333.6
42/654	M25R_100_100d	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	54.4 87.3 -30.6	92.5 340.6	1.0 0.0 0.75	54.2 86.7 -28.6	91.3 341.6 2.0	342	1.0 0.0 0.766	54.4 87.3 -30.6	92.5 340.6
43/653	M38R_100_100d	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	53.0 83.9 -13.6	85.0 350.7	1.0 0.0 0.625	53.0 83.6 -12.6	84.6 351.4 1.0	351	1.0 0.0 0.633	53.0 83.9 -13.6	85.0 350.7
44/652	M50R_100_100d	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	0.0	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
45/651	M63R_100_100d	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	51.3 79.3 22.7	82.5 16.0	1.0 0.0 0.375	51.3 79.2 21.6	82.1 15.2 1.1	368	1.0 0.0 0.366	51.3 79.3 22.7	82.5 16.0
46/650	M75R_100_100d	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8	1.0 0.0 0.25	50.8 77.9 39.2	87.2 26.6 2.0	377	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8
47/649	M88R_100_100d	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	50.5 77.2 55.6	95.1 35.7	1.0 0.0 0.125	50.6 77.2 54.9	94.8 35.4 0.6	383	1.0 0.0 0.116	50.5 77.2 55.6	95.1 35.7
48/648	R00Y_100_100d	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5	100.4 39.9 0.0	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
49/0	NW_000d	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0
50/91	NW_013d	0.125 0.125 0.125	0.125 0.0 0.125	360										

vedi file simili: <http://farbe.li.tu-berlin.de/PI31/PI31.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n/j	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsi_Md	rgb*Md	LabCh*Ma
0/648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.0	50.4	76.9	64.5	100.4	40.0
1/666	R25Y_100_100a	1.0	0.25	0.0	1.0	1.0	0.233	53.7	67.6	65.8	94.4	44.2
2/684	R50Y_100_100a	1.0	0.5	0.0	1.0	1.0	0.5	63.6	41.3	71.0	82.2	59.7
3/702	R75Y_100_100a	1.0	0.75	0.0	1.0	1.0	0.766	78.2	9.8	80.6	81.0	84.4
4/720	Y00G_100_100a	1.0	1.0	0.0	1.0	1.0	1.0	92.6	-20.6	90.7	93.0	102.8
5/558	Y25G_100_100a	0.75	1.0	0.0	1.0	1.0	0.5	88.7	-43.3	86.2	96.5	116.6
6/396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	85.7	-65.2	82.4	105.1	128.3
7/234	Y75G_100_100a	0.25	1.0	0.0	1.0	1.0	0.5	84.0	-78.7	80.4	112.2	134.1
8/72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	83.6	-82.7	79.8	115.0	136.0
9/72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	83.6	-82.7	79.8	115.0	136.0
10/76	G25B_100_100a	0.0	1.0	0.5	1.0	1.0	0.5	84.3	-73.7	44.9	86.4	148.6
11/80	G50B_100_100a	0.0	1.0	1.0	1.0	1.0	0.5	86.8	-46.1	-13.5	48.1	196.3
12/44	G75B_100_100a	0.0	0.5	1.0	1.0	1.0	0.5	51.7	18.3	-68.3	70.7	285.0
13/8	B00M_100_100a	0.0	0.0	1.0	1.0	1.0	0.5	30.3	76.0	-103.5	128.5	306.2
14/332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	38.5	79.8	-89.7	120.1	311.6
15/656	B50R_100_100a	1.0	0.0	1.0	1.0	1.0	0.5	57.2	94.3	-58.4	111.0	328.2
16/652	B75R_100_100a	1.0	0.0	0.5	1.0	1.0	0.5	52.0	81.1	4.1	81.2	2.9
17/648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.5	50.4	76.9	64.5	100.4	40.0
18/688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	72.9	38.4	32.2	50.2	40.0
19/706	R50Y_100_050a	1.0	0.75	0.5	1.0	0.5	0.75	79.5	20.6	35.5	41.1	59.7
20/724	Y00G_100_050a	1.0	1.0	0.5	1.0	0.5	0.75	94.0	-10.3	45.3	46.5	102.8
21/562	Y50G_100_050a	0.75	1.0	0.5	1.0	0.5	0.75	90.5	-32.6	41.2	52.5	128.3
22/400	G00B_100_050a	0.5	1.0	0.5	1.0	0.5	0.75	89.5	-41.3	39.9	57.5	136.0
23/404	G50B_100_050a	0.5	1.0	1.0	1.0	0.5	0.75	81.1	-23.0	-6.7	24.0	196.3
24/368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	62.8	38.0	-51.7	64.2	306.2
25/692	B50R_100_050a	1.0	0.5	1.0	1.0	0.5	0.75	76.3	47.1	-29.2	55.4	328.2
26/688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	72.9	38.4	32.2	50.2	40.0
27/506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	49.0	38.4	32.2	50.2	40.0
28/524	R50Y_075_050a	0.75	0.5	0.25	0.75	0.5	0.5	55.6	20.6	35.5	41.1	59.7
29/542	Y00G_075_050a	0.75	0.75	0.25	0.75	0.5	0.5	70.1	-12.3	45.3	46.5	102.8
30/380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	66.7	-32.6	41.2	52.5	128.3
31/218	G00B_075_050a	0.25	0.75	0.25	0.75	0.5	0.5	65.6	33.9	57.5	136.0	306.2
32/222	G50B_075_050a	0.25	0.75	0.75	0.75	0.5	0.5	67.5	-23.0	-6.7	24.0	196.3
33/186	B00R_075_050a	0.25	0.25	0.75	0.75	0.5	0.5	39.0	38.0	-51.7	64.2	306.2
34/510	B50R_075_050a	0.75	0.25	0.75	0.75	0.5	0.5	52.5	47.1	-29.2	55.4	328.2
35/506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	49.0	38.4	32.2	50.2	40.0
36/324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.25	25.2	38.4	32.2	50.2	40.0
37/342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	31.8	20.6	35.5	41.1	59.7
38/360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	46.8	-10.3	45.3	46.5	102.8
39/198	Y50G_050_050a	0.25	0.5	0.0	0.5	0.5	0.25	42.8	-32.6	41.2	52.5	128.3
40/36	G00B_050_050a	0.0	0.5	0.0	0.5	0.5	0.25	41.8	-41.3	39.9	57.5	136.0
41/40	G50B_050_050a	0.0	0.5	0.5	0.5	0.5	0.25	43.4	-23.0	-6.7	24.0	196.3
42/4	B00R_050_050a	0.0	0.0	0.5	0.5	0.5	0.25	15.1	38.0	-51.7	64.2	306.2
43/328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.25	28.6	47.1	-29.2	55.4	328.2
44/324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.25	25.2	38.4	32.2	50.2	40.0
45/0	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
46/91	NW_013a	0.125	0.125	0.125	0.125	0.0	0.125	11.9	0.0	0.0	0.0	0.0
47/182	NW_025a	0.25	0.25	0.25	0.25	0.0	0.25	23.8	0.0	0.0	0.0	0.0
48/273	NW_038a	0.375	0.375	0.375	0.375	0.0	0.375	35.7	0.0	0.0	0.0	0.0
49/364	NW_050a	0.5	0.5	0.5	0.5	0.0	0.5	47.7	0.0	0.0	0.0	0.0
50/455	NW_063a	0.625	0.625	0.625	0.625	0.0	0.625	59.6	0.0	0.0	0.0	0.0
51/546	NW_075a	0.75	0.75	0.75	0.75	0.0	0.75	71.5	0.0	0.0	0.0	0.0
52/637	NW_088a	0.875	0.875	0.875	0.875	0.0	0.875	83.4	0.0	0.0	0.0	0.0
53/728	NW_100a	1.0	1.0	1.0	1.0	0.0	1.0	95.4	0.0	0.0	0.0	0.0

delta E* = 6.5

Grafico TUB-PI31; riproduzione del colore
colori e la differenza, ΔE^* , 3D=0, de=0, sRGB

Input: $rgb/cmyk \rightarrow rgb_d$
Output: trasferire a rgb_d

iscrizione TUB: 20160501-PI31/PI31L0NP.PDF /.PS
Applicazione per la misura dell'output su display, nessuna separazione
TUB materiale: code=rha4ta

vedi file simili: <http://farbe.li.tu-berlin.de/PI31/PI31.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n\j	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md
0	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	1.0
1	B00R_012_012a	0.0	0.0	0.125	0.125	0.125	0.062	0.360	0.0	0.0	1.0	1.0
2	B00R_025_025a	0.0	0.0	0.25	0.25	0.25	0.125	0.270	0.0	0.0	1.0	1.0
3	B00R_037_037a	0.0	0.0	0.375	0.375	0.375	0.187	0.270	0.0	0.0	1.0	1.0
4	B00R_050_050a	0.0	0.0	0.5	0.5	0.5	0.25	0.270	0.0	0.0	1.0	1.0
5	B00R_062_062a	0.0	0.0	0.625	0.625	0.625	0.312	0.270	0.0	0.0	1.0	1.0
6	B00R_075_075a	0.0	0.0	0.75	0.75	0.75	0.375	0.270	0.0	0.0	1.0	1.0
7	B00R_087_087a	0.0	0.0	0.875	0.875	0.875	0.437	0.270	0.0	0.0	1.0	1.0
8	B00R_100_100a	0.0	0.0	1.0	1.0	1.0	0.5	0.270	0.0	0.0	1.0	1.0
9	G00B_012_012a	0.0	0.125	0.0	0.125	0.125	0.062	0.150	0.0	0.125	0.0	0.125
10	G50B_012_012a	0.0	0.125	0.125	0.125	0.125	0.062	0.210	0.0	0.125	0.125	0.125
11	G75B_025_025a	0.0	0.125	0.25	0.25	0.25	0.125	0.240	0.0	0.125	0.25	0.125
12	G84B_037_037a	0.0	0.125	0.375	0.375	0.375	0.187	0.251	0.0	0.125	0.375	0.187
13	G88B_050_050a	0.0	0.125	0.5	0.5	0.5	0.25	0.256	0.0	0.125	0.5	0.25
14	G90B_062_062a	0.0	0.125	0.625	0.625	0.625	0.312	0.259	0.0	0.125	0.625	0.312
15	G92B_075_075a	0.0	0.125	0.75	0.75	0.75	0.375	0.261	0.0	0.125	0.75	0.375
16	G93B_087_087a	0.0	0.125	0.875	0.875	0.875	0.437	0.262	0.0	0.125	0.875	0.437
17	G94B_100_100a	0.0	0.125	1.0	1.0	1.0	0.5	0.263	0.0	0.125	1.0	0.5
18	G00B_025_025a	0.0	0.25	0.0	0.25	0.25	0.125	0.150	0.0	0.25	0.0	0.25
19	G25B_025_025a	0.0	0.25	0.125	0.25	0.25	0.125	0.180	0.0	0.25	0.125	0.125
20	G50B_025_025a	0.0	0.25	0.25	0.25	0.25	0.125	0.210	0.0	0.25	0.25	0.125
21	G65B_037_037a	0.0	0.25	0.375	0.375	0.375	0.187	0.229	0.0	0.25	0.375	0.187
22	G75B_050_050a	0.0	0.25	0.5	0.5	0.5	0.25	0.240	0.0	0.25	0.5	0.25
23	G80B_062_062a	0.0	0.25	0.625	0.625	0.625	0.312	0.247	0.0	0.25	0.625	0.312
24	G84B_075_075a	0.0	0.25	0.75	0.75	0.75	0.375	0.251	0.0	0.25	0.75	0.375
25	G86B_087_087a	0.0	0.25	0.875	0.875	0.875	0.437	0.254	0.0	0.25	0.875	0.437
26	G88B_100_100a	0.0	0.25	1.0	1.0	1.0	0.5	0.256	0.0	0.25	1.0	0.5
27	G00B_037_037a	0.0	0.375	0.0	0.375	0.375	0.187	0.150	0.0	0.375	0.0	0.375
28	G15B_037_037a	0.0	0.375	0.125	0.375	0.375	0.187	0.169	0.0	0.375	0.125	0.187
29	G34B_037_037a	0.0	0.375	0.25	0.375	0.375	0.187	0.191	0.0	0.375	0.25	0.187
30	G50B_037_037a	0.0	0.375	0.375	0.375	0.375	0.187	0.210	0.0	0.375	0.375	0.187
31	G61B_050_050a	0.0	0.375	0.5	0.5	0.5	0.25	0.224	0.0	0.375	0.5	0.25
32	G69B_062_062a	0.0	0.375	0.625	0.625	0.625	0.312	0.233	0.0	0.375	0.625	0.312
33	G75B_075_075a	0.0	0.375	0.75	0.75	0.75	0.375	0.240	0.0	0.375	0.75	0.375
34	G79B_087_087a	0.0	0.375	0.875	0.875	0.875	0.437	0.245	0.0	0.375	0.875	0.437
35	G81B_100_100a	0.0	0.375	1.0	1.0	1.0	0.5	0.248	0.0	0.375	1.0	0.5
36	G00B_050_050a	0.0	0.5	0.0	0.5	0.5	0.25	0.150	0.0	0.5	0.0	0.5
37	G11B_050_050a	0.0	0.5	0.125	0.5	0.5	0.25	0.164	0.0	0.5	0.125	0.125
38	G25B_050_050a	0.0	0.5	0.25	0.5	0.5	0.25	0.180	0.0	0.5	0.25	0.125
39	G38B_050_050a	0.0	0.5	0.375	0.5	0.5	0.25	0.196	0.0	0.5	0.375	0.125
40	G50B_050_050a	0.0	0.5	0.5	0.5	0.5	0.25	0.210	0.0	0.5	0.5	0.25
41	G59B_062_062a	0.0	0.5	0.625	0.625	0.625	0.312	0.221	0.0	0.5	0.625	0.312
42	G65B_075_075a	0.0	0.5	0.75	0.75	0.75	0.375	0.229	0.0	0.5	0.75	0.375
43	G70B_087_087a	0.0	0.5	0.875	0.875	0.875	0.437	0.235	0.0	0.5	0.875	0.437
44	G75B_100_100a	0.0	0.5	1.0	1.0	1.0	0.5	0.240	0.0	0.5	1.0	0.5
45	G00B_062_062a	0.0	0.625	0.0	0.625	0.625	0.312	0.150	0.0	0.625	0.0	0.625
46	G09B_062_062a	0.0	0.625	0.125	0.625	0.625	0.312	0.161	0.0	0.625	0.125	0.125
47	G19B_062_062a	0.0	0.625	0.25	0.625	0.625	0.312	0.173	0.0	0.625	0.25	0.125
48	G30B_062_062a	0.0	0.625	0.375	0.625	0.625	0.312	0.187	0.0	0.625	0.375	0.125
49	G40B_062_062a	0.0	0.625	0.5	0.625	0.625	0.312	0.199	0.0	0.625	0.5	0.125
50	G50B_062_062a	0.0	0.625	0.625	0.625	0.625	0.312	0.210	0.0	0.625	0.625	0.125
51	G57B_075_075a	0.0	0.625	0.75	0.75	0.75	0.375	0.219	0.0	0.625	0.75	0.375
52	G63B_087_087a	0.0	0.625	0.875	0.875	0.875	0.437	0.226	0.0	0.625	0.875	0.437
53	G68B_100_100a	0.0	0.625	1.0	1.0	1.0	0.5	0.232	0.0	0.625	1.0	0.5
54	G00B_075_075a	0.0	0.75	0.0	0.75	0.75	0.375	0.150	0.0	0.75	0.0	0.75
55	G07B_075_075a	0.0	0.75	0.125	0.75	0.75	0.375	0.159	0.0	0.75	0.125	0.125
56	G15B_075_075a	0.0	0.75	0.25	0.75	0.75	0.375	0.169	0.0	0.75	0.25	0.125
57	G25B_075_075a	0.0	0.75	0.375	0.75	0.75	0.375	0.180	0.0	0.75	0.375	0.125
58	G34B_075_075a	0.0	0.75	0.5	0.75	0.75	0.375	0.191	0.0	0.75	0.5	0.125
59	G42B_075_075a	0.0	0.75	0.625	0.75	0.75	0.375	0.201	0.0	0.75	0.625	0.125
60	G50B_075_075a	0.0	0.75	0.75	0.75	0.75	0.375	0.210	0.0	0.75	0.75	0.125
61	G56B_087_087a	0.0	0.75	0.875	0.875	0.875	0.437	0.218	0.0	0.75	0.875	0.437
62	G61B_100_100a	0.0	0.75	1.0	1.0	1.0	0.5	0.224	0.0	0.75	1.0	0.5
63	G00B_087_087a	0.0	0.875	0.0	0.875	0.875	0.437	0.150	0.0	0.875	0.0	0.875
64	G06B_087_087a	0.0	0.875	0.125	0.875	0.875	0.437	0.158	0.0	0.875	0.125	0.125
65	G13B_087_087a	0.0	0.875	0.25	0.875	0.875	0.437	0.166	0.0	0.875	0.25	0.125
66	G20B_087_087a	0.0	0.875	0.375	0.875	0.875	0.437	0.175	0.0	0.875	0.375	0.125
67	G29B_087_087a	0.0	0.875	0.5	0.875	0.875	0.437	0.185	0.0	0.875	0.5	0.125
68	G36B_087_087a	0.0	0.875	0.625	0.875	0.875	0.437	0.194	0.0	0.875	0.625	0.125
69	G43B_087_087a	0.0	0.875	0.75	0.875	0.875	0.437	0.202	0.0	0.875	0.75	0.125
70	G50B_087_087a	0.0	0.875	0.875	0.875	0.875	0.437	0.210	0.0	0.875	0.875	0.125
71	G55B_100_100a	0.0	0.875	1.0	1.0	1.0	0.5	0.217	0.0	0.875	1.0	0.5
72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	0.150	0.0	1.0	0.0	1.0
73	G05B_100_100a	0.0	1.0	0.125	1.0	1.0	0.5	0.157	0.0	1.0	0.125	0.125
74	G11B_100_100a	0.0	1.0	0.25	1.0	1.0	0.5	0.164	0.0	1.0	0.25	0.125
75	G18B_100_100a	0.0	1.0	0.375	1.0	1.0	0.5	0.172	0.0	1.0	0.375	0.125
76	G25B_100_100a	0.0	1.0	0.5	1.0	1.0	0.5	0.180	0.0	1.0	0.5	0.125
77	G31B_100_100a	0.0	1.0	0.625	1.0	1.0	0.5	0.188	0.0	1.0	0.625	0.125
78	G38B_100_100a	0.0	1.0	0.75	1.0	1.0	0.5	0.196	0.0	1.0	0.75	0.125
79	G44B_100_100a	0.0	1.0	0.875	1.0	1.0	0.5	0.203	0.0	1.0	0.875	0.125
80	G50B_100_100a	0.0	1.0	1.0	1.0	1.0	0.5	0.210	0.0	1.0	1.0	0.5

delta E** = 4.6

Grafico TUB-PI31; riproduzione del colore
colori e la differenza, ΔE^* , 3D=0, de=0, sRGB

Input: $rgb/cmyk \rightarrow rgb_d$
Output: trasferire a rgb_d

iscrizione TUB: 20160501-PI31/PI31L0NP.PDF /PS
Applicazione per la misura dell'output su display, nessuna separazione
TUB materiale: code=rha4ra

vedi file simili: <http://farbe.li.tu-berlin.de/PI31/PI31.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
81	R00Y_012_012a	0.125 0.0 0.0	0.125 0.125 0.062	390	0.125 0.0 0.0	6.3 9.6 8.0	12.5 40.0	0.125 0.0 0.0	2.4 10.9 3.8	11.6 19.4 5.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
82	B50R_012_012a	0.125 0.0 0.125	0.125 0.125 0.062	330	0.125 0.0 0.125	7.1 11.7 -7.3	13.8 328.2	0.125 0.0 0.125	3.2 16.7 -11.6	20.4 325.1 7.6	330	1.0 0.0 0.0	57.2 94.3 -58.4	110.9 328.2
83	B25R_025_025a	0.125 0.0 0.25	0.25 0.25 0.125	300	0.125 0.0 0.25	9.6 19.9 -22.4	30.0 311.6	0.125 0.0 0.25	5.3 28.5 -31.2	42.3 312.3 13.0	300	0.5 0.0 0.0	38.5 79.8 -89.7	120.0 311.6
84	B15R_037_037a	0.125 0.0 0.375	0.375 0.375 0.187	289	0.118 0.0 0.375	12.7 29.0 -36.5	46.7 308.4	0.125 0.0 0.375	9.0 38.1 -46.3	60.0 309.4 13.8	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
85	B11R_050_050a	0.125 0.0 0.5	0.5 0.5 0.25	284	0.116 0.0 0.5	16.1 38.3 -50.0	63.1 307.4	0.125 0.0 0.5	13.4 46.1 -59.0	74.9 307.9 12.1	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
86	B09R_062_062a	0.125 0.0 0.625	0.625 0.625 0.312	281	0.114 0.0 0.625	19.8 47.8 -63.2	79.3 307.0	0.125 0.0 0.625	17.9 53.9 -70.7	88.9 307.3 9.8	279	0.183 0.0 1.0	31.7 76.5 -101.2	126.9 307.0
87	B07R_075_075a	0.125 0.0 0.75	0.75 0.75 0.375	279	0.112 0.0 0.75	23.5 57.2 -76.4	95.5 306.8	0.125 0.0 0.75	22.3 61.5 -81.7	102.3 306.9 6.9	278	0.15 0.0 1.0	31.3 76.3 -101.9	127.4 306.8
88	B06R_087_087a	0.125 0.0 0.875	0.875 0.875 0.437	278	0.110 0.0 0.875	27.2 66.7 -89.5	111.6 306.7	0.125 0.0 0.875	26.7 69.0 -92.3	115.2 306.7 3.6	277	0.133 0.0 1.0	31.1 76.3 -102.3	127.6 306.7
89	B05R_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	30.9 76.2 -102.5	127.8 306.6	0.125 0.0 1.0	31.0 76.2 -102.5	127.7 306.6 0.0	276	0.116 0.0 1.0	30.9 76.2 -102.5	127.8 306.6
90	Y00G_012_012a	0.125 0.125 0.0	0.125 0.125 0.062	90	0.125 0.125 0.0	11.5 -25.5 11.3	11.6 102.8	0.125 0.125 0.0	10.4 -5.0 15.4	16.2 108.0 4.8	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
91	NW_012a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	11.9 0.0 0.0	0.0 0.0	0.0 0.125 0.125	11.0 0.0 0.0	0.0 325.7 0.8	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0
92	B00R_025_012a	0.125 0.125 0.25	0.25 0.125 0.187	270	0.124 0.124 0.25	15.7 9.5 -12.9	16.0 306.2	0.125 0.125 0.25	12.6 9.6 -19.5	21.8 296.2 7.3	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
93	B00R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.124 0.375	19.5 19.0 -25.8	32.1 306.2	0.125 0.125 0.375	15.0 21.1 -36.5	42.1 300.0 11.6	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
94	B00R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.124 0.5	23.3 28.5 -38.8	48.1 306.2	0.125 0.125 0.5	18.1 32.4 -51.3	60.0 302.2 14.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
95	B00R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	27.1 38.0 -51.7	64.2 306.2	0.125 0.125 0.625	21.6 42.8 -64.6	77.5 303.5 14.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
96	B00R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	30.9 47.5 -64.7	80.3 306.2	0.125 0.125 0.75	25.3 52.5 -76.8	93.0 304.3 14.2	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
97	B00R_087_050a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	34.7 57.0 -77.6	96.3 306.2	0.125 0.125 0.875	29.1 61.5 -88.2	107.5 304.8 12.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
98	B00R_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	38.5 66.5 -90.6	112.4 306.2	0.125 0.125 1.0	33.0 69.9 -99.0	121.3 305.2 10.6	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
99	Y50G_025_025a	0.125 0.25 0.0	0.25 0.25 0.125	120	0.125 0.25 0.0	21.4 -16.3 20.6	26.2 128.3	0.125 0.25 0.0	21.9 -22.3 29.7	37.2 126.9 10.9	119	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
100	G00B_025_012a	0.125 0.25 0.125	0.25 0.125 0.187	150	0.124 0.25 0.124	22.3 -10.3 9.9	14.3 136.0	0.125 0.25 0.125	22.2 -18.8 15.2	24.2 141.0 10.0	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
101	G50B_025_012a	0.125 0.25 0.25	0.25 0.125 0.187	210	0.124 0.25 0.25	22.7 -5.7 -1.6	6.0 196.3	0.125 0.25 0.25	23.0 -11.2 -3.5	11.7 197.3 5.7	210	0.0 1.0 0.0	86.8 -46.1 -13.5	48.1 196.3
102	G75B_037_025a	0.125 0.25 0.375	0.375 0.25 0.25	240	0.124 0.25 0.375	24.8 4.5 -17.0	17.6 285.0	0.125 0.25 0.375	24.4 -0.5 -21.5	21.5 268.6 6.7	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
103	G84B_050_037a	0.125 0.25 0.5	0.5 0.375 0.312	251	0.124 0.243 0.5	27.2 17.1 -32.5	36.7 297.8	0.125 0.25 0.5	26.3 11.5 -37.9	39.6 286.9 7.8	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
104	G88B_062_050a	0.125 0.25 0.625	0.625 0.5 0.375	256	0.125 0.241 0.625	30.1 28.8 -46.7	54.8 301.6	0.125 0.25 0.625	28.7 23.7 -52.9	58.0 294.1 8.1	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6
105	G90B_075_062a	0.125 0.25 0.75	0.75 0.625 0.437	259	0.125 0.239 0.75	33.5 39.4 -60.3	72.1 303.1	0.125 0.25 0.75	31.4 35.4 -66.7	75.5 297.7 7.7	260	0.0 0.183 1.0	34.6 63.0 -96.6	115.3 303.1
106	G92B_087_050a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.237 0.875	36.9 50.0 -73.9	88.3 304.0	0.125 0.25 0.875	34.4 46.3 -79.5	92.0 300.2 7.1	262	0.0 0.15 1.0	33.4 66.7 -98.6	119.1 304.0
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.241 1.0	40.6 60.0 -87.1	105.8 304.5	0.125 0.25 1.0	37.6 56.5 -91.4	107.5 301.7 6.3	262	0.0 0.133 1.0	32.8 68.6 -99.6	120.9 304.5
108	Y68G_037_037a	0.125 0.375 0.0	0.375 0.375 0.187	131	0.118 0.375 0.0	31.6 -28.2 30.3	41.4 132.9	0.125 0.375 0.0	33.1 -35.2 39.6	53.0 131.5 11.7	131	0.316 1.0 0.0	84.4 -75.3 80.9	110.6 132.9
109	G00B_037_025a	0.125 0.375 0.125	0.375 0.25 0.25	150	0.124 0.375 0.124	32.8 -20.6 19.9	28.7 136.0	0.125 0.375 0.125	33.3 -22.9 28.6	43.6 138.9 14.9	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
110	G25B_037_025a	0.125 0.375 0.25	0.375 0.25 0.25	180	0.124 0.375 0.25	33.0 -18.4 11.2	21.6 148.6	0.125 0.375 0.25	33.8 -27.4 11.9	29.9 156.5 9.0	180	0.0 1.0 0.5	84.3 -73.7 44.9	86.4 148.6
111	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	33.6 -11.5 -3.3	12.0 196.3	0.125 0.375 0.375	34.7 -18.9 -5.7	19.8 196.8 7.8	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
112	G65B_050_037a	0.125 0.375 0.5	0.5 0.375 0.312	229	0.124 0.381 0.5	36.0 -3.4 -18.3	18.6 259.3	0.125 0.375 0.5	35.9 -8.3 -22.7	24.1 249.7 6.5	228	0.0 0.683 1.0	64.4 -9.2 -48.8	49.7 259.3
113	G75B_062_050a	0.125 0.375 0.625	0.625 0.5 0.375	240	0.125 0.375 0.625	37.8 9.1 -34.1	35.3 285.0	0.125 0.375 0.625	37.5 3.3 -38.6	38.7 274.9 7.3	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.364 0.75	39.6 22.6 -50.3	55.1 294.2	0.125 0.375 0.75	39.5 15.3 -53.5	55.6 285.9 7.9	247	0.0 0.383 1.0	44.3 36.2 -80.5	88.2 294.2
115	G84B_087_050a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.362 0.875	42.4 34.3 -65.0	73.5 297.8	0.125 0.375 0.875	41.7 27.1 -67.4	72.7 291.5 7.5	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
116	G86B_100_087a	0.125 0.375 1.0	1.0 0.875 0.562	254	0.125 0.358 1.0	45.2 46.6 -79.6	92.2 300.3	0.125 0.375 1.0	44.2 38.6 -80.5	89.3 295.5 8.1	255	0.0 0.266 1.0	38.0 53.3 -91.0	105.4 300.3
117	Y76G_050_050a	0.125 0.5 0.0	0.5 0.5 0.25	136	0.116 0.5 0.0	42.0 -39.3 40.2	56.2 134.3	0.125 0.5 0.0	43.9 -45.9 48.2	66.6 133.6 10.6	137	0.233 1.0 0.0	84.0 -78.7 80.4	112.5 134.3
118	G00B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.124	43.2 -31.0 29.9	43.1 136.0	0.125 0.5 0.125	44.1 -44.3 40.1	59.8 137.8 16.7	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.243	43.3 -29.7 23.6	38.0 141.4	0.125 0.5 0.25	44.4 -40.3 25.7	47.9 147.4 10.9	168	0.0 1.0 0.316	83.9 -79.2 63.1	101.3 141.4
120	G34B_050_037a	0.125 0.5 0.375	0.5 0.375 0.312	191	0.124 0.5 0.381	43.8 -24.7 8.7	26.2 160.4	0.125 0.5 0.375	45.0 -33.8 9.2	35.1 164.7 9.2	191	0.0 1.0 0.683	85.0 -65.8 23.4	69.9 160.4
121	G50B_050_037a	0.125 0.5 0.5	0.5 0.375 0.312	210	0.124 0.5 0.5	44.5 -17.3 -5.0	18.0 196.3	0.125 0.5 0.5	45.9 -25.2 -7.5	26.3 196.6 8.3	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
122	G61B_062_050a	0.125 0.5 0.625	0.625 0.5 0.375	224	0.125 0.508 0.625	47.0 -9.7 -19.6	21.9 243.6	0.125 0.5 0.625	47.0 -14.9 -23.7	28.0 237.7 6.6	222	0.0 0.766 1.0	70.2 -19.5 -39.3	43.9 243.6
123	G69B_075_062a	0.125 0.5 0.75	0.75 0.625 0.437	233	0.125 0.51 0.75	49.2 0.5 -34.8	34.8 270.8	0.125 0.5 0.75	48.4 -3.8 -39.2	39.3 264.4 6.2	232	0.0 0.616 1.0	59.7 0.8 -55.6	55.7 270.8
124	G75B_087_075a	0.125 0.5 0.875	0.875 0.75 0.5	240	0.125 0.5 0.875	50.7 13.7 -51.2	53.0 285.0	0.125 0.5 0.875	50.1 7.7 -53.8	54.4 278.2 6.5	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
125	G79B_100_087a	0.125 0.5 1.0	1.0 0.875 0.562	245	0.125 0.489 1.0	52.6 26.8 -67.7	72.8 291.5	0.125 0.5 1.0	52.0 19.4 -67.8	70.5 285.9 7.4	245	0.0 0.416 1.0	46.5 30.6 -77.4	83.2 291.5
126	Y81G_062_062a	0.125 0.625 0.0	0.625 0.625 0.312	139	0.114 0.625 0.0	52.4 -49.9 50.1	70.8 134.8	0.125 0.625 0.0	54.3 -55.6 56.5	79.3 134.5 8.7	149	0.183 1.0 0.0	83.9 -79.9 80.2	113.3 134.8
127	G00B_062_050a	0.125 0.625 0.125	0.625 0.5 0.375	150	0.125 0.625 0.125	53.7 -41.3 39.9	57.5 136.0	0.125 0.625 0.125	54.4 -54.4 50.5	74.1 137.2 16.7	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
128	G11B_062_050a	0.125 0.625 0.25	0.625 0.5 0.375	164	0.125 0.625 0.241	53.8 -40.4 35.0	53.4 139.0	0.125 0.625 0.25	54.7 -51.4 38.2	64.1 143.3 11.5	162	0.0 1.0 0.233	83.7 -80.8 70.1	106.9 139.0
129	G25B_062_050a	0.125 0.625 0.375	0.6											

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
162	R00Y_025_025a	0.25 0.0 0.0	0.25 0.25 0.125	390	0.25 0.0 0.0	12.6 19.2 16.1	25.1 40.0	0.25 0.0 0.0	8.6 28.5 13.6	31.6 25.5 10.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 100.4
163	R00Y_025_025a	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.125	13.0 20.2 10.1	20.3 2.9	0.25 0.0 0.125	9.4 30.5 -1.8	30.6 356.5 11.2	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
164	B50R_025_025a	0.25 0.0 0.25	0.25 0.25 0.125	330	0.25 0.0 0.25	14.3 23.5 -14.6	27.7 328.2	0.25 0.0 0.25	11.1 34.9 -21.6	41.1 328.2 13.7	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
165	B34R_037_037a	0.25 0.0 0.375	0.375 0.375 0.187	311	0.256 0.0 0.375	16.8 31.5 -29.7	43.3 316.7	0.25 0.0 0.375	13.8 41.1 -38.3	56.2 316.9 13.2	311	1.0 0.683 0.0	44.8 84.1 -79.2	115.5 316.7
166	B25R_050_050a	0.25 0.0 0.5	0.5 0.5 0.25	300	0.25 0.0 0.5	19.2 39.9 -44.8	60.0 311.6	0.25 0.0 0.5	17.1 48.0 -52.8	71.4 312.2 11.6	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
167	B19R_062_062a	0.25 0.0 0.625	0.625 0.625 0.312	293	0.239 0.0 0.625	22.1 48.8 -59.4	76.9 309.3	0.25 0.0 0.625	20.7 55.2 -65.9	86.0 309.9 9.2	292	0.383 0.0 1.0	35.3 78.1 -95.1	123.0 309.3
168	B15R_075_075a	0.25 0.0 0.75	0.75 0.75 0.375	289	0.237 0.0 0.75	25.4 58.1 -73.1	93.4 308.4	0.25 0.0 0.75	24.6 62.5 -77.8	99.8 308.7 6.5	288	0.316 0.0 1.0	33.9 77.4 -99.5	124.5 308.4
169	B13R_087_087a	0.25 0.0 0.875	0.875 0.875 0.437	286	0.233 0.0 0.875	28.8 67.3 -86.8	109.9 307.8	0.25 0.0 0.875	28.6 69.7 -89.1	113.1 308.0 3.2	284	0.266 0.0 1.0	32.9 77.0 -99.2	125.6 307.8
170	B11R_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4	0.25 0.0 1.0	32.6 76.8 -99.8	125.9 307.5 0.4	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
171	R50Y_025_025a	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.125 0.0	15.9 10.3 17.7	20.5 59.7	0.25 0.125 0.0	14.7 12.2 22.0	25.2 60.9 4.8	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
172	R00Y_025_012a	0.25 0.125 0.125	0.25 0.125 0.187	390	0.25 0.124 0.124	18.2 9.6 8.0	12.5 40.0	0.25 0.125 0.125	15.2 14.7 6.5	16.1 23.9 6.1	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
173	B50R_025_012a	0.25 0.125 0.25	0.25 0.125 0.187	330	0.25 0.124 0.25	19.0 11.7 -7.3	13.8 328.2	0.25 0.125 0.25	16.4 20.2 -13.2	24.2 326.7 10.6	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
174	B25R_037_025a	0.25 0.125 0.375	0.375 0.25 0.25	300	0.25 0.124 0.375	21.5 19.9 -22.4	30.0 311.6	0.25 0.125 0.375	18.4 28.0 -30.9	41.7 312.1 12.1	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
175	B15R_050_037a	0.25 0.125 0.5	0.5 0.375 0.312	289	0.243 0.124 0.5	24.6 29.0 -36.5	46.7 308.4	0.25 0.125 0.5	20.9 36.7 -46.5	59.3 308.3 13.1	288	0.316 0.0 1.0	33.9 77.4 -99.5	124.5 308.4
176	B11R_062_050a	0.25 0.125 0.625	0.625 0.5 0.375	284	0.241 0.125 0.625	28.1 38.3 -50.0	63.1 307.4	0.25 0.125 0.625	23.9 45.7 -60.5	75.9 307.0 13.4	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
177	B09R_075_062a	0.25 0.125 0.75	0.75 0.625 0.437	281	0.239 0.125 0.75	31.7 47.8 -63.2	79.3 307.0	0.25 0.125 0.75	27.3 54.4 -73.4	91.4 306.5 12.9	279	0.183 0.0 1.0	31.7 76.5 -101.2	126.9 307.0
178	B07R_087_075a	0.25 0.125 0.875	0.875 0.75 0.5	279	0.237 0.125 0.875	35.4 57.2 -76.4	95.5 306.8	0.25 0.125 0.875	30.8 62.8 -85.3	106.0 306.3 11.4	278	0.15 0.0 1.0	31.3 76.3 -101.9	127.4 306.8
179	B06R_100_087a	0.25 0.125 1.0	1.0 0.875 0.562	278	0.241 0.125 1.0	39.1 66.7 -89.5	111.6 306.7	0.25 0.125 1.0	34.5 70.9 -96.6	119.8 306.2 9.4	277	0.133 0.0 1.0	31.1 76.3 -102.3	127.6 306.7
180	Y00G_025_025a	0.25 0.25 0.0	0.25 0.25 0.125	90	0.25 0.25 0.0	23.1 -5.1 22.6	23.2 102.8	0.25 0.25 0.0	24.2 -7.6 32.9	33.7 103.1 10.5	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8
181	Y00G_025_012a	0.25 0.25 0.125	0.25 0.125 0.187	90	0.25 0.25 0.124	23.5 -2.5 11.3	11.6 102.8	0.25 0.25 0.125	24.5 -5.3 18.6	19.4 105.9 7.8	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8
182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	23.8 0.0 0.0	0.0 0.0	0.25 0.25 0.25	25.2 0.0 0.0	0.0 325.5 1.4	360	1.0 1.0 0.0	95.4 0.0 0.0	0.0 0.0
183	B00R_037_012a	0.25 0.25 0.375	0.375 0.125 0.312	270	0.249 0.249 0.375	27.6 9.5 -12.9	16.0 306.2	0.25 0.25 0.375	26.5 8.0 -18.0	19.8 294.0 5.4	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
184	B00R_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.249 0.5	31.4 19.0 -25.8	32.1 306.2	0.25 0.25 0.5	28.2 17.7 -34.7	39.0 297.0 9.5	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
185	B00R_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	35.2 28.5 -38.8	48.1 306.2	0.25 0.25 0.625	30.4 28.1 -50.7	57.4 299.3 12.2	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	39.0 38.0 -51.7	64.2 306.2	0.25 0.25 0.75	32.9 38.8 -64.1	74.8 301.0 13.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
187	B00R_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.8 47.5 -64.7	80.3 306.2	0.25 0.25 0.875	35.8 48.6 -77.1	91.2 302.1 14.3	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
188	B00R_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	46.6 57.0 -77.6	96.3 306.2	0.25 0.25 1.0	38.8 58.2 -89.4	106.7 303.0 14.1	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
189	Y31G_037_037a	0.25 0.375 0.0	0.375 0.375 0.187	109	0.256 0.375 0.0	32.8 -19.0 31.8	37.1 120.8	0.25 0.375 0.0	34.6 -24.3 41.4	48.0 120.4 11.0	108	0.683 1.0 0.0	87.6 -50.7	84.9 98.9 120.8
190	Y50G_037_025a	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	33.3 -16.3 20.6	26.2 128.3	0.25 0.375 0.125	34.8 -22.5 30.5	38.0 126.3 11.8	119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3
191	G00B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	34.3 -10.3 9.9	14.3 136.0	0.25 0.375 0.25	35.2 -18.1 14.0	22.9 142.2 8.8	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
192	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	34.7 -5.7 -1.6	6.0 196.3	0.25 0.375 0.375	36.0 -11.0 -3.5	11.6 197.8 5.8	210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3
193	G75B_050_025a	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.375 0.5	36.7 4.5 -17.0	17.6 285.0	0.25 0.375 0.5	37.2 -2.0 -20.5	20.6 264.3 7.4	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
194	G84B_062_037a	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.368 0.625	39.1 17.1 -32.5	36.7 297.8	0.25 0.375 0.625	38.7 8.2 -36.6	37.5 282.7 9.7	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
195	G88B_075_050a	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.366 0.75	42.1 28.8 -46.7	54.8 301.6	0.25 0.375 0.75	40.6 19.1 -51.6	55.0 290.3 10.9	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6
196	G90B_087_062a	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.364 0.875	45.5 39.4 -60.3	72.1 301.5	0.25 0.375 0.875	42.8 30.1 -65.7	72.2 294.6 11.0	260	0.0 0.183 1.0	34.6 63.0 -96.6	115.3 301.5
197	G92B_100_075a	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.362 1.0	48.9 50.0 -73.9	89.3 304.0	0.25 0.375 1.0	45.2 40.8 -78.9	88.9 297.3 11.1	262	0.0 0.15 1.0	33.4 66.7 -98.6	119.1 304.0
198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	42.8 -32.6 41.2	52.5 128.3	0.25 0.5 0.0	44.9 -37.9 49.4	62.3 127.5 10.0	119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3
199	Y68G_050_037a	0.25 0.5 0.125	0.5 0.375 0.312	131	0.243 0.5 0.124	43.6 -28.2 30.3	41.4 132.9	0.25 0.5 0.125	45.0 -36.5 41.4	55.2 131.4 13.9	131	0.316 1.0 0.0	84.4 -75.3	80.9 110.6 132.9
200	G00B_050_025a	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.249	44.7 -20.6 19.9	28.7 136.0	0.25 0.5 0.25	45.4 -33.0 27.2	42.8 140.5 14.3	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
201	G25B_050_025a	0.25 0.5 0.375	0.5 0.25 0.375	180	0.249 0.5 0.375	44.9 -18.4 11.2	21.6 146.6	0.25 0.5 0.375	45.9 -27.3 10.6	29.3 158.6 8.9	180	0.0 1.0 0.5	84.3 -73.7	44.9 86.4 148.6
202	G50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	45.5 -11.5 -3.3	12.0 196.3	0.25 0.5 0.5	46.8 -19.5 -6.0	20.4 197.2 8.5	210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3
203	G65B_062_037a	0.25 0.625 0.625	0.625 0.375 0.437	229	0.25 0.506 0.625	48.0 -3.4 -18.3	18.6 259.3	0.25 0.625 0.625	47.9 -10.2 -22.3	24.5 245.3 7.8	228	0.0 0.683 1.0	64.4 -9.2	-48.8 49.7 259.3
204	G75B_075_050a	0.25 0.5 0.75	0.75 0.5 0.5	240	0.25 0.5 0.75	49.7 9.1 -34.1	35.3 285.0	0.25 0.5 0.75	49.3 0.1 -37.8	37.8 270.1 9.7	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
205	G80B_087_062a	0.25 0.5 0.875	0.875 0.625 0.562	247	0.25 0.489 0.875	51.5 22.6 -50.3	55.1 294.2	0.25 0.5 0.875	50.9 10.9 -52.5	53.6 281.7 11.9	247	0.0 0.383 1.0	44.3 36.2 -80.5	88.2 294.2
206	G84B_100_075a	0.25 0.5 1.0	1.0 0.75 0.625	251	0.25 0.487 1.0	54.4 34.3 -65.0	73.5 297.8	0.25 0.5 1.0	52.8 21.9 -66.5	70.0 288.2 12.5	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
207	Y61G_062_062a	0.25 0.625 0.0	0.625 0.625 0.312	127	0.239 0.625 0.0	53.0 -45.2 50.8	68.0 131.6	0.25 0.625 0.0	55.1 -49.5 57.4	75.8 130.7 8.1	127	0.383 1.0 0.0	84.8 -72.3	81.3 108.8 131.6
208	Y76G_062_050a	0.25 0.625 0.125	0.625 0.5 0.375	136	0.241 0.625 0.125	53.9 -39.3 40.2	56.2 134.3	0.25 0.625 0.125	55.2 -48.4 51.2	70.5 133.3 14.3	137	0.233 1.0 0.0	84.0 -78.7	80.4 112.5 134.3
209	G00B_062_037a	0.25 0.625 0.25	0.625 0.375 0.437	150	0.25 0.625 0.25	55.2 -31.0 29.9	43.1 136.0	0.25 0.625 0.25	55.4 -45.7 39.2	60.2 139.3 17.3	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
210	G15B_062_037a	0.25 0.625 0.375	0.625 0.375 0.437	169	0.25 0.625 0.368	55.3 -29.7 23.6	38.0 141.4	0.25 0.625 0.375	55.8 -41.0 24.0	47.5 149.5 11.3	168	0.0 1.0 0.316	83.9 -79.2	63.1 101.3 141.4
211	G34B_062_037a	0.25 0.625 0.5	0.625 0.375 0.											

vedi file simili: <http://farbe.li.tu-berlin.de/PI31/PI31.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
243	R00Y_037_037a	0.375 0.0 0.0	0.375 0.375 0.187	390	0.375 0.0 0.0	18.9 28.8 24.2	37.6 40.0	0.375 0.0 0.0	16.4 37.5 25.4	45.3 34.1 9.1	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
244	R18Y_037_037a	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.118	19.1 26.6 11.1	31.7 20.6	0.375 0.0 0.125	16.8 38.7 9.7	39.9 14.1 9.4	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6
245	B65R_037_037a	0.375 0.0 0.25	0.375 0.375 0.187	349	0.375 0.0 0.256	20.0 32.0 -7.4	32.9 346.8	0.375 0.0 0.25	17.9 41.5 -10.4	42.8 345.8 10.2	348	1.0 0.0 0.683	53.5 85.4 -19.9	87.7 346.8
246	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	21.4 35.3 -21.9	41.6 328.2	0.375 0.0 0.375	19.7 46.0 -28.5	54.1 328.2 12.6	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
247	B38R_050_050a	0.375 0.0 0.5	0.5 0.5 0.25	316	0.383 0.0 0.5	23.9 43.2 -37.0	56.9 319.4	0.375 0.0 0.5	22.1 51.5 -44.4	68.1 319.2 11.3	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4
248	B30R_062_062a	0.375 0.0 0.625	0.625 0.625 0.312	307	0.385 0.0 0.625	25.5 51.4 -52.0	73.1 314.6	0.375 0.0 0.625	24.9 57.8 -58.7	82.4 314.5 9.4	307	0.616 0.0 1.0	42.4 82.3 -83.2	117.0 314.6
249	B25R_075_075a	0.375 0.0 0.75	0.75 0.75 0.375	300	0.375 0.0 0.75	28.9 59.8 -67.2	90.0 311.6	0.375 0.0 0.75	28.1 64.4 -71.9	96.5 311.8 6.5	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
250	B20R_087_087a	0.375 0.0 0.875	0.875 0.875 0.437	295	0.364 0.0 0.875	31.7 68.8 -81.8	106.9 310.0	0.375 0.0 0.875	31.6 71.2 -84.0	110.1 310.2 3.2	294	0.416 0.0 1.0	36.3 78.6 -93.5	122.2 310.0
251	B18R_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	34.9 77.9 -95.7	123.4 309.1	0.375 0.0 1.0	35.1 77.9 -95.5	123.3 309.2 0.3	291	0.366 0.0 1.0	34.9 77.9 -95.7	123.4 309.1
252	R31Y_037_037a	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.118 0.0	21.1 22.7 25.2	33.9 47.9	0.375 0.125 0.0	20.4 26.4 30.1	40.1 48.7 6.2	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9
253	R00Y_037_025a	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.124	24.5 19.2 16.1	25.1 40.0	0.375 0.125 0.125	20.7 27.8 14.8	31.5 28.0 9.5	389	1.0 0.0 0.0	50.4 76.9 64.5	100.0 40.0
254	R00Y_037_025a	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.25	24.9 20.2 1.0	20.3 2.9	0.375 0.125 0.25	21.6 31.1 -4.9	31.5 351.0 12.8	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
255	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.124 0.375	26.2 23.5 -14.6	27.7 328.2	0.375 0.125 0.375	23.1 36.3 -23.1	43.0 327.5 15.6	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
256	B34R_050_037a	0.375 0.125 0.5	0.5 0.375 0.312	311	0.381 0.124 0.5	28.7 31.5 -29.7	43.3 316.7	0.375 0.125 0.5	25.1 42.8 -39.5	58.3 317.2 15.3	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7
257	B25R_062_050a	0.375 0.125 0.625	0.625 0.5 0.375	300	0.375 0.125 0.625	31.2 39.9 -44.8	60.0 311.6	0.375 0.125 0.625	27.6 50.0 -54.4	73.9 312.5 14.4	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
258	B19R_075_062a	0.375 0.125 0.75	0.75 0.625 0.437	293	0.364 0.125 0.75	34.0 48.8 -59.4	76.9 309.3	0.375 0.125 0.75	30.4 57.5 -68.1	89.1 310.2 12.8	292	0.383 0.0 1.0	35.3 78.1 -95.1	120.9 309.3
259	B15R_087_075a	0.375 0.125 0.875	0.875 0.75 0.5	289	0.362 0.125 0.875	37.4 58.1 -73.1	93.4 308.4	0.375 0.125 0.875	33.6 65.1 -80.7	103.7 308.9 11.0	288	0.316 0.0 1.0	32.9 77.4 -97.5	124.5 308.4
260	B13R_100_087a	0.375 0.125 1.0	1.0 0.875 0.562	286	0.358 0.125 1.0	40.7 68.3 -86.8	109.9 307.8	0.375 0.125 1.0	36.9 72.6 -92.6	117.1 308.1 8.7	284	0.266 0.0 1.0	33.9 77.0 -99.2	125.6 307.8
261	R68Y_037_037a	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.256 0.0	27.5 6.9 29.1	29.9 76.5	0.375 0.25 0.0	27.8 8.3 37.5	38.4 77.4 8.5	71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5
262	R50Y_037_025a	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.25 0.124	27.8 10.3 17.7	20.5 59.7	0.375 0.25 0.125	28.1 9.8 23.7	25.7 67.5 6.0	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
263	R00Y_037_012a	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.249	30.1 9.6 8.0	12.5 40.0	0.375 0.25 0.25	28.7 13.3 5.4	14.4 22.0 4.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
264	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.249 0.375	31.0 11.7 -7.3	13.8 328.2	0.375 0.25 0.375	29.7 19.0 -12.7	22.9 326.1 9.1	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
265	B25R_050_025a	0.375 0.25 0.5	0.5 0.25 0.375	300	0.375 0.249 0.5	33.5 19.9 -22.4	30.0 311.6	0.375 0.25 0.5	31.2 26.3 -29.7	39.7 311.5 9.9	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
266	B15R_062_037a	0.375 0.25 0.625	0.625 0.375 0.437	289	0.368 0.25 0.625	36.5 29.0 -36.5	46.7 308.4	0.375 0.25 0.625	33.2 34.6 -45.4	57.0 307.3 10.9	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
267	B11R_075_050a	0.375 0.25 0.75	0.75 0.5 0.5	284	0.366 0.25 0.75	40.0 38.3 -50.0	63.1 307.4	0.375 0.25 0.75	35.4 43.3 -59.8	73.9 305.9 11.8	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
268	B09R_087_062a	0.375 0.25 0.875	0.875 0.625 0.562	281	0.364 0.25 0.875	43.7 47.8 -63.2	79.3 307.0	0.375 0.25 0.875	38.0 52.2 -73.3	90.0 305.4 12.3	279	0.183 0.0 1.0	31.7 76.5 -101.2	126.9 307.0
269	B07R_100_075a	0.375 0.25 1.0	1.0 0.75 0.625	279	0.362 0.25 1.0	47.3 57.2 -76.4	104.4 306.8	0.375 0.25 1.0	40.9 60.9 -86.0	105.4 305.3 12.0	278	0.15 0.0 1.0	31.3 76.3 -101.9	127.4 306.8
270	Y00G_037_037a	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.375 0.0	34.7 -7.7 34.0	34.9 102.8	0.375 0.375 0.0	36.9 10.0 -40.2	45.3 102.8 10.7	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
271	Y00G_037_025a	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.375 0.124	35.0 -5.1 22.6	23.2 102.8	0.375 0.375 0.125	37.1 -8.7 33.8	34.9 104.4 11.8	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
272	Y00G_037_012a	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.375 0.249	35.4 -2.5 11.3	11.6 102.8	0.375 0.375 0.25	37.5 -5.4 17.5	18.3 107.1 7.1	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
273	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	35.7 0.0 0.0	0.0 0.0	0.375 0.375 0.375	38.3 0.0 0.0	0.0 325.3 2.5	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0
274	B00R_050_012a	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.375 0.5	39.5 9.5 -12.9	16.0 306.2	0.375 0.375 0.5	39.4 7.2 -17.0	18.5 292.9 4.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
275	B00R_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	43.3 19.0 -25.8	32.1 306.2	0.375 0.375 0.625	40.8 15.7 -33.2	36.8 295.4 8.4	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
276	B00R_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	47.1 28.5 -38.8	48.1 306.2	0.375 0.375 0.75	42.5 25.1 -48.4	54.5 297.4 11.1	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
277	B00R_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	50.9 38.0 -51.7	64.2 306.2	0.375 0.375 0.875	44.6 34.8 -62.7	71.7 299.0 13.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
278	B00R_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	54.7 47.5 -64.7	80.3 306.2	0.375 0.375 1.0	46.8 44.5 -76.1	88.2 300.3 14.2	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
279	Y23G_050_050a	0.375 0.5 0.0	0.5 0.5 0.25	104	0.383 0.5 0.0	44.3 -21.6 43.1	48.2 116.6	0.375 0.5 0.0	46.6 -26.1 51.4	57.7 116.9 9.7	102	0.766 1.0 0.0	88.7 -43.3 86.2	96.5 116.6
280	Y31G_050_037a	0.375 0.5 0.125	0.5 0.375 0.312	109	0.381 0.5 0.124	44.8 -19.0 31.8	37.1 120.8	0.375 0.5 0.125	46.7 -25.0 43.6	50.2 119.8 13.3	108	0.683 1.0 0.0	87.6 -50.7 84.9	98.9 120.8
281	Y50G_050_025a	0.375 0.5 0.25	0.5 0.25 0.375	120	0.375 0.5 0.249	45.2 -16.3 20.6	26.2 128.3	0.375 0.5 0.25	47.0 -22.1 29.6	36.9 126.8 10.8	119	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
282	G00B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	46.2 -10.3 9.9	14.3 136.0	0.375 0.5 0.375	47.6 -17.3 13.1	21.8 142.8 7.8	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
283	G50B_050_012a	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.5	46.6 -5.7 -1.6	6.0 196.3	0.375 0.5 0.5	48.4 -10.7 -3.5	11.3 198.2 5.6	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
284	G75B_062_025a	0.375 0.5 0.625	0.625 0.25 0.5	240	0.375 0.5 0.625	48.7 4.5 -17.0	17.6 285.0	0.375 0.5 0.625	49.4 -2.7 -19.8	20.0 262.1 7.8	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
285	G84B_075_037a	0.375 0.5 0.75	0.75 0.375 0.562	251	0.375 0.493 0.75	51.0 17.1 -32.5	36.7 297.8	0.375 0.5 0.75	50.7 6.3 -35.4	35.9 280.2 11.1	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
286	G88B_087_050a	0.375 0.5 0.875	0.875 0.5 0.625	256	0.375 0.491 0.875	54.0 28.8 -46.7	54.8 301.6	0.375 0.5 0.875	52.3 16.1 -50.2	52.7 287.8 13.2	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6
287	G90B_100_062a	0.375 0.5 1.0	1.0 0.625 0.687	259	0.375 0.489 1.0	57.4 39.4 -60.3	72.1 303.1	0.375 0.5 1.0	54.1 26.2 -64.3	69.4 292.1 14.1	260	0.0 0.183 1.0	34.6 63.0 -96.6	115.3 303.1
288	Y38G_062_062a	0.375 0.625 0.0	0.625 0.625 0.312	113	0.385 0.625 0.0	54.2 -35.2 52.4	63.1 123.9	0.375 0.625 0.0	56.3 -39.9 58.9	71.2 124.1 8.3	112	0.616 1.0 0.0	86.8 -56.4 83.8	101.0 123.9
289	Y50G_062_050a	0.375 0.625 0.125	0.625 0.5 0.375	120	0.375 0.625 0.125	54.7 -32.6 41.4	52.5 128.3	0.375 0.625 0.125	56.4 -39.0 52.8	65.7 126.4 13.4	119	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
290	Y68G_062_037a	0.375 0.625 0.25	0.625 0.375 0.437	131	0.368 0.625 0.25	55.5 -28.2 30.3	41.4 132.9	0.375 0.625 0.25	56.6 -36.6 40.9	54.9 131.8 13.5	131	0.316 1.0 0.0	84.4 -75.3 80.9	110.6 132.9
291	G00B_062_025a	0.375 0.625												

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informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsi_d	rgb*Md	LabCh*Md				
324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.25	390	0.5	0.0	0.0	50.4	76.9	64.5	100.4	40.0
325	R26Y_050_050a	0.5	0.0	0.125	0.5	0.5	0.25	376	0.5	0.0	0.125	50.8	78.0	41.2	88.2	27.8
326	R00Y_050_050a	0.5	0.0	0.25	0.5	0.5	0.25	360	0.5	0.0	0.25	52.0	81.1	4.1	81.2	2.9
327	B61R_050_050a	0.5	0.0	0.375	0.5	0.5	0.25	344	0.5	0.0	0.375	54.4	87.3	-30.6	92.5	340.6
328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.25	330	0.5	0.0	0.5	57.2	94.3	-58.4	110.9	328.2
329	B40R_062_062a	0.5	0.0	0.625	0.625	0.625	0.312	319	0.51	0.0	0.625	31.0	48.8	-70.7	113.0	321.2
330	B34R_075_075a	0.5	0.0	0.75	0.75	0.75	0.375	311	0.512	0.0	0.75	31.6	44.8	-79.2	115.5	316.7
331	B29R_087_087a	0.5	0.0	0.875	0.875	0.875	0.437	305	0.51	0.0	0.875	31.6	41.3	-85.1	117.9	313.8
332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	300	0.5	0.0	1.0	38.5	79.8	-89.7	120.0	311.6
333	R23Y_050_050a	0.5	0.125	0.0	0.5	0.5	0.25	44	0.5	0.116	0.0	53.7	67.6	65.8	94.4	44.2
334	R00Y_050_037a	0.5	0.125	0.125	0.5	0.375	0.312	390	0.5	0.124	0.124	30.8	28.8	-24.2	37.6	40.0
335	R18Y_050_037a	0.5	0.125	0.25	0.5	0.375	0.312	371	0.5	0.124	0.243	31.0	29.6	11.1	31.7	20.6
336	B65R_050_037a	0.5	0.125	0.375	0.5	0.375	0.312	349	0.5	0.124	0.381	32.0	32.0	-7.4	32.9	346.8
337	B50R_050_037a	0.5	0.125	0.5	0.5	0.375	0.312	330	0.5	0.124	0.5	33.4	35.3	-21.9	41.6	328.2
338	B38R_062_050a	0.5	0.125	0.625	0.625	0.5	0.375	316	0.508	0.125	0.625	35.8	43.2	-37.0	56.9	319.4
339	B30R_075_062a	0.5	0.125	0.75	0.75	0.625	0.437	307	0.51	0.125	0.75	38.4	51.4	-52.0	73.1	314.6
340	B25R_087_075a	0.5	0.125	0.875	0.875	0.75	0.5	300	0.5	0.125	0.875	40.8	59.8	-67.2	90.0	311.6
341	B20R_100_087a	0.5	0.125	1.0	1.0	0.875	0.562	295	0.489	0.125	1.0	43.6	68.8	-81.8	106.9	310.0
342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	60	0.5	0.25	0.0	31.8	20.6	35.5	41.1	59.7
343	R31Y_050_037a	0.5	0.25	0.125	0.5	0.375	0.312	49	0.5	0.243	0.124	33.0	22.7	25.2	33.9	49.9
344	R00Y_050_025a	0.5	0.25	0.25	0.5	0.25	0.375	390	0.5	0.249	0.249	36.4	19.2	16.1	25.1	40.0
345	R00Y_050_025a	0.5	0.25	0.375	0.5	0.25	0.375	360	0.5	0.249	0.375	36.8	20.2	1.0	20.3	2.9
346	B50R_050_025a	0.5	0.25	0.5	0.5	0.25	0.375	330	0.5	0.249	0.5	38.1	23.5	-14.6	27.7	328.2
347	B34R_062_037a	0.5	0.25	0.625	0.625	0.375	0.437	311	0.506	0.25	0.625	40.6	31.5	-29.7	43.3	316.7
348	B25R_075_050a	0.5	0.25	0.75	0.75	0.5	0.5	300	0.5	0.25	0.75	43.1	39.9	-44.8	60.0	311.6
349	B19R_087_062a	0.5	0.25	0.875	0.875	0.625	0.562	293	0.489	0.25	0.875	45.9	48.8	-59.4	76.9	309.3
350	B15R_100_075a	0.5	0.25	1.0	1.0	0.75	0.625	289	0.487	0.25	1.0	49.3	58.1	-73.1	93.4	308.4
351	R76Y_050_050a	0.5	0.375	0.0	0.5	0.5	0.25	76	0.5	0.383	0.0	39.1	3.9	40.3	40.5	84.4
352	R68Y_050_037a	0.5	0.375	0.125	0.5	0.375	0.312	71	0.5	0.381	0.124	39.4	6.9	29.1	29.9	76.5
353	R50Y_050_025a	0.5	0.375	0.25	0.5	0.25	0.375	60	0.5	0.375	0.249	39.7	10.3	17.7	20.5	59.7
354	R00Y_050_012a	0.5	0.375	0.375	0.5	0.125	0.437	390	0.5	0.375	0.375	42.0	9.6	8.0	12.5	40.0
355	B50R_050_012a	0.5	0.375	0.5	0.5	0.125	0.437	330	0.5	0.375	0.5	42.9	11.7	-7.3	13.8	328.2
356	B25R_062_025a	0.5	0.375	0.625	0.625	0.25	0.5	300	0.5	0.375	0.625	45.4	19.9	-22.4	30.0	311.6
357	B15R_075_037a	0.5	0.375	0.75	0.75	0.375	0.562	289	0.493	0.375	0.75	48.5	29.0	-36.5	46.7	308.4
358	B11R_087_050a	0.5	0.375	0.875	0.875	0.5	0.625	284	0.491	0.375	0.875	51.9	38.3	-50.0	63.1	307.4
359	B09R_100_062a	0.5	0.375	1.0	1.0	0.625	0.687	281	0.489	0.375	1.0	55.6	47.8	-63.2	79.3	307.0
360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	90	0.5	0.5	0.0	46.3	-10.3	45.3	46.5	102.8
361	Y00G_050_037a	0.5	0.5	0.125	0.5	0.375	0.312	90	0.5	0.5	0.124	46.6	-7.7	34.0	34.9	102.8
362	Y00G_050_025a	0.5	0.5	0.25	0.5	0.25	0.375	90	0.5	0.5	0.249	47.0	-5.1	22.6	23.2	102.8
363	Y00G_050_012a	0.5	0.5	0.375	0.5	0.125	0.437	90	0.5	0.5	0.375	47.3	-2.5	11.3	11.6	102.8
364	NW_050a	0.5	0.5	0.5	0.5	0.0	0.5	360	0.5	0.5	0.5	47.7	0.0	0.0	0.0	0.0
365	B00R_062_012a	0.5	0.625	0.625	0.625	0.125	0.562	270	0.5	0.5	0.625	51.5	9.5	-12.9	16.0	306.2
366	B00R_075_025a	0.5	0.5	0.75	0.75	0.25	0.625	270	0.5	0.5	0.75	55.3	19.0	-25.8	32.1	306.2
367	B00R_087_037a	0.5	0.5	0.875	0.875	0.375	0.687	270	0.5	0.5	0.875	59.1	28.5	-38.8	48.1	306.2
368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	270	0.5	0.5	1.0	62.8	38.0	-51.7	64.2	306.2
369	Y18G_062_062a	0.5	0.625	0.0	0.625	0.625	0.312	101	0.51	0.625	0.0	55.9	-24.1	54.5	59.6	113.8
370	Y23G_062_050a	0.5	0.625	0.125	0.625	0.5	0.375	104	0.508	0.625	0.125	56.3	-21.6	43.1	48.2	116.6
371	Y31G_062_037a	0.5	0.625	0.25	0.625	0.375	0.437	109	0.506	0.625	0.25	56.7	-19.0	31.8	37.1	120.8
372	Y50G_062_025a	0.5	0.625	0.375	0.625	0.25	0.5	120	0.5	0.625	0.375	57.2	-16.3	20.6	26.2	128.3
373	G00B_062_012a	0.5	0.625	0.5	0.625	0.125	0.562	150	0.5	0.625	0.5	58.1	-10.3	9.9	14.3	136.0
374	G50B_062_012a	0.5	0.625	0.625	0.625	0.125	0.562	210	0.5	0.625	0.625	58.5	-5.7	-1.6	6.0	196.3
375	G75B_075_025a	0.5	0.625	0.75	0.75	0.25	0.625	240	0.5	0.625	0.75	60.6	4.5	-17.0	17.6	285.0
376	G84B_087_037a	0.5	0.625	0.875	0.875	0.375	0.687	251	0.5	0.618	0.875	62.9	17.1	-32.5	36.7	297.8
377	G88B_100_050a	0.5	0.625	1.0	1.0	0.5	0.75	256	0.5	0.616	1.0	65.9	28.8	-46.7	54.8	301.6
378	Y31G_075_075a	0.5	0.75	0.0	0.75	0.75	0.375	109	0.512	0.75	0.0	65.7	-38.0	63.7	74.2	120.8
379	Y38G_075_062a	0.5	0.75	0.125	0.75	0.625	0.437	113	0.51	0.75	0.125	66.2	-35.2	52.4	63.1	123.9
380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	120	0.5	0.75	0.25	66.7	-32.6	41.2	52.5	128.3
381	Y68G_075_037a	0.5	0.75	0.375	0.75	0.375	0.562	131	0.493	0.75	0.375	67.4	-28.2	30.3	41.4	132.9
382	G00B_075_025a	0.5	0.75	0.5	0.75	0.25	0.625	150	0.5	0.75	0.5	68.6	-20.6	19.9	28.7	136.0
383	G25B_075_025a	0.5	0.75	0.625	0.75	0.25	0.625	180	0.5	0.75	0.625	68.7	-18.4	11.2	21.6	148.6
384	G50B_075_025a	0.5	0.75	0.75	0.75	0.25	0.625	210	0.5	0.75	0.75	69.4	-11.5	-3.3	12.0	196.3
385	G65B_087_037a	0.5	0.75	0.875	0.875	0.375	0.687	229	0.5	0.756	0.875	71.8	-3.4	-18.3	18.6	259.3
386	G75B_100_050a	0.5	0.75	1.0	1.0	0.5	0.75	240	0.5	0.75	1.0	73.5	9.1	-34.1	35.3	285.0
387	Y41G_087_087a	0.5	0.875	0.0	0.875	0.875	0.437	115	0.51	0.875	0.0	75.7	-51.5	73.1	89.4	125.1
388	Y50G_087_075a	0.5	0.875	0.125	0.875	0.75	0.5	120	0.5	0.875	0.125	76.2	-48.9	61.8	78.8	128.3
389	Y61G_087_062a	0.5	0.875	0.25	0.875	0.625	0.562	127	0.489	0.875	0.25	76.8	-45.2	50.8	68.0	131.6
390	Y76G_087_050a	0.5	0.875	0.375	0.875	0.5	0.625	136	0.491	0.875	0.375	77.8	-39.3	40.2	56.2	134.3
391	G00B_087_037a	0.5	0.875	0.5	0.875	0.375	0.687	150	0.5	0.875	0.5	79.0	-31.0	29.9	43.1	136.0
392	G15B_087_037a	0.5	0.875	0.625	0.875	0.375	0.687	169	0.5	0.875	0.618	79.1	-29.7	23.6	38.0	141.4
393	G34B_087_037a	0.5	0.875	0.75	0.875	0.375	0.687	191	0.5	0.875	0.75	79.5	-24.7	8.7	26.2	160.0
394	G50B_087_037a	0.5	0.875	0.875	0.875	0.375	0.687	210	0.5	0.875	0.875	80.2	-17.3	-5.0	18.0	196.3
395	G61B_100_050a	0.5	0.875	1.0	1.0	0.5	0.75	224	0.5	0.883	1.0	82.8	-9.7	-19.6	21.9	243.6
396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	120	0.5	1.0	0.0	85.7	-65.2	82.4	105.1	128.3
397	Y58G_100_087a	0.5	1.0	0.125	1.0	0										

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informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsi_d	rgb*Md	LabCh*Md
405	R00Y_062_062a	0.625 0.0 0.0	0.625 0.625 0.312	390	0.625 0.0 0.0	31.5 48.0 40.3	62.7 40.0	0.625 0.0 0.0	30.7 54.1 44.5	70.1 39.4 7.4	389	50.4 76.9 64.5
406	R31Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.114	31.7 48.7 29.7	57.0 31.3	0.625 0.0 0.125	31.0 54.7 30.0	62.4 28.7 6.0	380	50.7 77.9 47.5
407	R11Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.239	32.1 49.6 12.8	51.3 14.4	0.625 0.0 0.25	31.5 56.2 10.9	57.2 11.0 6.7	367	51.4 79.5 20.4
408	B69R_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.385	33.0 52.2 -7.1	52.7 352.1	0.625 0.0 0.375	32.4 58.6 -7.7	59.1 352.5 6.4	352	52.9 83.6 -11.4
409	B59R_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.625 0.0 0.51	34.3 55.5 -22.8	60.1 337.6	0.625 0.0 0.5	33.8 62.1 -25.0	67.0 338.0 6.9	339	54.9 88.9 -36.6
410	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	35.8 58.9 -36.5	62.9 328.2	0.625 0.0 0.625	35.5 66.4 -41.1	78.1 328.2 8.7	330	57.2 94.3 -58.4
411	B42R_075_075a	0.625 0.0 0.75	0.75 0.75 0.375	321	0.637 0.0 0.75	38.4 66.8 -51.4	84.3 322.4	0.625 0.0 0.75	37.6 71.3 -55.9	90.6 312.8 6.4	322	51.2 89.1 -68.5
412	B36R_087_087a	0.625 0.0 0.875	0.875 0.875 0.375	314	0.641 0.0 0.875	40.8 74.7 -66.6	100.1 318.3	0.625 0.0 0.875	40.0 76.7 -69.8	103.7 317.8 3.8	315	46.6 85.4 -76.1
413	B31R_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	43.0 82.7 -82.2	116.6 315.1	0.625 0.0 1.0	42.7 82.5 -82.8	116.8 314.8 0.6	308	43.0 82.7 -82.2
414	R18Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	41	0.625 0.114 0.0	32.9 44.0 40.9	60.1 42.8	0.625 0.125 0.0	32.8 48.2 45.9	66.6 34.6 6.5	39	52.7 70.5 65.5
415	R00Y_062_050a	0.625 0.125 0.125	0.625 0.5 0.375	390	0.625 0.125 0.125	37.1 38.4 32.2	50.2 40.0	0.625 0.125 0.125	33.0 48.8 32.2	58.5 33.3 11.1	389	50.4 76.9 64.5
416	R26Y_062_050a	0.625 0.125 0.25	0.625 0.5 0.375	376	0.625 0.125 0.241	37.3 39.0 20.6	44.1 27.8	0.625 0.125 0.25	33.5 50.4 13.6	52.2 15.1 13.9	377	50.8 78.0 41.2
417	R00Y_062_050a	0.625 0.125 0.375	0.625 0.5 0.375	360	0.625 0.125 0.375	37.9 40.5 2.0	40.6 2.9	0.625 0.125 0.375	34.4 53.1 -4.8	53.3 354.8 14.7	360	52.0 81.1 4.1
418	B61R_062_050a	0.625 0.125 0.5	0.625 0.5 0.375	344	0.625 0.125 0.508	39.1 43.6 -15.3	46.2 340.6	0.625 0.125 0.5	35.6 56.7 -22.2	60.9 338.6 15.2	342	54.4 87.3 -30.6
419	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	40.5 47.1 -29.2	55.4 328.2	0.625 0.125 0.625	37.3 61.3 -38.3	72.3 327.9 17.2	330	57.2 94.3 -58.4
420	B40R_075_062a	0.625 0.125 0.75	0.75 0.625 0.375	319	0.635 0.125 0.75	43.1 55.0 -44.2	70.6 321.2	0.625 0.125 0.75	39.2 66.6 -53.4	85.3 321.2 15.2	320	49.8 88.1 -70.7
421	B34R_087_075a	0.625 0.125 0.875	0.875 0.75 0.5	311	0.637 0.125 0.875	45.5 63.1 -59.4	86.6 316.7	0.625 0.125 0.875	41.5 72.3 -67.4	98.9 317.0 12.9	311	44.8 84.1 -79.2
422	B29R_100_087a	0.625 0.125 1.0	1.0 0.875 0.562	305	0.635 0.125 1.0	48.0 71.4 -74.4	103.2 313.8	0.625 0.125 1.0	44.0 78.4 -80.5	112.4 314.2 10.1	305	41.3 81.6 -85.1
423	R38Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	53	0.625 0.239 0.0	36.6 34.0 42.6	54.6 51.3	0.625 0.25 0.0	37.4 35.7 48.5	60.2 53.5 6.1	52	58.5 54.5 68.2
424	R23Y_062_050a	0.625 0.25 0.125	0.625 0.5 0.375	44	0.625 0.241 0.125	38.8 33.8 32.9	47.2 44.2	0.625 0.25 0.125	37.6 36.4 36.8	51.8 45.2 4.8	42	57.3 67.6 65.8
425	R00Y_062_037a	0.625 0.25 0.25	0.625 0.375 0.437	390	0.625 0.25 0.25	42.7 28.8 24.2	37.6 40.0	0.625 0.25 0.25	38.0 38.2 19.6	42.9 27.1 11.4	389	50.4 76.9 64.5
426	R18Y_062_037a	0.625 0.25 0.375	0.625 0.375 0.437	371	0.625 0.25 0.368	43.0 29.6 11.1	31.7 20.6	0.625 0.25 0.375	38.7 41.1 1.5	41.1 21.5 15.5	371	51.1 79.1 29.7
427	B65R_062_037a	0.625 0.25 0.5	0.625 0.375 0.437	349	0.625 0.25 0.506	43.9 32.0 -7.4	32.9 346.8	0.625 0.25 0.5	39.8 45.1 -15.7	47.8 340.7 16.0	349	53.5 85.4 -19.9
428	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	45.3 35.3 -21.9	41.6 328.2	0.625 0.25 0.625	41.2 50.2 -32.1	59.6 327.4 18.5	330	57.2 94.3 -58.4
429	B38R_075_050a	0.625 0.25 0.75	0.75 0.5 0.5	316	0.633 0.25 0.75	47.8 43.2 -37.0	56.9 319.4	0.625 0.25 0.75	42.9 56.0 -47.4	73.4 319.7 17.2	317	47.9 86.4 -74.0
430	B30R_087_062a	0.625 0.25 0.875	0.875 0.625 0.562	307	0.635 0.25 0.875	50.3 51.4 -52.0	73.1 314.6	0.625 0.25 0.875	44.9 62.4 -61.8	87.9 315.2 15.7	307	42.4 82.3 -83.2
431	B25R_100_075a	0.625 0.25 1.0	1.0 0.75 0.625	300	0.625 0.25 1.0	52.8 59.8 -67.2	90.0 311.6	0.625 0.25 1.0	47.2 69.2 -75.4	102.3 312.5 13.5	300	38.5 79.8 -89.7
432	R61Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	67	0.625 0.385 0.0	43.5 16.7 46.8	49.7 70.2	0.625 0.375 0.0	44.1 19.3 52.4	55.9 69.7 6.2	67	69.6 26.8 74.8
433	R50Y_062_050a	0.625 0.375 0.125	0.625 0.5 0.375	60	0.625 0.375 0.125	43.7 20.6 35.5	41.1 59.7	0.625 0.375 0.125	44.2 20.0 43.2	47.6 65.1 7.7	59	63.6 41.3 71.0
434	R31Y_062_037a	0.625 0.375 0.25	0.625 0.375 0.437	49	0.625 0.368 0.25	44.9 22.7 25.2	33.9 47.9	0.625 0.375 0.25	44.5 21.8 27.8	35.4 51.9 2.8	48	56.2 60.6 67.2
435	R00Y_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	48.4 19.2 16.1	25.1 40.0	0.625 0.375 0.375	45.1 24.9 10.6	27.0 23.1 8.5	389	50.4 76.9 64.5
436	R00Y_062_025a	0.625 0.375 0.5	0.625 0.25 0.5	360	0.625 0.375 0.5	48.7 20.2 1.0	20.3 2.9	0.625 0.375 0.5	46.0 29.2 -6.4	29.9 347.5 11.9	360	52.0 81.1 4.1
437	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	50.1 23.5 -14.6	27.7 328.2	0.625 0.375 0.625	47.1 34.6 -22.9	41.5 326.5 14.1	330	57.2 94.3 -58.4
438	B34R_075_037a	0.625 0.375 0.75	0.75 0.375 0.562	311	0.631 0.375 0.75	52.5 31.5 -29.7	43.3 316.7	0.625 0.375 0.75	48.5 40.9 -38.5	56.2 316.7 13.5	311	44.8 84.1 -79.2
439	B25R_087_050a	0.625 0.375 0.875	0.875 0.5 0.625	300	0.625 0.375 0.875	55.0 39.9 -44.8	60.0 311.6	0.625 0.375 0.875	50.2 47.9 -53.3	71.7 311.9 12.6	300	38.5 79.8 -89.7
440	B19R_100_062a	0.625 0.375 1.0	1.0 0.625 0.687	293	0.614 0.375 1.0	57.8 48.8 -59.4	76.9 309.3	0.625 0.375 1.0	52.1 55.3 -67.3	87.1 309.4 11.7	292	35.3 78.1 -95.1
441	R81Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	79	0.625 0.51 0.0	50.8 1.0 51.8	51.8 88.7	0.625 0.5 0.0	51.9 1.9 57.7	57.8 88.0 6.0	80	81.2 1.7 82.9
442	R76Y_062_050a	0.625 0.5 0.125	0.625 0.5 0.375	76	0.625 0.508 0.125	51.0 3.9 40.3	40.5 84.4	0.625 0.5 0.125	52.0 2.6 50.5	50.6 86.9 10.3	77	78.2 7.8 80.6
443	R68Y_062_037a	0.625 0.5 0.25	0.625 0.375 0.437	71	0.625 0.506 0.25	51.3 6.9 29.1	29.9 76.5	0.625 0.5 0.25	52.3 4.4 37.1	37.4 83.2 8.4	71	73.4 18.5 77.6
444	R50Y_062_025a	0.625 0.5 0.375	0.625 0.25 0.5	60	0.625 0.5 0.375	51.6 10.3 17.7	20.5 59.7	0.625 0.5 0.375	52.8 7.4 21.1	22.3 70.5 4.5	59	63.6 41.3 71.0
445	R00Y_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	390	0.625 0.5 0.5	54.0 9.6 8.0	12.5 40.0	0.625 0.5 0.5	53.4 11.7 4.4	12.6 20.7 4.2	389	50.4 76.9 64.5
446	B50R_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	54.8 11.7 -7.3	13.8 328.2	0.625 0.5 0.625	54.4 17.2 -11.8	20.9 325.5 7.1	330	57.2 94.3 -58.4
447	B25R_075_025a	0.625 0.5 0.75	0.75 0.25 0.625	300	0.625 0.5 0.75	57.3 19.9 -22.4	30.0 311.6	0.625 0.5 0.75	55.5 23.7 -27.6	36.4 310.7 6.6	300	38.5 79.8 -89.7
448	B15R_087_037a	0.625 0.5 0.875	0.875 0.375 0.687	289	0.618 0.5 0.875	60.4 29.0 -36.5	46.7 308.4	0.625 0.5 0.875	56.9 31.0 -42.7	52.8 305.9 7.3	288	33.9 77.4 -97.5
449	B11R_100_050a	0.625 0.5 1.0	1.0 0.5 0.75	284	0.616 0.5 1.0	63.9 38.3 -50.0	63.1 307.4	0.625 0.5 1.0	58.5 38.8 -57.1	69.0 304.8 8.8	282	32.3 76.7 -100.1
450	Y00G_062_062a	0.625 0.625 0.0	0.625 0.625 0.312	90	0.625 0.625 0.0	57.9 -12.9 56.7	58.1 102.8	0.625 0.625 0.0	60.4 -14.5 63.8	65.4 102.8 7.7	89	92.6 -20.7
451	Y00G_062_050a	0.625 0.625 0.125	0.625 0.5 0.375	90	0.625 0.625 0.125	58.2 -10.3 45.3	46.5 102.8	0.625 0.625 0.125	60.5 -13.9 58.1	59.7 103.4 13.4	89	92.6 -20.7
452	Y00G_062_037a	0.625 0.625 0.25	0.625 0.375 0.437	90	0.625 0.625 0.25	58.5 -7.7 34.0	34.9 102.8	0.625 0.625 0.25	60.7 -12.2 46.6	48.2 104.7 13.5	89	92.6 -20.7
453	Y00G_062_025a	0.625 0.625 0.375	0.625 0.25 0.5	90	0.625 0.625 0.375	58.9 -5.1 22.6	23.2 102.8	0.625 0.625 0.375	61.1 -9.3 31.9	33.2 106.3 10.3	89	92.6 -20.7
454	Y00G_062_012a	0.625 0.625 0.5	0.625 0.125 0.562	90	0.625 0.625 0.5	59.2 -2.5 11.3	11.6 102.8	0.625 0.625 0.5	61.6 -5.2 16.0	16.8 108.2 5.8	89	92.6 -20.7
455	NW_062a	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	59.6 0.0 0.0	0.0 0.0	0.625 0.625 0.625	62.4 0.0 0.0	0.0 325.2 2.7	360	95.4 0.0 0.0
456	B00R_075_012a	0.625 0.625 0.75	0.75 0.125 0.687	270	0.625 0.625 0.75	63.4 9.5 -12.9	16.0 306.2	0.625 0.625 0.75	63.3 6.3 -15.7	16.9 292.0 4.1	270	30.3 76.0 -103.5
457	B00R_087_025a	0.625 0.625 0.875	0.875 0.25 0.75	270	0.625 0.625 0.875	67.2 19.0 -25.8	32.1 306.2	0.625 0.625 0.875	64.4 13.5 -30.9	33.8 293.6 7.9	270	30.3 76.0 -103.5
458	B00R_100_037a	0.625 0.625 1.0	1.0 0.375 0.812	270	0.625 0.625 1.0	71.0 28.5 -38.8	48.1 306.2	0.625 0.625 1.0	65.7 21.4 -45.6	50.4 295.1 11.1	270	30.3 76.0 -103.5
459	Y15G_075_075a	0.625 0.75 0.0	0.75 0.75 0.375	99	0.637 0.75 0.0	67.5 -26.5 65.8	71.0 111.9	0.625 0.75 0.0	69.2 -29.6 70.3	76.3 112.8 5.7	97	85.1 0.0 0.0
460	Y18G_075_062a	0.625 0.75 0.125	0.75 0.625 0.437	101	0.635 0.75 0.125	67.9 -24.1 54.5	59.6 113.					

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md
486	R00Y_075_075d	0.75	0.0	0.0	0.75	0.75	0.0	0.0	37.5	61.9	51.9	80.8
487	R35Y_075_075d	0.75	0.0	0.125	0.75	0.75	0.0	0.125	37.7	62.4	38.9	73.5
488	R18Y_075_075d	0.75	0.0	0.25	0.75	0.75	0.0	0.25	38.1	63.5	20.8	66.9
489	R00Y_075_075d	0.75	0.0	0.375	0.75	0.75	0.0	0.375	38.8	65.5	2.4	65.5
490	B65R_075_075d	0.75	0.0	0.5	0.75	0.75	0.0	0.5	39.9	68.2	-15.1	69.9
491	B57R_075_075d	0.75	0.0	0.625	0.75	0.75	0.0	0.625	41.3	71.8	-31.6	78.4
492	B50R_075_075d	0.75	0.0	0.75	0.75	0.75	0.0	0.75	43.0	76.0	-47.0	89.4
493	B43R_087_087d	0.75	0.0	0.875	0.875	0.875	0.0	0.875	45.0	80.7	-61.5	101.5
494	B38R_100_100d	0.75	0.0	1.0	1.0	1.0	0.5	1.0	47.2	85.8	-75.1	114.1
495	R15Y_075_075d	0.75	0.125	0.0	0.75	0.75	0.125	0.0	39.1	57.3	52.5	77.8
496	R00Y_075_062d	0.75	0.125	0.125	0.75	0.75	0.125	0.125	39.3	57.8	40.4	70.6
497	R31Y_075_062d	0.75	0.125	0.25	0.75	0.75	0.125	0.25	39.7	59.0	22.8	63.3
498	R11Y_075_062d	0.75	0.125	0.375	0.75	0.75	0.125	0.375	40.3	61.0	4.6	61.2
499	B69R_075_062d	0.75	0.125	0.5	0.75	0.75	0.125	0.5	41.4	64.0	-12.9	65.2
500	B59R_075_062d	0.75	0.125	0.625	0.75	0.75	0.125	0.625	42.7	67.7	-29.4	73.8
501	B50R_075_062d	0.75	0.125	0.75	0.75	0.75	0.125	0.75	44.3	72.1	-44.9	84.9
502	B42R_087_075d	0.75	0.125	0.875	0.875	0.75	0.125	0.875	46.2	77.0	-59.5	97.3
503	B36R_100_087d	0.75	0.125	1.0	1.0	0.875	0.562	314	48.4	82.4	-73.2	110.2
504	R31Y_075_075d	0.75	0.25	0.0	0.75	0.75	0.25	0.0	42.8	47.1	54.2	71.8
505	R18Y_075_062d	0.75	0.25	0.125	0.75	0.75	0.25	0.125	42.9	47.6	43.8	64.7
506	R00Y_075_050d	0.75	0.25	0.25	0.75	0.75	0.25	0.25	43.3	48.9	27.4	56.0
507	R26Y_075_050d	0.75	0.25	0.375	0.75	0.75	0.25	0.375	43.9	51.1	9.6	52.0
508	R00Y_075_050d	0.75	0.25	0.5	0.75	0.75	0.25	0.5	44.8	54.3	-7.7	54.8
509	B61R_075_050d	0.75	0.25	0.625	0.75	0.75	0.25	0.625	46.0	58.3	-24.3	63.1
510	B50R_075_050d	0.75	0.25	0.75	0.75	0.75	0.25	0.75	47.5	63.1	-39.9	74.6
511	B40R_087_062d	0.75	0.25	0.875	0.875	0.625	0.562	319	49.2	68.4	-54.7	87.6
512	B34R_100_075d	0.75	0.25	1.0	1.0	0.75	0.625	311	51.2	74.3	-68.7	101.2
513	R50Y_075_075d	0.75	0.375	0.0	0.75	0.75	0.375	0.0	48.5	32.5	57.4	65.9
514	R38Y_075_062d	0.75	0.375	0.125	0.75	0.75	0.375	0.125	48.6	33.0	48.8	59.0
515	R23Y_075_050d	0.75	0.375	0.25	0.75	0.75	0.375	0.25	48.9	34.4	34.1	48.4
516	R00Y_075_037d	0.75	0.375	0.375	0.75	0.75	0.375	0.375	49.4	36.7	17.1	40.5
517	R18Y_075_037d	0.75	0.375	0.5	0.75	0.75	0.375	0.5	50.1	40.1	0.1	40.1
518	B65R_075_037d	0.75	0.375	0.625	0.75	0.75	0.375	0.625	51.1	44.4	-16.4	47.4
519	B50R_075_037d	0.75	0.375	0.75	0.75	0.75	0.375	0.75	52.4	49.6	-32.2	59.1
520	B38R_087_050d	0.75	0.375	0.875	0.875	0.5	0.625	316	53.9	55.4	-47.2	72.8
521	B30R_100_062d	0.75	0.375	1.0	1.0	0.625	0.687	307	55.6	61.8	-61.5	87.2
522	R68Y_075_075d	0.75	0.5	0.0	0.75	0.75	0.5	0.0	55.4	15.9	61.8	87.2
523	R61Y_075_062d	0.75	0.5	0.125	0.75	0.75	0.5	0.125	55.5	16.4	54.9	57.3
524	R50Y_075_050d	0.75	0.5	0.25	0.75	0.75	0.5	0.25	55.8	17.8	42.0	45.6
525	R31Y_075_037d	0.75	0.5	0.375	0.75	0.75	0.5	0.375	56.2	20.2	26.2	33.1
526	R00Y_075_025d	0.75	0.5	0.5	0.75	0.75	0.5	0.5	56.8	23.7	9.7	25.6
527	R00Y_075_025d	0.75	0.5	0.625	0.75	0.75	0.5	0.625	57.6	28.2	-6.6	28.9
528	B50R_075_025d	0.75	0.5	0.75	0.75	0.75	0.5	0.75	58.7	33.5	-22.4	40.4
529	B34R_087_037d	0.75	0.5	0.875	0.875	0.375	0.687	311	60.0	39.7	-37.6	54.7
530	B25R_100_050d	0.75	0.5	1.0	1.0	0.5	0.75	300	61.4	46.4	-52.2	69.9
531	R85Y_075_075d	0.75	0.625	0.0	0.75	0.75	0.625	0.0	63.2	-0.7	67.1	67.1
532	R81Y_075_062d	0.75	0.625	0.125	0.75	0.75	0.625	0.125	63.3	-0.2	61.6	61.6
533	R76Y_075_050d	0.75	0.625	0.25	0.75	0.75	0.625	0.25	63.5	1.0	50.4	50.5
534	R68Y_075_037d	0.75	0.625	0.375	0.75	0.75	0.625	0.375	63.8	3.4	35.9	36.1
535	R50Y_075_025d	0.75	0.625	0.5	0.75	0.75	0.625	0.5	64.3	6.8	20.2	21.3
536	R00Y_075_012d	0.75	0.625	0.625	0.75	0.75	0.625	0.625	65.0	11.2	4.2	12.0
537	B50R_075_012d	0.75	0.625	0.75	0.75	0.75	0.625	0.75	65.9	16.6	-11.5	20.2
538	B25R_087_025d	0.75	0.625	0.875	0.875	0.25	0.75	300	67.0	22.8	-26.8	35.2
539	B15R_100_037d	0.75	0.625	1.0	1.0	0.375	0.812	289	68.2	29.7	-41.5	51.1
540	Y00G_075_075d	0.75	0.75	0.0	0.75	0.75	0.75	0.0	71.5	-16.6	73.0	74.9
541	Y00G_075_062d	0.75	0.75	0.125	0.75	0.75	0.75	0.125	71.5	-16.1	68.5	70.4
542	Y00G_075_050d	0.75	0.75	0.25	0.75	0.75	0.75	0.25	71.7	-14.8	58.9	68.8
543	Y00G_075_037d	0.75	0.75	0.375	0.75	0.75	0.75	0.375	72.0	-12.6	45.8	47.5
544	Y00G_075_025d	0.75	0.75	0.5	0.75	0.75	0.75	0.5	72.4	-9.4	30.9	32.3
545	Y00G_075_012d	0.75	0.75	0.625	0.75	0.75	0.75	0.625	73.0	-5.1	15.4	16.3
546	NW_075d	0.75	0.75	0.75	0.75	0.75	0.75	0.75	73.7	0.0	0.0	0.0
547	B00R_087_012d	0.75	0.75	0.875	0.875	0.125	0.812	270	74.6	6.0	-15.2	16.4
548	B00R_100_025d	0.75	0.75	1.0	1.0	0.25	0.875	270	75.6	-12.8	-30.0	32.7
549	Y13G_087_087d	0.75	0.875	0.0	0.875	0.875	0.437	98	79.9	-31.3	79.3	85.3
550	Y15G_087_075d	0.75	0.875	0.125	0.875	0.75	0.5	99	80.0	-30.9	75.6	81.7
551	Y18G_087_062d	0.75	0.875	0.25	0.875	0.625	0.562	101	80.1	-29.7	67.3	73.6
552	Y23G_087_050d	0.75	0.875	0.375	0.875	0.5	0.625	104	80.4	-27.6	55.5	62.0
553	Y31G_087_037d	0.75	0.875	0.5	0.875	0.375	0.687	109	80.7	-24.6	41.5	48.3
554	Y50G_087_025d	0.75	0.875	0.625	0.875	0.25	0.75	120	81.2	-20.7	26.7	33.8
555	G00B_087_012d	0.75	0.875	0.75	0.875	0.125	0.812	150	81.8	-15.7	11.6	19.6
556	G50B_087_012d	0.75	0.875	0.875	0.875	0.125	0.812	210	82.5	-10.0	-3.3	10.5
557	G75B_100_025d	0.75	0.875	1.0	1.0	0.25	0.875	240	83.4	-3.4	-18.1	18.5
558	Y23G_100_100d	0.75	1.0	0.0	1.0	0.5	0.75	104	88.5	-44.9	85.8	96.8
559	Y26G_100_087d	0.75	1.0	0.125	1.0	0.875	0.562	106	88.5	-44.2	82.7	93.9
560	Y31G_100_075d	0.75	1.0	0.25	1.0	0.75	0.625	109	88.6	-43.4	75.5	87.1
561	Y38G_100_062d	0.75	1.0	0.375	1.0	0.625	0.687	113	88.8	-41.5	64.8	77.0
562	Y50G_100_050d	0.75	1.0	0.5	1.0	0.5	0.75	120	89.1	-38.7	51.9	64.8
563	Y68G_100_037d	0.75	1.0	0.625	1.0	0.375	0.812	131	89.5	-35.1	37.8	51.6
564	G00B_100_025d	0.75	1.0	0.75	1.0	0.25	0.875	150	90.1	-30.5	23.2	38.3
565	G25B_100_025d	0.75	1.0	0.875	1.0	0.25	0.875	180	90.7	-25.1	8.4	26.5
566	G50B_100_025d	0.75	1.0	1.0	1.0	0.25	0.875	210	91.5	-18.9	-6.2	19.9

delta E* = 9.4

Grafico TUB-PI31; riproduzione del colore
colori e la differenza, ΔE^* , 3D=0, de=0, sRGB

Input: $rgb/cmyk \rightarrow rgb_d$
Output: trasferire a rgb_d

iscrizione TUB: 20160501-PI31/PI31L0NP.PDF /.PS
Applicazione per la misura dell'output su display, nessuna separazione

TUB materiale: code=rha4ta

vedi file simili: <http://farbe.li.tu-berlin.de/PI31/PI31.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md
567	R00Y_087_087a	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.0	44.1 67.3 56.4	87.8 40.0	0.875 0.0 0.0	44.1 69.5 58.3	39.9 2.9 389	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0
568	R36Y_087_087a	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.116	44.2 67.7 47.1	82.5 34.8	0.875 0.0 0.125	44.2 69.9 47.2 84.3	34.0 2.2 382	1.0 0.0 0.133	50.6 77.3 53.9 94.3 34.8
569	R23Y_087_087a	0.875 0.0 0.25	0.875 0.875 0.437	374	0.875 0.0 0.233	44.5 68.5 32.2	75.7 25.1	0.875 0.0 0.25	44.5 70.8 30.2 77.0	23.1 2.9 375	1.0 0.0 0.266	50.9 78.3 36.8 86.6 25.1
570	R08Y_087_087a	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.364	45.1 70.2 13.8	71.6 11.1	0.875 0.0 0.375	45.1 72.4 12.2 73.4	9.5 2.6 365	1.0 0.0 0.416	51.5 80.3 15.8 81.8 11.1
571	B70R_087_087a	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.51	46.1 72.8 -6.0	73.0 355.2	0.875 0.0 0.5	46.0 74.6 -5.3 74.8	355.8 1.9 354	1.0 0.0 0.583	52.7 83.2 -6.9 83.5 355.2
572	B30R_087_087a	0.875 0.0 0.625	0.875 0.875 0.437	346	0.875 0.0 0.641	47.2 75.6 -23.1	79.1 342.9	0.875 0.0 0.625	47.1 77.6 -22.1 80.7	344.0 2.2 344	1.0 0.0 0.733	54.0 86.5 -26.4 90.4 342.9
573	B56R_087_087a	0.875 0.0 0.75	0.875 0.875 0.437	338	0.875 0.0 0.758	48.6 78.8 -37.5	87.3 334.5	0.875 0.0 0.75	48.5 81.2 -37.9 89.6	334.9 2.3 337	1.0 0.0 0.866	55.5 90.1 -42.8 99.8 334.5
574	B63R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	50.1 82.5 -51.1	97.1 328.2	0.875 0.0 0.875	50.2 85.3 -52.8 100.3	328.2 3.2 330	1.0 0.0 1.0	57.2 94.3 -58.4 110.9 328.2
575	B44R_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	52.5 90.1 -66.3	111.9 323.6	0.875 0.0 1.0	52.1 89.8 -66.9 112.0	323.3 0.7 323	0.883 0.0 1.0	52.5 90.1 -66.3 111.9 323.6
576	R13Y_087_087a	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.116 0.0	45.2 64.2 56.9	85.8 41.5	0.875 0.125 0.0	45.3 65.8 58.8	88.3 41.7 2.5 37	1.0 0.133 0.0	51.7 73.4 65.0 98.0 41.5
577	R00Y_087_075a	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.125	49.7 57.7 48.4	75.3 40.0	0.875 0.125 0.125	45.5 66.2 48.3	81.9 36.0 9.5 389	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0
578	R35Y_087_075a	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.237	49.9 58.2 38.8	69.9 33.6	0.875 0.125 0.25	45.8 67.1 31.8	74.3 25.3 12.0 382	1.0 0.0 0.15	50.6 77.6 51.7 93.3 33.6
579	R18Y_087_075a	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.362	50.2 59.3 22.3	63.4 20.6	0.875 0.125 0.375	46.4 68.8 13.9	70.2 11.4 13.2 371	1.0 0.0 0.316	51.1 79.1 29.7 84.5 20.6
580	R00Y_087_075a	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.5	50.9 60.8 3.1	60.9 2.9	0.875 0.125 0.5	47.2 71.1 -3.6	71.2 357.1 12.8 360	1.0 0.0 0.5	52.0 81.1 4.1 81.2 2.9
581	B65R_087_075a	0.875 0.125 0.625	0.875 0.75 0.5	349	0.875 0.125 0.637	52.1 64.1 -14.9	65.8 346.8	0.875 0.125 0.625	48.3 74.2 -20.3	76.9 344.6 12.0 348	1.0 0.0 0.683	53.5 85.8 -19.9 87.7 346.8
582	B57R_087_075a	0.875 0.125 0.75	0.875 0.75 0.5	339	0.875 0.125 0.762	53.4 67.3 -30.5	73.9 335.5	0.875 0.125 0.75	49.6 77.9 -36.1	85.9 335.1 12.5 337	1.0 0.0 0.85	55.3 89.4 -40.7 98.6 335.5
583	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	54.9 70.7 -43.8	83.2 328.2	0.875 0.125 0.875	51.3 82.1 -51.1	96.7 328.1 14.0 330	1.0 0.0 1.0	57.2 94.3 -58.4 110.9 328.2
584	B43R_100_087a	0.875 0.125 1.0	1.0 1.0 0.875	322	0.883 0.125 1.0	57.3 78.4 -59.0	89.1 323.0	0.875 0.125 1.0	53.1 86.9 -65.3	108.7 323.0 11.3 322	0.866 0.0 1.0	51.8 89.6 -67.4 112.1 323.0
585	R26Y_087_087a	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.233 0.0	47.0 57.0 58.0	81.3 45.5	0.875 0.25 0.0	48.4 57.3 60.1	83.0 46.3 2.1 44	1.0 0.266 0.0	54.6 65.1 66.3 93.0 45.5
586	R15Y_087_075a	0.875 0.25 0.125	0.875 0.75 0.5	39	0.875 0.237 0.125	50.9 54.3 48.9	73.1 41.9	0.875 0.25 0.125	48.5 57.7 50.8	76.9 41.3 4.5 37	1.0 0.15 0.0	52.0 72.4 65.2 97.4 41.9
587	R00Y_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	55.4 48.0 40.3	62.7 40.0	0.875 0.25 0.25	48.8 58.7 35.3	68.5 31.0 13.4 389	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0
588	R31Y_087_062a	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.364	55.5 48.7 29.7	57.0 31.3	0.875 0.25 0.375	49.3 60.4 17.9	63.0 16.5 17.7 380	1.0 0.0 0.183	50.7 77.9 47.5 91.2 31.3
589	R11Y_087_062a	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.489	55.9 49.6 12.8	51.3 14.4	0.875 0.25 0.5	50.1 63.0 0.6	63.0 0.5 18.9 367	1.0 0.0 0.383	51.4 79.5 20.4 82.1 14.4
590	B69R_087_062a	0.875 0.25 0.625	0.875 0.625 0.562	353	0.875 0.25 0.635	56.9 52.2 -7.1	52.7 352.1	0.875 0.25 0.625	51.1 66.2 -16.0	68.1 346.3 17.5 352	1.0 0.0 0.616	52.9 83.6 -11.4 84.3 352.1
591	B59R_087_062a	0.875 0.25 0.75	0.875 0.625 0.562	341	0.875 0.25 0.76	58.2 55.5 -22.8	60.1 337.6	0.875 0.25 0.75	52.3 70.2 -31.9	77.1 335.5 18.1 339	1.0 0.0 0.816	54.9 88.9 -36.6 96.2 337.6
592	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	59.6 58.9 -36.5	69.3 328.2	0.875 0.25 0.875	53.8 74.7 -47.0	88.3 327.8 19.8 330	1.0 0.0 1.0	57.2 94.3 -58.4 110.9 328.2
593	B42R_100_075a	0.875 0.25 1.0	1.0 1.0 0.75	321	0.887 0.25 1.0	62.2 66.8 -51.4	84.3 324.4	0.875 0.25 1.0	55.6 79.8 -61.3	100.7 324.4 17.6 322	0.85 0.0 1.0	51.2 89.1 -68.5 112.4 322.4
594	R41Y_087_087a	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.364 0.0	52.5 44.4 60.6	75.1 53.7	0.875 0.375 0.0	53.2 44.5 62.6	76.8 54.5 2.1 54	1.0 0.416 0.0	60.0 50.7 69.3 85.9 53.7
595	R31Y_087_075a	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.362 0.125	54.1 45.5 50.4	67.9 47.9	0.875 0.375 0.125	53.3 44.9 54.7	70.8 50.6 4.1 48	1.0 0.316 0.0	56.2 60.6 67.2 90.5 47.9
596	R18Y_087_062a	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.364 0.25	56.8 44.0 40.9	60.1 42.8	0.875 0.375 0.25	53.5 45.9 40.7	61.4 41.5 3.7 39	1.0 0.183 0.0	52.7 70.5 65.5 96.2 42.8
597	R00Y_087_050a	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	61.0 38.4 32.2	50.2 40.0	0.875 0.375 0.375	54.0 47.8 24.1	53.6 26.8 14.2 389	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0
598	R26Y_087_050a	0.875 0.375 0.5	0.875 0.5 0.625	376	0.875 0.375 0.491	61.1 39.0 20.6	44.1 27.8	0.875 0.375 0.5	54.6 50.5 7.2	51.0 8.1 18.8 377	1.0 0.0 0.233	50.8 78.0 41.2 88.2 27.8
599	R00Y_087_050a	0.875 0.375 0.625	0.875 0.5 0.625	360	0.875 0.375 0.625	61.8 40.5 2.0	40.6 2.9	0.875 0.375 0.625	55.5 54.0 -9.3	54.8 350.2 18.7 360	1.0 0.0 0.5	52.0 81.1 4.1 81.2 2.9
600	B61R_087_050a	0.875 0.375 0.75	0.875 0.5 0.625	344	0.875 0.375 0.758	62.9 43.6 -15.3	46.2 340.6	0.875 0.375 0.75	56.6 58.3 -25.2	63.5 336.5 18.7 342	1.0 0.0 0.766	54.4 87.3 -30.6 92.5 340.6
601	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	64.4 47.1 -29.2	55.4 328.2	0.875 0.375 0.875	58.0 63.2 -40.5	75.0 327.3 20.6 330	1.0 0.0 1.0	57.2 94.3 -58.4 110.9 328.2
602	B40R_100_062a	0.875 0.375 1.0	1.0 1.0 0.625	318	0.885 0.375 1.0	66.9 55.0 -44.2	70.6 321.2	0.875 0.375 1.0	59.5 68.6 -55.0	88.0 321.2 18.8 320	0.816 0.0 1.0	49.8 88.1 -70.7 113.0 321.2
603	R58Y_087_087a	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.51 0.0	59.4 27.1 64.8	70.2 67.2	0.875 0.5 0.0	59.4 29.0 66.2	72.3 66.2 2.4 65	1.0 0.583 0.0	67.9 31.0 74.0 80.3 67.2
604	R50Y_087_075a	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.5 0.125	59.6 31.0 53.2	61.6 59.7	0.875 0.5 0.125	59.4 29.5 59.8	66.7 63.7 6.7 59	1.0 0.5 0.0	63.6 41.3 71.0 82.2 59.7
605	R38Y_087_062a	0.875 0.5 0.25	0.875 0.625 0.562	53	0.875 0.489 0.25	60.4 34.0 42.6	54.6 51.3	0.875 0.5 0.25	59.7 30.6 47.4	56.4 57.1 5.9 52	1.0 0.383 0.0	58.5 54.5 68.2 87.3 51.3
606	R23Y_087_050a	0.875 0.5 0.375	0.875 0.5 0.625	44	0.875 0.491 0.375	62.6 33.8 32.9	47.2 44.2	0.875 0.5 0.375	60.0 32.5 31.9	45.6 44.4 3.0 42	1.0 0.233 0.0	53.7 67.6 65.8 94.4 44.2
607	R00Y_087_037a	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.5	66.6 28.8 24.2	37.6 40.0	0.875 0.5 0.5	60.6 35.3 15.5	38.6 23.7 12.3 389	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0
608	R18Y_087_037a	0.875 0.5 0.625	0.875 0.375 0.687	371	0.875 0.5 0.618	66.8 29.6 11.1	31.7 20.6	0.875 0.5 0.625	61.3 39.0 -0.7	39.0 358.9 16.0 371	1.0 0.0 0.316	51.1 79.1 29.7 84.5 20.6
609	B65R_087_037a	0.875 0.5 0.75	0.875 0.375 0.687	349	0.875 0.5 0.756	67.7 32.0 -7.4	32.9 346.8	0.875 0.5 0.75	62.3 43.5 -16.6	46.5 339.0 15.6 348	1.0 0.0 0.683	53.5 85.4 -19.9 87.7 346.8
610	B50R_087_037a	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	69.1 35.3 -21.9	41.6 328.2	0.875 0.5 0.875	63.5 48.6 -31.9	58.2 326.7 17.6 330	1.0 0.0 1.0	57.2 94.3 -58.4 110.9 328.2
611	B38R_100_050a	0.875 0.5 1.0	1.0 1.0 0.5	316	0.883 0.5 1.0	71.6 43.2 -37.0	56.9 319.4	0.875 0.5 1.0	64.8 54.4 -46.6	71.7 319.3 16.3 317	0.766 0.0 1.0	47.9 86.4 -74.0 113.8 319.4
612	R73Y_087_087a	0.875 0.625 0.0	0.875 0.875 0.437	74	0.875 0.641 0.0	66.7 10.5 69.4	70.2 81.3	0.875 0.625 0.0	66.5 12.7 70.9	72.0 79.7 2.7 75	1.0 0.733 0.0	76.2 12.0 79.3 80.2 81.3
613	R68Y_087_075a	0.875 0.625 0.125	0.875 0.75 0.5	71	0.875 0.637 0.125	67.0 13.8 58.2	59.8 76.5	0.875 0.625 0.125	66.5 13.2 65.6	66.9 78.6 7.4 71	1.0 0.683 0.0	73.4 18.5 77.6 79.8 76.5
614	R61Y_087_062a	0.875 0.625 0.25	0.875 0.625 0.562	67	0.875 0.635 0.25	67.4 16.7 46.8	49.7 70.2	0.875 0.625 0.25	66.7 14.3 54.8	56.6 75.3 8.4 67	1.0 0.616 0.0	69.6 26.8 74.8 79.5 70.2
615	R50Y_087_050a	0.875 0.625 0.375	0.875 0.5 0.625	60	0.875 0.625 0.375	67.6 20.6 35.5	41.1 59.7	0.875 0.625 0.375	67.0 16.2 40.6	43.7 68.1 6.7 59	1.0 0.5 0.0	63.6 41.3 71.0 82.2 59.7
616	R31Y_087_037a	0.875 0.625 0.5	0.875 0.375 0.687	49	0.875 0.618 0.5	68.2 22.7 25.2	33.9 47.9	0.875 0.625 0.5	67.5 19.0 24.9	31.4 52.6 3.9 48	1.0 0.316 0.0	56.2 60.6 67.2 90.5 47.9

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md						
648	R00Y_100_100a	1.0	0.0	0.0	1.0	0.5	390	100.4	40.0	0.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0	
649	R38Y_100_100a	1.0	0.0	0.125	1.0	0.5	383	100.0	35.7	0.0	0.0	0.116	50.5	77.2	55.6	95.1	35.7	
650	R26Y_100_100a	1.0	0.0	0.25	1.0	0.5	376	100.0	27.8	0.0	0.0	0.233	50.8	78.0	41.2	88.2	27.8	
651	R13Y_100_100a	1.0	0.0	0.375	1.0	0.5	368	100.0	16.0	0.0	0.0	0.366	51.3	79.3	22.7	82.5	16.0	
652	R00Y_100_100a	1.0	0.0	0.5	1.0	0.5	360	100.0	2.9	0.0	0.0	0.5	52.0	81.1	4.1	81.2	2.9	
653	B68R_100_100a	1.0	0.0	0.625	1.0	0.5	352	100.0	350.7	1.0	0.0	0.633	53.0	83.9	-13.6	85.0	350.7	
654	B61R_100_100a	1.0	0.0	0.75	1.0	0.5	344	100.0	340.6	1.0	0.0	0.766	54.4	87.3	-30.6	92.5	340.6	
655	B55R_100_100a	1.0	0.0	0.875	1.0	0.5	337	100.0	333.6	1.0	0.0	0.883	55.7	90.6	-44.8	101.1	333.6	
656	B50R_100_100a	1.0	0.0	1.0	1.0	0.5	330	100.0	328.2	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	
657	R11Y_100_100a	1.0	0.125	0.0	1.0	0.5	37	100.0	41.2	0.125	0.0	0.116	51.4	74.1	64.9	98.5	41.2	
658	R00Y_100_087a	1.0	0.125	0.125	1.0	0.875	562	390	87.8	0.400	0.0	0.125	51.6	74.2	55.7	92.8	87.8	
659	R36Y_100_087a	1.0	0.125	0.25	1.0	0.875	562	382	34.8	0.125	0.0	0.133	50.6	77.3	53.9	94.3	34.8	
660	R23Y_100_087a	1.0	0.125	0.375	1.0	0.875	562	374	25.1	0.125	0.0	0.266	50.9	78.3	36.8	86.6	25.1	
661	R08Y_100_087a	1.0	0.125	0.5	1.0	0.875	562	365	11.1	0.125	0.0	0.416	51.5	80.3	15.8	81.8	11.1	
662	B70R_100_087a	1.0	0.125	0.625	1.0	0.875	562	355	355.2	0.125	0.0	0.583	52.7	83.2	-6.9	83.5	355.2	
663	B63R_100_087a	1.0	0.125	0.75	1.0	0.875	562	346	342.9	0.125	0.0	0.733	54.0	86.5	-26.4	90.4	342.9	
664	B56R_100_087a	1.0	0.125	0.875	1.0	0.875	562	338	334.5	0.125	0.0	0.866	55.5	90.1	-42.8	99.8	334.5	
665	B50R_100_087a	1.0	0.125	1.0	1.0	0.875	562	330	328.2	0.125	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	
666	R23Y_100_100a	1.0	0.25	0.0	1.0	0.5	44	100.0	44.2	0.25	0.0	0.233	0.0	53.7	67.6	65.8	44.2	
667	R13Y_100_100a	1.0	0.25	0.125	1.0	0.875	562	38	41.5	0.125	0.0	0.133	0.0	51.7	73.4	65.0	41.5	
668	R00Y_100_075a	1.0	0.25	0.25	1.0	0.75	625	390	75.0	0.25	0.0	0.0	50.4	76.9	64.5	100.4	75.0	
669	R35Y_100_075a	1.0	0.25	0.375	1.0	0.75	625	381	33.6	0.25	0.0	0.15	50.6	77.6	51.7	93.3	33.6	
670	R18Y_100_075a	1.0	0.25	0.5	1.0	0.75	625	371	20.6	0.25	0.0	0.316	51.1	79.1	29.7	84.5	20.6	
671	R00Y_100_075a	1.0	0.25	0.625	1.0	0.75	625	360	9.9	0.25	0.0	0.5	52.0	81.1	4.1	81.2	9.9	
672	B65R_100_075a	1.0	0.25	0.75	1.0	0.75	625	349	346.8	0.25	0.0	0.683	53.5	85.4	-19.9	87.7	346.8	
673	B57R_100_075a	1.0	0.25	0.875	1.0	0.75	625	339	335.5	0.25	0.0	0.85	55.3	89.8	-40.7	98.6	335.5	
674	B50R_100_075a	1.0	0.25	1.0	1.0	0.75	625	330	328.2	0.25	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	
675	R36Y_100_100a	1.0	0.375	0.0	1.0	0.5	52	100.0	50.3	0.375	0.0	0.366	0.0	57.9	56.2	67.9	50.3	
676	R15Y_100_100a	1.0	0.375	0.125	1.0	0.875	562	46	45.5	0.375	0.0	0.266	0.0	54.6	65.1	66.3	45.5	
677	R15Y_100_075a	1.0	0.375	0.25	1.0	0.75	625	39	41.9	0.375	0.0	0.15	50.2	72.4	65.2	97.4	41.9	
678	R00Y_100_062a	1.0	0.375	0.375	1.0	0.625	687	390	62.7	0.375	0.0	0.0	50.4	76.9	64.5	100.4	62.7	
679	R31Y_100_062a	1.0	0.375	0.5	1.0	0.625	687	379	31.3	0.375	0.0	0.183	50.7	77.9	47.5	91.2	31.3	
680	R11Y_100_062a	1.0	0.375	0.625	1.0	0.625	687	367	14.4	0.375	0.0	0.383	51.4	79.5	20.4	82.1	14.4	
681	B69R_100_062a	1.0	0.375	0.75	1.0	0.625	687	353	352.1	0.375	0.0	0.616	52.9	83.6	-11.4	84.3	352.1	
682	B59R_100_062a	1.0	0.375	0.875	1.0	0.625	687	341	337.6	0.375	0.0	0.816	54.9	88.9	-36.6	96.2	337.6	
683	B50R_100_062a	1.0	0.375	1.0	1.0	0.625	687	330	328.2	0.375	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	
684	R50Y_100_100a	1.0	0.5	0.0	1.0	0.5	60	100.0	59.7	0.5	0.0	0.5	63.6	41.3	71.0	82.2	59.7	
685	R41Y_100_087a	1.0	0.5	0.125	1.0	0.875	562	55	53.7	0.5	0.125	0.416	60.0	50.7	69.3	85.9	53.7	
686	R31Y_100_075a	1.0	0.5	0.25	1.0	0.75	625	49	47.9	0.5	0.25	0.316	50.6	60.6	67.2	90.5	47.9	
687	R18Y_100_062a	1.0	0.5	0.375	1.0	0.625	687	41	42.8	0.5	0.375	0.183	50.7	70.5	65.5	96.2	42.8	
688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	75	390	40.0	0.5	0.5	0.5	50.4	76.9	64.5	100.4	40.0	
689	R26Y_100_050a	1.0	0.5	0.625	1.0	0.5	75	376	27.8	0.5	0.625	0.233	50.8	78.0	41.2	88.2	27.8	
690	R00Y_100_050a	1.0	0.5	0.75	1.0	0.5	75	360	9.9	0.5	0.75	0.5	52.0	81.1	4.1	81.2	9.9	
691	B61R_100_050a	1.0	0.5	0.875	1.0	0.5	75	344	340.6	0.5	0.875	0.766	54.4	87.3	-30.6	92.5	340.6	
692	B50R_100_050a	1.0	0.5	1.0	1.0	0.5	75	330	328.2	0.5	1.0	1.0	57.2	94.3	-58.4	110.9	328.2	
693	R63Y_100_100a	1.0	0.625	0.0	1.0	0.5	68	100.0	68.0	0.625	0.0	0.633	0.0	70.5	24.7	75.4	79.4	71.8
694	R58Y_100_087a	1.0	0.625	0.125	1.0	0.875	562	65	61.2	0.625	0.125	0.583	0.0	67.9	31.0	74.0	80.3	67.2
695	R50Y_100_075a	1.0	0.625	0.25	1.0	0.75	625	60	59.7	0.625	0.25	0.5	63.6	41.3	71.0	82.2	59.7	
696	R38Y_100_062a	1.0	0.625	0.375	1.0	0.625	687	53	51.3	0.625	0.375	0.383	0.0	58.5	45.5	68.2	87.3	51.3
697	R23Y_100_050a	1.0	0.625	0.5	1.0	0.5	75	44	44.2	0.625	0.5	0.233	0.0	53.7	67.6	65.8	94.4	44.2
698	R00Y_100_037a	1.0	0.625	0.625	1.0	0.375	812	390	37.0	0.625	0.625	0.0	50.4	76.9	64.5	100.4	40.0	
699	R18Y_100_037a	1.0	0.625	0.75	1.0	0.375	812	371	20.6	0.625	0.75	0.316	51.1	79.1	29.7	84.5	20.6	
700	B65R_100_037a	1.0	0.625	0.875	1.0	0.375	812	349	346.8	0.625	0.875	0.683	53.5	85.4	-19.9	87.7	346.8	
701	B50R_100_037a	1.0	0.625	1.0	1.0	0.375	812	330	328.2	0.625	1.0	1.0	57.2	94.3	-58.4	110.9	328.2	
702	R76Y_100_100a	1.0	0.75	0.0	1.0	0.5	76	100.0	84.4	0.75	0.0	0.7	78.0	79.7	80.3	82.9	2.3	77
703	R73Y_100_087a	1.0	0.75	0.125	1.0	0.875	562	74	70.2	0.75	0.125	0.733	0.0	76.2	12.0	79.3	80.2	81.3
704	R68Y_100_075a	1.0	0.75	0.25	1.0	0.75	625	71	59.7	0.75	0.25	0.683	0.0	73.4	18.5	77.6	79.8	76.5
705	R61Y_100_062a	1.0	0.75	0.375	1.0	0.625	687	67	60.7	0.75	0.375	0.616	0.0	69.6	26.8	74.8	79.5	70.2
706	R50Y_100_050a	1.0	0.75	0.5	1.0	0.5	75	60	59.7	0.75	0.5	0.5	63.6	41.3	71.0	82.2	59.7	
707	R31Y_100_037a	1.0	0.75	0.625	1.0	0.375	812	49	47.9	0.75	0.625	0.316	50.6	60.6	67.2	90.5	47.9	
708	R00Y_100_025a	1.0	0.75	0.75	1.0	0.25	875	390	29.0	0.75	0.75	0.0	50.4	76.9	64.5	100.4	40.0	
709	R00Y_100_025a	1.0	0.75	0.875	1.0	0.25	875	360	20.3	0.75	0.875	0.0	50.4	76.9	64.5	100.4	20.3	
710	B50R_100_025a	1.0	0.75	1.0	1.0	0.25	875	330	27.7	0.75	1.0	1.0	57.2	94.3	-58.4	110.9	328.2	
711	R88Y_100_100a	1.0	0.875	0.0	1.0	0.5	83	100.0	84.4	0.875	0.0	0.883	0.0	85.3	-6.7	85.5	85.8	94.4
712	R86Y_100_087a	1.0	0.875	0.125	1.0	0.875	562	82	93.1	0.875	0.125	0.866	0.0	84.3	-4.6	84.8	84.9	93.1
713	R85Y_100_075a	1.0	0.875	0.25	1.0	0.75	625	81	91.7	0.875	0.25	0.85	0.0	83.3	-2.5	84.2	84.3	91.7
714	R81Y_100_062a	1.0	0.875	0.375	1.0	0.625	687	79	88.7	0.875	0.375	0.816	0.0	81.2	1.7	82.9	83.0	88.7
715	R76Y_100_050a	1.0	0.875	0.5	1.0	0.5	75	76	84.4	0.875	0.5	0.766	0.0	78.2	7.8	80.6	81.0	84.4
716	R68Y_100_037a	1.0	0.875	0.625	1.0	0.375	812	71	76.5	0.875	0.625	0.683	0.0	73.4	18.5	77.6	79.8	76.5
717	R50Y_100_025a	1.0	0.875	0.75	1.0	0.25	875	60	59.7	0.875	0.75	0.5	63.6	41.3	71.0	82.2	59.7	
718	R00Y_100_012a	1.0	0.875	0.875	1.0	0.125	937	390	12.5	0.875	0.875	0.0	50.4	76.9	64.5</			

vedi file simili: <http://farbe.li.tu-berlin.de/PI31/PI31.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

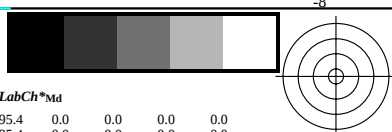
n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md
729	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0
730	G50B_100_012d	0.875 1.0 1.0	1.0 0.125 0.937	210	0.875 1.0 1.0	94.3 -5.7 -1.6	6.0 0.0 0.0	0.875 1.0 1.0	93.3 -9.7 -3.3	10.3 198.8 4.4	210 0.0 1.0 1.0	86.8 -46.1 -13.5
731	G50B_100_025d	0.75 1.0 1.0	1.0 0.25 0.875	210	0.75 1.0 1.0	93.2 -11.5 -3.3	12.0 196.3 0.0	0.75 1.0 1.0	91.5 -18.9 -6.2	19.9 198.1 8.0	210 0.0 1.0 1.0	86.8 -46.1 -13.5
732	G50B_100_037d	0.625 1.0 1.0	1.0 0.375 0.812	210	0.625 1.0 1.0	92.2 -17.3 -5.0	18.0 196.3 0.0	0.625 1.0 1.0	90.0 -27.0 -8.5	28.3 197.6 10.5	210 0.0 1.0 1.0	86.8 -46.1 -13.5
733	G50B_100_050d	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	91.1 -23.0 -6.7	24.0 196.3 0.0	0.5 1.0 1.0	88.8 -33.9 -10.4	35.4 197.1 11.6	210 0.0 1.0 1.0	86.8 -46.1 -13.5
734	G50B_100_062d	0.375 1.0 1.0	1.0 0.625 0.687	210	0.375 1.0 1.0	90.0 -28.8 -8.4	30.0 196.3 0.0	0.375 1.0 1.0	87.9 -39.3 -11.8	41.0 196.8 11.1	210 0.0 1.0 1.0	86.8 -46.1 -13.5
735	G50B_100_075d	0.25 1.0 1.0	1.0 0.75 0.625	210	0.25 1.0 1.0	89.0 -34.6 -10.1	36.1 196.3 0.0	0.25 1.0 1.0	87.3 -43.0 -12.8	44.9 196.5 9.0	210 0.0 1.0 1.0	86.8 -46.1 -13.5
736	G50B_100_087d	0.125 1.0 1.0	1.0 0.875 0.562	210	0.125 1.0 1.0	87.9 -40.4 -11.8	42.1 196.3 0.0	0.125 1.0 1.0	87.0 -45.2 -13.3	47.2 196.4 5.1	210 0.0 1.0 1.0	86.8 -46.1 -13.5
737	G50B_100_100d	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3 0.0	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3 0.0	210 0.0 1.0 1.0	86.8 -46.1 -13.5
738	RO0Y_100_012d	1.0 0.875 0.875	1.0 0.125 0.937	390	1.0 0.875 0.875	96.7 8.0 12.5	40.0 0.0 0.0	1.0 0.875 0.875	87.1 10.5 3.8	11.2 201.0 5.0	389 1.0 0.0 0.0	50.4 76.9 64.5
739	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	83.4 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	84.7 0.0 0.0	0.0 325.2 1.2	360 1.0 1.0 1.0	95.4 0.0 0.0
740	G50B_087_012d	0.75 0.875 0.875	0.875 0.125 0.812	210	0.75 0.875 0.875	82.4 -5.7 -1.6	6.0 196.3 0.0	0.75 0.875 0.875	82.5 -10.0 -3.3	10.5 198.7 4.5	210 0.0 1.0 1.0	86.8 -46.1 -13.5
741	G50B_087_025d	0.625 0.875 0.875	0.875 0.25 0.75	210	0.625 0.875 0.875	81.3 -11.5 -3.3	12.0 196.3 0.0	0.625 0.875 0.875	80.7 -19.1 -6.2	20.1 197.9 8.1	210 0.0 1.0 1.0	86.8 -46.1 -13.5
742	G50B_087_037d	0.5 0.875 0.875	0.875 0.375 0.687	210	0.5 0.875 0.875	80.2 -17.3 -5.0	18.0 196.3 0.0	0.5 0.875 0.875	79.3 -27.1 -8.5	28.4 197.4 10.4	210 0.0 1.0 1.0	86.8 -46.1 -13.5
743	G50B_087_050d	0.375 0.875 0.875	0.875 0.5 0.625	210	0.375 0.875 0.875	79.2 -23.0 -6.7	24.0 196.3 0.0	0.375 0.875 0.875	78.3 -33.4 -10.2	34.9 196.9 10.9	210 0.0 1.0 1.0	86.8 -46.1 -13.5
744	G50B_087_062d	0.25 0.875 0.875	0.875 0.625 0.562	210	0.25 0.875 0.875	78.1 -28.8 -8.4	30.0 196.3 0.0	0.25 0.875 0.875	77.5 -37.9 -11.3	39.6 196.6 9.5	210 0.0 1.0 1.0	86.8 -46.1 -13.5
745	G50B_087_075d	0.125 0.875 0.875	0.875 0.75 0.5	210	0.125 0.875 0.875	77.0 -34.6 -10.1	36.1 196.3 0.0	0.125 0.875 0.875	77.1 -40.6 -12.0	42.4 196.4 6.3	210 0.0 1.0 1.0	86.8 -46.1 -13.5
746	G50B_087_087d	0.0 0.875 0.875	0.875 0.875 0.437	210	0.0 0.875 0.875	76.0 -40.4 -11.8	42.1 196.3 0.0	0.0 0.875 0.875	77.0 -41.7 -12.2	43.5 196.3 1.7	210 0.0 1.0 1.0	86.8 -46.1 -13.5
747	RO0Y_100_025d	1.0 0.75 0.75	1.0 0.25 0.875	390	1.0 0.75 0.75	84.2 19.2 16.1	25.1 40.0 0.0	1.0 0.75 0.75	79.2 21.9 8.5	23.5 21.3 9.4	389 1.0 0.0 0.0	50.4 76.9 64.5
748	RO0Y_087_012d	0.875 0.75 0.75	0.875 0.125 0.812	390	0.875 0.75 0.75	77.8 9.6 8.0	12.5 40.0 0.0	0.875 0.75 0.75	76.2 10.8 4.0	11.6 20.3 4.5	389 1.0 0.0 0.0	50.4 76.9 64.5
749	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	71.5 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	73.7 0.0 0.0	0.0 325.2 2.1	360 1.0 1.0 1.0	95.4 0.0 0.0
750	G50B_075_012d	0.625 0.75 0.75	0.75 0.125 0.687	210	0.625 0.75 0.75	70.4 -5.7 -1.6	6.0 196.3 0.0	0.625 0.75 0.75	71.5 -10.2 -3.4	10.8 198.5 4.9	210 0.0 1.0 1.0	86.8 -46.1 -13.5
751	G50B_075_025d	0.5 0.75 0.75	0.75 0.25 0.625	210	0.5 0.75 0.75	69.4 -11.5 -3.3	12.0 196.3 0.0	0.5 0.75 0.75	69.8 -19.4 -6.2	20.3 197.8 8.3	210 0.0 1.0 1.0	86.8 -46.1 -13.5
752	G50B_075_037d	0.375 0.75 0.75	0.75 0.375 0.562	210	0.375 0.75 0.75	68.3 -17.3 -5.0	18.0 196.3 0.0	0.375 0.75 0.75	68.4 -26.9 -8.3	28.2 197.1 10.1	210 0.0 1.0 1.0	86.8 -46.1 -13.5
753	G50B_075_050d	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.75	67.2 -23.0 -6.7	24.0 196.3 0.0	0.25 0.75 0.75	67.5 -32.5 -9.7	33.9 196.7 9.8	210 0.0 1.0 1.0	86.8 -46.1 -13.5
754	G50B_075_062d	0.125 0.75 0.75	0.75 0.625 0.437	210	0.125 0.75 0.75	66.2 -28.8 -8.4	30.0 196.3 0.0	0.125 0.75 0.75	67.0 -35.8 -10.6	37.3 196.4 7.3	210 0.0 1.0 1.0	86.8 -46.1 -13.5
755	G50B_075_075d	0.0 0.75 0.75	0.75 0.75 0.375	210	0.0 0.75 0.75	65.1 -34.6 -10.1	36.1 196.3 0.0	0.0 0.75 0.75	66.8 -37.7 -10.9	38.7 196.3 3.1	210 0.0 1.0 1.0	86.8 -46.1 -13.5
756	RO0Y_100_037d	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	78.5 28.8 24.2	37.6 40.0 0.0	1.0 0.625 0.625	71.6 34.1 14.4	37.0 22.9 13.0	389 1.0 0.0 0.0	50.4 76.9 64.5
757	RO0Y_087_025d	0.875 0.625 0.625	0.875 0.25 0.75	390	0.875 0.625 0.625	72.2 19.2 16.1	25.1 40.0 0.0	0.875 0.625 0.625	68.1 22.7 9.0	24.5 21.7 8.8	389 1.0 0.0 0.0	50.4 76.9 64.5
758	RO0Y_075_012d	0.75 0.625 0.625	0.75 0.125 0.687	390	0.75 0.625 0.625	65.9 9.6 8.0	12.5 40.0 0.0	0.75 0.625 0.625	65.0 11.2 4.2	12.0 20.4 4.2	389 1.0 0.0 0.0	50.4 76.9 64.5
759	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	59.6 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	62.4 0.0 0.0	0.0 325.2 2.7	360 1.0 1.0 1.0	95.4 0.0 0.0
760	G50B_062_012d	0.5 0.625 0.625	0.625 0.125 0.562	210	0.5 0.625 0.625	58.5 -5.7 -1.6	6.0 196.3 0.0	0.5 0.625 0.625	60.1 -10.5 -3.5	11.0 198.4 5.3	210 0.0 1.0 1.0	86.8 -46.1 -13.5
761	G50B_062_025d	0.375 0.625 0.625	0.625 0.25 0.5	210	0.375 0.625 0.625	57.4 -11.5 -3.3	12.0 196.3 0.0	0.375 0.625 0.625	58.5 -19.5 -6.1	20.5 197.5 8.5	210 0.0 1.0 1.0	86.8 -46.1 -13.5
762	G50B_062_037d	0.25 0.625 0.625	0.625 0.375 0.437	210	0.25 0.625 0.625	56.4 -17.3 -5.0	18.0 196.3 0.0	0.25 0.625 0.625	57.3 -26.4 -8.0	27.6 196.9 9.6	210 0.0 1.0 1.0	86.8 -46.1 -13.5
763	G50B_062_050d	0.125 0.625 0.625	0.625 0.5 0.375	210	0.125 0.625 0.625	55.3 -23.0 -6.7	24.0 196.3 0.0	0.125 0.625 0.625	56.6 -30.7 -9.1	32.0 196.5 8.0	210 0.0 1.0 1.0	86.8 -46.1 -13.5
764	G50B_062_062d	0.0 0.625 0.625	0.625 0.625 0.312	210	0.0 0.625 0.625	54.2 -28.8 -8.4	30.0 196.3 0.0	0.0 0.625 0.625	56.3 -32.4 -9.5	33.8 196.3 4.3	210 0.0 1.0 1.0	86.8 -46.1 -13.5
765	RO0Y_100_050d	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	72.9 38.4 32.2	50.2 40.0 0.0	1.0 0.5 0.5	64.7 46.4 21.9	51.3 25.2 15.4	389 1.0 0.0 0.0	50.4 76.9 64.5
766	RO0Y_087_037d	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.5	66.6 28.8 24.2	37.6 40.0 0.0	0.875 0.5 0.5	60.6 35.3 15.5	38.6 23.7 12.3	389 1.0 0.0 0.0	50.4 76.9 64.5
767	RO0Y_075_025d	0.75 0.5 0.5	0.75 0.25 0.625	390	0.75 0.5 0.5	60.3 19.2 16.1	25.1 40.0 0.0	0.75 0.5 0.5	56.8 23.7 9.7	25.6 22.2 8.5	389 1.0 0.0 0.0	50.4 76.9 64.5
768	RO0Y_062_012d	0.625 0.5 0.5	0.625 0.125 0.562	390	0.625 0.5 0.5	54.0 9.6 8.0	12.5 40.0 0.0	0.625 0.5 0.5	53.4 11.7 4.4	12.6 20.7 4.2	389 1.0 0.0 0.0	50.4 76.9 64.5
769	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	47.7 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	50.6 0.0 0.0	0.0 325.3 2.9	360 1.0 1.0 1.0	95.4 0.0 0.0
770	G50B_050_012d	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.5	46.6 -5.7 -1.6	6.0 196.3 0.0	0.375 0.5 0.5	48.4 -10.7 -3.5	11.3 198.2 5.6	210 0.0 1.0 1.0	86.8 -46.1 -13.5
771	G50B_050_025d	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	45.5 -11.5 -3.3	12.0 196.3 0.0	0.25 0.5 0.5	46.8 -19.5 -6.0	20.4 197.2 8.5	210 0.0 1.0 1.0	86.8 -46.1 -13.5
772	G50B_050_037d	0.125 0.5 0.5	0.5 0.375 0.312	210	0.124 0.5 0.5	44.5 -17.3 -5.0	18.0 196.3 0.0	0.125 0.5 0.5	45.9 -25.2 -7.5	26.3 196.6 8.3	210 0.0 1.0 1.0	86.8 -46.1 -13.5
773	G50B_050_050d	0.0 0.5 0.5	0.5 0.5 0.25	210	0.0 0.5 0.5	43.4 -23.0 -6.7	24.0 196.3 0.0	0.0 0.5 0.5	45.5 -27.6 -8.1	28.7 196.3 5.1	210 0.0 1.0 1.0	86.8 -46.1 -13.5
774	RO0Y_100_062d	1.0 0.375 0.375	1.0 0.625 0.687	390	1.0 0.375 0.375	67.3 48.0 40.3	62.7 40.0 0.0	1.0 0.375 0.375	58.9 58.1 31.4	66.1 28.3 15.8	389 1.0 0.0 0.0	50.4 76.9 64.5
775	RO0Y_087_050d	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	61.0 38.4 32.2	50.2 40.0 0.0	0.875 0.375 0.375	54.0 47.8 24.1	53.6 26.8 14.2	389 1.0 0.0 0.0	50.4 76.9 64.5
776	RO0Y_075_037d	0.75 0.375 0.375	0.75 0.375 0.562	390	0.75 0.375 0.375	54.7 28.8 24.2	37.6 40.0 0.0	0.75 0.375 0.375	49.4 36.7 17.1	40.5 25.0 11.8	389 1.0 0.0 0.0	50.4 76.9 64.5
777	RO0Y_062_025d	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	48.4 19.2 16.1	25.1 40.0 0.0	0.625 0.375 0.375	45.1 24.9 10.6	27.0 23.1 8.5	389 1.0 0.0 0.0	50.4 76.9 64.5
778	RO0Y_050_012d	0.5 0.375 0.375	0.5 0.125 0.437	390	0.5 0.375 0.375	42.0 9.6 8.0	12.5 40.0 0.0	0.5 0.375 0.375	41.4 12.4 4.8	13.3 21.2 4.3	389 1.0 0.0 0.0	50.4 76.9 64.5
779	NW_037d	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	35.7 0.0 0.0	0.0 0.0 0.0	0.375 0.375 0.375	38.3 0.0 0.0	0.0 325.3 2.5	360 1.0 1.0 1.0	95.4 0.0 0.0
780	G50B_037_012d	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	34.7 -5.7 -1.6	6.0 196.3 0.0	0.25 0.375 0.375	36.0 -11.0 -3.5	11.6 197.8 5.8	210 0.0 1.0 1.0	86.8 -46.1 -13.5
781	G50B_037_025d	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	33.6 -11.5 -3.3	12.0 196.3 0.0	0.125 0.375 0.375	34.7 -18.9 -5.7	19.8 196.8 7.8	210 0.0 1.0 1.0	86.8 -46.1 -13.5

vedi file simili: <http://farbe.li.tu-berlin.de/PI31/PI31.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsi_d	rgb*Md	LabCh*Md
810	NW_100d	1.0 1.0 1.0	1.0 0.0 0.0	1.0 1.0 1.0	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
811	B00R_100_012d	0.875 0.875 1.0	1.0 0.125 0.937	270	0.875 0.875 1.0	87.2 9.5 -12.9	16.0 306.2	0.875 0.875 1.0	85.5 5.8 -14.8	15.9 291.5 4.4	270	0.0 0.0 1.0
812	B00R_100_025d	0.75 0.75 1.0	1.0 0.25 0.875	270	0.75 0.75 1.0	79.1 19.0 -25.8	32.1 306.2	0.75 0.75 1.0	75.6 12.8 -30.0	32.7 293.1 8.2	270	0.0 0.0 1.0
813	B00R_100_037d	0.625 0.625 1.0	1.0 0.375 0.812	270	0.625 0.625 1.0	71.0 28.5 -38.8	48.1 306.2	0.625 0.625 1.0	65.7 21.4 -45.6	50.4 295.1 11.1	270	0.0 0.0 1.0
814	B00R_100_050d	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	62.8 38.0 -51.7	64.2 306.2	0.5 0.5 1.0	56.0 31.9 -61.1	69.0 297.5 13.0	270	0.0 0.0 1.0
815	B00R_100_062d	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	54.7 47.5 -64.7	80.3 306.2	0.375 0.375 1.0	46.8 44.5 -76.1	88.2 300.3 14.2	270	0.0 0.0 1.0
816	B00R_100_075d	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	46.6 57.0 -77.6	96.3 306.2	0.25 0.25 1.0	38.8 58.2 -89.4	106.7 303.0 14.1	270	0.0 0.0 1.0
817	B00R_100_087d	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	38.5 66.5 -90.6	112.4 306.2	0.125 0.125 1.0	33.0 69.9 -99.0	121.3 305.2 10.6	270	0.0 0.0 1.0
818	B00R_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2 0.0	270	0.0 0.0 1.0
819	Y00G_100_012d	1.0 1.0 0.875	1.0 0.125 0.937	90	1.0 1.0 0.875	95.0 -2.5 11.3	11.6 102.8	1.0 1.0 0.875	94.7 -5.0 14.6	15.9 108.9 4.1	89	1.0 1.0 0.0
820	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	83.4 0.0 0.0	0.0 0.0	0.875 0.875 0.875	84.7 0.0 0.0	0.0 325.2 1.2	360	1.0 1.0 1.0
821	B00R_087_012d	0.75 0.75 0.875	0.875 0.125 0.812	270	0.75 0.75 0.875	75.3 9.5 -12.9	16.0 306.2	0.75 0.75 0.875	74.6 6.0 -15.2	16.4 291.7 4.1	270	0.0 0.0 1.0
822	B00R_087_025d	0.625 0.625 0.875	0.875 0.25 0.75	270	0.625 0.625 0.875	67.2 19.0 -25.8	32.1 306.2	0.625 0.625 0.875	64.4 13.5 -30.9	33.8 293.6 7.9	270	0.0 0.0 1.0
823	B00R_087_037d	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.5 0.875	59.1 28.5 -38.8	48.1 306.2	0.5 0.5 0.875	54.3 23.0 -46.9	52.2 296.1 10.8	270	0.0 0.0 1.0
824	B00R_087_050d	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	50.9 38.0 -51.7	64.2 306.2	0.375 0.375 0.875	44.6 34.8 -62.7	71.7 299.0 13.0	270	0.0 0.0 1.0
825	B00R_087_062d	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.8 47.5 -64.7	80.3 306.2	0.25 0.25 0.875	35.8 48.6 -77.1	91.2 302.1 14.3	270	0.0 0.0 1.0
826	B00R_087_075d	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	34.7 57.0 -77.6	96.3 306.2	0.125 0.125 0.875	29.1 61.5 -88.2	107.5 304.8 12.7	270	0.0 0.0 1.0
827	B00R_087_087d	0.0 0.0 0.875	0.875 0.875 0.437	270	0.0 0.0 0.875	26.5 66.5 -90.6	112.4 306.2	0.0 0.0 0.875	25.9 68.7 -93.6	116.1 306.2 3.7	270	0.0 0.0 1.0
828	Y00G_100_025d	1.0 1.0 0.75	1.0 0.25 0.875	90	1.0 1.0 0.75	94.7 -5.1 22.6	23.2 102.8	1.0 1.0 0.75	94.1 -9.3 29.3	30.8 107.7 7.9	89	1.0 1.0 0.0
829	Y00G_087_012d	0.875 0.875 0.75	0.875 0.125 0.812	90	0.875 0.875 0.75	83.1 -2.5 11.3	11.6 102.8	0.875 0.875 0.75	84.0 -5.1 15.0	15.8 108.7 4.5	89	1.0 1.0 0.0
830	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	71.5 0.0 0.0	0.0 0.0	0.75 0.75 0.75	73.7 0.0 0.0	0.0 325.2 2.1	360	1.0 1.0 1.0
831	B00R_075_012d	0.625 0.625 0.75	0.75 0.125 0.687	270	0.625 0.625 0.75	63.4 9.5 -12.9	16.0 306.2	0.625 0.625 0.75	63.3 6.3 -15.7	16.9 292.0 4.1	270	0.0 0.0 1.0
832	B00R_075_025d	0.5 0.5 0.75	0.75 0.25 0.625	270	0.5 0.5 0.75	55.3 19.0 -25.8	32.1 306.2	0.5 0.5 0.75	52.8 14.4 -31.9	35.1 294.3 7.9	270	0.0 0.0 1.0
833	B00R_075_037d	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	47.1 28.5 -38.8	48.1 306.2	0.375 0.375 0.75	42.5 25.1 -48.4	54.5 297.4 11.1	270	0.0 0.0 1.0
834	B00R_075_050d	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	39.0 38.0 -51.7	64.2 306.2	0.25 0.25 0.75	32.9 38.5 -64.1	74.8 301.0 13.7	270	0.0 0.0 1.0
835	B00R_075_062d	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	30.9 47.5 -64.7	80.3 306.2	0.125 0.125 0.75	25.3 52.5 -76.8	93.0 304.3 14.2	270	0.0 0.0 1.0
836	B00R_075_075d	0.0 0.0 0.75	0.75 0.75 0.375	270	0.0 0.0 0.75	22.7 57.0 -77.6	96.3 306.2	0.0 0.0 0.75	21.3 61.2 -83.4	103.5 306.2 7.2	270	0.0 0.0 1.0
837	Y00G_100_037d	1.0 1.0 0.625	1.0 0.375 0.812	90	1.0 1.0 0.625	94.3 -7.7 34.0	34.9 102.8	1.0 1.0 0.625	93.6 -13.0 43.8	45.7 106.5 11.1	89	1.0 1.0 0.0
838	Y00G_087_025d	0.875 0.875 0.625	0.875 0.25 0.75	90	0.875 0.875 0.625	82.7 -5.1 22.6	23.2 102.8	0.875 0.875 0.625	83.4 -9.4 30.0	31.5 107.3 8.5	89	1.0 1.0 0.0
839	Y00G_075_012d	0.75 0.75 0.625	0.75 0.125 0.687	90	0.75 0.75 0.625	71.2 -2.5 11.3	11.6 102.8	0.75 0.75 0.625	73.0 -5.1 15.4	16.3 108.5 5.2	89	1.0 1.0 0.0
840	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	59.6 0.0 0.0	0.0 0.0	0.625 0.625 0.625	62.4 0.0 0.0	0.0 325.2 2.7	360	1.0 1.0 1.0
841	B00R_062_012d	0.5 0.5 0.625	0.625 0.125 0.562	270	0.5 0.5 0.625	51.5 9.5 -12.9	16.0 306.2	0.5 0.5 0.625	51.6 6.7 -16.3	17.6 292.4 4.3	270	0.0 0.0 1.0
842	B00R_062_025d	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	43.3 19.0 -25.8	32.1 306.2	0.375 0.375 0.625	40.8 15.7 -33.2	36.8 295.4 8.4	270	0.0 0.0 1.0
843	B00R_062_037d	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	35.2 28.5 -38.8	48.1 306.2	0.25 0.25 0.625	30.4 28.1 -50.0	57.4 299.3 12.2	270	0.0 0.0 1.0
844	B00R_062_050d	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	27.1 38.0 -51.7	64.2 306.2	0.125 0.125 0.625	21.6 42.8 -64.6	77.5 305.3 14.7	270	0.0 0.0 1.0
845	B00R_062_062d	0.0 0.0 0.625	0.625 0.625 0.312	270	0.0 0.0 0.625	18.9 47.5 -64.7	80.3 306.2	0.0 0.0 0.625	16.6 53.5 -72.9	90.4 306.2 10.3	270	0.0 0.0 1.0
846	Y00G_100_050d	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	94.0 -10.3 45.3	46.5 102.8	1.0 1.0 0.5	93.2 -15.9 57.8	59.9 103.3 13.6	89	1.0 1.0 0.0
847	Y00G_087_037d	0.875 0.875 0.5	0.875 0.375 0.687	90	0.875 0.875 0.5	82.4 -7.7 34.0	34.9 102.8	0.875 0.875 0.5	82.9 -12.9 44.8	46.6 106.0 11.9	89	1.0 1.0 0.0
848	Y00G_075_025d	0.75 0.75 0.5	0.75 0.25 0.625	90	0.75 0.75 0.5	70.8 -5.1 22.6	23.2 102.8	0.75 0.75 0.5	72.4 -9.4 30.9	32.3 106.9 9.4	89	1.0 1.0 0.0
849	Y00G_062_012d	0.625 0.625 0.5	0.625 0.125 0.562	90	0.625 0.625 0.5	59.2 -2.5 11.3	11.6 102.8	0.625 0.625 0.5	61.6 -5.2 16.0	16.8 108.2 5.8	89	1.0 1.0 0.0
850	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	47.7 0.0 0.0	0.0 0.0	0.5 0.5 0.5	50.6 0.0 0.0	0.0 325.3 2.9	360	1.0 1.0 1.0
851	B00R_050_012d	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.375 0.5	39.5 9.5 -12.9	16.0 306.2	0.375 0.375 0.5	39.4 7.2 -17.0	18.5 292.9 4.7	270	0.0 0.0 1.0
852	B00R_050_025d	0.25 0.25 0.5	0.5 0.25 0.375	270	0.25 0.25 0.5	31.4 19.0 -25.8	32.1 306.2	0.25 0.25 0.5	28.2 17.7 -34.7	30.9 297.0 9.5	270	0.0 0.0 1.0
853	B00R_050_037d	0.125 0.125 0.5	0.5 0.375 0.312	270	0.125 0.125 0.5	23.3 28.5 -38.8	48.1 306.2	0.125 0.125 0.5	18.1 32.4 -51.3	60.6 302.2 14.0	270	0.0 0.0 1.0
854	B00R_050_050d	0.0 0.0 0.5	0.5 0.5 0.25	270	0.0 0.0 0.5	15.1 38.0 -51.7	64.2 306.2	0.0 0.0 0.5	11.7 45.5 -61.9	76.8 306.2 13.0	270	0.0 0.0 1.0
855	Y00G_100_062d	1.0 1.0 0.375	1.0 0.625 0.687	90	1.0 1.0 0.375	93.6 -12.9 56.7	58.1 102.8	1.0 1.0 0.375	92.9 -18.0 70.4	72.7 104.3 14.7	89	1.0 1.0 0.0
856	Y00G_087_050d	0.875 0.875 0.375	0.875 0.5 0.625	90	0.875 0.875 0.375	82.1 -10.3 45.3	46.5 102.8	0.875 0.875 0.375	82.6 -15.5 58.6	60.6 104.8 14.2	89	1.0 1.0 0.0
857	Y00G_075_037d	0.75 0.75 0.375	0.75 0.375 0.562	90	0.75 0.75 0.375	70.5 -7.7 34.0	34.9 102.8	0.75 0.75 0.375	72.0 -12.6 45.8	47.5 105.4 12.8	89	1.0 1.0 0.0
858	Y00G_062_025d	0.625 0.625 0.375	0.625 0.25 0.5	90	0.625 0.625 0.375	58.9 -5.1 22.6	23.2 102.8	0.625 0.625 0.375	61.1 -9.3 31.9	33.2 106.3 10.3	89	1.0 1.0 0.0
859	Y00G_050_012d	0.5 0.5 0.375	0.5 0.125 0.437	90	0.5 0.5 0.375	47.3 -2.5 11.3	11.6 102.8	0.5 0.5 0.375	49.8 -5.3 16.6	17.5 107.8 6.5	89	1.0 1.0 0.0
860	NW_037d	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	35.7 0.0 0.0	0.0 0.0	0.375 0.375 0.375	38.3 0.0 0.0	0.0 325.3 2.5	360	1.0 1.0 1.0
861	B00R_037_012d	0.25 0.25 0.375	0.375 0.125 0.312	270	0.25 0.25 0.375	27.6 9.5 -12.9	16.0 306.2	0.25 0.25 0.375	26.5 8.0 -18.0	19.8 294.0 5.4	270	0.0 0.0 1.0
862	B00R_037_025d	0.125 0.125 0.375	0.375 0.25 0.25	270	0.125 0.125 0.375	19.5 19.0 -25.8	32.1 306.2	0.125 0.125 0.375	15.0 21.1 -36.5	42.1 300.0 11.6	270	0.0 0.0 1.0
863	B00R_037_037d	0.0 0.0 0.375	0.375 0.375 0.187	270	0.0 0.0 0.375	11.3 28.5 -38.8	48.1 306.2	0.0 0.0 0.375	6.7 36.7 -50.3	62.3 306.1 14.9	270	0.0 0.0 1.0
864	Y00G_100_075d	1.0 1.0 0.25	1.0 0.75 0.625	90	1.0 1.0 0.25	93.3 -15.5 68.0	69.8 102.8	1.0 1.0 0.25	92.8 -19.5 80.8	83.1 103.5 13.4	89	1.0 1.0 0.0
865	Y00G_087_062d	0.875 0.875 0.25	0.875 0.625 0.562	90	0.875 0.875 0.25	81.7 -12.9 56.7	58.1 102.8	0.875 0.875 0.25	82.4 -17.2 70.3	72.4 103.8 14.2	89	1.0 1.0 0.0
866	Y00G_075_050d	0.75 0.75 0.25	0.75 0.5 0.5	90	0.75 0.75 0.25	70.1 -10.3 45.3	46.5 102.8	0.75 0.75 0.25	71.7 -14.8 58.9	60.8 104.1 14.4	89	1.0 1.0 0.0
867	Y00G_062_037d	0.625 0.625 0.25	0.6									

vedi file simili: <http://farbe.li.tu-berlin.de/PI31/PI31.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md
891	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	1.0 1.0 1.0	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
892	B50R_100_012d	1.0 0.875 1.0	1.0 0.875 1.0	1.0 0.125 0.937	330	1.0 0.875 1.0	90.6 11.7 -7.3	13.8 328.2	1.0 0.875 1.0	87.9 15.7 -10.9	19.1 325.1 5.9	330 1.0 0.875 1.0
893	B50R_100_025d	1.0 0.75 1.0	1.0 0.75 1.0	1.0 0.25 0.875	330	1.0 0.75 1.0	85.8 23.5 -14.6	27.7 328.2	1.0 0.75 1.0	80.9 31.7 -21.5	38.4 325.8 11.8	330 1.0 0.75 1.0
894	B50R_100_037d	1.0 0.625 1.0	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	81.1 35.3 -21.9	41.6 328.2	1.0 0.625 1.0	74.3 47.6 -31.5	57.1 326.4 17.0	330 1.0 0.625 1.0
895	B50R_100_050d	1.0 0.5 1.0	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	76.3 47.1 -29.2	55.4 328.2	1.0 0.5 1.0	68.6 62.6 -40.5	74.6 327.0 20.6	330 1.0 0.5 1.0
896	B50R_100_062d	1.0 0.375 1.0	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	71.5 58.9 -36.5	69.3 328.2	1.0 0.375 1.0	63.8 75.6 -48.0	89.6 327.5 21.6	330 1.0 0.375 1.0
897	B50R_100_075d	1.0 0.25 1.0	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	66.8 70.7 -43.8	83.2 328.2	1.0 0.25 1.0	60.2 85.6 -53.6	101.0 327.9 18.9	330 1.0 0.25 1.0
898	B50R_100_087d	1.0 0.125 1.0	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	62.0 82.5 -51.1	97.1 328.2	1.0 0.125 1.0	58.1 91.8 -57.0	108.0 328.1 11.6	330 1.0 0.125 1.0
899	B50R_100_100d	1.0 0.0 1.0	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	1.0 0.0 1.0	57.2 94.3 -58.4	111.0 328.2 0.0	330 1.0 0.0 1.0
900	GO0B_100_012d	0.875 1.0 0.875	1.0 1.0 0.875	1.0 1.25 0.937	150	0.875 1.0 0.875	93.9 -10.3 9.9	14.3 136.0	0.875 1.0 0.875	92.5 -15.4 11.3	19.1 143.6 5.4	149 0.0 1.0 0.875 1.0
901	NW_087d	0.875 0.875 0.875	0.875 0.875 0.875	0.875 0.875 0.875	360	0.875 0.875 0.875	83.4 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	84.7 0.0 0.0	0.0 325.2 1.2	360 1.0 1.0 0.875 1.0
902	B50R_087_012d	0.875 0.75 0.875	0.875 0.75 0.875	0.875 0.125 0.812	330	0.875 0.75 0.875	78.7 11.7 -7.3	13.8 328.2	0.875 0.75 0.875	77.1 16.1 -11.2	19.6 325.2 6.0	330 1.0 0.75 0.875 1.0
903	B50R_087_025d	0.875 0.625 0.875	0.875 0.625 0.875	0.875 0.25 0.75	330	0.875 0.625 0.875	73.9 23.5 -14.6	27.7 328.2	0.875 0.625 0.875	69.9 32.6 -22.0	39.3 325.9 12.3	330 1.0 0.625 0.875 1.0
904	B50R_087_037d	0.875 0.5 0.875	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	69.1 35.3 -21.9	41.6 328.2	0.875 0.5 0.875	63.5 48.6 -31.9	58.2 326.7 17.6	330 1.0 0.5 0.875 1.0
905	B50R_087_050d	0.875 0.375 0.875	0.875 0.375 0.875	0.875 0.625 0.330	330	0.875 0.375 0.875	64.4 47.1 -29.2	55.4 328.2	0.875 0.375 0.875	58.0 63.2 -40.5	75.0 327.3 20.6	330 1.0 0.375 0.875 1.0
906	B50R_087_062d	0.875 0.25 0.875	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	59.6 58.9 -36.5	69.3 328.2	0.875 0.25 0.875	53.8 74.7 -47.0	88.3 327.8 19.8	330 1.0 0.25 0.875 1.0
907	B50R_087_075d	0.875 0.125 0.875	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	54.9 70.7 -43.8	83.2 328.2	0.875 0.125 0.875	51.3 82.1 -51.1	96.7 328.1 14.0	330 1.0 0.125 0.875 1.0
908	B50R_087_087d	0.875 0.0 0.875	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	50.1 82.5 -51.1	97.1 328.2	0.875 0.0 0.875	50.2 85.3 -52.8	100.3 328.2 3.2	330 1.0 0.0 0.875 1.0
909	GO0B_100_025d	0.75 1.0 0.75	1.0 0.75 1.0	1.0 0.25 0.875	150	0.75 1.0 0.75	92.4 -20.6 19.9	28.7 136.0	0.75 1.0 0.75	90.1 -20.5 23.2	38.3 142.7 10.6	149 0.0 1.0 0.75 1.0
910	GO0B_087_012d	0.75 0.875 0.75	0.875 0.875 0.75	0.875 0.125 0.812	150	0.75 0.875 0.75	82.0 -10.3 9.9	14.3 136.0	0.75 0.875 0.75	81.8 -15.7 11.6	19.6 143.5 5.6	149 0.0 1.0 0.875 1.0
911	NW_075d	0.75 0.75 0.75	0.75 0.75 0.75	0.75 0.75 0.75	360	0.75 0.75 0.75	71.5 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	73.7 0.0 0.0	0.0 325.2 2.1	360 1.0 1.0 0.75 1.0
912	B50R_075_012d	0.75 0.625 0.75	0.75 0.625 0.75	0.75 0.125 0.687	330	0.75 0.625 0.75	66.7 11.7 -7.3	13.8 328.2	0.75 0.625 0.75	65.9 16.6 -11.5	20.2 325.3 6.4	330 1.0 0.625 0.75 1.0
913	B50R_075_025d	0.75 0.5 0.75	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	62.0 23.5 -14.6	27.7 328.2	0.75 0.5 0.75	58.7 33.5 -22.4	40.4 326.2 13.1	330 1.0 0.5 0.75 1.0
914	B50R_075_037d	0.75 0.375 0.75	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	57.2 35.3 -21.9	41.6 328.2	0.75 0.375 0.75	52.4 49.6 -32.2	59.1 327.0 18.2	330 1.0 0.375 0.75 1.0
915	B50R_075_050d	0.75 0.25 0.75	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	52.5 47.1 -29.2	55.4 328.2	0.75 0.25 0.75	47.5 63.1 -39.9	74.6 327.6 19.8	330 1.0 0.25 0.75 1.0
916	B50R_075_062d	0.75 0.125 0.75	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	47.7 58.9 -36.5	69.3 328.2	0.75 0.125 0.75	44.3 72.1 -44.9	84.9 328.0 15.9	330 1.0 0.125 0.75 1.0
917	B50R_075_075d	0.75 0.0 0.75	0.75 0.0 0.75	0.75 0.375 0.330	330	0.75 0.0 0.75	42.9 70.7 -43.8	83.2 328.2	0.75 0.0 0.75	43.0 76.0 -47.0	89.4 328.2 6.1	330 1.0 0.0 0.75 1.0
918	GO0B_100_037d	0.625 1.0 0.625	1.0 0.625 1.0	1.0 0.375 0.812	150	0.625 1.0 0.625	90.9 -31.0 29.9	43.1 136.0	0.625 1.0 0.625	88.0 -44.8 35.5	57.2 141.6 15.1	149 0.0 1.0 0.625 1.0
919	GO0B_087_025d	0.625 0.875 0.625	0.875 0.25 0.75	1.0 0.625 0.687	150	0.625 0.875 0.625	80.5 -20.6 19.9	28.7 136.0	0.625 0.875 0.625	79.3 -31.1 23.9	39.3 142.4 11.2	149 0.0 1.0 0.875 1.0
920	GO0B_075_012d	0.625 0.75 0.625	0.75 0.125 0.687	1.0 0.625 0.687	150	0.625 0.75 0.625	70.0 -10.3 9.9	14.3 136.0	0.625 0.75 0.625	70.8 -16.2 12.0	20.2 143.4 6.2	149 0.0 1.0 0.75 1.0
921	NW_062d	0.625 0.625 0.625	0.625 0.625 0.625	0.625 0.625 0.625	360	0.625 0.625 0.625	59.6 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	62.4 0.0 0.0	0.0 325.2 2.7	360 1.0 1.0 0.625 1.0
922	B50R_062_012d	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	54.8 11.7 -7.3	13.8 328.2	0.625 0.5 0.625	54.4 17.2 -11.8	20.9 325.5 7.1	330 1.0 0.5 0.625 1.0	
923	B50R_062_025d	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	50.1 23.5 -14.6	27.7 328.2	0.625 0.375 0.625	47.1 34.6 -22.9	41.5 326.5 14.1	330 1.0 0.375 0.625 1.0	
924	B50R_062_037d	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	45.3 35.3 -21.9	41.6 328.2	0.625 0.25 0.625	41.2 50.2 -32.1	59.6 327.4 18.5	330 1.0 0.25 0.625 1.0	
925	B50R_062_050d	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	40.5 47.1 -29.2	55.4 328.2	0.625 0.125 0.625	37.3 61.3 -38.3	72.3 327.9 17.2	330 1.0 0.125 0.625 1.0	
926	B50R_062_062d	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	35.8 58.9 -36.5	69.3 328.2	0.625 0.0 0.625	35.5 66.4 -41.1	78.1 328.2 8.7	330 1.0 0.0 0.625 1.0	
927	GO0B_100_050d	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	89.5 -51.3 39.9	57.5 136.0	0.5 1.0 0.5	86.3 -57.6 47.9	75.0 140.2 18.4	149 0.0 1.0 0.5 1.0	
928	GO0B_087_037d	0.5 0.875 0.5	0.875 0.375 0.687	150	0.5 0.875 0.5	79.0 -31.0 29.9	43.1 136.0	0.5 0.875 0.5	77.4 -45.4 36.6	58.3 141.0 15.9	149 0.0 1.0 0.875 1.0	
929	GO0B_075_025d	0.5 0.75 0.5	0.75 0.25 0.625	150	0.5 0.75 0.5	68.6 -20.6 19.9	28.7 136.0	0.5 0.75 0.5	68.3 -31.8 24.8	40.4 142.0 12.1	149 0.0 1.0 0.75 1.0	
930	GO0B_062_012d	0.5 0.625 0.5	0.625 0.125 0.562	150	0.5 0.625 0.5	58.1 -10.3 9.9	14.3 136.0	0.5 0.625 0.5	59.4 -16.7 12.5	20.9 143.1 7.0	149 0.0 1.0 0.625 1.0	
931	NW_050d	0.5 0.5 0.5	0.5 0.5 0.5	360	0.5 0.5 0.5	47.7 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	50.6 0.0 0.0	0.0 325.3 2.9	360 1.0 1.0 0.5 1.0	
932	B50R_050_012d	0.5 0.375 0.5	0.5 0.125 0.437	330	0.5 0.375 0.5	42.9 11.7 -7.3	13.8 328.2	0.5 0.375 0.5	42.3 18.0 -12.2	21.8 325.7 8.0	330 1.0 0.375 0.5 1.0	
933	B50R_050_025d	0.5 0.25 0.5	0.5 0.25 0.375	330	0.5 0.25 0.5	38.1 23.5 -14.6	27.7 328.2	0.5 0.25 0.5	35.2 35.7 -23.2	42.6 326.9 15.2	330 1.0 0.25 0.5 1.0	
934	B50R_050_037d	0.5 0.125 0.5	0.5 0.375 0.312	330	0.5 0.125 0.5	33.4 35.3 -21.9	41.6 328.2	0.5 0.125 0.5	30.1 49.6 -31.2	58.6 327.8 17.3	330 1.0 0.125 0.5 1.0	
935	B50R_050_050d	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	28.6 47.1 -29.2	55.4 328.2	0.5 0.0 0.5	27.8 56.4 -34.9	66.3 328.2 10.9	330 1.0 0.0 0.5 1.0	
936	GO0B_100_062d	0.375 1.0 0.375	1.0 0.625 0.687	150	0.375 1.0 0.375	88.0 -51.7 49.9	71.9 136.0	0.375 1.0 0.375	85.1 -68.3 59.7	90.7 138.8 19.4	149 0.0 1.0 0.375 1.0	
937	GO0B_087_050d	0.375 0.875 0.375	0.875 0.5 0.625	150	0.375 0.875 0.375	77.5 -41.3 39.9	57.5 136.0	0.375 0.875 0.375	75.9 -57.6 49.1	75.7 139.5 18.7	149 0.0 1.0 0.875 1.0	
938	GO0B_075_037d	0.375 0.75 0.375	0.75 0.375 0.562	150	0.375 0.75 0.375	67.1 -31.0 29.9	43.1 136.0	0.375 0.75 0.375	66.5 -45.8 37.9	59.4 140.3 16.7	149 0.0 1.0 0.75 1.0	
939	GO0B_062_025d	0.375 0.625 0.375	0.625 0.25 0.5	150	0.375 0.625 0.375	56.6 -20.6 19.9	28.7 136.0	0.375 0.625 0.375	57.0 -32.5 25.9	41.6 141.4 13.2	149 0.0 1.0 0.625 1.0	
940	GO0B_050_012d	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	46.2 -10.3 9.9	14.3 136.0	0.375 0.5 0.375	47.6 -17.3 13.1	21.8 142.8 7.8	149 0.0 1.0 0.5 1.0	
941	NW_037d	0.375 0.375 0.375	0.375 0.375 0.375	360	0.375 0.375 0.375	35.7 0.0 0.0	0.0 0.0 0.0	0.375 0.375 0.375	38.3 0.0 0.0	0.0 325.3 2.5	360 1.0 1.0 0.375 1.0	
942	B50R_037_012d	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.25 0.375	31.0 11.7 -7.3	13.8 328.2	0.375 0.25 0.375	29.7 19.0 -12.7	22.9 326.1 9.1	330 1.0 0.25 0.375 1.0	
943	B50R_037_025d	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.125 0.375	26.2 23.5 -14.6	27.7 328.2	0.375 0.125 0.375	23.1 36.3 -23.1	43.0 327.5 15.6	330 1.0 0.125 0.375 1.0	
944	B50R_037_037d	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	21.4 35.3 -21.9	41.6 328.2	0.375 0.0 0.375	19.7 46.0 -28.5	54		



TUB materiale: code=rha4ta.

$$\Delta E^* = 1.6$$

Input: $rgb/cmyk \rightarrow rgb_d$
Output: trasferire a rgb_d



vedi file simili: <http://farbe.li.tu-berlin.de/PI31/PI31.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd			ict_Fd			hsi_Fd			rgb**Fd			LabCh**Fd			rgb*Fd			LabCh*Fd			DE**Fd			hsiM,d	rgb**Md			LabCh**Md					
1053	NW_086d	0.866	0.866	0.866	0.866	0.0	0.866	360	0.866	0.866	0.866	82.6	0.0	0.0	0.0	0.0	0.866	0.866	0.866	83.9	0.0	0.0	0.0	0.0	325.2	1.3	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1054	NW_093d	0.933	0.933	0.933	0.933	0.0	0.933	360	0.933	0.933	0.933	89.0	0.0	0.0	0.0	0.0	0.933	0.933	0.933	89.7	0.0	0.0	0.0	0.0	325.2	0.6	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1055	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0	325.2	0.0	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1056	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1057	NW_006d	0.066	0.066	0.066	0.066	0.0	0.066	360	0.066	0.066	0.066	6.2	0.0	0.0	0.0	0.0	0.066	0.066	0.066	4.4	0.0	0.0	0.0	0.0	326.3	1.8	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1058	NW_013d	0.133	0.133	0.133	0.133	0.0	0.133	360	0.133	0.133	0.133	12.6	0.0	0.0	0.0	0.0	0.133	0.133	0.133	12.0	0.0	0.0	0.0	0.0	325.6	0.6	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1059	NW_020d	0.2	0.2	0.2	0.2	0.0	0.2	360	0.2	0.2	0.2	19.0	0.0	0.0	0.0	0.0	0.2	0.2	0.2	19.7	0.0	0.0	0.0	0.0	325.5	0.6	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1060	NW_026d	0.266	0.266	0.266	0.266	0.0	0.266	360	0.266	0.266	0.266	25.3	0.0	0.0	0.0	0.0	0.266	0.266	0.266	27.0	0.0	0.0	0.0	0.0	325.4	1.6	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1061	NW_033d	0.333	0.333	0.333	0.333	0.0	0.333	360	0.333	0.333	0.333	31.7	0.0	0.0	0.0	0.0	0.333	0.333	0.333	34.0	0.0	0.0	0.0	0.0	325.3	2.2	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1062	NW_040d	0.4	0.4	0.4	0.4	0.0	0.4	360	0.4	0.4	0.4	38.1	0.0	0.0	0.0	0.0	0.4	0.4	0.4	40.8	0.0	0.0	0.0	0.0	325.3	2.6	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1063	NW_046d	0.466	0.466	0.466	0.466	0.0	0.466	360	0.466	0.466	0.466	44.4	0.0	0.0	0.0	0.0	0.466	0.466	0.466	47.3	0.0	0.0	0.0	0.0	325.4	2.8	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1064	NW_053d	0.533	0.533	0.533	0.533	0.0	0.533	360	0.533	0.533	0.533	50.8	0.0	0.0	0.0	0.0	0.533	0.533	0.533	53.7	0.0	0.0	0.0	0.0	325.3	2.9	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1065	NW_060d	0.6	0.6	0.6	0.6	0.0	0.6	360	0.6	0.6	0.6	57.2	0.0	0.0	0.0	0.0	0.6	0.6	0.6	60.0	0.0	0.0	0.0	0.0	325.3	2.8	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1066	NW_066d	0.666	0.666	0.666	0.666	0.0	0.666	360	0.666	0.666	0.666	63.5	0.0	0.0	0.0	0.0	0.666	0.666	0.666	66.5	0.0	0.0	0.0	0.0	325.2	2.6	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1067	NW_073d	0.734	0.734	0.734	0.734	0.0	0.734	360	0.734	0.734	0.734	70.0	0.0	0.0	0.0	0.0	0.734	0.734	0.734	72.3	0.0	0.0	0.0	0.0	325.2	2.2	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1068	NW_080d	0.8	0.8	0.8	0.8	0.0	0.8	360	0.8	0.8	0.8	76.3	0.0	0.0	0.0	0.0	0.8	0.8	0.8	78.1	0.0	0.0	0.0	0.0	325.2	1.8	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1069	NW_086d	0.866	0.866	0.866	0.866	0.0	0.866	360	0.866	0.866	0.866	82.6	0.0	0.0	0.0	0.0	0.866	0.866	0.866	83.9	0.0	0.0	0.0	0.0	325.2	1.3	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1070	NW_093d	0.933	0.933	0.933	0.933	0.0	0.933	360	0.933	0.933	0.933	89.0	0.0	0.0	0.0	0.0	0.933	0.933	0.933	89.7	0.0	0.0	0.0	0.0	325.2	0.6	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1071	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0	325.2	0.0	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1072	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1073	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0	325.2	0.0	360	1.0	1.0	1.0	95.4	0.0	0.0	0.0	0.0
1074	R00Y_100_100d	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0	1.0	0.0	0.0	50.4	76.9	64.5	100.4	39.9	0.0	389	1.0	0.0	0.0	50.4	76.9	64.5	100.4	40.0	
1075	G50B_100_100d	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	1.0	86.8	-46.1	-13.5	48.1	196.3	0.0	1.0	1.0	86.8	-46.1	-13.5	48.1	196.3	0.0	210	0.0	0.0	1.0	86.8	-46.1	-13.5	48.1	196.3	
1076	Y00G_100_100d	1.0	1.0	0.0	1.0	1.0	0.5	90	1.0	1.0	0.0	92.6	-20.7	90.7	93.0	102.8	1.0	1.0	0.0	92.6	-20.6	90.7	93.0	102.8	0.0	89	1.0	1.0	0.0	92.6	-20.7	90.7	93.0	102.8	
1077	B00R_100_100d	0.0	0.0	1.0	1.0	1.0	0.5	270	0.0	0.0	1.0	30.3	76.0	-103.5	128.5	306.2	0.0	0.0	1.0	30.3	76.0	-103.5	128.5	306.2	0.0	270	0.0	0.0	1.0	30.3	76.0	-103.5	128.5	306.2	
1078	G00B_100_100d	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	83.6	-82.7	79.8	115.0	136.0	0.0	1.0	0.0	83.6	-82.7	79.8	115.0	136.0	0.0	149	0.0	1.0	0.0	83.6	-82.7	79.8	115.0	136.0	
1079	B50R_100_100d	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	1.0	0.0	1.0	57.2	94.3	-58.4	111.0	328.2	0.0	330	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328.2	

delta E* = 1.0

Grafico TUB-PI31; riproduzione del colore
colori e la differenza, ΔE^* , 3D=0, de=0, sRGB

Input: $rgb/cmyk \rightarrow rgb_d$
Output: trasferire a rgb_d

iscrizione TUB: 20160501-PI31/PI31L0NP.PDF /.PS
Applicazione per la misura dell'output su display, nessuna separazione
TUB materiale: code=rha4ta