

Immettere y uscita: Offset Reflective System ORS18a for relative CIELAB hue $h_{ab,a,rel} = h_{ab}/360 = 102/360 = 0.28$

$H^*_ = Y25G_$

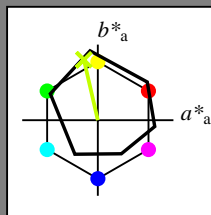
Dati del dispositivo (d) o colori elementari (e):

$HIC^*_$

codice di tonalità per i colori questa pagina:

$H^*_ = Y25G_$

triangolo chiarezza T^*



ORS18a; dati atti CIELAB (a)

name	$L^*=L^*_a a^*_a$	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
R_.,Ma	47.9	65.3	50.5	82.6
Y_.,Ma	90.3	-10.2	91.7	92.3
G_.,Ma	50.9	-62.8	34.9	71.9
C_.,Ma	58.6	-30.3	-45.0	54.2
B_.,Ma	25.7	31.0	-44.4	54.2
M_.,Ma	48.1	75.2	-8.3	75.7
N_.,Ma	18.0	0.0	0.0	0.0
W_.,Ma	95.4	0.0	0.0	0.0
R_.,CIE	39.9	58.7	27.9	65.0
Y_.,CIE	81.2	-2.8	71.5	71.6
G_.,CIE	52.2	-42.4	13.6	44.5
B_.,CIE	30.5	1.4	-46.4	46.4

Il dati per il massimo colore (Ma):

$LabCh^*_{-,Ma}$: 83 -18 79 81 102

$HIC^*_{-,Ma}$: Y25G_100_100_

$rgbic^*_{-,Ma}$:

0.76 1.0 0.0 1.0 1.0

triangolo chiarezza T^*

%Gamma

$u^*_{rel} = 92$

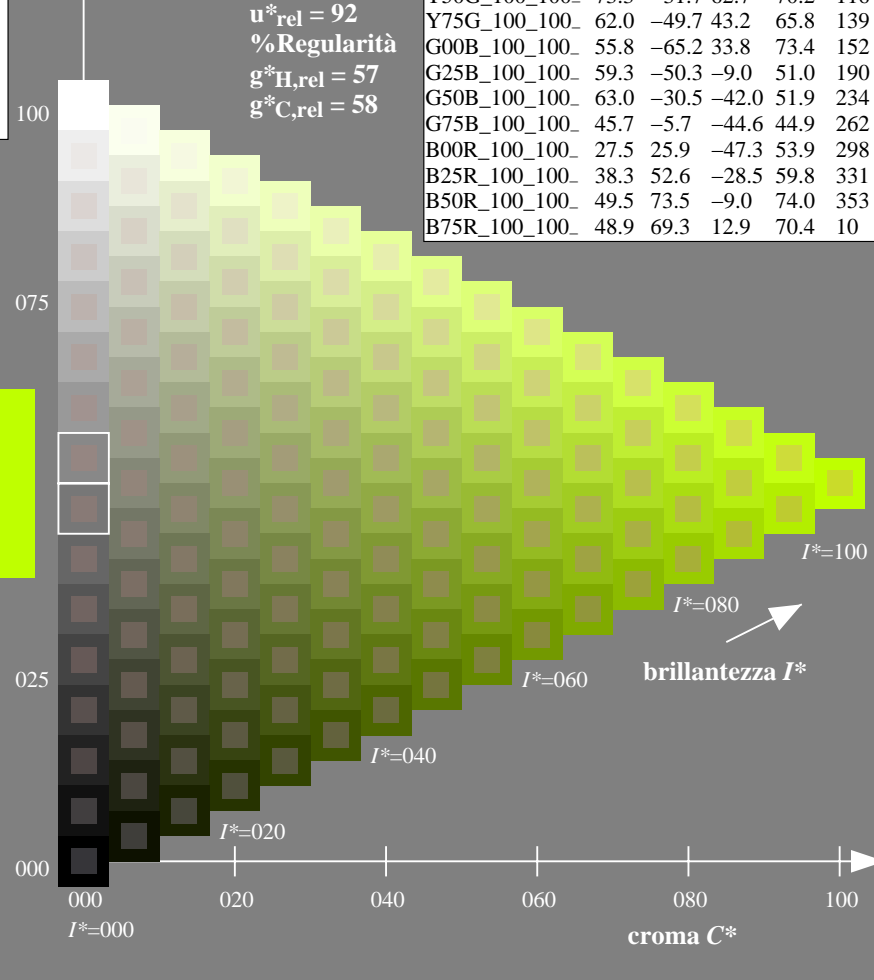
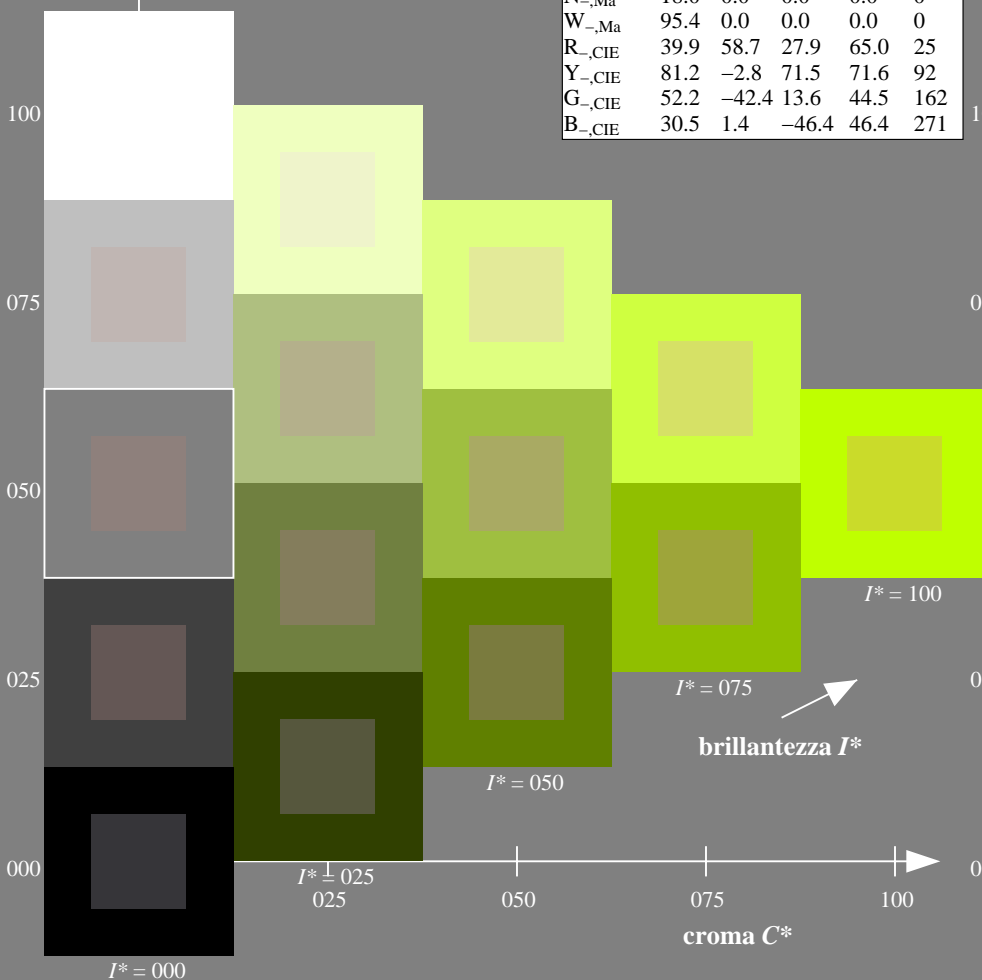
%Regularità

$g^*_{H,rel} = 57$

$g^*_{C,rel} = 58$

ORS20a; dati atti CIELAB (a)

$H^*_$	$L^*=L^*_a a^*_a$	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
R00Y_100_100_	48.4	66.1	40.2	77.3
R25Y_100_100_	56.8	48.0	50.5	69.6
R50Y_100_100_	68.6	25.0	63.9	68.6
R75Y_100_100_	80.6	4.8	77.2	77.3
Y00G_100_100_	90.2	-9.6	88.2	88.7
Y25G_100_100_	83.2	-18.4	79.9	81.9
Y50G_100_100_	73.3	-31.7	62.7	70.2
Y75G_100_100_	62.0	-49.7	43.2	65.8
G00B_100_100_	55.8	-65.2	33.8	73.4
G25B_100_100_	59.3	-50.3	-9.0	51.0
G50B_100_100_	63.0	-30.5	-42.0	51.9
G75B_100_100_	45.7	-5.7	-44.6	44.9
B00R_100_100_	27.5	25.9	-47.3	53.9
B25R_100_100_	38.3	52.6	-28.5	59.8
B50R_100_100_	49.5	73.5	-9.0	74.0
B75R_100_100_	48.9	69.3	12.9	70.4



vedere dei file simili: http://130.149.60.45/~farbmetrik/QI41/QI41.HTM
 informazioni tecniche: http://www.ps.bam.de o http://130.149.60.45/~farbmetrik

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
 la domanda per la misura di stampa di display

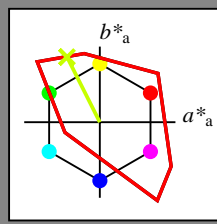
TUB materiale: code=rh4ta

Immettere y uscita: Television Luminous System TLS00a for relative CIELAB hue $h_{ab,a,rel} = h_{ab}/360 = 116/360 = 0.32$

$H^*_d = Y25G_d$

Dati del dispositivo (d) o colori elementari (e):

HIC^*_d
codice di tonalità per i colori questa pagina:
 $H^*_d = Y25G_d$
triangolo chiarezza T^*



TLS00a; dati atti CIELAB (a)

name	$L^*=L^*_a$	a^*_a	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
R _{d,Ma}	50.4	76.9	64.5	100.4	40
Y _{d,Ma}	92.6	-20.7	90.7	93.0	102
G _{d,Ma}	83.6	-82.7	79.8	115.0	136
C _{d,Ma}	86.8	-46.1	-13.5	48.1	196
B _{d,Ma}	30.3	76.0	-103.5	128.5	306
M _{d,Ma}	57.2	94.3	-58.4	110.9	328
N _{d,Ma}	0.0	0.0	0.0	0.0	0
W _{d,Ma}	95.4	0.0	0.0	0.0	0
R _{d,CIE}	39.9	58.7	27.9	65.0	25
Y _{d,CIE}	81.2	-2.8	71.5	71.6	92
G _{d,CIE}	52.2	-42.4	13.6	44.5	162
B _{d,CIE}	30.5	1.4	-46.4	46.4	271

Il dati per il massimo colore (Ma):

$LabCh^*_d, Ma: 88 -43 86 96 116$

$HIC^*_d, Ma: Y25G_{100_{100d}}$

$rgbic^*_d, Ma:$

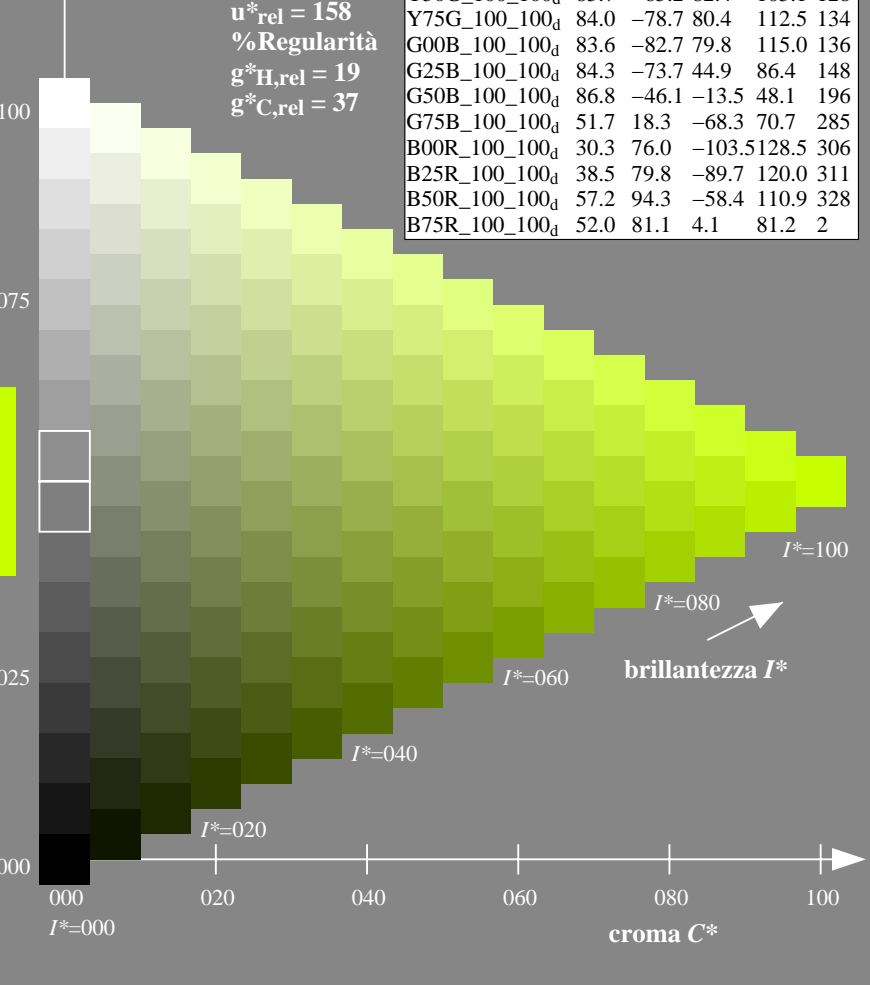
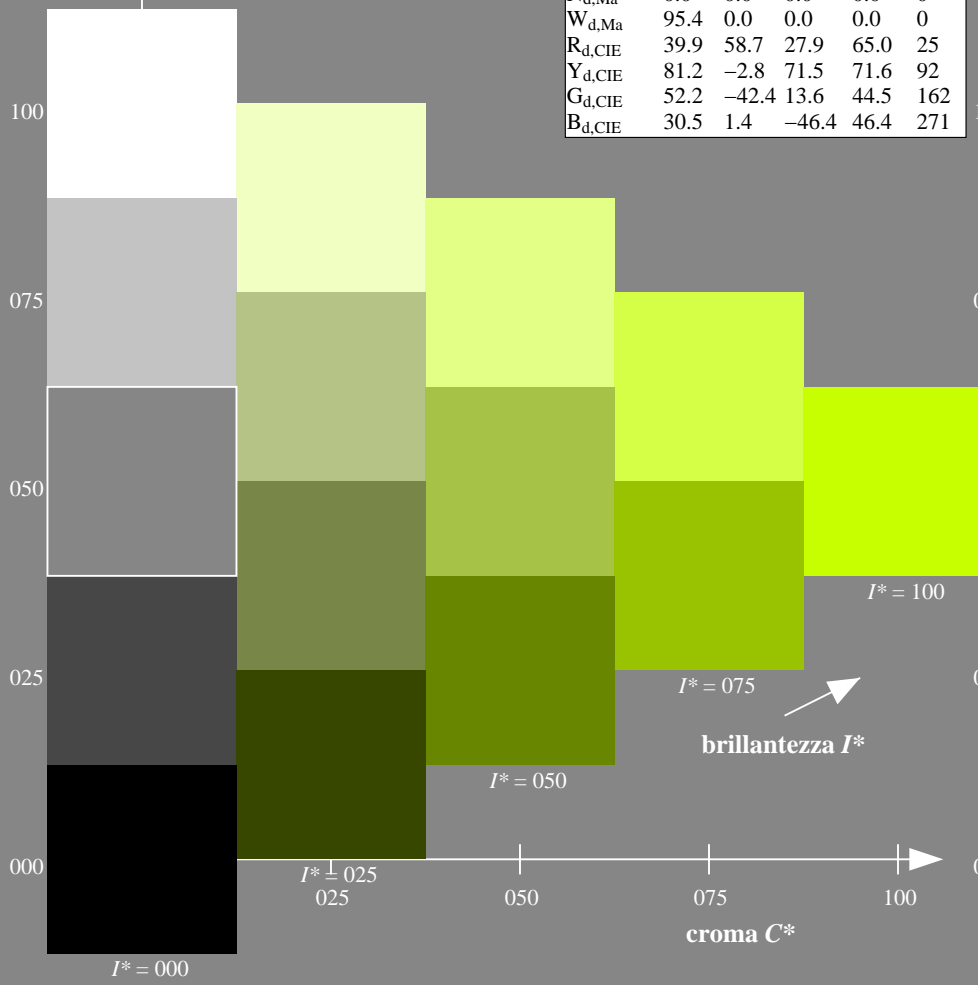
0.76 1.0 0.0 1.0 1.0

triangolo chiarezza T^*

TLS00a; dati atti CIELAB (a)

H^*_d	$L^*=L^*_a$	a^*_a	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
R00Y_100_100 _d	50.4	76.9	64.5	100.4	40
R25Y_100_100 _d	53.7	67.6	65.8	94.4	44
R50Y_100_100 _d	63.6	41.3	71.0	82.2	59
R75Y_100_100 _d	78.2	7.8	80.6	81.0	84
Y00G_100_100 _d	92.6	-20.7	90.7	93.0	102
Y25G_100_100 _d	88.7	-43.3	86.2	96.5	116
Y50G_100_100 _d	85.7	-65.2	82.4	105.1	128
Y75G_100_100 _d	84.0	-78.7	80.4	112.5	134
G00B_100_100 _d	83.6	-82.7	79.8	115.0	136
G25B_100_100 _d	84.3	-73.7	44.9	86.4	148
G50B_100_100 _d	86.8	-46.1	-13.5	48.1	196
G75B_100_100 _d	51.7	18.3	-68.3	70.7	285
B00R_100_100 _d	30.3	76.0	-103.5	128.5	306
B25R_100_100 _d	38.5	79.8	-89.7	120.0	311
B50R_100_100 _d	57.2	94.3	-58.4	110.9	328
B75R_100_100 _d	52.0	81.1	4.1	81.2	2

%Gamma
 $u^*_{rel} = 158$
%Regularità
 $g^*_{H,rel} = 19$
 $g^*_{C,rel} = 37$



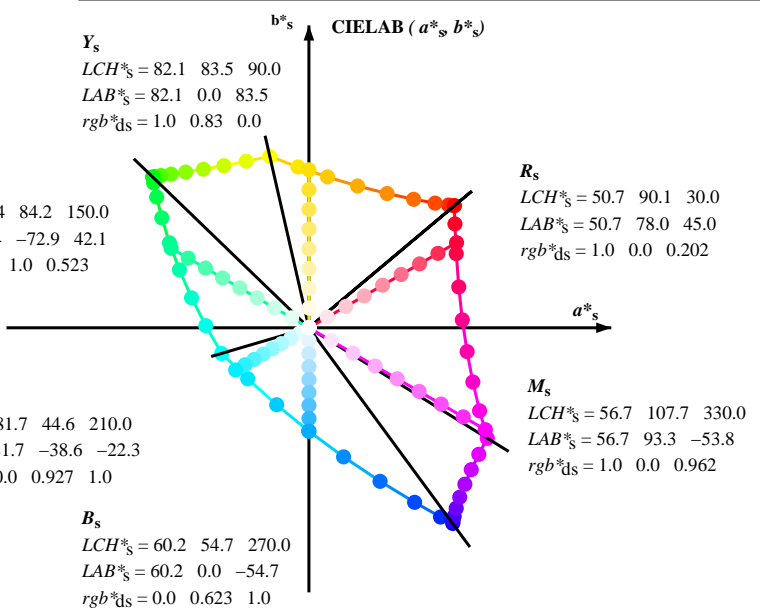
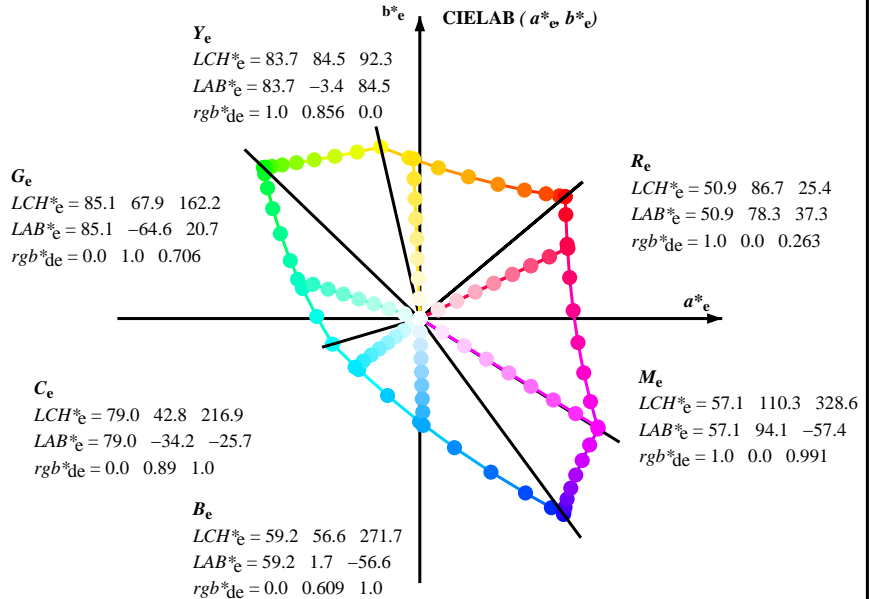
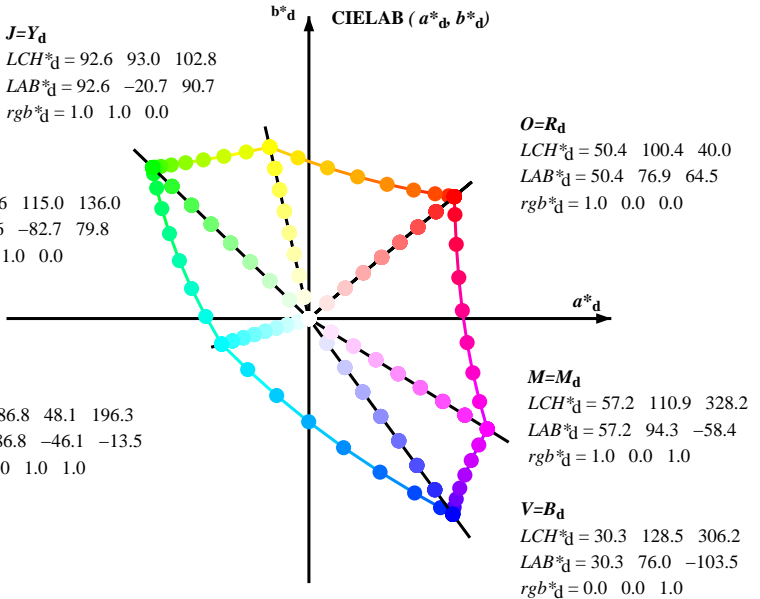
vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione

TUB materiale: code=rh4ta



Data of Maximum color M in colorimetric system sRGB standard device; no separation, D65 for input or output; Six hue angles of the 60 degree standard colours $RYGCBM_s$: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$; Six hue angles of the device colours $RYGCBM_d$: $h_{ab,d} = 40.0, 102.9, 136.0, 196.4, 306.3, 328.2$; Six hue angles of the elementary colours $RYGCBM_e$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$



$(a^*_d, b^*_d), (a^*_s, b^*_s), (a^*_e, b^*_e)$
 $rgb^*_d, LCH^*_d, LAB^*_d$
 h_{ab}, rgb^*_d
 $h_{ab,s} = atan [r^*_d \ cos(30) + g^*_d \ cos(150)] / [r^*_d \ sin(30) + g^*_d \ sin(150) + b^*_d \ sin(270)]$ (1)
 $h_{ab,s}$
 $s: h_{ab,s} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0, 390.0 (i=0,6)$
 $h_{48ab,sij} = h_{ab,si} + j [h_{ab,si+1} - h_{ab,si}] / 8 (i = 0, 1, \dots, 5; j = 0, 1, \dots, 7)$ (2)
 $h_{360ab,sij} = h_{ab,si} + j [h_{ab,si+1} - h_{ab,si}] / 60 (i = 0, 1, \dots, 5; j = 0, 1, \dots, 59)$ (3)
 $h_{ab,e}$
 $e: h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6, 385.5 (i=0,6)$
 $h_{48ab,eij} = h_{ab,ei} + j [h_{ab,ei+1} - h_{ab,ei}] / 8 (i = 0, 1, \dots, 5; j = 0, 1, \dots, 7)$ (4)
 $h_{360ab,eij} = h_{ab,ei} + j [h_{ab,ei+1} - h_{ab,ei}] / 60 (i = 0, 1, \dots, 5; j = 0, 1, \dots, 59)$ (5)
 $h_{ab}, h_{ab,d}$
 rgb^*_{de}

vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rh4ta

Data of maximum color M in colorimetric system sRGB standard device; no separation, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM_s: h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;

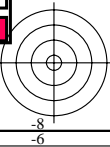
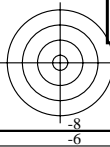
Six hue angles of the device colours RYGBM_d: h_{ab,d} = 40.0, 102.9, 136.0, 196.4, 306.3, 328.2; Six hue angles of the elementary colours RYGBM_e: h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6

Table with 12 columns of colorimetric data (h_{ab,d}, h_{ab,s}, h_{ab,e}, r_{gb}^a, d_{64M}, LAB*, d_{361M}, LAB*, d_{361M}, r_{gb}^a, d_{361M}, LAB*, d_{361M}, r_{gb}^a, d_{361M}, LAB*, d_{361M}) and 12 corresponding color patches (rgb^a_{dd}, rgb^a_{ds}, rgb^a_{de}) on the right.

vedere dei file simili: http://130.149.60.45/~farbmetrik/QI41/QI41.HTM
informazioni tecniche: http://www.ps.bam.de o http://130.149.60.45/~farbmetrik

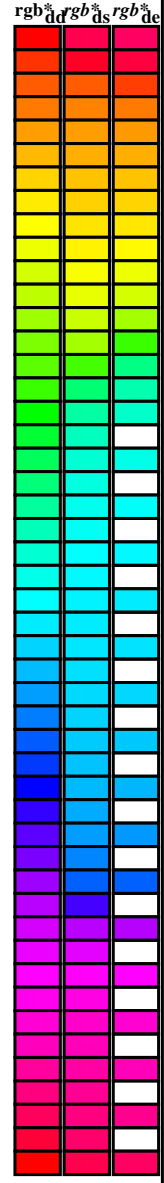
TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione

TUB materiale: code=rhatha



Data of Maximum color M in colorimetric system sRGB standard device; no separation, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM_s: h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;
Six hue angles of the device colours RYGBM_d: h_{ab,d} = 40.0, 102.9, 136.0, 196.4, 306.3, 328.2; Six hue angles of the elementary colours RYGBM_e: h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6

h _{ab,d}	h _{ab,s}	h _{ab,e}	rgb* dd64M	LAB* ddx64M (x=LabCh)	rgb* dex361M	LAB* dex361M
40.0	30.0	25.4	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0	1.0 0.0 0.263 50.9	78.3 37.3 86.7 25
41.3	37.5	33.8	1.0 0.125 0.0	51.5 73.9 64.9 98.3 41.3	1.0 0.0 0.156 50.7	77.7 51.0 92.9 33
44.6	45.0	42.1	1.0 0.25 0.0	54.0 66.7 65.9 93.8 44.6	1.0 0.157 0.0	52.2 72.0 65.3 97.2 42
50.7	52.5	50.5	1.0 0.375 0.0	58.2 55.4 67.9 87.7 50.7	1.0 0.358 0.0	57.7 56.9 67.8 88.6 49
59.7	60.0	58.8	1.0 0.5 0.0	63.6 41.3 71.0 82.2 59.7	1.0 0.488 0.0	63.1 42.8 70.9 82.8 58
71.0	67.5	67.2	1.0 0.625 0.0	70.1 25.7 75.0 79.3 71.0	1.0 0.577 0.0	67.6 31.8 73.9 80.5 66
82.9	75.0	75.6	1.0 0.75 0.0	77.2 9.8 79.7 80.4 82.9	1.0 0.673 0.0	72.8 19.8 77.3 79.8 75
93.8	82.5	83.9	1.0 0.875 0.0	84.8 -5.7 85.0 85.2 93.8	1.0 0.755 0.0	77.5 9.3 80.1 80.6 83
102.8	90.0	92.3	1.0 1.0 0.0	92.6 -20.7 90.7 93.0 102.8	1.0 0.857 0.0	83.7 -3.3 84.5 84.6 92
110.5	97.5	101.0	0.875 1.0 0.0	90.4 -33.1 88.1 94.1 110.5	1.0 0.967 0.0	90.6 -16.4 89.5 91.0 100
117.6	105.0	109.7	0.75 1.0 0.0	88.5 -44.9 85.8 96.8 117.6	0.888 1.0 0.0	90.7 -31.7 88.5 94.0 109
123.6	112.5	118.5	0.625 1.0 0.0	86.9 -55.8 83.9 100.7 123.6	0.743 1.0 0.0	88.5 -45.4 85.8 97.1 117
128.3	120.0	127.2	0.5 1.0 0.0	85.7 -65.2 82.4 105.1 128.3	0.529 1.0 0.0	86.0 -62.9 82.9 104.1 127
131.8	127.5	136.0	0.375 1.0 0.0	84.7 -72.8 81.2 109.1 131.8	0.132 1.0 0.0	83.8 -81.2 80.1 114.1 135
134.1	135.0	144.7	0.25 1.0 0.0	84.1 -78.2 80.5 112.2 134.1	0.0 1.0 0.41	84.1 -76.8 54.3 94.1 144
135.5	142.5	153.4	0.125 1.0 0.0	83.7 -81.4 80.0 114.2 135.5	0.0 1.0 0.573	84.6 -70.9 36.3 79.8 152
136.0	150.0	162.2	0.0 1.0 0.0	83.6 -82.7 79.8 115.0 136.0	0.0 1.0 0.706	85.2 -64.6 20.7 67.9 162
137.0	157.5	169.0	0.0 1.0 0.125	83.6 -82.1 76.6 112.3 137.0	0.0 1.0 0.778	85.5 -60.6 12.2 61.9 168
139.3	165.0	175.9	0.0 1.0 0.25	83.8 -80.5 69.1 106.1 139.3	0.0 1.0 0.847	85.9 -56.4 4.0 56.7 175
143.2	172.5	182.7	0.0 1.0 0.375	84.0 -77.8 58.1 97.1 143.2	0.0 1.0 0.9	86.2 -53.2 -2.0 53.3 182
148.6	180.0	189.6	0.0 1.0 0.5	84.3 -73.7 44.9 86.4 148.6	0.0 1.0 0.952	86.6 -49.8 -8.3 50.6 189
155.8	187.5	196.4	0.0 1.0 0.625	84.7 -68.5 30.6 75.0 155.8	0.0 1.0 0.997	86.9 -46.3 -13.2 48.3 195
165.6	195.0	203.2	0.0 1.0 0.75	85.3 -62.0 15.9 64.0 165.6	0.0 0.963	1.0 84.3 -42.5 -18.2 46.4 203
178.8	202.5	210.1	0.0 1.0 0.875	86.0 -54.5 1.0 54.5 178.8	0.0 0.929	1.0 81.8 -38.8 -22.1 44.7 209
196.3	210.0	216.9	0.0 1.0 1.0	86.8 -46.1 -13.5 48.1 196.3	0.0 0.89	1.0 79.1 -34.2 -25.7 42.9 216
219.8	217.5	223.8	0.0 0.875 1.0	77.9 -32.3 -27.0 42.1 219.8	0.0 0.859	1.0 76.9 -30.7 -29.0 42.4 223
247.2	225.0	230.6	0.0 0.75 1.0	69.1 -17.0 -40.7 44.1 247.2	0.0 0.826	1.0 74.5 -27.1 -33.1 43.0 230
269.8	232.5	237.5	0.0 0.625 1.0	60.3 -0.1 -54.6 54.6 269.8	0.0 0.797	1.0 72.4 -23.5 -36.3 43.4 237
285.0	240.0	244.3	0.0 0.5 1.0	51.7 18.3 -68.3 70.7 285.0	0.0 0.763	1.0 70.1 -18.9 -39.5 44.0 244
294.8	247.5	251.2	0.0 0.375 1.0	43.8 37.6 -81.2 89.5 294.8	0.0 0.731	1.0 67.8 -15.0 -43.1 45.8 250
301.1	255.0	258.0	0.0 0.25 1.0	37.1 55.9 -92.3 107.9 301.1	0.0 0.69	1.0 64.9 -10.1 -48.0 49.2 258
304.8	262.5	264.8	0.0 0.125 1.0	32.4 69.5 -100.0 121.8 304.8	0.0 0.655	1.0 62.4 -5.0 -51.8 52.1 264
306.2	270.0	271.7	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2	0.0 0.609	1.0 59.3 1.7 -56.5 56.6 271
306.6	277.5	278.8	0.125 0.0 1.0	31.0 76.2 -102.4 127.7 306.6	0.0 0.555	1.0 55.5 9.3 -62.9 63.7 278
307.5	285.0	285.9	0.25 0.0 1.0	32.6 76.8 -99.8 125.9 307.5	0.0 0.488	1.0 51.0 19.9 -69.6 72.5 285
309.2	292.5	293.0	0.375 0.0 1.0	35.1 77.9 -95.5 123.3 309.2	0.0 0.404	1.0 45.7 32.7 -78.5 85.2 292
311.6	300.0	300.1	0.5 0.0 1.0	38.5 79.8 -89.7 120.0 311.6	0.0 0.27	1.0 38.2 52.8 -90.6 105.0 300
314.8	307.5	307.2	0.625 0.0 1.0	42.7 82.5 -82.7 116.8 314.8	0.0 0.146	0.0 31.3 76.4 -102.0 127.5 306
318.8	315.0	314.3	0.75 0.0 1.0	47.2 85.8 -75.1 114.0 318.8	0.0 0.605	0.0 42.1 82.1 -83.8 117.4 314
323.3	322.5	321.4	0.875 0.0 1.0	52.1 89.8 -66.9 112.0 323.3	0.0 0.811	0.0 49.7 87.9 -71.0 113.1 321
328.2	330.0	328.6	1.0 0.0 1.0	57.2 94.3 -58.4 110.9 328.2	0.0 0.992	0.0 57.2 94.2 -57.4 110.3 328
334.0	337.5	335.7	1.0 0.0 0.875	55.6 90.3 -43.9 100.4 334.0	0.0 0.856	0.0 55.4 89.9 -41.4 99.0 335
341.6	345.0	342.8	1.0 0.0 0.75	54.2 86.7 -28.6 91.3 341.6	0.0 0.735	0.0 54.1 86.5 -26.6 90.6 342
351.4	352.5	349.9	1.0 0.0 0.625	53.0 83.6 -12.6 84.6 351.4	0.0 0.65	0.0 53.3 84.5 -15.6 86.0 349
362.9	360.0	357.0	1.0 0.0 0.5	52.0 81.1 4.1 81.2 362.9	0.0 0.618	0.0 53.0 83.6 -11.6 84.4 352
375.2	367.5	364.1	1.0 0.0 0.375	51.3 79.2 21.6 82.1 375.2	0.0 0.533	0.0 52.3 82.2 -0.1 82.2 359
386.7	375.0	371.2	1.0 0.0 0.25	50.8 77.9 39.2 87.2 386.7	0.0 0.441	0.0 51.7 80.7 12.5 81.7 368
395.4	382.5	378.3	1.0 0.0 0.125	50.6 77.2 54.9 94.8 395.4	0.0 0.361	0.0 51.3 79.3 23.6 82.8 376
400.0	390.0	385.4	1.0 0.0 0.0	50.4 76.9 64.5 100.4 400.0	0.0 0.263	0.0 50.9 78.3 37.3 86.7 385



vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF> / .PS
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rh4ta

Data of Maximum color M in colorimetric system sRGB standard device; no separation, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM_s: h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;
Six hue angles of the device colours RYGBM_d: h_{ab,d} = 40.0, 102.9, 136.0, 196.4, 306.3, 328.2; Six hue angles of the elementary colours RYGBM_e: h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6

h _{ab,d}	h _{ab,s}	h _{ab,e}	rgb ^{ab} * dd361M	LAB ^{ab} * ddx361Mi (x=LabCh)	R _d	rgb ^{ab} * ds361Mi	LAB ^{ab} * dsx361Mi (x=LabCh)	R _s	rgb ^{ab} * dd361Mi	LAB ^{ab} * dex361Mi (x=LabCh)	R _e	rgb ^{ab} * dd361Mi	rgb ^{ab} * dd	rgb ^{ab} * ds	rgb ^{ab} * de
40	30	25	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40	1.0	1.0 0.0 0.203 50.8 78.0 45.1 90.1 30	1.0	1.0 0.0 0.0	1.0 0.0 0.263 50.9 78.3 37.3 86.7 25	1.0	1.0 0.0 0.0				
40	31	26	1.0 0.016 0.0	50.6 76.5 64.6 100.1 40	1.0	1.0 0.0 0.189 50.7 78.0 46.9 91.0 31	1.0	1.0 0.017 0.0	1.0 0.0 0.251 50.9 78.0 39.0 87.2 26	1.0	1.0 0.017 0.0				
40	32	27	1.0 0.033 0.0	50.7 76.1 64.6 99.8 40	1.0	1.0 0.0 0.174 50.7 77.9 48.7 91.8 32	1.0	1.0 0.033 0.0	1.0 0.0 0.236 50.8 78.0 41.0 88.1 27	1.0	1.0 0.033 0.0				
40	33	28	1.0 0.05 0.0	50.9 75.7 64.7 99.6 40	1.0	1.0 0.0 0.16 50.7 77.7 50.5 92.7 33	1.0	1.0 0.05 0.0	1.0 0.0 0.22 50.8 78.1 43.0 89.1 28	1.0	1.0 0.05 0.0				
40	34	29	1.0 0.066 0.0	51.0 75.3 64.7 99.3 40	1.0	1.0 0.0 0.146 50.6 77.6 52.3 93.6 34	1.0	1.0 0.067 0.0	1.0 0.0 0.204 50.8 78.0 44.9 90.1 29	1.0	1.0 0.067 0.0				
40	35	31	1.0 0.083 0.0	51.1 74.9 64.8 99.0 40	1.0	1.0 0.0 0.131 50.6 77.3 54.2 94.4 35	1.0	1.0 0.083 0.0	1.0 0.0 0.188 50.7 78.0 46.9 91.0 31	1.0	1.0 0.083 0.0				
41	36	32	1.0 0.1 0.0	51.3 74.5 64.8 98.7 41	1.0	1.0 0.0 0.11 50.6 77.3 56.1 95.5 36	1.0	1.0 0.1 0.0	1.0 0.0 0.172 50.7 77.9 49.0 92.0 32	1.0	1.0 0.1 0.0				
41	37	33	1.0 0.116 0.0	51.4 74.1 64.9 98.5 41	1.0	1.0 0.0 0.082 50.6 77.2 58.2 96.7 37	1.0	1.0 0.117 0.0	1.0 0.0 0.156 50.7 77.7 51.0 92.9 33	1.0	1.0 0.117 0.0				
41	38	34	1.0 0.133 0.0	51.7 73.4 65.0 98.0 41	1.0	1.0 0.0 0.055 50.5 77.2 60.3 98.0 38	1.0	1.0 0.133 0.0	1.0 0.0 0.14 50.6 77.5 53.0 93.9 34	1.0	1.0 0.133 0.0				
41	39	35	1.0 0.15 0.0	52.0 72.4 65.2 97.4 41	1.0	1.0 0.0 0.028 50.5 77.1 62.4 99.2 39	1.0	1.0 0.15 0.0	1.0 0.0 0.123 50.6 77.2 55.1 94.9 35	1.0	1.0 0.15 0.0				
42	40	36	1.0 0.166 0.0	52.3 71.4 65.3 96.8 42	1.0	1.0 0.0 0.0 50.5 76.9 64.6 100.4 40	1.0	1.0 0.167 0.0	1.0 0.0 0.093 50.6 77.3 57.4 96.3 36	1.0	1.0 0.167 0.0				
42	41	37	1.0 0.183 0.0	52.7 70.5 65.5 96.2 42	1.0	1.0 0.095 0.0 51.3 74.6 64.9 98.9 41	1.0	1.0 0.183 0.0	1.0 0.0 0.062 50.5 77.2 59.7 97.6 37	1.0	1.0 0.183 0.0				
43	42	38	1.0 0.2 0.0	53.0 69.5 65.6 95.6 43	1.0	1.0 0.151 0.0 52.1 72.4 65.2 97.5 42	1.0	1.0 0.2 0.0	1.0 0.0 0.032 50.5 77.1 62.1 99.0 38	1.0	1.0 0.2 0.0				
43	43	39	1.0 0.216 0.0	53.4 68.6 65.7 95.0 43	1.0	1.0 0.188 0.0 52.8 70.3 65.5 96.1 43	1.0	1.0 0.217 0.0	1.0 0.0 0.001 50.5 76.9 64.5 100.4 39	1.0	1.0 0.217 0.0				
44	44	41	1.0 0.233 0.0	53.7 67.6 65.8 94.4 44	1.0	1.0 0.225 0.0 53.6 68.2 65.8 94.8 44	1.0	1.0 0.233 0.0	1.0 0.102 0.0 51.4 74.4 64.9 98.8 41	1.0	1.0 0.233 0.0				
44	45	42	1.0 0.25 0.0	54.0 66.7 65.9 93.8 44	1.0	1.0 0.256 0.0 54.3 66.1 66.1 93.5 45	1.0	1.0 0.25 0.0	1.0 0.157 0.0 52.2 72.0 65.3 97.2 42	1.0	1.0 0.25 0.0				
45	46	43	1.0 0.266 0.0	54.6 65.1 66.3 93.0 45	1.0	1.0 0.277 0.0 55.0 64.3 66.6 92.5 46	1.0	1.0 0.267 0.0	1.0 0.199 0.0 53.0 69.6 65.6 95.7 43	1.0	1.0 0.267 0.0				
46	47	44	1.0 0.283 0.0	55.1 63.6 66.6 92.2 46	1.0	1.0 0.297 0.0 55.6 62.4 66.9 91.5 47	1.0	1.0 0.283 0.0	1.0 0.24 0.0 53.9 67.3 65.9 94.2 44	1.0	1.0 0.283 0.0				
47	48	45	1.0 0.3 0.0	55.7 62.1 66.9 91.3 47	1.0	1.0 0.318 0.0 56.3 60.6 67.3 90.5 48	1.0	1.0 0.3 0.0	1.0 0.267 0.0 54.7 65.1 66.4 93.0 45	1.0	1.0 0.3 0.0				
47	49	46	1.0 0.316 0.0	56.2 60.6 67.2 90.5 47	1.0	1.0 0.338 0.0 57.0 58.7 67.6 89.5 49	1.0	1.0 0.317 0.0	1.0 0.29 0.0 55.4 63.1 66.8 91.9 46	1.0	1.0 0.317 0.0				
48	50	47	1.0 0.333 0.0	56.8 59.1 67.5 89.7 48	1.0	1.0 0.359 0.0 57.7 56.9 67.8 88.5 50	1.0	1.0 0.333 0.0	1.0 0.313 0.0 56.2 61.0 67.2 90.8 47	1.0	1.0 0.333 0.0				
49	51	48	1.0 0.35 0.0	57.3 57.6 67.7 88.9 49	1.0	1.0 0.378 0.0 58.3 55.1 68.1 87.6 51	1.0	1.0 0.35 0.0	1.0 0.336 0.0 56.9 59.0 67.5 89.7 48	1.0	1.0 0.35 0.0				
50	52	49	1.0 0.366 0.0	57.9 56.2 67.9 88.1 50	1.0	1.0 0.392 0.0 58.9 53.6 68.6 87.0 52	1.0	1.0 0.367 0.0	1.0 0.358 0.0 57.7 56.9 67.8 88.6 49	1.0	1.0 0.367 0.0				
51	53	51	1.0 0.383 0.0	58.5 54.5 68.2 87.3 51	1.0	1.0 0.406 0.0 59.6 52.0 69.0 86.4 53	1.0	1.0 0.383 0.0	1.0 0.379 0.0 58.4 55.0 68.1 87.6 51	1.0	1.0 0.383 0.0				
52	54	52	1.0 0.4 0.0	59.3 52.6 68.8 86.6 52	1.0	1.0 0.42 0.0 60.2 50.4 69.4 85.8 54	1.0	1.0 0.4 0.0	1.0 0.395 0.0 59.1 53.2 68.7 86.9 52	1.0	1.0 0.4 0.0				
53	55	53	1.0 0.416 0.0	60.0 50.7 69.3 85.9 53	1.0	1.0 0.433 0.0 60.8 48.8 69.8 85.2 55	1.0	1.0 0.417 0.0	1.0 0.41 0.0 59.7 51.5 69.1 86.2 53	1.0	1.0 0.417 0.0				
54	56	54	1.0 0.433 0.0	60.7 48.8 69.7 85.1 54	1.0	1.0 0.447 0.0 61.4 47.3 70.1 84.5 56	1.0	1.0 0.433 0.0	1.0 0.426 0.0 60.4 49.7 69.6 85.5 54	1.0	1.0 0.433 0.0				
56	57	55	1.0 0.45 0.0	61.4 46.9 70.1 84.4 56	1.0	1.0 0.461 0.0 62.0 45.7 70.4 83.9 57	1.0	1.0 0.45 0.0	1.0 0.441 0.0 61.1 48.0 69.9 84.8 55	1.0	1.0 0.45 0.0				
57	58	56	1.0 0.466 0.0	62.2 45.1 70.4 83.6 57	1.0	1.0 0.475 0.0 62.6 44.1 70.7 83.3 58	1.0	1.0 0.467 0.0	1.0 0.457 0.0 61.8 46.2 70.3 84.1 56	1.0	1.0 0.467 0.0				
58	59	57	1.0 0.483 0.0	62.9 43.2 70.7 82.9 58	1.0	1.0 0.489 0.0 63.2 42.6 70.9 82.7 59	1.0	1.0 0.483 0.0	1.0 0.472 0.0 62.5 44.5 70.6 83.4 57	1.0	1.0 0.483 0.0				
59	60	58	1.0 0.5 0.0	63.6 41.3 71.0 82.2 59	1.0	1.0 0.502 0.0 63.8 41.1 71.2 82.2 60	1.0	1.0 0.5 0.0	1.0 0.488 0.0 63.1 42.8 70.9 82.8 58	1.0	1.0 0.5 0.0				
61	61	60	1.0 0.516 0.0	64.5 39.3 71.7 81.8 61	1.0	1.0 0.513 0.0 64.4 39.7 71.6 81.9 61	1.0	1.0 0.517 0.0	1.0 0.502 0.0 63.8 41.1 71.2 82.2 60	1.0	1.0 0.517 0.0				
62	62	61	1.0 0.533 0.0	65.3 37.2 72.4 81.4 62	1.0	1.0 0.525 0.0 64.9 38.3 72.1 81.7 62	1.0	1.0 0.533 0.0	1.0 0.515 0.0 64.4 39.5 71.7 81.9 61	1.0	1.0 0.533 0.0				
64	63	62	1.0 0.55 0.0	66.2 35.1 73.0 81.0 64	1.0	1.0 0.536 0.0 65.5 37.0 72.5 81.4 63	1.0	1.0 0.55 0.0	1.0 0.527 0.0 65.1 38.0 72.2 81.6 62	1.0	1.0 0.55 0.0				
65	64	63	1.0 0.566 0.0	67.1 33.0 73.5 80.6 65	1.0	1.0 0.547 0.0 66.1 35.6 72.9 81.1 64	1.0	1.0 0.567 0.0	1.0 0.54 0.0 65.7 36.5 72.7 81.3 63	1.0	1.0 0.567 0.0				
67	65	64	1.0 0.583 0.0	67.9 31.0 74.0 80.3 67	1.0	1.0 0.558 0.0 66.7 34.2 73.3 80.9 65	1.0	1.0 0.583 0.0	1.0 0.552 0.0 66.4 34.9 73.1 81.0 64	1.0	1.0 0.583 0.0				
68	66	65	1.0 0.6 0.0	68.8 28.9 74.5 79.9 68	1.0	1.0 0.569 0.0 67.2 32.8 73.7 80.6 66	1.0	1.0 0.6 0.0	1.0 0.564 0.0 67.0 33.4 73.5 80.7 65	1.0	1.0 0.6 0.0				
70	67	66	1.0 0.616 0.0	69.6 26.8 74.8 79.5 70	1.0	1.0 0.58 0.0 67.8 31.4 74.0 80.4 67	1.0	1.0 0.617 0.0	1.0 0.577 0.0 67.6 31.8 73.9 80.5 66	1.0	1.0 0.617 0.0				
71	68	67	1.0 0.633 0.0	70.5 24.7 75.4 79.4 71	1.0	1.0 0.591 0.0 68.4 30.0 74.3 80.1 68	1.0	1.0 0.633 0.0	1.0 0.589 0.0 68.3 30.3 74.2 80.2 67	1.0	1.0 0.633 0.0				
73	69	68	1.0 0.65 0.0	71.5 22.7 76.2 79.5 73	1.0	1.0 0.602 0.0 69.0 28.6 74.6 79.9 69	1.0	1.0 0.65 0.0	1.0 0.602 0.0 68.9 28.7 74.5 79.9 68	1.0	1.0 0.65 0.0				
75	70	70	1.0 0.666 0.0	72.4 20.6 76.9 79.7 75	1.0	1.0 0.614 0.0 69.5 27.2 74.8 79.6 70	1.0	1.0 0.667 0.0	1.0 0.614 0.0 69.5 27.2 74.8 79.6 70	1.0	1.0 0.667 0.0				
76	71	71	1.0 0.683 0.0	73.4 18.5 77.6 79.8 76	1.0	1.0 0.625 0.0 70.1 25.8 75.0 79.4 71	1.0	1.0 0.683 0.0	1.0 0.626 0.0 70.2 25.6 75.1 79.4 71	1.0	1.0 0.683 0.0				
78	72	72	1.0 0.7 0.0	74.3 16.3 78.2 79.9 78	1.0	1.0 0.635 0.0 70.7 24.5 75.6 79.4 72	1.0	1.0 0.7 0.0	1.0 0.638 0.0 70.9 24.2 75.7 79.5 72	1.0	1.0 0.7 0.0				
79	73	73	1.0 0.716 0.0	75.3 14.2 78.8 80.1 79	1.0	1.0 0.646 0.0 71.3 23.3 76.1 79.5 73	1.0	1.0 0.717 0.0	1.0 0.65 0.0 71.5 22.8 76.2 79.6 73	1.0	1.0 0.717 0.0				
81	74	74	1.0 0.733 0.0	76.2 12.0 79.3 80.2 81	1.0	1.0 0.656 0.0 71.9 21.9 76.5 79.6 74	1.0	1.0 0.733 0.0	1.0 0.661 0.0 72.2 21.3 76.8 79.7 74	1.0	1.0 0.733 0.0				
82	75	75	1.0 0.75 0.0	77.2 9.8 79.7 80.4 82	1.0	1.0 0.667 0.0 72.5 20.6 77.0 79.7 75	1.0	1.0 0.75 0.0	1.0 0.673 0.0 72.8 19.8 77.3 79.8 75	1.0	1.0 0.75 0.0				

vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rh4ta

Data of Maximum color M in colorimetric system sRGB standard device; no separation, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM_s; h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;
Six hue angles of the device colours RYGBM_d; h_{ab,d} = 40.0, 102.9, 136.0, 196.4, 306.3, 328.2; Six hue angles of the elementary colours RYGBM_e; h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6

h _{ab,d}	h _{ab,s}	h _{ab,e}	rgb [*] _{dd361M}	LAB [*] _{ddx361Mi (x=LabCh)}	rgb [*] _{ds361Mi}	LAB [*] _{dsx361Mi (x=LabCh)}	rgb [*] _{dd361Mi}	LAB [*] _{de361Mi}	rgb [*] _{dex361Mi (x=LabCh)}	rgb [*] _{dd361Mi}	LAB [*] _{de361Mi}	rgb [*] _{dd361Mi}	rgb [*] _{dd}	rgb [*] _{ds}	rgb [*] _{de}
82	75	75	1.0 0.75 0.0	77.2 9.8 79.7 80.4 82	1.0 0.667 0.0	72.5 20.6 77.0 79.7 75	1.0 0.75 0.0	1.0 0.673 0.0	72.8 19.8 77.3 79.8 75	1.0 0.75 0.0	1.0 0.673 0.0	1.0 0.75 0.0			
84	76	76	1.0 0.766 0.0	78.2 7.8 80.6 81.0 84	1.0 0.677 0.0	73.1 19.3 77.4 79.8 76	1.0 0.767 0.0	1.0 0.685 0.0	73.5 18.3 77.7 79.9 76	1.0 0.767 0.0	1.0 0.685 0.0	1.0 0.767 0.0			
85	77	77	1.0 0.783 0.0	79.2 5.8 81.4 81.7 85	1.0 0.688 0.0	73.7 18.0 77.8 79.9 77	1.0 0.783 0.0	1.0 0.696 0.0	74.2 16.9 78.2 80.0 77	1.0 0.783 0.0	1.0 0.696 0.0	1.0 0.783 0.0			
87	78	78	1.0 0.8 0.0	80.2 3.8 82.2 82.3 87	1.0 0.698 0.0	74.3 16.6 78.2 80.0 78	1.0 0.8 0.0	1.0 0.708 0.0	74.8 15.3 78.6 80.1 78	1.0 0.8 0.0	1.0 0.708 0.0	1.0 0.8 0.0			
88	79	80	1.0 0.816 0.0	81.2 1.7 82.9 83.0 88	1.0 0.708 0.0	74.9 15.3 78.6 80.1 79	1.0 0.817 0.0	1.0 0.72 0.0	75.5 13.8 78.9 80.1 80	1.0 0.817 0.0	1.0 0.72 0.0	1.0 0.817 0.0			
90	80	81	1.0 0.833 0.0	82.2 -0.3 83.6 83.6 90	1.0 0.719 0.0	75.5 13.9 78.9 80.1 80	1.0 0.833 0.0	1.0 0.731 0.0	76.2 12.3 79.3 80.2 81	1.0 0.833 0.0	1.0 0.731 0.0	1.0 0.833 0.0			
91	81	82	1.0 0.85 0.0	83.3 -2.5 84.2 84.3 91	1.0 0.729 0.0	76.1 12.6 79.2 80.2 81	1.0 0.85 0.0	1.0 0.743 0.0	76.8 10.8 79.6 80.3 82	1.0 0.85 0.0	1.0 0.743 0.0	1.0 0.85 0.0			
93	82	83	1.0 0.866 0.0	84.3 -4.6 84.8 84.9 93	1.0 0.74 0.0	76.7 11.2 79.5 80.3 82	1.0 0.867 0.0	1.0 0.755 0.0	77.5 9.3 80.1 80.6 83	1.0 0.867 0.0	1.0 0.755 0.0	1.0 0.867 0.0			
94	83	84	1.0 0.883 0.0	85.3 -6.7 85.5 85.8 94	1.0 0.75 0.0	77.3 9.8 79.8 80.4 83	1.0 0.883 0.0	1.0 0.768 0.0	78.3 7.8 80.7 81.1 84	1.0 0.883 0.0	1.0 0.768 0.0	1.0 0.883 0.0			
95	84	85	1.0 0.9 0.0	86.3 -8.5 86.4 86.8 95	1.0 0.762 0.0	78.0 8.5 80.4 80.9 84	1.0 0.9 0.0	1.0 0.78 0.0	79.1 6.2 81.4 81.6 85	1.0 0.9 0.0	1.0 0.78 0.0	1.0 0.9 0.0			
96	85	86	1.0 0.916 0.0	87.4 -10.5 87.2 87.8 96	1.0 0.773 0.0	78.7 7.1 81.0 81.3 85	1.0 0.917 0.0	1.0 0.793 0.0	79.9 4.7 82.0 82.1 86	1.0 0.917 0.0	1.0 0.793 0.0	1.0 0.917 0.0			
98	86	87	1.0 0.933 0.0	88.4 -12.4 88.0 88.9 98	1.0 0.785 0.0	79.3 5.7 81.6 81.8 86	1.0 0.933 0.0	1.0 0.806 0.0	80.6 3.1 82.5 82.6 87	1.0 0.933 0.0	1.0 0.806 0.0	1.0 0.933 0.0			
99	87	88	1.0 0.95 0.0	89.5 -14.4 88.7 89.9 99	1.0 0.796 0.0	80.0 4.3 82.1 82.2 87	1.0 0.95 0.0	1.0 0.819 0.0	81.4 1.5 83.1 83.1 88	1.0 0.95 0.0	1.0 0.819 0.0	1.0 0.95 0.0			
100	88	90	1.0 0.966 0.0	90.5 -16.5 89.4 91.0 100	1.0 0.808 0.0	80.7 2.9 82.6 82.7 88	1.0 0.967 0.0	1.0 0.831 0.0	82.2 0.0 83.6 83.6 90	1.0 0.967 0.0	1.0 0.831 0.0	1.0 0.967 0.0			
101	89	91	1.0 0.983 0.0	91.6 -18.5 90.1 92.0 101	1.0 0.819 0.0	81.4 1.5 83.1 83.1 89	1.0 0.983 0.0	1.0 0.844 0.0	83.0 -1.7 84.1 84.1 91	1.0 0.983 0.0	1.0 0.844 0.0	1.0 0.983 0.0			
102	90	92	1.0 1.0 0.0	92.6 -20.7 90.7 93.0 102	Y _d 1.0 0.831 0.0	82.1 0.0 83.5 83.5 90	Y _s 1.0 1.0 0.0	1.0 0.857 0.0	83.7 -3.3 84.5 84.6 92	Y _e 1.0 1.0 0.0	1.0 0.857 0.0	1.0 1.0 0.0			
103	91	93	0.983 1.0 0.0	92.3 -22.3 90.5 93.2 103	1.0 0.842 0.0	82.8 -1.4 84.0 84.0 91	0.983 1.0 0.0	1.0 0.87 0.0	84.5 -5.1 84.9 85.1 93	0.983 1.0 0.0	1.0 0.87 0.0	0.983 1.0 0.0			
104	92	94	0.966 1.0 0.0	92.0 -24.0 90.2 93.3 104	1.0 0.853 0.0	83.5 -2.8 84.4 84.4 92	0.967 1.0 0.0	1.0 0.886 0.0	85.5 -6.9 85.7 85.9 94	0.967 1.0 0.0	1.0 0.886 0.0	0.967 1.0 0.0			
105	93	95	0.95 1.0 0.0	91.7 -25.6 89.9 93.5 105	1.0 0.865 0.0	84.2 -4.3 84.8 84.9 93	0.95 1.0 0.0	1.0 0.902 0.0	86.5 -8.7 86.5 87.0 95	0.95 1.0 0.0	1.0 0.902 0.0	0.95 1.0 0.0			
106	94	96	0.933 1.0 0.0	91.4 -27.3 89.5 93.6 106	1.0 0.877 0.0	84.9 -5.9 85.2 85.4 94	0.933 1.0 0.0	1.0 0.918 0.0	87.5 -10.6 87.3 88.0 96	0.933 1.0 0.0	1.0 0.918 0.0	0.933 1.0 0.0			
108	95	98	0.916 1.0 0.0	91.1 -28.9 89.1 93.7 108	1.0 0.891 0.0	85.8 -7.4 85.9 86.3 95	0.917 1.0 0.0	1.0 0.934 0.0	88.5 -12.5 88.1 89.0 98	0.917 1.0 0.0	1.0 0.934 0.0	0.917 1.0 0.0			
109	96	99	0.9 1.0 0.0	90.8 -30.6 88.7 93.9 109	1.0 0.904 0.0	86.7 -9.0 86.6 87.1 96	0.9 1.0 0.0	1.0 0.951 0.0	89.6 -14.4 88.8 90.0 99	0.9 1.0 0.0	1.0 0.951 0.0	0.9 1.0 0.0			
110	97	100	0.883 1.0 0.0	90.5 -32.2 88.3 94.0 110	1.0 0.918 0.0	87.5 -10.6 87.3 88.0 97	0.883 1.0 0.0	1.0 0.967 0.0	90.6 -16.4 89.5 91.0 100	0.883 1.0 0.0	1.0 0.967 0.0	0.883 1.0 0.0			
111	98	101	0.866 1.0 0.0	90.3 -33.8 88.0 94.3 111	1.0 0.932 0.0	88.4 -12.3 88.0 88.9 98	0.867 1.0 0.0	1.0 0.983 0.0	91.6 -18.5 90.1 92.0 101	0.867 1.0 0.0	1.0 0.983 0.0	0.867 1.0 0.0			
111	99	102	0.85 1.0 0.0	90.0 -35.4 87.7 94.6 111	1.0 0.946 0.0	89.3 -13.9 88.6 89.7 99	0.85 1.0 0.0	1.0 0.999 0.0	92.6 -20.5 90.7 93.0 102	0.85 1.0 0.0	1.0 0.999 0.0	0.85 1.0 0.0			
112	100	103	0.833 1.0 0.0	89.8 -37.0 87.5 95.0 112	1.0 0.96 0.0	90.2 -15.6 89.2 90.6 100	0.833 1.0 0.0	0.982 1.0 0.0	92.3 -22.4 90.5 93.2 103	0.833 1.0 0.0	0.982 1.0 0.0	0.833 1.0 0.0			
113	101	105	0.816 1.0 0.0	89.5 -38.6 87.2 95.4 113	1.0 0.974 0.0	91.0 -17.4 89.8 91.5 101	0.817 1.0 0.0	0.963 1.0 0.0	92.0 -24.3 90.2 93.4 105	0.817 1.0 0.0	0.963 1.0 0.0	0.817 1.0 0.0			
114	102	106	0.8 1.0 0.0	89.3 -40.1 86.9 95.7 114	1.0 0.988 0.0	91.9 -19.1 90.3 92.3 102	0.8 1.0 0.0	0.944 1.0 0.0	91.7 -26.1 89.8 93.6 106	0.8 1.0 0.0	0.944 1.0 0.0	0.8 1.0 0.0			
115	103	107	0.783 1.0 0.0	89.0 -41.7 86.6 96.1 115	0.998 1.0 0.0	92.6 -20.8 90.7 93.1 103	0.783 1.0 0.0	0.926 1.0 0.0	91.3 -28.0 89.4 93.7 107	0.783 1.0 0.0	0.926 1.0 0.0	0.783 1.0 0.0			
116	104	108	0.766 1.0 0.0	88.7 -43.3 86.2 96.5 116	0.981 1.0 0.0	92.3 -22.5 90.5 93.2 104	0.767 1.0 0.0	0.907 1.0 0.0	91.0 -29.9 89.0 93.9 108	0.767 1.0 0.0	0.907 1.0 0.0	0.767 1.0 0.0			
117	105	109	0.75 1.0 0.0	88.5 -44.9 85.8 96.8 117	0.965 1.0 0.0	92.0 -24.1 90.2 93.4 105	0.75 1.0 0.0	0.888 1.0 0.0	90.7 -31.7 88.5 94.0 109	0.75 1.0 0.0	0.888 1.0 0.0	0.75 1.0 0.0			
118	106	110	0.733 1.0 0.0	88.3 -46.3 85.6 97.4 118	0.949 1.0 0.0	91.8 -25.7 89.9 93.5 106	0.733 1.0 0.0	0.868 1.0 0.0	90.3 -33.6 88.0 94.3 110	0.733 1.0 0.0	0.868 1.0 0.0	0.733 1.0 0.0			
119	107	112	0.716 1.0 0.0	88.1 -47.8 85.4 97.9 119	0.933 1.0 0.0	91.5 -27.3 89.6 93.6 107	0.717 1.0 0.0	0.848 1.0 0.0	90.0 -35.6 87.8 94.7 112	0.717 1.0 0.0	0.848 1.0 0.0	0.717 1.0 0.0			
120	108	113	0.7 1.0 0.0	87.9 -49.2 85.2 98.4 120	0.917 1.0 0.0	91.2 -28.9 89.2 93.8 108	0.7 1.0 0.0	0.827 1.0 0.0	89.7 -37.5 87.4 95.2 113	0.7 1.0 0.0	0.827 1.0 0.0	0.7 1.0 0.0			
120	109	114	0.683 1.0 0.0	87.6 -50.7 84.9 98.9 120	0.901 1.0 0.0	90.9 -30.5 88.8 93.9 109	0.683 1.0 0.0	0.806 1.0 0.0	89.4 -39.5 87.1 95.7 114	0.683 1.0 0.0	0.806 1.0 0.0	0.683 1.0 0.0			
121	110	115	0.666 1.0 0.0	87.4 -52.1 84.7 99.4 121	0.884 1.0 0.0	90.6 -32.1 88.4 94.1 110	0.667 1.0 0.0	0.786 1.0 0.0	89.1 -41.5 86.7 96.1 115	0.667 1.0 0.0	0.786 1.0 0.0	0.667 1.0 0.0			
122	111	116	0.65 1.0 0.0	87.2 -53.6 84.4 100.0 122	0.868 1.0 0.0	90.3 -33.7 88.0 94.3 111	0.65 1.0 0.0	0.765 1.0 0.0	88.8 -43.4 86.2 96.6 116	0.65 1.0 0.0	0.765 1.0 0.0	0.65 1.0 0.0			
123	112	117	0.633 1.0 0.0	87.0 -55.0 84.1 100.5 123	0.85 1.0 0.0	90.1 -35.4 87.8 94.7 112	0.633 1.0 0.0	0.743 1.0 0.0	88.5 -45.4 85.8 97.1 117	0.633 1.0 0.0	0.743 1.0 0.0	0.633 1.0 0.0			
123	113	119	0.616 1.0 0.0	86.8 -56.4 83.8 101.0 123	0.832 1.0 0.0	89.8 -37.1 87.5 95.1 113	0.617 1.0 0.0	0.719 1.0 0.0	88.2 -47.5 85.5 97.9 119	0.617 1.0 0.0	0.719 1.0 0.0	0.617 1.0 0.0			
124	114	120	0.6 1.0 0.0	86.7 -57.6 83.7 101.6 124	0.814 1.0 0.0	89.5 -38.7 87.2 95.5 114	0.6 1.0 0.0	0.695 1.0 0.0	87.8 -49.6 85.2 98.6 120	0.6 1.0 0.0	0.695 1.0 0.0	0.6 1.0 0.0			
125	115	121	0.583 1.0 0.0	86.5 -58.9 83.5 102.2 125	0.797 1.0 0.0	89.3 -40.4 86.9 95.9 115	0.583 1.0 0.0	0.67 1.0 0.0	87.5 -51.7 84.8 99.4 121	0.583 1.0 0.0	0.67 1.0 0.0	0.583 1.0 0.0			
125	116	122	0.566 1.0 0.0	86.3 -60.1 83.3 102.8 125	0.779 1.0 0.0	89.0 -42.1 86.5 96.3 116	0.567 1.0 0.0	0.646 1.0 0.0	87.2 -53.9 84.4 100.1 122	0.567 1.0 0.0	0.646 1.0 0.0	0.567 1.0 0.0			
126	117	123	0.55 1.0 0.0	86.2 -61.4 83.1 103.3 126	0.761 1.0 0.0	88.7 -43.8 86.1 96.6 117	0.55 1.0 0.0	0.621 1.0 0.0	86.9 -56.0 83.9 100.9 123	0.55 1.0 0.0	0.621 1.0 0.0	0.55 1.0 0.0			
127	118	124	0.533 1.0 0.0	86.0 -62.7 82.9 103.9 127	0.742 1.0 0.0	88.4 -45.5 85.8 97.1 118	0.533 1.0 0.0	0.59 1.0 0.0	86.6 -58.3 83.6 102.0 124	0.533 1.0 0.0	0.59 1.0 0.0	0.533 1.0 0.0			
127	119	126	0.516 1.0 0.0	85.8 -63.9 82.6 104.5 127	0.721 1.0 0.0	88.2 -47.3 85.5 97.8 119	0.517 1.0 0.0	0.56 1.0 0.0	86.3 -60.6 83.3 103.1 126	0.517 1.0 0.0	0.56 1.0 0.0	0.517 1.0 0.0			
128	120	127	0.5 1.0 0.0	85.7 -65.2 82.4 105.1 128	0.7 1.0 0.0	87.9 -49.1 85.3 98.4 120	0.5 1.0 0.0	0.529 1.0 0.0	86.0 -62.9 82.9 104.1 127	0.5 1.0 0.0	0.529 1.0 0.0	0.5 1.0 0.0			

vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF> /PS
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /PS
la domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rh4ta

Data of Maximum color M in colorimetric system sRGB standard device; no separation, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM_s; h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;

Six hue angles of the device colours RYGBM_d; h_{ab,d} = 40.0, 102.9, 136.0, 196.4, 306.3, 328.2; Six hue angles of the elementary colours RYGBM_e; h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6

h _{ab,d}	h _{ab,s}	h _{ab,e}	rgb* dd361M	LAB* ddx361Mi (x=LabCh)	rgb* ds361Mi	LAB* dsx361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)																			
128	120	127	0.5	1.0	0.0	85.7	-65.2	82.4	105.1	128	0.7	1.0	0.0	87.9	-49.1	85.3	98.4	120	0.5	1.0	0.0	0.529	1.0	0.0	86.0	-62.9	82.9	104.1	127	0.5	1.0	0.0					
128	121	128	0.483	1.0	0.0	85.5	-66.2	82.3	105.6	128	0.68	1.0	0.0	87.7	-50.9	84.9	99.1	121	0.483	1.0	0.0	0.498	1.0	0.0	85.7	-65.3	82.4	105.2	128	0.483	1.0	0.0					
129	122	129	0.466	1.0	0.0	85.4	-67.2	82.1	106.1	129	0.659	1.0	0.0	87.4	-52.8	84.6	99.7	122	0.466	1.0	0.0	0.456	1.0	0.0	85.4	-67.8	82.1	106.5	129	0.466	1.0	0.0					
129	123	130	0.45	1.0	0.0	85.3	-68.2	82.0	106.7	129	0.638	1.0	0.0	87.1	-54.6	84.2	100.4	123	0.45	1.0	0.0	0.414	1.0	0.0	85.1	-70.3	81.7	107.9	130	0.45	1.0	0.0					
130	124	131	0.433	1.0	0.0	85.0	-69.2	81.8	107.2	130	0.615	1.0	0.0	86.9	-56.5	83.9	101.1	124	0.433	1.0	0.0	0.372	1.0	0.0	84.7	-72.9	81.3	109.2	131	0.433	1.0	0.0					
130	125	133	0.416	1.0	0.0	85.2	-70.2	81.7	107.8	130	0.589	1.0	0.0	86.6	-58.4	83.6	102.1	125	0.416	1.0	0.0	0.309	1.0	0.0	84.4	-75.6	80.9	110.8	133	0.416	1.0	0.0					
131	126	134	0.4	1.0	0.0	84.9	-71.3	81.5	108.3	131	0.562	1.0	0.0	86.3	-60.4	83.3	103.0	126	0.4	1.0	0.0	0.244	1.0	0.0	84.1	-78.3	80.5	112.4	134	0.4	1.0	0.0					
131	127	135	0.383	1.0	0.0	84.8	-72.3	81.3	108.8	131	0.536	1.0	0.0	86.1	-62.4	83.0	103.9	127	0.383	1.0	0.0	0.132	1.0	0.0	83.8	-81.2	80.1	114.1	135	0.383	1.0	0.0					
132	128	136	0.366	1.0	0.0	84.7	-73.2	81.2	109.3	132	0.51	1.0	0.0	85.8	-64.4	82.6	104.8	128	0.366	1.0	0.0	0.0	1.0	0.0	0.073	83.7	-82.3	78.0	113.5	136	0.366	1.0	0.0				
132	129	137	0.35	1.0	0.0	84.6	-73.9	81.1	109.7	132	0.477	1.0	0.0	85.5	-66.5	82.3	105.8	129	0.35	1.0	0.0	0.0	1.0	0.0	0.165	83.7	-81.6	74.2	110.4	137	0.35	1.0	0.0				
132	130	138	0.333	1.0	0.0	84.5	-74.6	81.0	110.1	132	0.442	1.0	0.0	85.3	-68.7	82.0	107.0	130	0.333	1.0	0.0	0.0	1.0	0.0	0.227	83.8	-80.8	70.5	107.3	138	0.333	1.0	0.0				
132	131	140	0.316	1.0	0.0	84.4	-75.3	80.9	110.6	132	0.406	1.0	0.0	85.0	-70.9	81.6	108.1	131	0.316	1.0	0.0	0.0	1.0	0.0	0.273	83.8	-80.0	67.0	104.5	140	0.316	1.0	0.0				
133	132	141	0.3	1.0	0.0	84.3	-76.0	80.8	111.0	133	0.368	1.0	0.0	84.7	-73.1	81.2	109.3	132	0.3	1.0	0.0	0.0	1.0	0.0	0.311	83.9	-79.3	63.7	101.8	141	0.3	1.0	0.0				
133	133	142	0.283	1.0	0.0	84.2	-76.8	80.7	111.4	133	0.314	1.0	0.0	84.5	-75.4	80.9	110.7	133	0.283	1.0	0.0	0.0	1.0	0.0	0.349	84.0	-78.4	60.4	99.0	142	0.283	1.0	0.0				
133	134	143	0.266	1.0	0.0	84.2	-77.5	80.6	111.8	133	0.261	1.0	0.0	84.2	-77.7	80.6	112.0	134	0.266	1.0	0.0	0.0	1.0	0.0	0.383	84.0	-77.5	57.3	96.4	143	0.266	1.0	0.0				
134	135	144	0.25	1.0	0.0	84.1	-78.2	80.5	112.2	134	0.173	1.0	0.0	83.9	-80.2	80.3	113.5	135	0.25	1.0	0.0	0.0	1.0	0.0	0.41	84.1	-76.8	54.3	94.1	144	0.25	1.0	0.0				
134	136	145	0.233	1.0	0.0	84.0	-78.7	80.4	112.5	134	0.004	1.0	0.0	83.6	-82.6	79.9	115.0	136	0.233	1.0	0.0	0.0	1.0	0.0	0.437	84.2	-75.9	51.5	91.8	145	0.233	1.0	0.0				
134	137	147	0.216	1.0	0.0	84.0	-79.1	80.4	112.8	134	0.0	1.0	0.0	0.125	83.7	-82.1	76.6	112.3	137	0.216	1.0	0.0	0.0	1.0	0.0	0.464	84.2	-75.0	48.7	89.5	147	0.216	1.0	0.0			
134	138	148	0.2	1.0	0.0	83.9	-79.5	80.3	113.0	134	0.0	1.0	0.0	0.178	83.7	-81.4	73.4	109.7	138	0.2	1.0	0.0	0.0	1.0	0.0	0.491	84.3	-74.1	45.9	87.2	148	0.2	1.0	0.0			
134	139	149	0.183	1.0	0.0	83.9	-79.9	80.2	113.3	134	0.0	1.0	0.0	0.231	83.8	-80.7	70.3	107.1	139	0.183	1.0	0.0	0.0	1.0	0.0	0.513	84.4	-73.3	43.4	85.2	149	0.183	1.0	0.0			
135	140	150	0.166	1.0	0.0	83.8	-80.4	80.2	113.5	135	0.0	1.0	0.0	0.271	83.8	-80.1	67.3	104.7	140	0.166	1.0	0.0	0.0	1.0	0.0	0.533	84.5	-72.5	41.0	83.4	150	0.166	1.0	0.0			
135	141	151	0.15	1.0	0.0	83.8	-80.8	80.1	113.8	135	0.0	1.0	0.0	0.303	83.9	-79.4	64.4	102.3	141	0.15	1.0	0.0	0.0	1.0	0.0	0.553	84.5	-71.7	38.6	81.6	151	0.15	1.0	0.0			
135	142	152	0.133	1.0	0.0	83.7	-81.2	80.1	114.1	135	0.0	1.0	0.0	0.335	83.9	-78.7	61.6	100.0	142	0.133	1.0	0.0	0.0	1.0	0.0	0.573	84.6	-70.9	36.3	79.8	152	0.133	1.0	0.0			
135	143	154	0.116	1.0	0.0	83.7	-81.5	80.0	114.2	135	0.0	1.0	0.0	0.368	84.0	-77.9	58.8	97.7	143	0.116	1.0	0.0	0.0	1.0	0.0	0.593	84.7	-70.0	34.1	77.9	154	0.116	1.0	0.0			
135	144	155	0.1	1.0	0.0	83.7	-81.7	80.0	114.4	135	0.0	1.0	0.0	0.393	84.1	-77.3	56.2	95.6	144	0.1	1.0	0.0	0.0	1.0	0.0	0.614	84.7	-69.0	31.9	76.1	155	0.1	1.0	0.0			
135	145	156	0.083	1.0	0.0	83.7	-81.9	80.0	114.5	135	0.0	1.0	0.0	0.416	84.1	-76.6	53.7	93.6	145	0.083	1.0	0.0	0.0	1.0	0.0	0.631	84.8	-68.2	29.8	74.5	156	0.083	1.0	0.0			
135	146	157	0.066	1.0	0.0	83.7	-82.0	79.9	114.6	135	0.0	1.0	0.0	0.439	84.2	-75.9	51.3	91.7	146	0.066	1.0	0.0	0.0	1.0	0.0	0.646	84.9	-67.5	27.9	73.2	157	0.066	1.0	0.0			
135	147	158	0.049	1.0	0.0	83.6	-82.2	79.9	114.7	135	0.0	1.0	0.0	0.462	84.2	-75.1	48.8	89.7	147	0.049	1.0	0.0	0.0	1.0	0.0	0.661	85.0	-66.9	26.1	71.9	158	0.049	1.0	0.0			
135	148	159	0.033	1.0	0.0	83.6	-82.4	79.9	114.8	135	0.0	1.0	0.0	0.485	84.3	-74.3	46.5	87.7	148	0.033	1.0	0.0	0.0	1.0	0.0	0.676	85.0	-66.2	24.3	70.6	159	0.033	1.0	0.0			
135	149	161	0.016	1.0	0.0	83.6	-82.6	79.9	114.9	135	0.0	1.0	0.0	0.506	84.4	-73.5	44.2	85.9	149	0.016	1.0	0.0	0.0	1.0	0.0	0.691	85.1	-65.4	22.5	69.2	161	0.016	1.0	0.0			
136	150	162	0.0	1.0	0.0	83.6	-82.7	79.8	115.0	136	G _d	0.0	1.0	0.0	0.523	84.4	-72.9	42.1	84.3	150	G _s	0.0	1.0	0.0	0.0	1.0	0.0	0.706	85.2	-64.6	20.7	67.9	162	G _e	0.0	1.0	0.0
136	151	163	0.0	1.0	0.016	83.6	-82.7	79.4	114.6	136	0.0	1.0	0.0	0.541	84.5	-72.3	40.1	82.7	151	0.0	1.0	0.017	0.0	1.0	0.0	0.718	85.2	-63.9	19.4	66.9	163	0.0	1.0	0.017			
136	152	164	0.0	1.0	0.033	83.6	-82.6	79.0	114.3	136	0.0	1.0	0.0	0.558	84.5	-71.6	38.1	81.2	152	0.0	1.0	0.033	0.0	1.0	0.0	0.73	85.3	-63.2	18.1	65.9	164	0.0	1.0	0.033			
136	153	164	0.0	1.0	0.05	83.6	-82.5	78.5	113.9	136	0.0	1.0	0.0	0.575	84.6	-70.8	36.1	79.6	153	0.0	1.0	0.05	0.0	1.0	0.0	0.741	85.3	-62.5	16.8	64.8	164	0.0	1.0	0.05			
136	154	165	0.0	1.0	0.066	83.6	-82.4	78.1	113.5	136	0.0	1.0	0.0	0.592	84.7	-70.0	34.2	78.0	154	0.0	1.0	0.067	0.0	1.0	0.0	0.752	85.4	-61.9	15.6	63.9	165	0.0	1.0	0.067			
136	155	166	0.0	1.0	0.083	83.6	-82.3	77.6	113.2	136	0.0	1.0	0.0	0.61	84.7	-69.2	32.3	76.5	155	0.0	1.0	0.083	0.0	1.0	0.0	0.761	85.4	-61.5	14.5	63.2	166	0.0	1.0	0.083			
136	156	167	0.0	1.0	0.1	83.6	-82.2	77.2	112.8	136	0.0	1.0	0.0	0.629	84.8	-68.4	30.3	74.9	156	0.0	1.0	0.1	0.0	1.0	0.0	0.77	85.5	-61.1	13.3	62.6	167	0.0	1.0	0.1			
136	157	168	0.0	1.0	0.116	83.6	-82.1	76.8	112.5	136	0.0	1.0	0.0	0.639	84.9	-67.8	28.8	73.8	157	0.0	1.0	0.117	0.0	1.0	0.0	0.778	85.5	-60.6	12.2	61.9	168	0.0	1.0	0.117			
137	158	169	0.0	1.0	0.133</																																

Data of Maximum color M in colorimetric system sRGB standard device; no separation, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM_s; h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;
Six hue angles of the device colours RYGBM_d; h_{ab,d} = 40.0, 102.9, 136.0, 196.4, 306.3, 328.2; Six hue angles of the elementary colours RYGBM_e; h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6

h _{ab,d}	h _{ab,s}	h _{ab,e}	rgb* dd361M	LAB* d361Mi (x=LabCh)	rgb* ds361Mi	LAB* ds361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)	rgb* dd361Mi	rgb* dd361Mi	rgb* ds361Mi	rgb* ds361Mi	rgb* ds361Mi																			
196	210	216	0.0	1.0	1.0	86.8	-46.1	-13.5	48.1	196	C _d	0.0	0.922	1.0	81.7	-38.6	-22.2	44.7	210	C _s	0.0	0.983	1.0	0.0	0.885	1.0	78.7	-33.6	-26.1	42.7	217	0.0	0.983	1.0
199	211	217	0.0	0.983	1.0	85.6	-44.6	-15.8	47.3	199		0.0	0.922	1.0	81.3	-38.0	-22.8	44.4	211		0.0	0.983	1.0	0.0	0.885	1.0	78.7	-33.6	-26.1	42.7	217	0.0	0.983	1.0
202	212	218	0.0	0.966	1.0	84.5	-42.9	-17.9	46.5	202		0.0	0.917	1.0	81.0	-37.3	-23.3	44.2	212		0.0	0.967	1.0	0.0	0.881	1.0	78.4	-33.0	-26.5	42.4	218	0.0	0.967	1.0
205	213	219	0.0	0.95	1.0	83.3	-41.1	-19.8	45.7	205		0.0	0.911	1.0	80.6	-36.7	-23.8	43.9	213		0.0	0.95	1.0	0.0	0.876	1.0	78.0	-32.3	-26.9	42.2	219	0.0	0.95	1.0
208	214	220	0.0	0.933	1.0	82.1	-39.3	-21.7	44.9	208		0.0	0.906	1.0	80.2	-36.1	-24.3	43.6	214		0.0	0.933	1.0	0.0	0.871	1.0	77.7	-31.9	-27.4	42.2	220	0.0	0.933	1.0
212	215	221	0.0	0.916	1.0	80.9	-37.4	-23.4	44.1	212		0.0	0.901	1.0	79.8	-35.4	-24.8	43.4	215		0.0	0.917	1.0	0.0	0.867	1.0	77.4	-31.5	-27.9	42.3	221	0.0	0.917	1.0
215	216	222	0.0	0.9	1.0	79.7	-35.4	-24.9	43.3	215		0.0	0.895	1.0	79.5	-34.8	-25.3	43.1	216		0.0	0.9	1.0	0.0	0.863	1.0	77.2	-31.1	-28.5	42.3	222	0.0	0.9	1.0
218	217	223	0.0	0.883	1.0	78.5	-33.4	-26.3	42.5	218		0.0	0.89	1.0	79.1	-34.1	-25.7	42.9	217		0.0	0.883	1.0	0.0	0.859	1.0	76.9	-30.7	-29.0	42.4	223	0.0	0.883	1.0
221	218	224	0.0	0.866	1.0	77.4	-31.5	-28.1	42.2	221		0.0	0.885	1.0	78.7	-33.5	-26.1	42.6	218		0.0	0.867	1.0	0.0	0.855	1.0	76.6	-30.3	-29.6	42.5	224	0.0	0.867	1.0
225	219	225	0.0	0.85	1.0	76.2	-29.9	-30.2	42.5	225		0.0	0.879	1.0	78.3	-32.8	-26.6	42.4	219		0.0	0.85	1.0	0.0	0.851	1.0	76.3	-29.9	-30.1	42.6	225	0.0	0.85	1.0
228	220	226	0.0	0.833	1.0	75.0	-28.1	-32.3	42.8	228		0.0	0.874	1.0	77.9	-32.2	-27.0	42.2	220		0.0	0.833	1.0	0.0	0.846	1.0	76.0	-29.4	-30.6	42.6	226	0.0	0.833	1.0
232	221	227	0.0	0.816	1.0	73.8	-26.1	-34.2	43.1	232		0.0	0.87	1.0	77.6	-31.8	-27.6	42.2	221		0.0	0.817	1.0	0.0	0.842	1.0	75.7	-29.0	-31.1	42.7	227	0.0	0.817	1.0
236	222	227	0.0	0.8	1.0	72.6	-24.0	-36.0	43.3	236		0.0	0.865	1.0	77.3	-31.3	-28.2	42.3	222		0.0	0.8	1.0	0.0	0.838	1.0	75.4	-28.5	-31.6	42.8	227	0.0	0.8	1.0
239	223	228	0.0	0.783	1.0	71.4	-21.8	-37.7	43.6	239		0.0	0.861	1.0	77.0	-30.9	-28.8	42.4	223		0.0	0.783	1.0	0.0	0.834	1.0	75.1	-28.1	-32.1	42.8	228	0.0	0.783	1.0
243	224	229	0.0	0.766	1.0	70.2	-19.5	-39.3	43.9	243		0.0	0.856	1.0	76.7	-30.4	-29.4	42.5	224		0.0	0.767	1.0	0.0	0.83	1.0	74.8	-27.6	-32.6	42.9	229	0.0	0.767	1.0
247	225	230	0.0	0.75	1.0	69.1	-17.0	-40.7	44.1	247		0.0	0.851	1.0	76.3	-30.0	-30.0	42.5	225		0.0	0.75	1.0	0.0	0.826	1.0	74.5	-27.1	-33.1	43.0	230	0.0	0.75	1.0
250	226	231	0.0	0.733	1.0	67.9	-15.3	-42.9	45.5	250		0.0	0.847	1.0	76.0	-29.5	-30.6	42.6	226		0.0	0.733	1.0	0.0	0.821	1.0	74.2	-26.6	-33.6	43.0	231	0.0	0.733	1.0
253	227	232	0.0	0.716	1.0	66.7	-13.5	-44.9	46.9	253		0.0	0.842	1.0	75.7	-29.0	-31.1	42.7	227		0.0	0.717	1.0	0.0	0.817	1.0	73.9	-26.1	-34.1	43.1	232	0.0	0.717	1.0
256	228	233	0.0	0.7	1.0	65.5	-11.4	-46.9	48.3	256		0.0	0.838	1.0	75.4	-28.5	-31.7	42.8	228		0.0	0.7	1.0	0.0	0.813	1.0	73.6	-25.6	-34.6	43.2	233	0.0	0.7	1.0
259	229	234	0.0	0.683	1.0	64.4	-9.2	-48.8	49.7	259		0.0	0.833	1.0	75.0	-28.0	-32.2	42.8	229		0.0	0.683	1.0	0.0	0.809	1.0	73.3	-25.1	-35.0	43.2	234	0.0	0.683	1.0
262	230	235	0.0	0.666	1.0	63.2	-6.8	-50.6	51.1	262		0.0	0.829	1.0	74.7	-27.5	-32.8	42.9	230		0.0	0.667	1.0	0.0	0.805	1.0	73.0	-24.6	-35.5	43.3	235	0.0	0.667	1.0
265	231	236	0.0	0.65	1.0	62.0	-4.2	-52.3	52.5	265		0.0	0.824	1.0	74.4	-26.9	-33.3	43.0	231		0.0	0.65	1.0	0.0	0.801	1.0	72.7	-24.1	-35.9	43.4	236	0.0	0.65	1.0
268	232	237	0.0	0.633	1.0	60.9	-1.5	-53.9	53.9	268		0.0	0.82	1.0	74.1	-26.4	-33.8	43.1	232		0.0	0.633	1.0	0.0	0.797	1.0	72.4	-23.5	-36.3	43.4	237	0.0	0.633	1.0
270	233	237	0.0	0.616	1.0	59.7	0.8	-55.6	55.7	270		0.0	0.815	1.0	73.7	-25.9	-34.3	43.1	233		0.0	0.617	1.0	0.0	0.792	1.0	72.1	-23.0	-36.8	43.5	237	0.0	0.617	1.0
272	234	238	0.0	0.6	1.0	58.6	2.9	-57.7	57.8	272		0.0	0.81	1.0	73.4	-25.3	-34.9	43.2	234		0.0	0.6	1.0	0.0	0.788	1.0	71.8	-22.4	-37.2	43.6	238	0.0	0.6	1.0
274	235	239	0.0	0.583	1.0	57.4	5.1	-59.7	59.9	274		0.0	0.806	1.0	73.1	-24.7	-35.4	43.3	235		0.0	0.583	1.0	0.0	0.784	1.0	71.5	-21.8	-37.6	43.6	239	0.0	0.583	1.0
276	236	240	0.0	0.566	1.0	56.3	7.4	-61.6	62.1	276		0.0	0.801	1.0	72.8	-24.1	-35.8	43.4	236		0.0	0.567	1.0	0.0	0.78	1.0	71.2	-21.3	-38.0	43.7	240	0.0	0.567	1.0
278	237	241	0.0	0.55	1.0	55.2	10.0	-63.5	64.2	278		0.0	0.797	1.0	72.4	-23.6	-36.3	43.4	237		0.0	0.55	1.0	0.0	0.776	1.0	70.9	-20.7	-38.4	43.8	241	0.0	0.55	1.0
280	238	242	0.0	0.533	1.0	54.0	12.6	-65.2	66.4	280		0.0	0.792	1.0	72.1	-23.0	-36.8	43.5	238		0.0	0.533	1.0	0.0	0.772	1.0	70.6	-20.1	-38.8	43.8	242	0.0	0.533	1.0
283	239	243	0.0	0.516	1.0	52.9	15.4	-66.8	68.5	283		0.0	0.788	1.0	71.8	-22.3	-37.2	43.6	239		0.0	0.517	1.0	0.0	0.767	1.0	70.3	-19.5	-39.2	43.9	243	0.0	0.517	1.0
285	240	244	0.0	0.5	1.0	51.7	18.3	-68.3	70.7	285		0.0	0.783	1.0	71.5	-21.7	-37.7	43.6	240		0.0	0.5	1.0	0.0	0.763	1.0	70.1	-18.9	-39.5	44.0	244	0.0	0.5	1.0
286	241	245	0.0	0.483	1.0	50.7	20.6	-70.2	73.2	286		0.0	0.779	1.0	71.1	-21.1	-38.1	43.7	241		0.0	0.483	1.0	0.0	0.759	1.0	69.8	-18.3	-39.9	44.0	245	0.0	0.483	1.0
287	242	246	0.0	0.466	1.0	49.6	22.9	-72.1	75.7	287		0.0	0.774	1.0	70.8	-20.5	-38.6	43.8	242		0.0	0.467	1.0	0.0	0.755	1.0	69.5	-17.7	-40.2	44.1	246	0.0	0.467	1.0
288	243	247	0.0	0.45	1.0	48.6	25.4	-74.0	78.2	288		0.0	0.769	1.0	70.5	-19.8	-39.0	43.9	243		0.0	0.45	1.0	0.0	0.751	1.0	69.2	-17.1	-40.6	44.2	247	0.0	0.45	1.0
290	244	248	0.0	0.433	1.0	47.5	28.0	-75.7	80.7	290		0.0	0.765	1.0	70.2	-19.2	-39.4	43.9	244		0.0	0.433	1.0	0.0	0.746	1.0	68.8	-16.6	-41.2	44.5	248	0.0	0.433	1.0
291	245	248	0.0	0.416	1.0	46.5	30.6	-77.4	83.2	291		0.0	0.76	1.0	69.8	-18.5	-39.8	44.0	245		0.0	0.417	1.0	0.0	0.741	1.0	68.5	-16.1	-41.8	45.0	248	0.0	0.417	1.0
292	246	249	0.0	0.4	1.0	45.4	33.3	-79.0	85.7	292		0.0	0.756	1.0	69.5	-17.8	-40.2	44.1	246		0.0	0.4	1.0	0.0	0.736	1.0	68.1	-15.5	-42.5	45.4	249	0.0	0.4	1.0
294	247	250	0.0	0.383	1.0	44.3	36.2	-80.5	88.2	294		0.0	0.751	1.0	69.2	-17.2	-40.6	44.2	247		0.0	0.383	1.0	0.0	0.731	1.0	67.8	-15.0	-43.1	45.8	250	0.0	0.383	1.0
295	248	251	0.0	0.366	1.0	43.4	38.7	-82.0	90.7	295		0.0	0.746	1.0	68.8	-16.6	-41.2	44.5	248		0.0	0.367	1.0	0.0	0.726	1.0	67.4	-14.4	-43.8	46.2	251	0.0	0.367	1.0
296	249	252	0.0	0.35	1.0</																													

Data of Maximum color M in colorimetric system sRGB standard device; no separation, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM_s: h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;
Six hue angles of the device colours RYGBM_d: h_{ab,d} = 40.0, 102.9, 136.0, 196.4, 306.3, 328.2; Six hue angles of the elementary colours RYGBM_e: h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6

h _{ab,d}	h _{ab,s}	h _{ab,e}	rgb [*] _{dd361M}	LAB [*] _{dsx361Mi (x=LabCh)}	rgb [*] _{ds361Mi}	LAB [*] _{dsx361Mi (x=LabCh)}	rgb [*] _{dd361Mi}	LAB [*] _{de361Mi (x=LabCh)}	rgb [*] _{de361Mi}	LAB [*] _{de361Mi (x=LabCh)}	rgb [*] _{dd361Mi}	rgb [*] _{dd361Mi}	rgb [*] _{ds361Mi}	rgb [*] _{de361Mi}																					
301	255	258	0.0	0.25	1.0	37.1	55.9	-92.3	107.9	301	0.0	0.702	1.0	65.7	-11.6	-46.7	48.2	256	0.0	0.233	1.0	0.0	0.685	1.0	64.6	-9.4	-48.6	49.6	258	0.0	0.233	1.0			
302	257	259	0.0	0.216	1.0	35.9	59.4	-94.5	111.6	302	0.0	0.696	1.0	65.3	-10.9	-47.3	48.7	257	0.0	0.217	1.0	0.0	0.68	1.0	64.2	-8.7	-49.1	50.0	259	0.0	0.217	1.0			
302	258	260	0.0	0.2	1.0	35.2	61.2	-95.5	113.5	302	0.0	0.691	1.0	64.9	-10.1	-48.0	49.1	258	0.0	0.2	1.0	0.0	0.675	1.0	63.8	-8.0	-49.7	50.4	260	0.0	0.2	1.0			
303	259	261	0.0	0.183	1.0	34.6	63.0	-96.6	115.3	303	0.0	0.685	1.0	64.5	-9.4	-48.6	49.6	259	0.0	0.183	1.0	0.0	0.67	1.0	63.5	-7.2	-50.2	50.9	261	0.0	0.183	1.0			
303	260	262	0.0	0.166	1.0	34.0	64.8	-97.6	117.2	303	0.0	0.679	1.0	64.2	-8.6	-49.2	50.1	260	0.0	0.167	1.0	0.0	0.665	1.0	63.1	-6.5	-50.8	51.3	262	0.0	0.167	1.0			
304	261	263	0.0	0.15	1.0	33.4	66.7	-98.6	119.1	304	0.0	0.674	1.0	63.8	-7.8	-49.8	50.5	261	0.0	0.15	1.0	0.0	0.66	1.0	62.8	-5.7	-51.3	51.7	263	0.0	0.15	1.0			
304	262	264	0.0	0.133	1.0	32.8	68.6	-99.6	120.9	304	0.0	0.668	1.0	63.4	-7.0	-50.4	51.0	262	0.0	0.133	1.0	0.0	0.655	1.0	62.4	-5.0	-51.8	52.1	264	0.0	0.133	1.0			
304	263	265	0.0	0.116	1.0	32.3	70.0	-100.3	122.3	304	0.0	0.663	1.0	63.0	-6.2	-51.0	51.5	263	0.0	0.117	1.0	0.0	0.65	1.0	62.1	-4.2	-52.3	52.5	265	0.0	0.117	1.0			
305	264	266	0.0	0.1	1.0	32.0	70.8	-100.8	123.2	305	0.0	0.657	1.0	62.6	-5.3	-51.5	51.9	264	0.0	0.1	1.0	0.0	0.645	1.0	61.7	-3.4	-52.8	53.0	266	0.0	0.1	1.0			
305	265	267	0.0	0.083	1.0	31.7	71.7	-101.2	124.1	305	0.0	0.652	1.0	62.2	-4.5	-52.1	52.4	265	0.0	0.083	1.0	0.0	0.64	1.0	61.4	-2.5	-53.2	53.4	267	0.0	0.083	1.0			
305	266	268	0.0	0.066	1.0	31.5	72.5	-101.7	124.9	305	0.0	0.646	1.0	61.8	-3.6	-52.6	52.8	266	0.0	0.067	1.0	0.0	0.635	1.0	61.0	-1.7	-53.7	53.8	268	0.0	0.067	1.0			
305	267	269	0.0	0.049	1.0	31.2	73.4	-102.2	125.8	305	0.0	0.641	1.0	61.4	-2.7	-53.1	53.3	267	0.0	0.05	1.0	0.0	0.63	1.0	60.6	-0.8	-54.1	54.2	269	0.0	0.05	1.0			
305	268	269	0.0	0.033	1.0	30.9	74.3	-102.6	126.7	305	0.0	0.635	1.0	61.0	-1.8	-53.6	53.8	268	0.0	0.033	1.0	0.0	0.624	1.0	60.3	0.0	-54.6	54.7	269	0.0	0.033	1.0			
306	269	270	0.0	0.016	1.0	30.6	75.1	-103.1	127.6	306	0.0	0.63	1.0	60.6	-0.8	-54.1	54.2	269	0.0	0.017	1.0	0.0	0.617	1.0	59.8	0.8	-55.6	55.7	270	0.0	0.017	1.0			
306	270	271	0.0	0.0	1.0	30.3	76.0	-103.5	128.5	306	B _d	0.0	0.624	1.0	60.2	0.0	-54.7	54.8	270	B _s	0.0	0.0	1.0	0.0	0.609	1.0	59.3	1.7	-56.5	56.6	271	B _e	0.0	0.0	1.0
306	271	272	0.016	0.0	1.0	30.4	76.0	-103.4	128.4	306	0.0	0.615	1.0	59.7	1.0	-55.7	55.9	271	0.0	0.017	0.0	1.0	0.0	0.602	1.0	58.7	2.7	-57.5	57.6	272	0.0	0.017	0.0	1.0	
306	272	273	0.033	0.0	1.0	30.5	76.1	-103.3	128.3	306	0.0	0.607	1.0	59.1	2.0	-56.8	56.9	272	0.033	0.0	1.0	0.0	0.594	1.0	58.2	3.7	-58.4	58.6	273	0.033	0.0	1.0			
306	273	274	0.05	0.0	1.0	30.6	76.1	-103.1	128.2	306	0.0	0.599	1.0	58.5	3.0	-57.8	58.0	273	0.05	0.0	1.0	0.0	0.586	1.0	57.7	4.8	-59.4	59.7	274	0.05	0.0	1.0			
306	274	275	0.066	0.0	1.0	30.7	76.1	-103.0	128.1	306	0.0	0.591	1.0	58.0	4.1	-58.8	59.0	274	0.067	0.0	1.0	0.0	0.578	1.0	57.1	5.8	-60.3	60.7	275	0.067	0.0	1.0			
306	275	276	0.083	0.0	1.0	30.8	76.2	-102.8	128.0	306	0.0	0.583	1.0	57.4	5.2	-59.8	60.1	275	0.083	0.0	1.0	0.0	0.57	1.0	56.6	7.0	-61.2	61.7	276	0.083	0.0	1.0			
306	276	277	0.1	0.0	1.0	30.9	76.2	-102.7	127.9	306	0.0	0.574	1.0	56.9	6.4	-60.7	61.2	276	0.1	0.0	1.0	0.0	0.563	1.0	56.1	8.1	-62.0	62.7	277	0.1	0.0	1.0			
306	277	278	0.116	0.0	1.0	30.9	76.2	-102.5	127.8	306	0.0	0.566	1.0	56.3	7.6	-61.7	62.2	277	0.117	0.0	1.0	0.0	0.555	1.0	55.5	9.3	-62.9	63.7	278	0.117	0.0	1.0			
306	278	279	0.133	0.0	1.0	31.1	76.3	-102.3	127.6	306	0.0	0.558	1.0	55.7	8.8	-62.6	63.3	278	0.133	0.0	1.0	0.0	0.547	1.0	55.0	10.5	-63.7	64.7	279	0.133	0.0	1.0			
306	279	280	0.15	0.0	1.0	31.3	76.3	-101.9	127.4	306	0.0	0.55	1.0	55.2	10.1	-63.5	64.3	279	0.15	0.0	1.0	0.0	0.539	1.0	54.5	11.7	-64.5	65.7	280	0.15	0.0	1.0			
306	280	281	0.166	0.0	1.0	31.5	76.4	-101.6	127.1	306	0.0	0.541	1.0	54.6	11.4	-64.3	65.4	280	0.167	0.0	1.0	0.0	0.531	1.0	53.9	13.0	-65.3	66.7	281	0.167	0.0	1.0			
307	281	282	0.183	0.0	1.0	31.7	76.5	-101.2	126.9	307	0.0	0.533	1.0	54.1	12.7	-65.1	66.5	281	0.183	0.0	1.0	0.0	0.524	1.0	53.4	14.3	-66.1	67.7	282	0.183	0.0	1.0			
307	282	283	0.2	0.0	1.0	31.9	76.6	-100.9	126.7	307	0.0	0.525	1.0	53.5	14.0	-66.0	67.5	282	0.2	0.0	1.0	0.0	0.516	1.0	52.9	15.6	-66.8	68.7	283	0.2	0.0	1.0			
307	283	284	0.216	0.0	1.0	32.1	76.6	-100.5	126.4	307	0.0	0.517	1.0	52.9	15.4	-66.7	68.6	283	0.217	0.0	1.0	0.0	0.508	1.0	52.3	16.9	-67.5	69.7	284	0.217	0.0	1.0			
307	284	285	0.233	0.0	1.0	32.3	76.7	-100.1	126.2	307	0.0	0.508	1.0	52.4	16.9	-67.5	69.7	284	0.233	0.0	1.0	0.0	0.5	1.0	51.8	18.3	-68.2	70.7	285	0.233	0.0	1.0			
307	285	285	0.25	0.0	1.0	32.6	76.8	-99.8	125.9	307	0.0	0.5	1.0	51.8	18.3	-68.2	70.7	285	0.25	0.0	1.0	0.0	0.488	1.0	51.0	19.9	-69.6	72.5	285	0.25	0.0	1.0			
307	286	286	0.266	0.0	1.0	32.9	77.0	-99.2	125.6	307	0.0	0.488	1.0	51.0	20.0	-69.7	72.6	286	0.267	0.0	1.0	0.0	0.476	1.0	50.3	21.6	-71.0	74.3	286	0.267	0.0	1.0			
308	287	287	0.283	0.0	1.0	33.2	77.1	-98.6	125.2	308	0.0	0.475	1.0	50.2	21.8	-71.2	74.5	287	0.283	0.0	1.0	0.0	0.464	1.0	49.5	23.3	-72.4	76.1	287	0.283	0.0	1.0			
308	288	288	0.3	0.0	1.0	33.6	77.3	-98.1	124.9	308	0.0	0.462	1.0	49.4	23.6	-72.6	76.4	288	0.3	0.0	1.0	0.0	0.452	1.0	48.8	25.1	-73.7	77.9	288	0.3	0.0	1.0			
308	289	289	0.316	0.0	1.0	33.9	77.4	-97.5	124.5	308	0.0	0.45	1.0	48.6	25.5	-74.0	78.3	289	0.317	0.0	1.0	0.0	0.44	1.0	48.0	26.9	-75.0	79.8	289	0.317	0.0	1.0			
308	290	290	0.333	0.0	1.0	34.3	77.6	-96.9	124.1	308	0.0	0.437	1.0	47.8	27.4	-75.3	80.2	290	0.333	0.0	1.0	0.0	0.428	1.0	47.2	28.8	-76.8	81.6	290	0.333	0.0	1.0			
308	291	291	0.35	0.0	1.0	34.6	77.7	-96.3	123.8	308	0.0	0.424	1.0	47.0	29.4	-76.6	82.1	291	0.35	0.0	1.0	0.0	0.416	1.0	46.5	30.7	-77.4	83.4	291	0.35	0.0	1.0			
309	292	292	0.366	0.0	1.0	34.9	77.9	-95.7	123.4	309	0.0	0.412	1.0	46.2	31.5	-77.8	84.1	292	0.367	0.0	1.0	0.0	0.404	1.0	45.7	32.7	-78.5	85.2	292	0.367	0.0	1.0			
309	293	293	0.383	0.0	1.0	35.3	78.1	-95.1	123.0	309	0.0	0.399	1.0	45.4	33.6	-79.0	86.0	293	0.383	0.0	1.0	0.0	0.392	1.0	44.9	34.7	-79.7	87.0	293	0.383	0.0	1.0			
309	294	294	0.4	0.0	1.0	35.8	78.3	-94.3	122.6	309	0.0	0.386	1.0	44.6	35.7	-80.2	87.9	294	0.4	0.0	1.0	0.0	0.38	1.0	44.2	36.8	-80.7	88.8	294	0.4	0.0	1.0			
310	295	295	0.416	0.0	1.0	36.3	78.6	-93.5	122.2	310	0.0	0.373	1.0	43.7	38.0	-81.4	89.9	295	0.417	0.0	1.0	0.0	0.364	1.0	43.3	39.2	-82.2	91.2	295	0.					

Data of Maximum color M in colorimetric system sRGB standard device; no separation, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM_s: h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;
Six hue angles of the device colours RYGBCM_d: h_{ab,d} = 40.0, 102.9, 136.0, 196.4, 306.3, 328.2; Six hue angles of the elementary colours RYGBM_e: h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6

h _{ab,d}	h _{ab,s}	h _{ab,e}	rgb* dd361M	LAB* dxx361Mi (x=LabCh)	rgb* ds361Mi	LAB* dsx361Mi (x=LabCh)	rgb* dd361Mi	LAB* de361Mi	rgb* dex361Mi (x=LabCh)	rgb* dd361Mi	LAB* de361Mi	rgb* dd361Mi	rgb* dd361Mi	rgb* ds361Mi	rgb* de361Mi																				
311	300	300	0.5	0.0	1.0	38.5	79.8	-89.7	120.0	311	0.0	0.274	1.0	38.4	52.2	-90.4	104.5	300	0.5	0.0	1.0	0.0	0.27	1.0	38.2	52.8	-90.6	105.0	300	0.5	0.0	1.0			
312	301	301	0.516	0.0	1.0	39.1	80.2	-88.7	119.6	312	0.0	0.254	1.0	37.4	55.3	-91.9	107.4	301	0.517	0.0	1.0	0.0	0.251	1.0	37.2	55.7	-92.1	107.7	301	0.517	0.0	1.0			
312	302	302	0.533	0.0	1.0	39.6	80.6	-87.8	119.2	312	0.0	0.222	1.0	36.1	58.8	-94.1	111.0	302	0.533	0.0	1.0	0.0	0.22	1.0	36.0	59.1	-94.2	111.3	302	0.533	0.0	1.0			
312	303	303	0.55	0.0	1.0	40.2	80.9	-86.9	118.8	312	0.0	0.188	1.0	34.8	62.6	-96.3	114.9	303	0.55	0.0	1.0	0.0	0.187	1.0	34.8	62.6	-96.3	115.0	303	0.55	0.0	1.0			
313	304	304	0.566	0.0	1.0	40.7	81.3	-86.0	118.3	313	0.0	0.153	1.0	33.5	66.4	-98.4	118.8	304	0.567	0.0	1.0	0.0	0.154	1.0	33.6	66.3	-98.3	118.6	304	0.567	0.0	1.0			
313	305	305	0.583	0.0	1.0	41.3	81.6	-85.1	117.9	313	0.0	0.109	1.0	32.2	70.4	-100.4	122.7	305	0.583	0.0	1.0	0.0	0.117	1.0	32.4	70.0	-100.2	122.3	304	0.583	0.0	1.0			
314	306	305	0.6	0.0	1.0	41.8	82.0	-84.1	117.5	314	0.0	0.024	1.0	30.8	74.8	-102.8	127.2	306	0.6	0.0	1.0	0.0	0.036	1.0	31.0	74.2	-102.5	126.6	305	0.6	0.0	1.0			
314	307	306	0.616	0.0	1.0	42.4	82.3	-83.2	117.0	314	0.172	0.0	1.0	31.6	76.5	-101.4	127.1	307	0.617	0.0	1.0	0.146	0.0	1.0	31.3	76.4	-102.0	127.5	306	0.617	0.0	1.0			
315	308	307	0.633	0.0	1.0	43.0	82.7	-82.2	116.6	315	0.282	0.0	1.0	33.2	77.2	-98.6	125.3	308	0.633	0.0	1.0	0.263	0.0	1.0	32.9	77.0	-99.3	125.7	307	0.633	0.0	1.0			
315	309	308	0.65	0.0	1.0	43.6	83.2	-81.2	116.3	315	0.357	0.0	1.0	34.8	77.8	-96.0	123.7	309	0.65	0.0	1.0	0.335	0.0	1.0	34.3	77.6	-96.8	124.2	308	0.65	0.0	1.0			
316	310	309	0.666	0.0	1.0	44.2	83.7	-80.2	115.9	316	0.414	0.0	1.0	36.2	78.6	-93.6	122.3	310	0.667	0.0	1.0	0.396	0.0	1.0	35.8	78.3	-94.4	122.8	309	0.667	0.0	1.0			
316	311	310	0.683	0.0	1.0	44.8	84.1	-79.2	115.5	316	0.465	0.0	1.0	37.6	79.4	-91.2	121.0	311	0.683	0.0	1.0	0.445	0.0	1.0	37.1	79.1	-92.2	121.5	310	0.683	0.0	1.0			
317	312	311	0.7	0.0	1.0	45.4	84.6	-78.1	115.2	317	0.513	0.0	1.0	39.0	80.1	-88.9	119.8	312	0.7	0.0	1.0	0.493	0.0	1.0	38.4	79.8	-89.9	120.3	311	0.7	0.0	1.0			
317	313	312	0.716	0.0	1.0	46.0	85.0	-77.1	114.8	317	0.551	0.0	1.0	40.3	81.0	-86.8	118.8	313	0.717	0.0	1.0	0.532	0.0	1.0	39.6	80.6	-87.9	119.3	312	0.717	0.0	1.0			
318	314	313	0.733	0.0	1.0	46.6	85.4	-76.1	114.4	318	0.59	0.0	1.0	41.6	81.8	-84.6	117.8	314	0.733	0.0	1.0	0.569	0.0	1.0	40.8	81.4	-85.8	118.3	313	0.733	0.0	1.0			
318	315	314	0.75	0.0	1.0	47.2	85.8	-75.1	114.0	318	0.628	0.0	1.0	42.8	82.6	-82.5	116.8	315	0.75	0.0	1.0	0.605	0.0	1.0	42.1	82.1	-83.8	117.4	314	0.75	0.0	1.0			
319	316	315	0.766	0.0	1.0	47.9	86.4	-74.0	113.8	319	0.66	0.0	1.0	44.0	83.5	-80.6	116.1	316	0.767	0.0	1.0	0.639	0.0	1.0	43.2	82.9	-81.8	116.6	315	0.767	0.0	1.0			
320	317	316	0.783	0.0	1.0	48.5	87.0	-72.9	113.5	320	0.692	0.0	1.0	45.2	84.4	-78.6	115.4	317	0.783	0.0	1.0	0.669	0.0	1.0	44.3	83.8	-80.0	115.9	316	0.783	0.0	1.0			
320	318	317	0.8	0.0	1.0	49.2	87.5	-71.8	113.2	320	0.724	0.0	1.0	46.3	85.2	-76.6	114.7	318	0.8	0.0	1.0	0.699	0.0	1.0	45.4	84.6	-78.1	115.2	317	0.8	0.0	1.0			
321	319	318	0.816	0.0	1.0	49.8	88.1	-70.7	113.0	321	0.755	0.0	1.0	47.5	86.0	-74.7	114.0	319	0.817	0.0	1.0	0.729	0.0	1.0	46.5	85.4	-76.3	114.5	318	0.817	0.0	1.0			
321	320	319	0.833	0.0	1.0	50.5	88.6	-69.6	112.7	321	0.783	0.0	1.0	48.6	87.0	-72.9	113.6	320	0.833	0.0	1.0	0.758	0.0	1.0	47.6	86.2	-74.5	114.0	319	0.833	0.0	1.0			
322	321	320	0.85	0.0	1.0	51.2	89.1	-68.5	112.4	322	0.81	0.0	1.0	49.7	87.9	-71.1	113.1	321	0.85	0.0	1.0	0.785	0.0	1.0	48.6	87.1	-72.8	113.5	320	0.85	0.0	1.0			
323	322	321	0.866	0.0	1.0	51.8	89.6	-67.4	112.1	323	0.838	0.0	1.0	50.7	88.8	-69.3	112.7	322	0.867	0.0	1.0	0.811	0.0	1.0	49.7	87.9	-71.0	113.1	321	0.867	0.0	1.0			
323	323	321	0.883	0.0	1.0	52.5	90.1	-66.3	111.9	323	0.866	0.0	1.0	51.8	89.6	-67.4	112.2	323	0.883	0.0	1.0	0.837	0.0	1.0	50.7	88.8	-69.3	112.7	321	0.883	0.0	1.0			
324	324	322	0.9	0.0	1.0	53.2	90.8	-65.2	111.8	324	0.892	0.0	1.0	52.9	90.5	-65.7	111.9	324	0.9	0.0	1.0	0.864	0.0	1.0	51.7	89.5	-67.6	112.2	322	0.9	0.0	1.0			
324	325	323	0.916	0.0	1.0	53.8	91.4	-64.1	111.6	324	0.918	0.0	1.0	53.9	91.5	-64.0	111.7	325	0.917	0.0	1.0	0.889	0.0	1.0	52.8	90.4	-65.9	111.9	323	0.917	0.0	1.0			
325	326	324	0.933	0.0	1.0	54.5	92.0	-62.9	111.5	325	0.943	0.0	1.0	55.0	92.4	-62.2	111.5	326	0.933	0.0	1.0	0.913	0.0	1.0	53.7	91.3	-64.3	111.7	324	0.933	0.0	1.0			
326	327	325	0.95	0.0	1.0	55.2	92.6	-61.8	111.4	326	0.969	0.0	1.0	56.0	93.3	-60.5	111.3	327	0.95	0.0	1.0	0.937	0.0	1.0	54.7	92.2	-62.6	111.5	325	0.95	0.0	1.0			
326	328	326	0.966	0.0	1.0	55.9	93.2	-60.7	111.2	326	0.994	0.0	1.0	57.1	94.2	-58.7	111.0	328	0.967	0.0	1.0	0.961	0.0	1.0	55.7	93.1	-61.0	111.3	326	0.967	0.0	1.0			
327	329	327	0.983	0.0	1.0	56.6	93.8	-59.5	111.1	327	1.0	0.0	1.0	0.984	57.1	93.9	-56.4	109.6	329	0.983	0.0	1.0	0.985	0.0	1.0	56.7	93.9	-59.3	111.1	327	0.983	0.0	1.0		
328	330	328	1.0	0.0	1.0	57.2	94.3	-58.4	110.9	328	M _d	1.0	0.0	0.962	56.8	93.4	-53.8	107.8	330	M _s	1.0	0.0	1.0	1.0	0.0	0.992	57.2	94.2	-57.4	110.3	328	M _e	1.0	0.0	1.0
329	331	329	1.0	0.0	0.983	57.0	93.9	-56.4	109.5	329	1.0	0.0	0.941	56.5	92.7	-51.3	106.0	331	1.0	0.0	0.983	1.0	0.0	0.972	56.9	93.6	-54.9	108.6	329	1.0	0.0	0.983			
329	332	330	1.0	0.0	0.966	56.8	93.4	-54.4	108.1	329	1.0	0.0	0.919	56.2	92.0	-48.8	104.2	332	1.0	0.0	0.967	1.0	0.0	0.951	56.7	93.0	-52.5	106.9	330	1.0	0.0	0.967			
330	333	331	1.0	0.0	0.95	56.6	92.9	-52.4	106.7	330	1.0	0.0	0.898	55.9	91.2	-46.4	102.4	333	1.0	0.0	0.95	1.0	0.0	0.931	56.4	92.4	-50.2	105.2	331	1.0	0.0	0.95			
331	334	332	1.0	0.0	0.933	56.4	92.4	-50.5	105.3	331	1.0	0.0	0.876	55.7	90.4	-44.0	100.5	334	1.0	0.0	0.933	1.0	0.0	0.911	56.1	91.7	-47.8	103.4	332	1.0	0.0	0.933			
332	335	333	1.0	0.0	0.916	56.1	91.8	-48.6	103.9	332	1.0	0.0	0.86	55.5	90.0	-41.9	99.3	335	1.0	0.0	0.917	1.0	0.0	0.89	55.8	90.9	-45.5	101.7	333	1.0	0.0	0.917			
332	336	334	1.0	0.0	0.9	55.9	91.2	-46.7	102.5	332	1.0	0.0	0.843	55.3	89.2	-39.8	98.3	336	1.0	0.0	0.9	1.0	0.0	0.871	55.6	90.2	-43.3	100.2	334	1.0	0.0	0.9			
333	337	335	1.0	0.0	0.883	55.7	90.6	-44.8	101.1	333	1.0	0.0	0.827	55.1	89.6	-37.8	96.9	337	1.0	0.0	0.883	1.0	0.0	0.856	55.4	89.9	-41.4	99.0	335	1.0	0.0	0.883			
334	338	336	1.0	0.0	0.866	55.5	90.1	-42.8	99.8	334	1.0	0.0	0.811	54.9	88.8	-35.8	95.8	338	1.0	0.0	0.867	1.0	0.0	0.84	55.2	89.6	-39.4	97.9	336	1.0	0.0	0.867			
335	339	337	1.0	0.0	0.85	55.3	89.8	-40.7	98.6	335	1.0	0.0	0.794	54.7	88.3	-33.8	94.6	339	1.0	0.0	0.85	1.0	0.0	0.825	5										

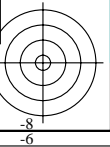
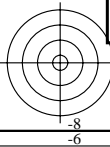
Data of Maximum color M in colorimetric system sRGB standard device; no separation, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM_s; h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;

Six hue angles of the device colours RYGBM_d; h_{ab,d} = 40.0, 102.9, 136.0, 196.4, 306.3, 328.2; Six hue angles of the elementary colours RYGBM_e; h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6

h _{ab,d}	h _{ab,s}	h _{ab,e}	rgb* dd361M	LAB* ddx361Mi (x=LabCh)	rgb* ds361Mi	LAB* dsx361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)	rgb* dd361Mi	rgb* dd	rgb* ds	rgb* de
341	345	342	1.0	0.0	0.75	54.2	86.7	-28.6	91.3	341	1.0	0.0	0.75	
342	346	343	1.0	0.0	0.733	54.0	86.5	-26.4	90.4	342	1.0	0.0	0.733	
344	347	344	1.0	0.0	0.716	53.8	86.2	-24.2	89.5	344	1.0	0.0	0.716	
345	348	345	1.0	0.0	0.7	53.7	85.8	-22.0	88.6	345	1.0	0.0	0.7	
346	349	346	1.0	0.0	0.683	53.5	85.4	-19.9	87.7	346	1.0	0.0	0.683	
348	350	347	1.0	0.0	0.666	53.4	85.0	-17.8	86.8	348	1.0	0.0	0.666	
349	351	348	1.0	0.0	0.65	53.2	84.5	-15.7	85.9	349	1.0	0.0	0.65	
350	352	349	1.0	0.0	0.633	53.0	83.9	-13.6	85.0	350	1.0	0.0	0.633	
352	353	350	1.0	0.0	0.616	52.9	83.6	-11.4	84.3	352	1.0	0.0	0.616	
353	354	351	1.0	0.0	0.6	52.8	83.4	-9.1	83.9	353	1.0	0.0	0.6	
355	355	352	1.0	0.0	0.583	52.7	83.2	-6.9	83.5	355	1.0	0.0	0.583	
356	356	353	1.0	0.0	0.566	52.5	82.9	-4.6	83.0	356	1.0	0.0	0.566	
358	357	354	1.0	0.0	0.55	52.4	82.5	-2.4	82.6	358	1.0	0.0	0.55	
359	358	355	1.0	0.0	0.533	52.3	82.1	-0.1	82.1	359	1.0	0.0	0.533	
361	359	356	1.0	0.0	0.516	52.1	81.6	2.0	81.7	361	1.0	0.0	0.516	
362	360	352	1.0	0.0	0.5	52.0	81.1	4.1	81.2	362	1.0	0.0	0.5	
364	361	353	1.0	0.0	0.483	51.9	81.1	6.5	81.3	364	1.0	0.0	0.483	
366	362	354	1.0	0.0	0.466	51.8	81.0	8.8	81.5	366	1.0	0.0	0.466	
367	363	355	1.0	0.0	0.45	51.7	80.8	11.1	81.6	367	1.0	0.0	0.45	
369	364	356	1.0	0.0	0.433	51.6	80.6	13.5	81.7	369	1.0	0.0	0.433	
371	365	357	1.0	0.0	0.416	51.5	80.3	15.8	81.8	371	1.0	0.0	0.416	
372	366	358	1.0	0.0	0.4	51.4	79.9	18.1	81.9	372	1.0	0.0	0.4	
374	367	359	1.0	0.0	0.383	51.4	79.5	20.4	82.1	374	1.0	0.0	0.383	
376	368	360	1.0	0.0	0.366	51.3	79.3	22.7	82.5	376	1.0	0.0	0.366	
377	369	362	1.0	0.0	0.35	51.2	79.3	25.1	83.2	377	1.0	0.0	0.35	
379	370	363	1.0	0.0	0.333	51.1	79.2	27.4	83.8	379	1.0	0.0	0.333	
380	371	364	1.0	0.0	0.316	51.1	79.1	29.7	84.5	380	1.0	0.0	0.316	
382	372	365	1.0	0.0	0.3	51.0	78.9	32.1	85.2	382	1.0	0.0	0.3	
383	373	366	1.0	0.0	0.283	51.0	78.7	34.4	85.9	383	1.0	0.0	0.283	
385	374	367	1.0	0.0	0.266	50.9	78.3	36.8	86.6	385	1.0	0.0	0.266	
386	375	368	1.0	0.0	0.25	50.8	77.9	39.2	87.2	386	1.0	0.0	0.25	
387	376	369	1.0	0.0	0.233	50.8	78.0	41.2	88.2	387	1.0	0.0	0.233	
389	377	370	1.0	0.0	0.216	50.8	78.0	43.3	89.2	389	1.0	0.0	0.216	
390	378	372	1.0	0.0	0.2	50.7	78.0	45.4	90.2	390	1.0	0.0	0.2	
391	379	373	1.0	0.0	0.183	50.7	77.9	47.5	91.2	391	1.0	0.0	0.183	
392	380	374	1.0	0.0	0.166	50.6	77.8	49.6	92.2	392	1.0	0.0	0.166	
393	381	375	1.0	0.0	0.15	50.6	77.6	51.9	93.3	393	1.0	0.0	0.15	
394	382	376	1.0	0.0	0.133	50.6	77.3	53.9	94.3	394	1.0	0.0	0.133	
395	383	377	1.0	0.0	0.116	50.5	77.2	55.6	95.1	395	1.0	0.0	0.116	
396	384	378	1.0	0.0	0.1	50.5	77.2	56.8	95.9	396	1.0	0.0	0.1	
396	385	379	1.0	0.0	0.083	50.5	77.2	58.1	96.6	396	1.0	0.0	0.083	
397	386	381	1.0	0.0	0.066	50.5	77.2	59.4	97.4	397	1.0	0.0	0.066	
398	387	382	1.0	0.0	0.049	50.5	77.1	60.6	98.1	398	1.0	0.0	0.049	
398	388	383	1.0	0.0	0.033	50.5	77.1	61.9	98.9	398	1.0	0.0	0.033	
399	389	384	1.0	0.0	0.016	50.5	77.0	63.2	99.6	399	1.0	0.0	0.016	
400	390	385	1.0	0.0	0.0	50.4	76.9	64.5	100.4	400	1.0	0.0	0.0	

vedere dei file simili: http://130.149.60.45/~farbmetrik/QI41/QI41.HTM
informazioni tecniche: http://www.ps.bam.de o http://130.149.60.45/~farbmetrik

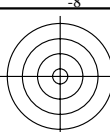
TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rh4ta



vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
La domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rh4ta

nj	HIC*Fa	rgb_Fa	ief_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMa	rgb*Ma	LabCh*Ma		
0/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5	100.4 39.9 0.0	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
1/657	R13Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	51.4 74.1 64.9	98.5 41.2	1.0 0.125 0.0	51.5 73.9 64.9	98.3 41.3 0.2	36	1.0 0.116 0.0	51.4 74.1 64.9	98.5 41.2
2/666	R25Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.7 67.6 65.8	94.4 44.2	1.0 0.25 0.0	54.0 66.7 65.9	93.8 44.6 1.0	42	1.0 0.233 0.0	53.7 67.6 65.8	94.4 44.2
3/675	R38Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	57.9 56.2 67.9	88.1 50.3	1.0 0.375 0.0	58.2 55.4 67.9	87.7 50.7 0.7	51	1.0 0.366 0.0	57.9 56.2 67.9	88.1 50.3
4/684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7 0.0	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
5/693	R63Y_100_100a	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	70.5 24.7 75.4	79.4 71.8	1.0 0.625 0.0	70.1 25.8 75.0	79.3 71.0 1.2	68	1.0 0.633 0.0	70.5 24.7 75.4	79.4 71.8
6/702	R75Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	78.2 7.8 80.6	81.0 84.4	1.0 0.75 0.0	77.2 9.8 79.7	80.3 82.9 2.3	77	1.0 0.766 0.0	78.2 7.8 80.6	81.0 84.4
7/711	R88Y_100_100a	1.0 0.875 0.0	1.0 1.0 0.5	83	1.0 0.883 0.0	85.3 -6.7 85.5	85.8 94.4	1.0 0.875 0.0	84.8 -5.7 85.0	85.2 93.8 1.1	83	1.0 0.883 0.0	85.3 -6.7 85.5	85.8 94.4
8/720	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8	1.0 1.0 0.0	92.6 -20.6 90.7	93.0 102.8 0.0	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
9/639	Y13G_100_100a	0.875 1.0 0.0	1.0 1.0 0.5	97	0.883 1.0 0.0	90.5 -32.2 88.3	94.0 110.0	0.875 1.0 0.0	90.4 -33.0 88.1	94.1 110.5 0.8	96	0.883 1.0 0.0	90.5 -32.2 88.3	94.0 110.0
10/558	Y25G_100_100a	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	88.7 -43.3 86.2	96.5 116.6	0.75 1.0 0.0	88.5 -44.9 85.8	96.8 117.6 1.6	102	0.766 1.0 0.0	88.7 -43.3 86.2	96.5 116.6
11/477	Y38G_100_100a	0.625 1.0 0.0	1.0 1.0 0.5	112	0.633 1.0 0.0	87.0 -55.0 84.1	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_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	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_100a	0.375 1.0 0.0	1.0 1.0 0.5	128	0.366 1.0 0.0	84.7 -73.2 81.2	109.3 132.0	0.375 1.0 0.0	84.7 -72.8 81.2	109.1 131.8 0.3	128	0.366 1.0 0.0	84.7 -73.2 81.2	109.3 132.0
14/234	Y75G_100_100a	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	84.0 -78.7 80.4	112.5 134.3	0.25 1.0 0.0	84.1 -78.2 80.4	112.2 134.1 0.4	137	0.233 1.0 0.0	84.0 -78.7 80.4	112.5 134.3
15/153	Y88G_100_100a	0.125 1.0 0.0	1.0 1.0 0.5	143	0.116 1.0 0.0	83.7 -81.5 80.0	114.2 135.5	0.125 1.0 0.0	83.7 -81.4 80.0	114.2 135.5 0.1	143	0.116 1.0 0.0	83.7 -81.5 80.0	114.2 135.5
16/72	G00C_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0 0.0	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
17/73	G13C_100_100a	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.116	83.6 -82.1 76.8	112.5 136.9	0.0 1.0 0.125	83.6 -82.1 76.5	112.3 137.0 0.2	156	0.0 1.0 0.116	83.6 -82.1 76.8	112.5 136.9
18/74	G25C_100_100a	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.233	83.7 -80.8 70.1	106.9 139.0	0.0 1.0 0.25	83.8 -80.5 69.1	106.1 139.3 1.0	162	0.0 1.0 0.233	83.7 -80.8 70.1	106.9 139.0
19/75	G38C_100_100a	0.0 1.0 0.375	1.0 1.0 0.5	172	0.0 1.0 0.366	84.0 -78.0 58.8	97.7 142.9	0.0 1.0 0.375	84.0 -77.7 58.1	97.1 143.2 0.7	171	0.0 1.0 0.366	84.0 -78.0 58.8	97.7 142.9
20/76	G50C_100_100a	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	84.3 -73.7 44.9	86.4 148.6	0.0 1.0 0.5	84.3 -73.7 44.9	86.3 148.6 0.0	180	0.0 1.0 0.5	84.3 -73.7 44.9	86.4 148.6
21/77	G63C_100_100a	0.0 1.0 0.625	1.0 1.0 0.5	188	0.0 1.0 0.633	84.8 -68.1 29.5	74.3 156.5	0.0 1.0 0.625	84.7 -68.5 30.6	75.0 155.9 1.1	188	0.0 1.0 0.633	84.8 -68.1 29.5	74.3 156.5
22/78	G75C_100_100a	0.0 1.0 0.75	1.0 1.0 0.5	196	0.0 1.0 0.766	85.4 -61.2 13.7	62.8 167.3	0.0 1.0 0.75	85.3 -62.0 15.8	64.0 165.6 2.3	197	0.0 1.0 0.766	85.4 -61.2 13.7	62.8 167.3
23/79	G88C_100_100a	0.0 1.0 0.875	1.0 1.0 0.5	203	0.0 1.0 0.883	86.1 -54.1 0.0	54.1 180.0	0.0 1.0 0.875	86.0 -54.5 1.0	54.5 178.8 1.1	203	0.0 1.0 0.883	86.1 -54.1 0.0	54.1 180.0
24/80	C00B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3 0.0	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
25/71	C13B_100_100a	0.0 0.875 1.0	1.0 1.0 0.5	217	0.0 0.883 1.0	78.5 -33.4 -26.3	42.5 218.2	0.0 0.875 1.0	77.9 -32.3 -27.0	42.1 219.8 1.3	216	0.0 0.883 1.0	78.5 -33.4 -26.3	42.5 218.2
26/62	C25B_100_100a	0.0 0.75 1.0	1.0 1.0 0.5	224	0.0 0.766 1.0	70.2 -19.5 -39.3	43.9 243.6	0.0 0.75 1.0	69.1 -17.0 -40.7	44.1 247.2 3.0	222	0.0 0.766 1.0	70.2 -19.5 -39.3	43.9 243.6
27/53	C38B_100_100a	0.0 0.625 1.0	1.0 1.0 0.5	232	0.0 0.633 1.0	60.9 -1.5 -53.9	53.9 268.3	0.0 0.625 1.0	60.3 -0.1 -54.6	54.6 269.8 1.7	231	0.0 0.633 1.0	60.9 -1.5 -53.9	53.9 268.3
28/44	C50B_100_100a	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0 0.0	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
29/35	C63B_100_100a	0.0 0.375 1.0	1.0 1.0 0.5	248	0.0 0.366 1.0	43.4 38.7 -82.0	90.7 295.3	0.0 0.375 1.0	43.8 37.6 -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_100a	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6	0.0 0.25 1.0	37.1 55.9 -92.3	107.9 301.1 2.1	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6
31/17	C88B_100_100a	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.116 1.0	32.3 70.0 -100.3	122.3 304.9	0.0 0.125 1.0	32.4 69.6 -100.0	121.9 304.8 0.5	263	0.0 0.116 1.0	32.3 70.0 -100.3	122.3 304.9
32/8	B00M_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2 0.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
33/89	B13M_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	30.9 76.2 -102.5	127.8 306.6	0.125 0.0 1.0	31.0 76.2 -102.5	127.7 306.6 0.0	276	0.116 0.0 1.0	30.9 76.2 -102.5	127.8 306.6
34/170	B25M_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4	0.25 0.0 1.0	32.6 76.8 -99.8	125.9 307.5 0.4	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
35/251	B38M_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	34.9 77.9 -95.7	123.4 309.1	0.375 0.0 1.0	35.1 77.9 -95.5	123.3 309.2 0.3	291	0.366 0.0 1.0	34.9 77.9 -95.7	123.4 309.1
36/332	B50M_100_100a	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6	0.5 0.0 1.0	38.5 79.8 -89.7	120.1 311.6 0.0	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
37/413	B63M_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	43.0 82.7 -82.2	116.6 315.1	0.625 0.0 1.0	42.7 82.5 -82.8	116.8 314.8 0.6	308	0.633 0.0 1.0	43.0 82.7 -82.2	116.6 315.1
38/494	B75M_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4	0.75 0.0 1.0	47.2 85.8 -75.1	114.1 318.8 1.3	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4
39/575	B88M_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	52.5 90.1 -66.3	111.9 323.6	0.875 0.0 1.0	52.1 89.8 -66.9	112.0 323.3 0.7	323	0.883 0.0 1.0	52.5 90.1 -66.3	111.9 323.6
40/656	M00R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	1.0 0.0 1.0	57.2 94.3 -58.4	111.0 328.2 0.0	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
41/655	M13R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	55.7 90.6 -44.8	101.1 333.6	1.0 0.0 0.875	55.6 90.3 -43.9	100.4 334.0 0.9	336	1.0 0.0 0.883	55.7 90.6 -44.8	101.1 333.6
42/654	M25R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	54.4 87.3 -30.6	92.5 340.6	1.0 0.0 0.75	54.2 86.7 -28.6	91.3 341.6 2.0	342	1.0 0.0 0.766	54.4 87.3 -30.6	92.5 340.6
43/653	M38R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	53.0 83.9 -13.6	85.0 350.7	1.0 0.0 0.625	53.0 83.6 -12.6	84.6 351.4 1.0	351	1.0 0.0 0.633	53.0 83.9 -13.6	85.0 350.7
44/652	M50R_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	0.0	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
45/651	M63R_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	51.3 79.3 22.7	82.5 16.0	1.0 0.0 0.375	51.3 79.2 21.6	82.1 15.2 1.1	368	1.0 0.0 0.366	51.3 79.3 22.7	82.5 16.0
46/650	M75R_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8	1.0 0.0 0.25	50.8 77.9 39.2	87.2 26.6 2.0	377	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8
47/649	M88R_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	50.5 77.2 55.6	95.1 35.7	1.0 0.0 0.125	50.6 77.2 54.9	94.8 35.4 0.6	383	1.0 0.0 0.116	50.5 77.2 55.6	95.1 35.7
48/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5	100.4 39.9 0.0	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
49/0	NW_000a	0.0 0.0 0.0	0.0											



vedere dei file simili: http://130.149.60.45/~farbmetrik/QI41/QI41.HTM
informazioni tecniche: http://www.ps.bam.de o http://130.149.60.45/~farbmetrik

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rh4ta

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

delta E* = 6.5

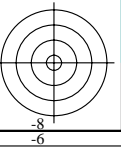
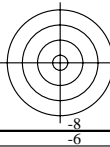
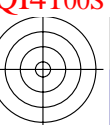
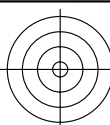


grafico TUB-QI41; codice di tinte: H*d=Y25Gd
colori e la differenza, ΔE*

immettere: rgb/cmyk -> rgbd
uscita: trasferire a rgbd



vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF> / .PS
 informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF / .PS
 la domanda per la misura di stampa di display, nessuna separazione
 TUB materiale: code=rh4t4

n=j	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	rgb*Md	LabCh*Md
0	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1	BOOR_012_012a	0.0	0.0	0.125	0.125	0.125	0.062	0.062	270	0.0	0.0	1.0
2	BOOR_025_025a	0.0	0.0	0.25	0.25	0.25	0.125	0.125	270	0.0	0.0	1.0
3	BOOR_037_037a	0.0	0.0	0.375	0.375	0.375	0.187	0.187	270	0.0	0.0	1.0
4	BOOR_050_050a	0.0	0.0	0.5	0.5	0.5	0.25	0.25	270	0.0	0.0	1.0
5	BOOR_062_062a	0.0	0.0	0.625	0.625	0.625	0.312	0.312	270	0.0	0.0	1.0
6	BOOR_075_075a	0.0	0.0	0.75	0.75	0.75	0.375	0.375	270	0.0	0.0	1.0
7	BOOR_087_087a	0.0	0.0	0.875	0.875	0.875	0.437	0.437	270	0.0	0.0	1.0
8	BOOR_100_100a	0.0	0.0	1.0	1.0	1.0	0.5	0.5	270	0.0	0.0	1.0
9	GOOB_012_012a	0.0	0.125	0.125	0.125	0.062	0.150	0.150	270	0.0	1.0	0.0
10	G50B_012_012a	0.0	0.125	0.125	0.125	0.062	0.210	0.210	270	0.0	1.0	0.0
11	G75B_025_025a	0.0	0.125	0.25	0.25	0.125	0.240	0.240	270	0.0	0.5	1.0
12	G84B_037_037a	0.0	0.125	0.375	0.375	0.187	0.251	0.251	270	0.0	0.316	1.0
13	G88B_050_050a	0.0	0.125	0.5	0.5	0.25	0.256	0.256	270	0.0	0.233	1.0
14	G90B_062_062a	0.0	0.125	0.625	0.625	0.312	0.259	0.259	270	0.0	0.183	1.0
15	G92B_075_075a	0.0	0.125	0.75	0.75	0.375	0.261	0.261	270	0.0	0.15	1.0
16	G93B_087_087a	0.0	0.125	0.875	0.875	0.437	0.262	0.262	270	0.0	0.133	1.0
17	G94B_100_100a	0.0	0.125	1.0	1.0	0.5	0.263	0.263	270	0.0	0.116	1.0
18	GOOB_025_025a	0.0	0.25	0.25	0.25	0.125	0.150	0.150	270	0.0	1.0	0.0
19	G25B_025_025a	0.0	0.25	0.125	0.25	0.125	0.180	0.180	270	0.0	1.0	0.5
20	G50B_025_025a	0.0	0.25	0.25	0.25	0.125	0.210	0.210	270	0.0	1.0	0.0
21	G65B_037_037a	0.0	0.25	0.375	0.375	0.187	0.229	0.229	270	0.0	0.683	1.0
22	G75B_050_050a	0.0	0.25	0.5	0.5	0.25	0.240	0.240	270	0.0	0.5	1.0
23	G80B_062_062a	0.0	0.25	0.625	0.625	0.312	0.247	0.247	270	0.0	0.383	1.0
24	G84B_075_075a	0.0	0.25	0.75	0.75	0.375	0.251	0.251	270	0.0	0.316	1.0
25	G86B_087_087a	0.0	0.25	0.875	0.875	0.437	0.254	0.254	270	0.0	0.266	1.0
26	G88B_100_100a	0.0	0.25	1.0	1.0	0.5	0.256	0.256	270	0.0	0.233	1.0
27	GOOB_037_037a	0.0	0.375	0.375	0.375	0.187	0.150	0.150	270	0.0	1.0	0.0
28	G15B_037_037a	0.0	0.375	0.125	0.375	0.187	0.169	0.169	270	0.0	0.316	0.836
29	G34B_037_037a	0.0	0.375	0.25	0.375	0.187	0.191	0.191	270	0.0	1.0	0.683
30	G50B_037_037a	0.0	0.375	0.375	0.375	0.187	0.210	0.210	270	0.0	1.0	0.0
31	G61B_050_050a	0.0	0.375	0.5	0.5	0.25	0.224	0.224	270	0.0	0.766	1.0
32	G69B_062_062a	0.0	0.375	0.625	0.625	0.312	0.233	0.233	270	0.0	0.616	1.0
33	G75B_075_075a	0.0	0.375	0.75	0.75	0.375	0.240	0.240	270	0.0	0.5	1.0
34	G79B_087_087a	0.0	0.375	0.875	0.875	0.437	0.245	0.245	270	0.0	0.416	1.0
35	G81B_100_100a	0.0	0.375	1.0	1.0	0.5	0.248	0.248	270	0.0	0.366	1.0
36	GOOB_050_050a	0.0	0.5	0.0	0.5	0.25	0.150	0.150	270	0.0	1.0	0.0
37	G11B_050_050a	0.0	0.5	0.125	0.5	0.25	0.164	0.164	270	0.0	1.0	0.233
38	G25B_050_050a	0.0	0.5	0.25	0.5	0.25	0.180	0.180	270	0.0	1.0	0.5
39	G38B_050_050a	0.0	0.5	0.375	0.5	0.25	0.196	0.196	270	0.0	1.0	0.766
40	G50B_050_050a	0.0	0.5	0.5	0.5	0.25	0.210	0.210	270	0.0	1.0	0.0
41	G59B_062_062a	0.0	0.5	0.625	0.625	0.312	0.221	0.221	270	0.0	0.816	1.0
42	G65B_075_075a	0.0	0.5	0.75	0.75	0.375	0.229	0.229	270	0.0	0.683	1.0
43	G70B_087_087a	0.0	0.5	0.875	0.875	0.437	0.235	0.235	270	0.0	0.583	1.0
44	G75B_100_100a	0.0	0.5	1.0	1.0	0.5	0.240	0.240	270	0.0	0.5	1.0
45	GOOB_062_062a	0.0	0.625	0.0	0.625	0.312	0.150	0.150	270	0.0	1.0	0.0
46	GO9B_062_062a	0.0	0.625	0.125	0.625	0.312	0.161	0.161	270	0.0	1.0	0.183
47	G19B_062_062a	0.0	0.625	0.25	0.625	0.312	0.173	0.173	270	0.0	1.0	0.383
48	G30B_062_062a	0.0	0.625	0.375	0.625	0.312	0.187	0.187	270	0.0	1.0	0.616
49	G40B_062_062a	0.0	0.625	0.5	0.625	0.312	0.199	0.199	270	0.0	1.0	0.816
50	G50B_062_062a	0.0	0.625	0.625	0.625	0.312	0.210	0.210	270	0.0	1.0	0.0
51	G57B_075_075a	0.0	0.625	0.75	0.75	0.375	0.219	0.219	270	0.0	0.85	1.0
52	G63B_087_087a	0.0	0.625	0.875	0.875	0.437	0.226	0.226	270	0.0	0.733	1.0
53	G68B_100_100a	0.0	0.625	1.0	1.0	0.5	0.232	0.232	270	0.0	0.633	1.0
54	GOOB_075_075a	0.0	0.75	0.0	0.75	0.375	0.150	0.150	270	0.0	1.0	0.0
55	G07B_075_075a	0.0	0.75	0.125	0.75	0.375	0.159	0.159	270	0.0	1.0	0.15
56	G15B_075_075a	0.0	0.75	0.25	0.75	0.375	0.169	0.169	270	0.0	1.0	0.316
57	G25B_075_075a	0.0	0.75	0.375	0.75	0.375	0.180	0.180	270	0.0	1.0	0.5
58	G34B_075_075a	0.0	0.75	0.5	0.75	0.375	0.191	0.191	270	0.0	1.0	0.683
59	G42B_075_075a	0.0	0.75	0.625	0.75	0.375	0.201	0.201	270	0.0	1.0	0.85
60	G50B_075_075a	0.0	0.75	0.75	0.75	0.375	0.210	0.210	270	0.0	1.0	0.0
61	G56B_087_087a	0.0	0.75	0.875	0.875	0.437	0.218	0.218	270	0.0	0.866	1.0
62	G61B_100_100a	0.0	0.75	1.0	1.0	0.5	0.224	0.224	270	0.0	0.766	1.0
63	GOOB_087_087a	0.0	0.875	0.0	0.875	0.437	0.150	0.150	270	0.0	1.0	0.0
64	G06B_087_087a	0.0	0.875	0.125	0.875	0.437	0.158	0.158	270	0.0	1.0	0.133
65	G13B_087_087a	0.0	0.875	0.25	0.875	0.437	0.166	0.166	270	0.0	1.0	0.266
66	G20B_087_087a	0.0	0.875	0.375	0.875	0.437	0.175	0.175	270	0.0	1.0	0.416
67	G29B_087_087a	0.0	0.875	0.5	0.875	0.437	0.185	0.185	270	0.0	1.0	0.583
68	G36B_087_087a	0.0	0.875	0.625	0.875	0.437	0.194	0.194	270	0.0	1.0	0.733
69	G43B_087_087a	0.0	0.875	0.75	0.875	0.437	0.202	0.202	270	0.0	1.0	0.866
70	G50B_087_087a	0.0	0.875	0.875	0.875	0.437	0.210	0.210	270	0.0	1.0	0.0
71	G55B_100_100a	0.0	0.875	1.0	1.0	0.5	0.217	0.217	270	0.0	0.883	1.0
72	GOOB_100_100a	0.0	1.0	0.0	1.0	0.5	0.150	0.150	270	0.0	1.0	0.0
73	G05B_100_100a	0.0	1.0	0.125	1.0	0.5	0.157	0.157	270	0.0	1.0	0.116
74	G11B_100_100a	0.0	1.0	0.25	1.0	0.5	0.164	0.164	270	0.0	1.0	0.233
75	G18B_100_100a	0.0	1.0	0.375	1.0	0.5	0.172	0.172	270	0.0	1.0	0.366
76	G25B_100_100a	0.0	1.0	0.5	1.0	0.5	0.180	0.180	270	0.0	1.0	0.5
77	G31B_100_100a	0.0	1.0	0.625	1.0	0.5	0.188	0.188	270	0.0	1.0	0.633
78	G38B_100_100a	0.0	1.0	0.75	1.0	0.5	0.196	0.196	270	0.0	1.0	0.766
79	G44B_100_100a	0.0	1.0	0.875	1.0	0.5	0.203	0.203	270	0.0	1.0	0.883
80	G50B_100_100a	0.0	1.0	1.0	1.0	0.5	0.210	0.210	270	0.0	1.0	0.0

delta E** = 4.6

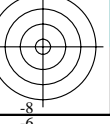
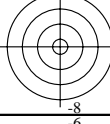
4-0031530-F0

QI410-7N, 16,29-F

grafico TUB-QI41; codice di tinte: H*d=Y25Gd
 colori e la differenza, ΔE*

immettere: rgb/cmyk -> rgbd
 uscita: trasferire a rgbd

4-0031530-F0



vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF> / .PS
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF / .PS
La domanda per la misura di stampa di display, nessuna separazione

TUB materiale: code=rh4t4

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Ma	rgb*Ma	LabCh*Ma		
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	20.4 325.7 7.6	330	1.0 0.0 1.0	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 1.0	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	0.0 0.0 0.0	0.0 325.7 0.8	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0
92	B00R_025_012a	0.125 0.125 0.25	0.25 0.125 0.187	270	0.124 0.124 0.25	15.7 9.5 -12.9	16.0 306.2	0.125 0.125 0.25	12.6 9.6 -19.5	21.8 296.2 7.3	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
93	B00R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.124 0.375	19.5 19.0 -25.8	32.1 306.2	0.125 0.125 0.375	15.0 21.1 -36.5	42.1 300.0 11.6	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
94	B00R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.124 0.5	23.3 28.5 -38.8	48.1 306.2	0.125 0.125 0.5	18.1 32.4 -51.3	60.6 302.2 14.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
95	B00R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	27.1 38.0 -51.7	64.2 306.2	0.125 0.125 0.625	21.6 42.8 -64.6	77.5 303.5 14.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
96	B00R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	30.9 47.5 -64.7	80.3 306.2	0.125 0.125 0.75	25.3 52.5 -76.8	93.0 304.3 14.2	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
97	B00R_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	34.7 57.0 -77.6	96.3 306.2	0.125 0.125 0.875	29.1 61.5 -88.2	107.5 304.8 12.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
98	B00R_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	38.5 66.5 -90.6	112.4 306.2	0.125 0.125 1.0	33.0 69.9 -99.0	121.3 305.2 10.6	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
99	Y50G_025_012a	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	51.7 183.3 -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	59.0 294.1 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_075a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.237 0.875	36.9 50.0 -73.9	89.3 304.0	0.125 0.25 0.875	34.4 46.3 -79.5	92.0 300.2 7.1	262	0.0 0.15 1.0	33.4 66.7 -98.6	119.1 304.0
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.241 1.0	40.6 60.0 -87.1	105.8 304.5	0.125 0.25 1.0	37.6 56.5 -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	51.7 183.3 -68.3	70.7 285.0
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.364 0.75	39.6 22.6 -50.3	55.1 294.2	0.125 0.375 0.75	39.5 15.3 -53.5	55.6 285.9 7.9	247	0.0 0.383 1.0	44.3 36.2 -80.5	88.2 294.2
115	G84B_087_075a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.362 0.875	42.4 34.3 -65.0	73.5 297.8	0.125 0.375 0.875	41.7 27.1 -67.4	72.7 291.9 7.5	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
116	G86B_100_087a	0.125 0.375 1.0	1.0 0.875 0.562	254	0.125 0.358 1.0	45.2 46.6 -79.6	92.2 300.3	0.125 0.375 1.0	44.2 38.6 -80.5	89.3 295.5 8.1	255	0.0 0.266 1.0	38.0 53.3 -91.0	105.4 300.3
117	Y76G_050_050a	0.125 0.5 0.0	0.5 0.5 0.25	136	0.116 0.5 0.0	42.0 -39.3 40.2	56.2 134.3	0.125 0.5 0.0	43.9 -45.9 48.2	66.6 133.6 10.6	137	0.233 1.0 0.0	84.0 -78.7	80.4 112.5 134.3
118	G00B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.124	43.2 -31.0 29.9	43.1 136.0	0.125 0.5 0.125	44.1 -44.3 40.1	59.8 137.8 16.7	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.243	43.3 -29.9 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 8.8 -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	51.7 183.3 -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 50.3	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.								

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Ma	rgb*Ma	LabCh*Ma			
162	R00Y_025_025a	0.25 0.0 0.0	0.25 0.25 0.125	390	0.25 0.0 0.0	12.6 19.2 16.1	25.1 25.1 4.0	0.25 0.0 0.0	8.6 28.5 13.6	31.6 31.6	25.5 10.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
163	R00Y_025_025a	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.125	13.0 20.2 1.0	20.3 2.9	0.25 0.0 0.125	9.4 30.5 -1.8	30.6 35.6	11.2 36.0	1.0 0.0 0.5	52.0 81.1 4.1	81.2 29.0	
164	B50R_025_025a	0.25 0.0 0.25	0.25 0.25 0.125	330	0.25 0.0 0.25	14.3 23.5 -14.6	27.7 328.2	0.25 0.0 0.25	11.1 34.9 -21.6	41.1 32.8	13.7 330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	
165	B34R_037_037a	0.25 0.0 0.375	0.375 0.375 0.187	311	0.256 0.0 0.375	16.8 31.5 -29.7	43.3 316.7	0.25 0.0 0.375	13.8 41.1 -38.3	56.2 316.9	13.2 311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7	
166	B25R_050_050a	0.25 0.0 0.5	0.5 0.5 0.25	300	0.25 0.0 0.5	19.2 39.9 -44.8	60.0 311.6	0.25 0.0 0.5	17.1 48.0 -52.8	71.4 312.2	11.6 300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6	
167	B19R_062_062a	0.25 0.0 0.625	0.625 0.625 0.312	293	0.239 0.0 0.625	22.1 48.8 -59.9	76.9 309.3	0.25 0.0 0.625	20.7 55.2 -65.9	86.0 309.9	9.2 292	0.383 0.0 1.0	35.3 78.1 -95.1	123.0 309.3	
168	B15R_075_075a	0.25 0.0 0.75	0.75 0.75 0.375	289	0.237 0.0 0.75	25.4 58.1 -73.1	93.4 308.4	0.25 0.0 0.75	24.6 62.5 -77.8	99.8 308.7	6.5 288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4	
169	B13R_087_087a	0.25 0.0 0.875	0.875 0.875 0.437	286	0.233 0.0 0.875	28.8 67.3 -86.8	109.9 307.8	0.25 0.0 0.875	28.6 69.7 -89.1	113.1 308.0	3.2 284	0.266 0.0 1.0	32.9 77.0 -99.2	125.6 307.8	
170	B11R_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4	0.25 0.0 1.0	32.6 76.8 -99.8	125.9 307.5	0.4 282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4	
171	R50Y_025_025a	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.125 0.0	15.9 10.3 17.7	20.5 59.7	0.25 0.125 0.0	14.7 12.2 22.0	25.2 60.9	4.8 59	1.0 0.5 0.0	63.6 46.3 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 0.312	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 0.25	289	0.243 0.124 0.5	24.6 29.0 -36.5	46.7 308.4	0.25 0.125 0.5	20.9 36.7 -46.5	59.3 308.3	13.1 288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4	
176	B11R_062_050a	0.25 0.125 0.625	0.625 0.5 0.375	284	0.241 0.125 0.625	28.1 38.3 -50.0	63.1 307.4	0.25 0.125 0.625	23.9 45.7 -60.5	75.9 307.0	13.4 282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4	
177	B09R_075_062a	0.25 0.125 0.75	0.75 0.625 0.437	281	0.239 0.125 0.75	31.7 47.8 -63.2	79.3 307.0	0.25 0.125 0.75	27.3 54.4 -73.4	91.4 306.5	12.9 279	0.233 0.0 1.0	31.7 76.5 -101.2	126.9 307.0	
178	B07R_087_075a	0.25 0.125 0.875	0.875 0.75 0.5	279	0.237 0.125 0.875	35.4 57.2 -76.4	95.5 306.8	0.25 0.125 0.875	30.8 62.8 -85.3	106.0 306.3	11.4 278	0.15 0.0 1.0	31.1 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	111.6 306.2	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 32.5	1.4 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	
183	B00R_037_012a	0.25 0.25 0.375	0.375 0.125 0.312	270	0.249 0.249 0.375	27.6 9.5 -12.9	16.0 306.2	0.25 0.25 0.375	26.5 8.0 -18.0	19.8 294.0	5.4 270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
184	B00R_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.249 0.5	31.4 19.0 -25.8	32.1 306.2	0.25 0.25 0.5	28.2 17.7 -34.7	39.0 297.0	9.5 270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
185	B00R_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	35.2 28.5 -38.8	48.1 306.2	0.25 0.25 0.625	30.4 28.1 -50.0	57.4 299.3	12.2 270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	39.0 38.0 -51.7	64.2 306.2	0.25 0.25 0.75	32.9 38.5 -64.1	74.8 301.0	13.7 270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
187	B00R_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.8 47.5 -64.7	80.3 306.2	0.25 0.25 0.875	35.8 48.6 -77.1	91.2 302.1	14.3 270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
188	B00R_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	46.6 57.0 -77.6	96.3 306.2	0.25 0.25 1.0	38.8 58.2 -89.4	106.7 303.0	14.1 270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
189	Y31G_037_037a	0.25 0.375 0.0	0.375 0.375 0.187	109	0.256 0.375 0.0	32.8 -19.0 31.8	37.1 120.8	0.25 0.375 0.0	34.6 -24.3 41.4	48.0 120.4	11.0 108	0.683 1.0 0.0	87.6 -50.7	84.9 98.9 120.8	
190	Y50G_037_025a	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	33.3 -16.3 20.6	26.2 128.3	0.25 0.375 0.125	34.8 -22.5 30.5	38.0 126.3	11.8 119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3	
191	G00B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	34.3 -10.3 9.9	14.3 136.0	0.25 0.375 0.25	35.2 -18.1 14.0	22.9 142.2	8.8 149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0	
192	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	34.7 -5.7 -1.6	6.0 196.3	0.25 0.375 0.375	36.0 -11.0 -3.5	11.6 197.8	5.8 210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3	
193	G75B_050_025a	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.375 0.5	36.7 4.5 -17.0	17.6 285.0	0.25 0.375 0.5	37.2 -2.0 -20.5	20.6 264.3	7.4 240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0	
194	G84B_062_037a	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.368 0.625	39.1 17.1 -32.5	36.7 297.8	0.25 0.375 0.625	38.7 8.2 -36.6	37.5 282.7	9.7 251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8	
195	G88B_075_050a	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.366 0.75	42.1 28.8 -46.7	54.8 301.6	0.25 0.375 0.75	40.6 19.1 -51.6	55.0 290.3	10.9 257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6	
196	G90B_087_062a	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.364 0.875	45.5 39.4 -60.3	72.1 303.1	0.25 0.375 0.875	42.8 30.1 -65.7	72.2 294.6	11.0 260	0.0 0.183 1.0	34.6 63.0 -96.6	115.3 303.1	
197	G92B_100_075a	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.362 1.0	48.9 50.0 -73.9	89.3 304.0	0.25 0.375 1.0	45.2 40.8 -78.9	88.9 297.3	11.1 262	0.0 0.15 1.0	33.4 66.7	-98.6 119.1 304.0	
198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.25 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 -19.3 10.6	29.3 158.6	8.9 180	0.0 1.0 0.5	84.3 -73.7	44.9 86.4 148.6	
202	G50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	45.5 -11.5 -3.3	12.0 196.3	0.25 0.5 0.5	46.8 -27.5 -6.0	20.4 197.2	8.5 210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3	
203	G65B_062_037a	0.25 0.5 0.625	0.625 0.375 0.437	229	0.25 0.506 0.625	48.0 -3.4 -18.3	18.6 259.3	0.25 0.5 0.625	47.9 -10.2 -22.3	24.5 245.3	7.8 228	0.0 0.683 1.0	64.4 -9.2	-48.8 49.7 259.3	
204	G75B_075_050a	0.25 0.5 0.75	0.75 0.5 0.5	240	0.25 0.5 0.75	49.7 9.1 -34.1	35.3 285.0	0.25 0.5 0.75	49.3 0.1 -37.8	37.8 270.1	9.7 240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0	
205	G80B_087_062a	0.25 0.5 0.875	0.875 0.625 0.562	247	0.25 0.489 0.875	51.5 22.6 -50.3	55.1 294.2	0.25 0.5 0.875	50.9 10.9 -52.5	53.6 281.7	11.9 247	0.0 0.383 1.0	44.3 36.2 -80.5	88.2 294.2	
206	G84B_100_075a	0.25 0.5 1.0	1.0 0.75 0.625	251	0.25 0.487 1.0	54.4 34.3 -60.5	73.5 297.8	0.25 0.5 1.0	52.8 21.9 -66.5	70.0 288.2	12.5 251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8	
207	Y61G_062_062a	0.25 0.625 0.0	0.625 0.625 0.312	127	0.239 0.625 0.0	53.0 -45.2 50.8	68.0 131.6	0.25 0.625 0.0	55.1 -49.5 57.4	75.8 130.7	8.1 127	0.383 1.0 0.0	84.8 -72.3	81.3 108.8 131.6	
208	Y76G_062_050a	0.25 0.625 0.125	0.625 0.5 0.375	136	0.241 0.625 0.125	53.9 -39.3 40.2	56.2 134.3	0.25 0.625 0.125	55.2 -48.4 51.2	70.5 133.3	14.3 137	0.233 1.0 0.0	84.0 -78.7	80.4 112.5 134.3	
209	G00B_062_037a	0.25 0.625 0.25	0.625 0.375 0.437	150	0.25 0.625 0.25	55.2 -31.0 29.9	43.1 136.0	0.25 0.625 0.25	55.4 -45.7 39.2	60.2 139.3	17.3 149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0	
210	G15B_062_037a	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 1	

vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF> / .PS
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF / .PS
la domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rhatha

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md			
243	R00Y_037_037a	0.375 0.0 0.0	0.375 0.375 0.187	390	0.375 0.0 0.0	18.9 28.8 24.2	37.6 40.0	0.375 0.0 0.0	16.4 37.5 25.4	45.3 34.1 9.1	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
244	R18Y_037_037a	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.118	19.1 29.6 11.1	31.7 20.6	0.375 0.0 0.125	16.8 38.7 9.7	39.9 14.1 9.4	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6	
245	B65R_037_037a	0.375 0.0 0.25	0.375 0.375 0.187	349	0.375 0.0 0.256	20.0 32.0 -7.4	32.9 346.8	0.375 0.0 0.25	17.9 41.5 -10.4	42.8 345.8 10.2	348	1.0 0.0 0.683	53.5 85.4 -19.9	87.7 346.8	
246	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	21.4 35.5 -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	0.383 0.0 0.625	26.5 51.4 -52.0	73.1 314.6	0.375 0.0 0.625	24.9 57.8 -58.7	82.4 314.5 9.4	307	0.616 0.0 1.0	42.4 82.3 -83.2	117.0 314.6	
249	B25R_075_075a	0.375 0.0 0.75	0.75 0.75 0.375	300	0.375 0.0 0.75	28.9 59.8 -67.2	90.0 311.6	0.375 0.0 0.75	28.1 64.4 -71.9	96.5 311.8 6.5	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6	
250	B20R_087_087a	0.375 0.0 0.875	0.875 0.875 0.437	295	0.364 0.0 0.875	31.7 68.8 -81.8	106.9 310.0	0.375 0.0 0.875	31.6 71.2 -84.0	110.1 310.2 3.2	294	0.416 0.0 1.0	36.3 78.6 -93.5	122.2 310.0	
251	B18R_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	34.9 77.9 -95.7	123.4 309.1	0.375 0.0 1.0	35.1 77.9 -95.5	123.3 309.2 0.3	291	0.366 0.0 1.0	34.9 77.9 -95.7	123.4 309.1	
252	R31Y_037_037a	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.118 0.0	21.1 22.7 25.2	33.9 47.9	0.375 0.125 0.0	20.4 26.4 30.1	40.1 48.7 6.2	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9	
253	R00Y_037_025a	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.124	24.5 19.2 16.1	25.1 40.0	0.375 0.125 0.125	20.7 27.8 14.8	31.5 28.0 9.5	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
254	R00Y_037_025a	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.25	24.9 20.2 1.0	20.3 2.9	0.375 0.125 0.25	21.6 31.1 -4.9	31.5 25.0 12.8	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	
255	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.124 0.375	26.2 23.5 -14.6	27.7 328.2	0.375 0.125 0.375	23.1 36.3 -23.1	43.0 327.5 15.6	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	
256	B34R_050_037a	0.375 0.125 0.5	0.5 0.5 0.375	312	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	123.0 309.3	
259	B15R_087_075a	0.375 0.125 0.875	0.875 0.75 0.5	289	0.362 0.125 0.875	37.4 58.1 -76.1	93.4 308.4	0.375 0.125 0.875	33.6 65.1 -80.7	103.7 308.9 11.0	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4	
260	B13R_100_087a	0.375 0.125 1.0	1.0 0.875 0.562	286	0.358 0.125 1.0	40.7 67.3 -83.8	109.9 307.8	0.375 0.125 1.0	36.9 72.6 -92.6	117.7 308.1 8.7	284	0.266 0.0 1.0	32.9 77.0 -99.2	125.6 307.8	
261	R68Y_037_037a	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.256 0.0	27.5 6.9 29.1	29.9 76.5	0.375 0.25 0.0	27.8 8.3 37.5	38.4 77.4 8.5	71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5	
262	R50Y_037_025a	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.25 0.124	27.8 10.3 17.7	20.5 59.7	0.375 0.25 0.125	28.1 9.8 23.7	25.7 67.5 6.0	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7	
263	R00Y_037_012a	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.249	30.1 9.6 8.0	12.5 40.0	0.375 0.25 0.25	28.7 13.3 5.4	14.4 22.0 4.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
264	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.249 0.375	31.0 11.7 -7.3	13.8 328.2	0.375 0.25 0.375	29.7 19.0 -12.7	22.9 326.1 9.1	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	
265	B25R_050_025a	0.375 0.25 0.5	0.5 0.25 0.375	300	0.375 0.249 0.5	33.5 19.9 -22.4	30.0 311.6	0.375 0.25 0.5	31.2 26.3 -29.7	39.7 311.5 9.9	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6	
266	B15R_062_037a	0.375 0.25 0.625	0.625 0.375 0.437	289	0.368 0.25 0.625	36.5 29.0 -36.5	46.7 308.4	0.375 0.25 0.625	33.2 34.6 -45.4	57.0 307.3 10.9	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4	
267	B11R_075_050a	0.375 0.25 0.75	0.75 0.5 0.5	284	0.366 0.25 0.75	40.0 38.3 -50.0	63.1 307.4	0.375 0.25 0.75	35.4 43.3 -59.8	73.9 305.9 11.8	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4	
268	B09R_087_062a	0.375 0.25 0.875	0.875 0.625 0.562	281	0.364 0.25 0.875	43.7 47.8 -63.2	79.3 307.0	0.375 0.25 0.875	38.0 52.2 -73.3	90.0 305.4 12.3	279	0.183 0.0 1.0	31.7 76.5 -101.2	126.9 307.0	
269	B07R_100_075a	0.375 0.25 1.0	1.0 0.75 0.625	279	0.362 0.25 1.0	47.3 -76.4	95.5 306.8	0.375 0.25 1.0	40.9 60.9 -86.0	105.4 305.3 12.0	278	0.15 0.0 1.0	31.3 76.3 -101.9	127.4 306.8	
270	Y00G_037_037a	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.375 0.0	34.7 -7.7	34.0 34.9	0.375 0.375 0.0	36.9 -10.0	44.2 45.3 10.2	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8	
271	Y00G_037_025a	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.375 0.124	35.0 -5.1	22.6 23.2	0.375 0.375 0.125	37.1 -8.7	33.8 34.9 10.4	11.8	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8
272	Y00G_037_012a	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.375 0.249	35.4 -2.5	11.3 11.6	0.375 0.375 0.25	37.5 -5.4	17.5 18.3 10.7	7.1	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8
273	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	35.7 0.0 0.0	0.0 0.0	0.375 0.375 0.375	38.3 0.0 0.0	0.0 0.0 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	
274	B00R_050_012a	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.375 0.5	39.5 9.5 -12.9	16.0 306.2	0.375 0.375 0.5	39.4 7.2 -17.0	18.5 292.9 4.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
275	B00R_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	43.3 19.0 -25.8	32.1 306.2	0.375 0.375 0.625	40.8 15.7 -33.2	36.8 295.4 8.4	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
276	B00R_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	47.1 28.5 -38.8	48.1 306.2	0.375 0.375 0.75	42.5 25.1 -48.4	54.5 297.4 11.1	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
277	B00R_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	50.9 38.0 -51.7	64.2 306.2	0.375 0.375 0.875	44.6 34.8 -62.7	71.7 299.0 13.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
278	B00R_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	54.7 47.5 -64.7	80.3 306.2	0.375 0.375 1.0	46.8 44.5 -76.1	88.2 300.3 14.2	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
279	Y23G_050_050a	0.375 0.5 0.0	0.5 0.5 0.25	104	0.383 0.5 0.0	44.3 -21.6	43.1 48.2	0.375 0.5 0.0	46.6 -26.1	51.4 57.7 116.9	9.7	102	0.766 1.0 0.0	88.7 -43.3	86.2 96.5 116.6
280	Y31G_050_037a	0.375 0.5 0.125	0.5 0.375 0.312	109	0.381 0.5 0.124	44.8 -19.0	31.8 37.1	0.375 0.5 0.125	46.7 -25.0	43.6 50.2 119.8	13.3	108	0.683 1.0 0.0	87.6 -50.7	84.9 98.9 120.8
281	Y50G_050_025a	0.375 0.5 0.25	0.5 0.25 0.375	120	0.375 0.5 0.249	45.2 -16.3	20.6 26.2	0.375 0.5 0.25	47.0 -22.1	29.6 36.9 126.8	10.8	119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3
282	G00B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	46.2 -10.3	9.9 14.3	0.375 0.5 0.375	47.6 -17.3	13.1 21.8 142.8	7.8	149	0.0 1.0 1.0	83.6 -82.7	79.8 115.0 136.0
283	G50B_050_012a	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.5	46.6 -5.7	-1.6 6.0	0.375 0.5 0.5	48.4 -10.7	-3.5 11.3 198.2	5.6	210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3
284	G75B_062_025a	0.375 0.5 0.625	0.625 0.25 0.5	240	0.375 0.5 0.625	48.7 4.5 -17.0	17.6 285.0	0.375 0.5 0.625	49.4 -2.7	-19.8 20.0 262.1	7.8	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
285	G84B_075_037a	0.375 0.5 0.75	0.75 0.375 0.562	251	0.375 0.493 0.75	51.0 17.1 -32.5	36.7 297.8	0.375 0.5 0.75	50.7 6.3 -35.4	35.9 280.2 11.1	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8	
286	G88B_087_050a	0.375 0.5 0.875	0.875 0.5 0.625	256	0.375 0.491 0.875	54.0 28.8 -46.7	54.8 301.6	0.375 0.5 0.875	52.3 16.1 -50.2	52.7 287.8 13.2	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6	
287	G90B_100_062a	0.375 0.5 1.0	1.0 0.625 0.687	259	0.375 0.489 1.0	57.4 39.4 -60.3	72.1 303.1	0.375 0.5 1.0	54.1 26.2 -64.3	69.4 292.1 14.1	260	0.0 0.183 1.0	34.6 63.0 -96.6	115.3 303.1	
288	Y38G_062_062a	0.375 0.625 0.0	0.625 0.625 0.312	113	0.385 0.625 0.0	54.2 -35.2	52.4 63.1	0.375 0.625 0.0	56.3 -39.9	58.9 71.7	12.4 8.3	112	0.616 1.0 0.0	86.8 -56.4	83.8 101.0 123.9
289	Y50G_062_050a	0.375 0.625 0.125	0.625 0.5 0.375	120	0.375 0.625 0.125	54.7 -32.6	41.1 52.5	0.375 0.625 0.125	56.4 -39.0	52.8 65.2	12.6 13.4	119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3
290	Y68G_062_037a	0.375 0.625 0.25	0.625 0.375 0.437	131	0.368 0.625 0.25	55.5 -28.2	30.3 41.4	0.375 0.625 0.25	56.6 -3						

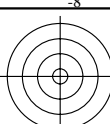
vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF> / .PS
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md			
324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.0	25.2 38.4 32.2	50.2 40.0	0.5 0.0 0.0	23.7 46.0 35.7	58.2 37.8 8.4	389	1.0 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 0.116	25.4 39.0 20.6	44.1 27.8	0.5 0.0 0.125	24.0 46.8 20.3	51.0 23.5 7.9	377	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8	
326	R00Y_050_050a	0.5 0.0 0.25	0.5 0.5 0.25	360	0.5 0.0 0.25	26.0 40.5 2.0	40.6 2.9	0.5 0.0 0.25	24.8 48.8 0.4	48.8 0.5 8.4	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	
327	B61R_050_050a	0.5 0.0 0.375	0.5 0.5 0.25	344	0.5 0.0 0.383	27.2 43.6	-15.3 46.2 340.6	0.5 0.0 0.375	26.0 52.0	-18.0 55.1 340.8	8.9 342	1.0 0.0 0.766	54.4 87.3	-30.6 92.5 340.6	
328	B50R_050_050a	0.5 0.0 0.5	0.5 0.5 0.25	330	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 1.0	57.2 94.3	-58.4 110.9 328.2	
329	B40R_062_062a	0.5 0.0 0.625	0.625 0.625 0.312	319	0.51 0.0 0.625	31.1 55.0	-44.2 70.6 321.2	0.5 0.0 0.625	30.0 61.6	-50.3 79.5 320.7	9.0 320	0.816 0.0 1.0	49.8 88.1	-70.7 113.0 321.2	
330	B34R_075_075a	0.5 0.0 0.75	0.75 0.75 0.375	311	0.512 0.0 0.75	33.6 63.1	-59.4 86.6 316.7	0.5 0.0 0.75	32.6 67.4	-64.4 93.2 316.3	6.6 311	0.683 0.0 1.0	44.8 84.1	-79.2 115.5 316.7	
331	B29R_087_087a	0.5 0.0 0.875	0.875 0.875 0.437	305	0.51 0.0 0.875	36.1 71.4	-74.4 103.2 313.8	0.5 0.0 0.875	35.5 73.5	-77.4 106.8 313.5	3.7 305	0.583 0.0 1.0	41.3 81.6	-85.1 117.9 313.8	
332	B25R_100_100a	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	38.5 79.8	-89.7 120.0 311.6	0.5 0.0 1.0	38.5 79.8	-89.7 120.1	311.6 0.0 300	0.5 0.0 1.0	38.5 79.8	-89.7 120.0 311.6	
333	R23Y_050_050a	0.5 0.125 0.0	0.5 0.5 0.25	44	0.5 0.116 0.0	26.8 33.8	32.9 47.2 44.2	0.5 0.125 0.0	26.5 38.1	38.3 54.1 6.9	42	1.0 0.233 0.0	53.7 67.6	65.8 94.4 44.2	
334	R00Y_050_037a	0.5 0.125 0.125	0.5 0.375 0.312	390	0.5 0.124 0.124	30.8 28.8	24.2 37.6 40.0	0.5 0.125 0.125	26.8 39.0	23.5 45.6 31.1	10.9 389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0	
335	R18Y_050_037a	0.5 0.125 0.25	0.5 0.375 0.312	371	0.5 0.124 0.243	31.0 29.6	11.1 31.7 20.6	0.5 0.125 0.25	27.4 41.2 4.3	41.4 5.9 13.9	371	1.0 0.0 0.316	51.1 79.1	29.7 84.5 20.6	
336	B63R_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	0.5 0.125 0.375	28.5 44.8	-14.1 47.0 342.4	14.8 348	1.0 0.0 0.683	53.5 85.4	-19.9 87.7 346.8	
337	B50R_050_037a	0.5 0.125 0.5	0.5 0.375 0.312	330	0.5 0.124 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 0.0 1.0	57.2 94.3	-58.4 110.9 328.2	
338	B38R_062_050a	0.5 0.125 0.625	0.625 0.5 0.375	316	0.508 0.125 0.625	35.8 43.2	-37.0 56.9 319.4	0.5 0.125 0.625	32.1 55.3	-46.8 72.5 319.7	16.0 317	0.766 0.0 1.0	47.9 86.4	-74.0 113.8 319.4	
339	B30R_075_062a	0.5 0.125 0.75	0.75 0.625 0.437	307	0.51 0.125 0.75	38.4 51.4	-52.0 73.1 314.6	0.5 0.125 0.75	34.5 61.7	-61.2 82.9 315.2	14.3 307	0.616 0.0 1.0	42.4 82.3	-83.2 117.0 314.6	
340	B25R_087_075a	0.5 0.125 0.875	0.875 0.75 0.5	300	0.5 0.125 0.875	40.8 59.8	-67.2 90.0 311.6	0.5 0.125 0.875	37.2 68.3	-74.6 101.2 312.4	11.8 300	0.5 0.0 1.0	38.5 79.8	-89.7 120.0 311.6	
341	B20R_100_087a	0.5 0.125 1.0	1.0 0.875 0.562	295	0.489 0.125 1.0	43.6 68.8	-81.8 106.9 310.0	0.5 0.125 1.0	40.1 75.2	-87.1 115.1 310.7	9.0 294	0.416 0.0 1.0	36.3 78.6	-93.5 122.2 310.0	
342	R50Y_050_050a	0.5 0.25 0.0	0.5 0.5 0.25	60	0.5 0.25 0.0	31.8 20.6	35.5 41.1 59.7	0.5 0.25 0.0	32.3 22.9	42.9 48.6 61.8	7.7 59	1.0 0.5 0.0	63.6 41.3	71.0 82.2 59.7	
343	R31Y_050_037a	0.5 0.25 0.125	0.5 0.375 0.312	49	0.5 0.243 0.124	33.0 22.7	25.2 33.9 47.9	0.5 0.25 0.125	32.5 23.9	30.0 38.4 51.4	4.9 48	1.0 0.316 0.0	56.2 60.6	67.2 90.5 47.9	
344	R00Y_050_025a	0.5 0.25 0.25	0.5 0.25 0.375	390	0.5 0.249 0.249	36.4 19.2	16.1 25.1 40.0	0.5 0.25 0.25	33.0 26.3	12.1 29.0 24.7	8.8 389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0	
345	R00Y_050_025a	0.5 0.25 0.375	0.5 0.25 0.375	360	0.5 0.249 0.375	36.8 20.2	1.0 20.3 2.9	0.5 0.25 0.375	33.9 20.3	-6.0 30.9 348.7	12.6 360	1.0 0.0 0.5	52.0 81.1	4.1 81.2 2.9	
346	B50R_050_025a	0.5 0.25 0.5	0.5 0.25 0.375	330	0.5 0.249 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 0.0 1.0	57.2 94.3	-58.4 110.9 328.2	
347	B34R_062_037a	0.5 0.25 0.625	0.625 0.375 0.437	311	0.506 0.25 0.625	40.6 31.1	-29.7 43.3 316.7	0.5 0.25 0.625	36.8 42.2	-33.9 57.6 317.0	14.8 311	0.683 0.0 1.0	44.8 84.1	-79.2 115.5 316.7	
348	B25R_075_050a	0.5 0.25 0.75	0.75 0.5 0.300	0.5	0.5 0.25 0.75	43.1 39.9	-44.8 60.0 311.6	0.5 0.25 0.75	38.8 49.3	-54.2 73.3 312.3	13.9 300	0.5 0.0 1.0	38.5 79.8	-89.7 120.0 311.6	
349	B19R_087_062a	0.5 0.25 0.875	0.875 0.625 0.293	0.489	0.489 0.25 0.875	45.9 48.8	59.4 76.9 309.3	0.5 0.25 0.875	41.1 56.9	-68.1 88.8 309.8	12.8 292	0.383 0.0 1.0	35.3 78.1	-95.1 123.0 309.3	
350	B15R_100_075a	0.5 0.25 1.0	1.0 0.75 0.625	289	0.487 0.25 1.0	49.3 58.1	-73.1 93.4 308.4	0.5 0.25 1.0	43.7 64.7	-81.2 103.8 308.5	11.8 288	0.316 0.0 1.0	33.9 77.4	-97.5 124.5 308.4	
351	R76Y_050_050a	0.5 0.375 0.0	0.5 0.5 0.25	76	0.5 0.383 0.0	39.1 3.9	40.3 40.5 84.4	0.5 0.375 0.0	40.2 4.9	48.0 48.2 84.1	7.8 77	1.0 0.766 0.0	78.2 7.8	80.6 81.0 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	0.5 0.375 0.125	40.3 5.9	38.1 38.6 81.1	9.1 71	1.0 0.683 0.0	73.4 18.5	77.6 79.8 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	0.5 0.375 0.25	40.7 8.3	22.2 23.7 69.3	5.0 59	1.0 0.5 0.0	63.6 41.3	71.0 82.2 59.7	
354	R00Y_050_012a	0.5 0.375 0.375	0.5 0.125 0.437	390	0.5 0.375 0.375	42.0 9.6	8.0 12.5 40.0	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	64.5 100.4 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	0.5 0.375 0.5	42.3 18.0	-12.2 21.8 325.7	8.0 330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2	
356	B25R_062_025a	0.5 0.375 0.625	0.625 0.25 0.5	300	0.5 0.375 0.625	45.4 19.9	-22.4 30.0 311.6	0.5 0.375 0.625	43.6 24.8	-28.6 37.9 311.0	8.0 300	0.5 0.0 1.0	38.5 79.8	-89.7 120.0 311.6	
357	B15R_075_037a	0.5 0.375 0.75	0.75 0.375 0.562	289	0.493 0.375 0.75	48.5 29.0	-36.5 46.7 308.4	0.5 0.375 0.75	45.2 32.6	-44.0 54.7 306.5	8.8 288	0.316 0.0 1.0	33.9 77.4	-97.5 124.5 308.4	
358	B11R_087_050a	0.5 0.375 0.875	0.875 0.5 0.625	284	0.491 0.375 0.875	51.9 38.3	-50.0 63.1 307.4	0.5 0.375 0.875	47.1 40.9	-58.5 71.4 304.9	10.0 282	0.233 0.0 1.0	32.3 76.7	-100.1 126.2 307.4	
359	B09R_100_062a	0.5 0.375 1.0	1.0 0.625 0.687	281	0.489 0.375 1.0	55.6 47.8	-63.2 79.3 307.0	0.5 0.375 1.0	49.2 49.5	-72.2 87.6 304.4	11.1 279	0.183 0.0 1.0	31.7 76.5	-101.2 126.9 307.0	
360	Y00G_050_050a	0.5 0.5 0.0	0.5 0.5 0.25	90	0.5 0.5 0.0	46.3 10.3	45.3 46.5 102.8	0.5 0.5 0.0	48.9	-12.3 54.2 55.6	102.8 9.5 89	1.0 1.0 0.0	92.6	-20.7 90.7 93.0 102.8	
361	Y00G_050_037a	0.5 0.5 0.125	0.5 0.375 0.312	90	0.5 0.5 0.124	46.6	-7.7 34.0 34.9	102.8	0.5 0.5 0.125	49.1	-11.4 46.7 48.0	103.7 13.4 89	1.0 1.0 0.0	92.6	-20.7 90.7 93.0 102.8
362	Y00G_050_025a	0.5 0.5 0.25	0.5 0.25 0.375	90	0.5 0.5 0.249	47.0	-5.1 22.6 23.2	102.8	0.5 0.5 0.25	49.3	-9.2 32.9 34.2	105.6 11.3 89	1.0 1.0 0.0	92.6	-20.7 90.7 93.0 102.8
363	Y00G_050_012a	0.5 0.5 0.375	0.5 0.125 0.437	90	0.5 0.5 0.375	47.3	-2.5 11.3 11.6	102.8	0.5 0.5 0.375	49.8	-5.3 16.6 17.5	107.8 6.5 89	1.0 1.0 0.0	92.6	-20.7 90.7 93.0 102.8
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	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 0.0 0.0	
365	B00R_062_012a	0.5 0.625 0.625	0.625 0.125 0.625	270	0.5 0.5 0.625	51.5 9.5	-12.9 16.0 306.2	0.5 0.625 0.5	51.6 6.7	-16.3 17.6 292.4	4.3 270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5 306.2	
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	0.5 0.5 0.75	52.8 14.4	-31.9 35.1 294.3	7.9 270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5 306.2	
367	B00R_087_037a	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.5 0.875	59.1 28.5	-38.8 48.1 306.2	0.5 0.5 0.875	54.3 23.0	-46.9 52.2 296.5	10.8 270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5 306.2	
368	B00R_100_050a	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 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 69.0 297.1	13.0 270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5 306.2	
369	Y18G_062_062a	0.5 0.625 0.0	0.625 0.625 0.312	101	0.51 0.625 0.0	55.9	-24.1 54.5 59.6	113.8	0.5 0.625 0.0	58.1	-27.9 61.0 67.1	114.5 7.8 99	0.816 1.0 0.0	89.5	-38.6 87.2 95.4 113.8
370	Y23G_062_050a	0.5 0.625 0.125	0.625 0.5 0.375	104	0.508 0.625 0.125	56.3	-21.6 43.1 48.2	116.6	0.5 0.625 0.125	58.2	-27.1 55.1 61.4	116.1 13.3 102	0.766 1.0 0.0	88.7	-43.3 86.2 96.5 116.6
371	Y31G_062_037a	0.5 0.625 0.25	0.625 0.375 0.437	109	0.506 0.625 0.25	56.7	-19.0 31.8 37.1	120.8	0.5 0.625 0.25	58.4	-25.1 43.4 50.2	120.0 13.2 108	0.683 1.0 0.0	87.6	-50.7 84.9 98.9 120.8
372	Y50G_062_025a	0.5 0.625 0.375	0.625 0.25 0.5	120	0.5 0.625 0.375	57.2	-16.3 20.6 26.2	128.3	0.5 0.625 0.375	58.8	-21.6 28.5 35.8	127.2 9.7 119	0.5 1.0 0.0	85.7	-65.2 82.4 105.1 128.3
373	G00B_062_012a														

vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF /.PS>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md				
405	R00Y_062_062a	0.625 0.0 0.0	0.625 0.625 0.312	390	0.625 0.0 0.0	31.5 48.0 40.3	62.7 40.0	0.625 0.0 0.0	30.7 54.1 44.5	70.1 39.4 7.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0		
406	R31Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.114	31.7 48.7 29.7	57.0 31.3	0.625 0.1125 0.114	31.0 54.7 30.0	62.4 28.7 6.0	380	1.0 0.0 0.183	50.7 70.7 47.5	91.2 31.3		
407	R11Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.239	32.1 49.6 12.8	51.3 14.4	0.625 0.0 0.235	31.5 56.2 10.9	57.2 11.0 6.7	367	1.0 0.0 0.383	51.4 79.5 20.4	82.1 14.4		
408	B69R_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.385	33.0 52.2 -7.1	52.7 35.2	0.625 0.0 0.375	32.4 58.6 -7.7	59.1 35.2 6.4	352	1.0 0.0 0.616	52.9 83.6 -11.4	84.3 35.2		
409	B59R_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.625 0.0 0.51	34.3 55.5 -22.8	60.1 337.6	0.625 0.0 0.5	33.8 62.1 -25.0	67.0 338.0 6.9	339	1.0 0.0 0.816	54.9 88.9 -36.6	96.2 337.6		
410	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	35.8 58.9 -36.5	69.3 328.2	0.625 0.0 0.625	35.5 66.4 -41.1	78.1 328.2 8.7	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2		
411	B42R_075_075a	0.625 0.0 0.75	0.75 0.75 0.375	321	0.637 0.0 0.75	38.4 66.8 -51.4	84.3 322.4	0.625 0.0 0.75	37.6 71.3 -55.9	90.6 312.8 6.4	322	0.85 0.0 1.0	51.2 89.1 -68.5	112.4 322.4		
412	B36R_087_087a	0.625 0.0 0.875	0.875 0.875 0.437	314	0.641 0.0 0.875	40.8 74.7 -66.6	100.1 318.3	0.625 0.0 0.875	40.0 76.7 -69.8	103.7 317.7 3.8	315	0.733 0.0 1.0	46.6 85.4 -76.1	114.4 318.3		
413	B31R_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	43.0 82.7 -82.2	116.6 315.1	0.625 0.0 1.0	42.7 82.5 -82.8	116.8 314.8 0.6	308	0.633 0.0 1.0	43.0 82.7 -82.2	116.6 315.1		
414	R18Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	41	0.625 0.114 0.0	32.9 44.0 40.9	60.1 42.8	0.625 0.125 0.0	32.8 48.2 45.9	66.6 6.5 3.9	379	1.0 0.183 0.0	52.7 70.5 65.5	96.2 42.8		
415	R00Y_062_050a	0.625 0.125 0.125	0.625 0.5 0.375	390	0.625 0.125 0.125	37.1 38.4 32.2	50.2 40.0	0.625 0.125 0.125	33.0 48.8 32.2	58.5 33.3 11.1	379	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0		
416	R26Y_062_050a	0.625 0.125 0.25	0.625 0.5 0.375	376	0.625 0.125 0.241	37.3 39.0 20.6	44.1 27.8	0.625 0.125 0.25	33.5 50.4 13.6	52.2 15.1 13.9	387	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8		
417	R00Y_062_050a	0.625 0.125 0.375	0.625 0.5 0.375	360	0.625 0.125 0.375	37.9 40.5 2.0	40.6 2.9	0.625 0.125 0.375	34.4 53.1 -4.8	53.3 354.8 14.7	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9		
418	B61R_062_050a	0.625 0.125 0.5	0.625 0.5 0.375	344	0.625 0.125 0.508	39.1 43.6 -15.3	46.2 340.6	0.625 0.125 0.5	35.6 56.7 -22.2	60.9 338.6 15.2	342	1.0 0.0 0.766	54.4 87.3 -30.6	92.5 340.6		
419	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	40.5 47.1 -29.2	55.4 328.2	0.625 0.125 0.625	37.3 61.3 -38.3	72.3 327.9 17.2	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2		
420	B40R_075_062a	0.625 0.125 0.75	0.75 0.625 0.437	319	0.635 0.125 0.75	43.1 55.0 -44.2	70.6 321.2	0.625 0.125 0.75	39.2 66.6 -53.4	85.3 321.2 15.2	320	0.816 0.0 1.0	49.8 88.1 -70.7	113.0 321.2		
421	B34R_087_075a	0.625 0.125 0.875	0.875 0.75 0.5	311	0.637 0.125 0.875	45.5 63.1 -59.4	86.6 316.7	0.625 0.125 0.875	41.5 72.3 -67.4	98.9 317.0 12.9	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7		
422	B29R_100_087a	0.625 0.125 1.0	1.0 0.875 0.562	305	0.635 0.125 1.0	48.0 71.4 -74.4	103.2 313.8	0.625 0.125 1.0	44.0 78.4 -80.5	112.4 314.2 10.1	305	0.583 0.0 1.0	41.3 81.6 -85.1	117.9 313.8		
423	R38Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	53	0.625 0.239 0.0	36.6 34.0 42.6	54.6 51.3	0.625 0.25 0.0	37.4 35.7 48.5	60.2 53.5 6.1	52	1.0 0.233 0.0	58.5 54.5 68.2	87.3 51.3		
424	R23Y_062_050a	0.625 0.25 0.125	0.625 0.5 0.375	44	0.625 0.241 0.125	38.8 33.8 32.9	47.2 44.2	0.625 0.25 0.125	37.6 36.4 36.8	51.8 45.2 4.8	42	1.0 0.383 0.0	57.7 67.6 65.8	94.4 44.2		
425	R00Y_062_037a	0.625 0.25 0.25	0.625 0.375 0.437	390	0.625 0.25 0.25	42.7 28.8 24.2	37.6 40.0	0.625 0.25 0.25	38.0 38.2 19.6	42.9 27.1 11.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0		
426	R18Y_062_037a	0.625 0.25 0.375	0.625 0.375 0.437	371	0.625 0.25 0.368	43.0 29.6 11.1	31.7 20.6	0.625 0.25 0.375	38.7 41.1 1.5	41.1 21.1 15.5	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6		
427	B65R_062_037a	0.625 0.25 0.5	0.625 0.375 0.437	349	0.625 0.25 0.506	43.9 32.0 -7.4	32.9 346.8	0.625 0.25 0.5	39.8 45.1 -15.7	47.8 340.7 16.0	348	1.0 0.0 0.683	53.5 85.4 -19.9	87.7 346.8		
428	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	45.3 35.3 -21.9	41.6 328.2	0.625 0.25 0.625	41.2 50.2 -32.1	59.6 327.4 18.5	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2		
429	B38R_075_050a	0.625 0.25 0.75	0.75 0.5 0.5	316	0.633 0.25 0.75	47.8 43.2 -37.0	56.9 319.4	0.625 0.25 0.75	42.9 56.0 -47.4	73.4 319.7 17.2	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4		
430	B30R_087_062a	0.625 0.25 0.875	0.875 0.625 0.562	307	0.635 0.25 0.875	50.3 51.4 -52.0	73.1 314.6	0.625 0.25 0.875	44.9 62.4 -61.8	87.9 315.2 15.7	307	0.616 0.0 1.0	42.4 82.3 -83.2	117.0 314.6		
431	B25R_100_075a	0.625 0.25 1.0	1.0 0.75 0.625	300	0.625 0.25 1.0	52.8 59.8 -67.2	90.0 314.6	0.625 0.25 1.0	47.2 69.2 -75.4	102.3 312.5 13.5	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 314.6		
432	R61Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	67	0.625 0.385 0.0	43.5 16.7 46.8	49.7 70.2	0.625 0.375 0.0	44.1 19.3 52.4	55.9 69.7 6.2	67	1.0 0.616 0.0	69.6 26.8 74.8	79.5 70.2		
433	R50Y_062_050a	0.625 0.375 0.125	0.625 0.5 0.375	60	0.625 0.375 0.125	43.7 20.6 35.5	41.1 59.7	0.625 0.375 0.125	44.2 20.0 43.2	47.6 65.1 7.7	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7		
434	R31Y_062_037a	0.625 0.375 0.25	0.625 0.375 0.437	49	0.625 0.368 0.25	44.9 22.7 25.2	33.9 47.9	0.625 0.375 0.25	44.5 21.8 27.8	35.4 51.9 2.8	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9		
435	R00Y_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	48.4 19.2 16.1	25.1 20.1	0.625 0.375 0.375	45.1 24.9 10.6	27.0 23.1 8.5	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0		
436	R00Y_062_025a	0.625 0.375 0.5	0.625 0.25 0.5	360	0.625 0.375 0.5	48.7 20.2 1.0	20.3 2.9	0.625 0.375 0.5	46.0 29.2 -6.4	29.9 347.5 11.9	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9		
437	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	50.1 23.5 -14.6	27.7 328.2	0.625 0.375 0.625	47.1 34.6 -22.9	41.5 326.5 14.1	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2		
438	B34R_075_037a	0.625 0.375 0.75	0.75 0.375 0.562	311	0.631 0.375 0.75	52.5 31.5 -29.7	43.3 316.7	0.625 0.375 0.75	48.5 40.9 -38.5	56.2 316.7 13.5	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7		
439	B25R_087_050a	0.625 0.375 0.875	0.875 0.5 0.625	300	0.625 0.375 0.875	55.0 39.9 -44.8	60.0 311.6	0.625 0.375 0.875	50.2 47.9 -53.3	71.7 311.9 12.6	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6		
440	B19R_100_062a	0.625 0.375 1.0	1.0 0.625 0.687	293	0.614 0.375 1.0	57.8 48.8 -59.4	76.9 309.3	0.625 0.375 1.0	52.1 55.3 -67.3	87.1 309.4 11.7	292	0.383 0.0 1.0	35.3 78.1 -95.1	123.0 309.3		
441	R81Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	79	0.625 0.51 0.0	50.8 1.0 51.8	51.8 88.7	0.625 0.5 0.0	51.9 1.9 57.7	57.8 88.0 6.0	80	1.0 0.816 0.0	81.2 1.7 82.9	83.0 88.7		
442	R76Y_062_050a	0.625 0.5 0.125	0.625 0.5 0.375	76	0.625 0.508 0.125	51.0 3.9 40.3	40.5 84.4	0.625 0.5 0.125	52.0 2.6 50.5	50.6 86.9 10.3	77	1.0 0.766 0.0	73.2 78.8 80.6	81.0 84.4		
443	R68Y_062_037a	0.625 0.5 0.25	0.625 0.375 0.437	71	0.625 0.506 0.25	51.3 6.9 29.1	29.9 76.5	0.625 0.5 0.25	52.3 4.4 37.1	37.4 83.2 8.4	71	1.0 0.683 0.0	72.4 18.5 77.6	79.8 76.5		
444	R50Y_062_025a	0.625 0.5 0.375	0.625 0.25 0.5	60	0.625 0.5 0.375	51.6 10.3 17.7	20.5 59.7	0.625 0.5 0.375	52.8 7.4 21.1	22.3 70.5 4.5	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7		
445	R00Y_062_012a	0.625 0.5 0.625	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	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0		
446	B50R_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	54.8 11.7 -7.3	13.8 328.2	0.625 0.5 0.625	54.4 17.2 -11.8	20.9 325.5 7.1	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2		
447	B25R_075_025a	0.625 0.5 0.75	0.75 0.25 0.625	300	0.625 0.5 0.75	57.3 19.9 -22.4	30.0 311.6	0.625 0.5 0.75	55.5 23.7 -27.6	36.4 310.7 6.6	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6		
448	B15R_087_037a	0.625 0.5 0.875	0.875 0.375 0.687	289	0.618 0.5 0.875	60.4 29.0 -36.5	46.7 308.4	0.625 0.5 0.875	56.9 31.0 -42.7	52.8 305.9 7.3	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4		
449	B11R_100_050a	0.625 0.5 1.0	1.0 0.5 0.75	284	0.616 0.5 1.0	63.9 38.3 -50.0	63.1 307.4	0.625 0.5 1.0	58.5 38.8 -57.1	69.0 304.2 8.8	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4		
450	Y00G_062_062a	0.625 0.625 0.0	0.625 0.625 0.312	90	0.625 0.625 0.0	57.9 -12.9	56.7 58.1	0.625 0.625 0.0	60.4 -14.5	63.8 65.4	102.8 7.7	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
451	Y00G_062_050a	0.625 0.625 0.125	0.625 0.5 0.375	90	0.625 0.625 0.125	58.2 -10.3	45.3 46.5	0.625 0.625 0.125	60.5 -13.9	58.1 59.7	103.4 13.4	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
452	Y00G_062_037a	0.625 0.625 0.25	0.625 0.375 0.437	90	0.625 0.625 0.25	58.5 -7.7	34.0 34.9	0.625 0.625 0.25	60.7 -12.2	46.6 48.2	104.7 13.5	89	1.0 1.0 0.0	92.6 -20		

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Ma	rgb*Ma	LabCh*Ma		
486	R00Y_075_075a	0.75 0.0 0.0	0.75 0.75 0.375	390	0.75 0.0 0.0	37.8 57.7 48.4	75.3 40.0	0.75 0.0 0.0	37.5 61.9 51.9	80.8 39.9 5.5	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
487	R35Y_075_075a	0.75 0.0 0.125	0.75 0.75 0.375	381	0.75 0.0 0.112	37.9 58.2 38.8	69.9 33.6	0.75 0.0 0.125	37.7 62.4 38.9	73.5 31.9 4.2	382	1.0 0.0 0.15	50.6 77.6 51.7	93.3 33.6
488	R18Y_075_075a	0.75 0.0 0.25	0.75 0.75 0.375	371	0.75 0.0 0.237	38.3 59.3 22.3	63.4 20.6	0.75 0.0 0.25	38.1 63.5 20.8	66.9 18.1 4.4	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6
489	R00Y_075_075a	0.75 0.0 0.375	0.75 0.75 0.375	360	0.75 0.0 0.375	39.0 60.8 3.1	60.9 2.9	0.75 0.0 0.375	38.8 65.5 2.4	65.5 2.1 4.6	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
490	B65R_075_075a	0.75 0.0 0.5	0.75 0.75 0.375	349	0.75 0.0 0.512	40.1 64.1 -14.9	65.8 346.8	0.75 0.0 0.5	39.9 68.2 -15.1	69.9 347.4 4.1	348	1.0 0.0 0.683	53.5 85.4 -19.9	87.7 346.8
491	B57R_075_075a	0.75 0.0 0.625	0.75 0.75 0.375	339	0.75 0.0 0.637	41.5 67.3 -30.5	73.9 335.5	0.75 0.0 0.625	41.3 71.8 -31.6	78.4 336.2 4.5	337	1.0 0.0 0.85	55.3 89.8 -40.7	98.6 335.5
492	B50R_075_075a	0.75 0.0 0.75	0.75 0.75 0.375	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 1.0	57.2 84.3 -58.4	110.9 328.2
493	B43R_087_087a	0.75 0.0 0.875	0.875 0.875 0.437	322	0.758 0.0 0.875	45.3 78.4 -59.0	98.1 323.0	0.75 0.0 0.875	45.0 80.7 -61.5	101.5 326.4 3.4	322	0.866 0.0 1.0	51.8 94.6 -67.4	112.1 323.0
494	B38R_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4	0.75 0.0 1.0	47.2 85.8 -75.1	114.1 318.8 1.3	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4
495	R15Y_075_075a	0.75 0.125 0.0	0.75 0.75 0.375	39	0.75 0.112 0.0	39.0 54.3 48.9	73.1 41.9	0.75 0.125 0.0	39.1 57.3 52.5	77.8 42.5 4.7	37	1.0 0.15 0.0	52.0 72.4 65.2	97.4 41.9
496	R00Y_075_062a	0.75 0.125 0.125	0.75 0.625 0.437	390	0.75 0.125 0.125	43.4 48.0 40.3	62.7 40.0	0.75 0.125 0.125	39.3 57.8 40.4	60.6 34.9 10.6	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
497	R31Y_075_062a	0.75 0.125 0.25	0.75 0.625 0.437	379	0.75 0.125 0.239	43.6 48.7 29.7	57.0 31.3	0.75 0.125 0.25	39.7 59.0 22.8	73.3 21.1 12.9	380	1.0 0.0 0.183	50.7 77.9 47.5	91.2 31.3
498	R11Y_075_062a	0.75 0.125 0.375	0.75 0.625 0.437	367	0.75 0.125 0.364	44.0 49.6 12.8	51.3 14.4	0.75 0.125 0.375	40.3 61.0 4.6	61.2 4.3 14.5	367	1.0 0.0 0.383	51.4 79.5 20.4	82.1 14.4
499	B69R_075_062a	0.75 0.125 0.5	0.75 0.625 0.437	353	0.75 0.125 0.51	45.0 52.2 -7.1	52.7 352.1	0.75 0.125 0.5	41.4 64.0 -12.9	65.2 348.5 13.5	352	1.0 0.0 0.616	52.9 83.6 -11.4	84.3 352.1
500	B59R_075_062a	0.75 0.125 0.625	0.75 0.625 0.437	341	0.75 0.125 0.635	46.2 55.5 -22.8	60.1 337.6	0.75 0.125 0.625	42.7 67.7 -29.4	73.8 336.4 14.2	339	1.0 0.0 0.816	54.9 88.9 -36.6	96.2 337.6
501	B50R_075_062a	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 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
502	B42R_087_075a	0.75 0.125 0.875	0.875 0.75 0.5	321	0.762 0.125 0.875	50.3 66.8 -51.4	84.3 322.4	0.75 0.125 0.875	46.2 77.0 -59.5	97.3 322.3 13.6	322	0.85 0.0 1.0	51.2 89.1 -68.5	112.4 322.4
503	B36R_100_087a	0.75 0.125 1.0	1.0 0.875 0.562	314	0.766 0.125 1.0	52.7 74.7 -66.6	100.1 318.3	0.75 0.125 1.0	48.4 82.4 -73.2	112.0 318.3 10.9	315	0.733 0.0 1.0	46.6 85.4 -76.1	114.4 318.3
504	R31Y_075_075a	0.75 0.25 0.0	0.75 0.75 0.375	49	0.75 0.237 0.0	42.2 45.5 50.4	67.9 47.9	0.75 0.25 0.0	42.8 47.1 54.2	71.8 49.0 4.2	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9
505	R18Y_075_062a	0.75 0.25 0.125	0.75 0.625 0.437	41	0.75 0.239 0.125	44.8 44.0 40.9	60.1 42.8	0.75 0.25 0.125	42.9 47.6 43.8	64.7 42.6 4.9	39	1.0 0.183 0.0	52.7 70.5 65.5	96.2 42.8
506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	49.0 38.4 32.2	50.2 40.0	0.75 0.25 0.25	43.3 48.9 27.4	56.0 29.2 12.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
507	K26Y_075_050a	0.75 0.25 0.375	0.75 0.5 0.5	376	0.75 0.25 0.366	49.2 39.0 20.6	44.1 27.8	0.75 0.25 0.375	43.9 51.1 9.6	52.0 10.6 17.1	377	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8
508	R00Y_075_050a	0.75 0.25 0.5	0.75 0.5 0.5	360	0.75 0.25 0.5	49.8 40.5 2.0	40.6 2.9	0.75 0.25 0.5	44.8 54.3 -7.7	54.8 35.1 17.6	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
509	B61R_075_050a	0.75 0.25 0.625	0.75 0.5 0.5	344	0.75 0.25 0.633	51.0 43.6 -15.3	46.2 340.6	0.75 0.25 0.625	46.0 58.3 -24.3	63.1 337.3 17.9	342	1.0 0.0 0.766	54.4 87.3 -30.6	92.5 340.6
510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	52.5 47.1 -29.2	55.4 328.7	0.75 0.25 0.75	47.5 63.1 -30.9	74.6 327.2 19.8	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
511	B40R_087_062a	0.75 0.25 0.875	0.875 0.625 0.562	319	0.76 0.25 0.875	55.0 55.0 -44.2	70.6 321.2	0.75 0.25 0.875	49.2 68.4 -54.7	87.6 321.3 18.0	320	0.816 0.0 1.0	42.8 88.1 -70.7	113.0 321.2
512	B34R_100_075a	0.75 0.25 1.0	1.0 0.75 0.625	311	0.762 0.25 1.0	57.4 63.1 -59.4	86.6 311.7	0.75 0.25 1.0	51.2 74.3 -68.7	101.2 317.2 15.0	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7
513	R50Y_075_075a	0.75 0.375 0.0	0.75 0.75 0.375	60	0.75 0.375 0.0	47.7 31.0 53.2	61.6 51.7	0.75 0.375 0.0	48.5 32.5 57.4	65.9 60.4 4.4	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
514	R38Y_075_062a	0.75 0.375 0.125	0.75 0.625 0.437	53	0.75 0.364 0.125	48.5 34.0 42.6	54.6 51.3	0.75 0.375 0.125	48.6 33.0 48.8	59.0 55.9 6.2	52	1.0 0.383 0.0	58.5 54.5 68.2	87.3 51.3
515	R23Y_075_050a	0.75 0.375 0.25	0.75 0.5 0.5	44	0.75 0.366 0.25	50.7 33.8 32.9	47.2 44.2	0.75 0.375 0.25	48.9 34.4 34.1	48.4 44.7 2.2	42	1.0 0.233 0.0	53.7 67.6 65.8	94.4 44.2
516	R00Y_075_037a	0.75 0.375 0.375	0.75 0.375 0.562	390	0.75 0.375 0.375	54.7 28.8 24.2	37.6 40.0	0.75 0.375 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	100.4 40.0
517	R18Y_075_037a	0.75 0.375 0.5	0.75 0.375 0.562	371	0.75 0.375 0.493	54.9 29.6 11.1	31.7 20.6	0.75 0.375 0.5	50.1 40.1 0.1	40.1 0.1 15.9	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6
518	B65R_075_037a	0.75 0.375 0.625	0.75 0.375 0.562	349	0.75 0.375 0.631	55.8 32.0 -7.4	32.9 346.8	0.75 0.375 0.625	51.1 44.4 -16.4	47.4 339.7 16.0	348	1.0 0.0 0.683	53.5 85.4 -19.9	87.7 346.8
519	B50R_075_037a	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	57.2 35.3 -21.9	41.6 328.2	0.75 0.375 0.75	52.4 49.6 -32.2	59.1 327.0 18.2	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
520	B38R_087_050a	0.75 0.375 0.875	0.875 0.5 0.625	316	0.758 0.375 0.875	59.7 43.2 -37.0	56.9 319.4	0.75 0.375 0.875	53.9 55.4 -47.2	72.8 319.5 16.9	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4
521	B30R_100_062a	0.75 0.375 1.0	1.0 0.625 0.687	307	0.76 0.375 1.0	62.2 51.4 -52.0	73.1 314.6	0.75 0.375 1.0	55.6 61.8 -61.5	87.2 315.1 15.5	307	0.616 0.0 1.0	42.4 82.3 -83.2	117.0 314.6
522	R68Y_075_075a	0.75 0.5 0.0	0.75 0.75 0.375	71	0.75 0.512 0.0	55.0 13.8 58.2	59.8 76.5	0.75 0.5 0.0	55.4 15.9 61.8	63.8 75.5 4.1	71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5
523	R61Y_075_062a	0.75 0.5 0.125	0.75 0.625 0.437	67	0.75 0.51 0.125	55.4 16.7 46.8	49.7 70.2	0.75 0.5 0.125	55.5 16.4 54.9	57.3 73.3 8.1	67	1.0 0.616 0.0	69.6 26.8 74.8	79.5 70.2
524	R50Y_075_050a	0.75 0.5 0.25	0.75 0.5 0.5	60	0.75 0.5 0.25	55.6 20.6 35.5	41.1 59.7	0.75 0.5 0.25	55.8 17.8 42.0	45.6 66.9 7.1	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
525	R31Y_075_037a	0.75 0.5 0.375	0.75 0.375 0.562	49	0.75 0.493 0.375	56.8 22.7 25.2	33.9 47.9	0.75 0.5 0.375	56.2 20.2 26.2	33.1 52.3 2.7	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9
526	R00Y_075_025a	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	0.75 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	100.4 40.0
527	R00Y_075_025a	0.75 0.5 0.625	0.75 0.25 0.625	360	0.75 0.5 0.625	60.7 20.2 1.0	20.3 2.9	0.75 0.5 0.625	57.6 28.2 -6.6	28.9 346.7 11.4	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
528	B50R_075_025a	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	62.0 23.5 -14.6	27.7 328.2	0.75 0.5 0.75	58.7 33.5 -22.4	40.4 326.2 13.1	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
529	B34R_087_037a	0.75 0.5 0.875	0.875 0.375 0.687	311	0.756 0.5 0.875	64.5 31.5 -29.7	43.3 316.7	0.75 0.5 0.875	60.0 39.7 -37.6	54.7 316.5 12.2	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7
530	B25R_100_050a	0.75 0.5 1.0	1.0 0.5 0.75	300	0.75 0.5 1.0	67.0 39.9 -44.8	60.0 311.6	0.75 0.5 1.0	61.4 46.4 -52.2	69.9 316.6 11.3	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
531	R85Y_075_075a	0.75 0.625 0.0	0.75 0.75 0.375	81	0.75 0.637 0.0	62.4 -1.8 63.2	63.2 91.7	0.75 0.625 0.0	63.2 -0.7 67.1	67.1 90.6 4.1	81	1.0 0.85 0.0	83.3 -2.5 84.2	84.3 91.7
532	R81Y_075_062a	0.75 0.625 0.125	0.75 0.625 0.437	79	0.75 0.635 0.125	62.7 1.0 51.8	51.8 88.7	0.75 0.625 0.125	63.3 -0.2 61.6	61.6 90.2 9.8	80	1.0 0.816 0.0	81.2 1.7 82.9	83.0 88.7
533	R76Y_075_050a	0.75 0.625 0.25	0.75 0.5 0.5	76	0.75 0.633 0.25	62.9 3.9 40.3	40.5 84.4	0.75 0.625 0.25	63.5 1.0 50.4	50.5 88.7 10.5	77	1.0 0.766 0.0	78.2 7.8 80.6	81.0 84.4
534	R68Y_075_037a	0.75 0.625 0.375	0.75 0.375 0.562	71	0.75 0.631 0.375	63.3 6.9 29.1	29.9 76.5	0.75 0.625 0.375	63.8 3.4 35.9	36.1 84.5 7.7	71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5
535	R50Y_075_025a	0.75 0.625 0.5	0											



vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF /.PS>
 informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
 la domanda per la misura di stampa di display, nessuna separazione
 TUB materiale: code=rh4t4

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Ma	rgb*Ma	LabCh*Ma		
567	R00Y_087_087a	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.0	44.1 67.3 56.4	87.8 40.0	0.875 0.0 0.0	44.1 69.5 58.3	90.8 39.9 2.9	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
568	R36Y_087_087a	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.116	44.2 67.7 47.1	82.5 34.8	0.875 0.0 0.125	44.2 69.9 47.2	84.3 34.0 2.2	382	1.0 0.0 0.133	50.6 77.3 53.9	94.4 34.8
569	R23Y_087_087a	0.875 0.0 0.25	0.875 0.875 0.437	374	0.875 0.0 0.233	44.5 68.5 32.2	75.7 25.1	0.875 0.0 0.25	44.5 70.8 30.2	77.0 23.1 2.9	375	1.0 0.0 0.266	50.9 78.3 36.8	86.6 25.1
570	R08Y_087_087a	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.364	45.1 70.2 13.8	71.6 11.1	0.875 0.0 0.375	45.1 72.4 12.2	73.4 9.5 2.6	365	1.0 0.0 0.416	51.5 80.3 15.8	81.8 11.1
571	B70R_087_087a	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.51	46.1 72.8 7.6	60.0 73.0 355.2	0.875 0.0 0.5	46.0 74.6 5.3	74.8 355.8 1.9	354	1.0 0.0 0.583	52.7 83.2 6.9	83.5 355.2
572	B63R_087_087a	0.875 0.0 0.625	0.875 0.875 0.437	346	0.875 0.0 0.641	47.2 75.6 23.1	79.1 342.9	0.875 0.0 0.625	47.1 77.6 22.1	80.7 344.0 2.2	344	1.0 0.0 0.753	54.0 86.5 26.4	90.4 342.9
573	B56R_087_087a	0.875 0.0 0.75	0.875 0.875 0.437	338	0.875 0.0 0.758	48.6 78.8 37.5	87.3 334.5	0.875 0.0 0.75	48.5 81.2 37.9	89.6 334.9 2.3	337	1.0 0.0 0.866	55.5 90.1 42.8	99.8 334.5
574	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	50.1 82.5 31.1	97.1 328.2	0.875 0.0 0.875	50.2 85.3 32.0	100.3 328.2 3.2	330	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2
575	B44R_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	52.5 90.1 66.3	111.9 323.6	0.875 0.0 1.0	52.1 89.8 66.9	112.0 323.3 0.7	323	0.883 0.0 1.0	52.5 90.1 66.3	111.9 323.6
576	R13Y_087_087a	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.116 0.0	45.2 64.2 56.9	85.8 41.5	0.875 0.125 0.0	45.3 65.8 58.8	88.3 41.7 2.5	37	1.0 0.133 0.0	51.7 73.4 65.0	98.0 41.5
577	R00Y_087_075a	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.125	49.7 57.7 48.4	75.3 40.0	0.875 0.125 0.125	45.5 66.2 48.3	81.9 36.0 9.5	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
578	R35Y_087_075a	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.237	49.9 58.2 38.8	69.9 33.6	0.875 0.125 0.25	45.8 67.1 31.8	74.3 25.3 12.0	382	1.0 0.0 0.15	50.6 77.6 51.7	93.3 33.6
579	R18Y_087_075a	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.362	50.2 59.3 22.3	63.4 20.6	0.875 0.125 0.375	46.4 68.8 13.9	70.2 11.4 13.2	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6
580	R00Y_087_075a	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.5	50.9 60.8 3.1	60.9 2.9	0.875 0.125 0.5	47.2 71.1 3.6	71.2 357.1 12.8	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
581	B65R_087_075a	0.875 0.125 0.625	0.875 0.75 0.5	349	0.875 0.125 0.637	52.1 64.1 14.9	65.8 346.8	0.875 0.125 0.625	48.3 74.2 20.3	76.9 344.6 12.0	348	1.0 0.0 0.683	53.5 85.4 19.9	87.7 346.8
582	B57R_087_075a	0.875 0.125 0.75	0.875 0.75 0.5	339	0.875 0.125 0.762	53.4 67.3 30.5	73.9 335.5	0.875 0.125 0.75	49.6 77.9 36.1	85.9 335.1 12.5	337	1.0 0.0 0.85	55.3 89.8 40.7	98.6 335.5
583	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	54.9 70.7 43.8	83.2 328.2	0.875 0.125 0.875	51.3 82.1 51.1	96.7 328.1 14.0	330	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2
584	B43R_100_087a	0.875 0.125 1.0	1.0 1.0 0.875	322	0.883 0.125 1.0	57.3 78.4 59.0	98.1 323.0	0.875 0.125 1.0	53.1 86.9 65.3	107.7 323.0 11.3	322	0.866 0.0 1.0	51.8 89.6 67.4	112.1 323.0
585	R26Y_087_087a	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.233 0.0	47.8 57.0 58.0	81.3 45.5	0.875 0.25 0.0	48.4 57.3 60.1	83.0 46.3 2.1	44	1.0 0.266 0.0	54.6 65.1 66.3	93.0 45.5
586	R15Y_087_075a	0.875 0.25 0.125	0.875 0.75 0.5	39	0.875 0.237 0.125	50.9 54.3 48.9	73.1 41.9	0.875 0.25 0.125	48.5 57.7 50.8	76.9 41.3 4.5	37	1.0 0.15 0.0	52.0 72.4 65.2	97.4 41.9
587	R00Y_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	55.4 48.0 40.3	62.7 40.0	0.875 0.25 0.25	48.8 58.7 35.3	68.5 31.0 13.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
588	R31Y_087_062a	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.364	55.5 48.7 29.7	57.0 31.3	0.875 0.25 0.375	49.3 60.4 17.9	63.0 16.5 17.7	380	1.0 0.0 0.183	50.7 77.9 47.5	91.2 31.3
589	R11Y_087_062a	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.489	55.9 49.6 12.8	51.3 14.4	0.875 0.25 0.5	50.1 63.0 0.6	63.0 0.5 18.9	367	1.0 0.0 0.383	51.4 79.5 20.4	82.1 14.4
590	B69R_087_062a	0.875 0.25 0.625	0.875 0.625 0.562	353	0.875 0.25 0.635	56.9 52.2 7.1	52.7 352.1	0.875 0.25 0.625	51.1 66.2 16.0	68.1 346.3 17.5	352	1.0 0.0 0.616	52.9 83.6 11.4	84.3 352.1
591	B59R_087_062a	0.875 0.25 0.75	0.875 0.625 0.562	341	0.875 0.25 0.76	58.2 55.5 22.8	60.1 337.6	0.875 0.25 0.75	52.3 70.2 31.9	77.1 335.5 18.1	339	1.0 0.0 0.816	54.9 88.9 36.6	96.2 337.6
592	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	59.6 58.9 36.5	69.3 328.2	0.875 0.25 0.875	53.8 74.7 47.0	88.3 327.8 19.8	330	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2
593	B42R_100_075a	0.875 0.25 1.0	1.0 1.0 0.75	325	0.887 0.25 1.0	62.2 66.8 51.4	84.1 324.4	0.875 0.25 1.0	55.6 79.8 67.3	100.7 322.4 17.6	322	0.85 0.0 1.0	51.2 89.1 68.5	112.4 322.4
594	R41Y_087_087a	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.364 0.0	52.5 44.4 60.6	75.1 53.7	0.875 0.375 0.0	53.2 44.5 62.6	76.8 54.5 2.1	54	1.0 0.416 0.0	60.0 50.7 69.3	85.9 53.7
595	R31Y_087_075a	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.362 0.125	54.1 45.5 50.4	67.9 47.9	0.875 0.375 0.125	53.3 44.9 54.7	70.8 50.6 4.4	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9
596	R18Y_087_062a	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.364 0.25	56.8 44.0 40.9	60.1 42.8	0.875 0.375 0.25	53.5 45.9 40.7	61.4 41.5 3.7	39	1.0 0.183 0.0	57.2 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	319	0.885 0.375 1.0	66.9 55.0 44.2	70.6 321.2	0.875 0.375 1.0	59.5 68.6 55.0	88.0 321.2 18.8	320	0.816 0.0 1.0	49.8 88.1 70.7	113.0 321.2
603	R58Y_087_087a	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.51 0.0	59.4 27.1 64.8	70.2 67.2	0.875 0.5 0.0	59.4 29.0 66.2	72.3 66.2 2.4	65	1.0 0.583 0.0	67.9 31.0 74.0	80.3 67.2
604	R50Y_087_075a	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.5 0.125	59.6 31.0 53.2	61.6 59.7	0.875 0.5 0.125	59.4 29.5 59.8	66.7 63.7 6.7	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	B63R_087_037a	0.875 0.5 0.75	0.875 0.375 0.687	349	0.875 0.5 0.756	67.7 32.0 7.4	32.9 346.8	0.875 0.5 0.75	62.3 43.5 16.6	46.5 339.0 15.6	348	1.0 0.0 0.683	53.5 85.4 19.9	87.7 346.8
610	B50R_087_037a	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	69.1 35.3 21.9	41.6 328.2	0.875 0.5 0.875	63.5 48.6 31.9	58.2 326.7 17.6	330	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2
611	B38R_100_050a	0.875 0.5 1.0	1.0 0.5 0.75	316	0.883 0.5 1.0	71.6 43.2 37.0	56.9 319.4	0.875 0.5 1.0	64.8 54.4 46.6	71.7 319.3 16.3	317	0.766 0.0 1.0	47.9 86.4 74.0	113.8 319.4
612	R73Y_087_087a	0.875 0.625 0.0	0.875 0.875 0.437	74	0.875 0.641 0.0	66.7 10.5 69.4	70.2 81.3	0.875 0.625 0.0	66.5 12.7 70.9	72.0 79.7 2.7	75	1.0 0.733 0.0	76.2 12.0 79.3	80.2 81.3
613	R68Y_087_075a	0.875 0.625 0.125	0.875 0.75 0.5	71	0.875 0.637 0.125	67.0 13.8 58.2	59.8 76.5	0.875 0.625 0.125	66.5 13.2 65.6	66.9 78.6 7.4	71	1.0 0.683 0.0	73.4 18.5 77.6	79.8

vedere dei file simili: http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF /.PS
informazioni tecniche: http://www.ps.bam.de o http://130.149.60.45/~farbmetrik

Table with columns: n, HIC*Fa, rgb_Fa, icf_Fa, hsi_Fa, rgb*Fa, LabCh*Fa, DE*Fa, hsiMd, rgb*Md, LabCh*Md. It contains a large grid of numerical data for various color and density measurements.

delta E** = 9.3

grafico TUB-QI41; codice di tinte: H*d=Y25Gd
colori e la differenza, ΔE*

immettere: rgb/cmyk -> rgbd
uscita: trasferire a rgbd

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione

TUB materiale: code=rhatha

vedere dei file simili: http://130.149.60.45/~farbmetrik/QI41/QI41.HTM informazioni tecniche: http://www.ps.bam.de o http://130.149.60.45/~farbmetrik

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS La domanda per la misura di stampa di display, nessuna separazione TUB materiale: code=rhath4

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

4-0032430-F0

QI410-7N, 25/29-F

delta E** = 7.3

grafico TUB-QI41; codice di tinte: H*d=Y25Gd colori e la differenza, ΔE*

immettere: rgb/cmyk -> rgbd uscita: trasferire a rgbd

vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF> / .PS
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
La domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rh4t4

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

delta E** = 8.7

grafico TUB-QI41; codice di tinte: H*d=Y25Gd
colori e la differenza, ΔE*

immettere: rgb/cmyk -> rgbd
uscita: trasferire a rgbd

http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF /.PS; uscita di trasferimento
N: nessun 3D-linearizzazione (OL) nel file (F) o PS-startup (S), pagina 27/29

vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41L0NP.PDF> / .PS
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

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

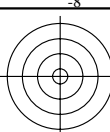
delta E** = 11.4

grafico TUB-QI41; codice di tinte: H*d=Y25Gd
colori e la differenza, ΔE*

immettere: rgb/cmyk -> rgbd
uscita: trasferire a rgbd

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione

TUB materiale: code=rh4ta

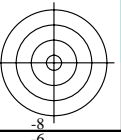
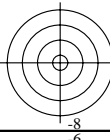


vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rh4ta

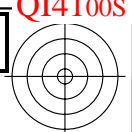
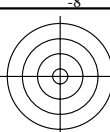
Table with columns: n, HIC*Fa, rgb*Fa, icf*Fa, hsi*Fa, rgbb*Fa, LabCh*Fa, DE*Fa, hsiMa, rgbb*Ma, LabCh*Ma. It contains a large grid of numerical data for various color and registration points.

delta E* = 1.6



colori e la differenza, ΔE^*
grafico TUB-QI41; codice di tinte: H*d=Y25Gd

immettere: $rgb/cmyk \rightarrow rgb_d$
uscita: trasferire a rgb_d



vedere dei file simili: <http://130.149.60.45/~farbmetrik/QI41/QI41.HTM>
informazioni tecniche: <http://www.ps.bam.de> o <http://130.149.60.45/~farbmetrik>

TUB iscrizione: 20130201-QI41/QI41L0NP.PDF /.PS
la domanda per la misura di stampa di display, nessuna separazione
TUB materiale: code=rh4ta

n	HIC*Fd	rgb_Fd	icf_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md	
1053	NW_086a	0.866 0.866 0.866	0.866 0.0	0.866 360	0.866 0.866 0.866	82.6 0.0 0.0	0.866 0.866 0.866	83.9 0.0 0.0	325.2 1.3	360	1.0 1.0 1.0	95.4 0.0 0.0	
1054	NW_093a	0.933 0.933 0.933	0.933 0.0	0.933 360	0.933 0.933 0.933	89.0 0.0 0.0	0.933 0.933 0.933	89.7 0.0 0.0	325.2 0.6	360	1.0 1.0 1.0	95.4 0.0 0.0	
1055	NW_100a	1.0 1.0 1.0	1.0 0.0	1.0 360	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	325.2 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0	
1056	NW_000a	0.0 0.0 0.0	0.0 0.0	0.0 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0	
1057	NW_006a	0.066 0.066 0.066	0.066 0.0	0.066 360	0.066 0.066 0.066	6.2 0.0 0.0	0.066 0.066 0.066	4.4 0.0 0.0	326.3 1.8	360	1.0 1.0 1.0	95.4 0.0 0.0	
1058	NW_013a	0.133 0.133 0.133	0.133 0.0	0.133 360	0.133 0.133 0.133	12.6 0.0 0.0	0.133 0.133 0.133	12.0 0.0 0.0	325.6 0.6	360	1.0 1.0 1.0	95.4 0.0 0.0	
1059	NW_020a	0.2 0.2 0.2	0.2 0.0	0.2 360	0.2 0.2 0.2	19.0 0.0 0.0	0.2 0.2 0.2	19.7 0.0 0.0	325.5 0.6	360	1.0 1.0 1.0	95.4 0.0 0.0	
1060	NW_026a	0.266 0.266 0.266	0.266 0.0	0.266 360	0.266 0.266 0.266	25.3 0.0 0.0	0.266 0.266 0.266	27.0 0.0 0.0	325.4 1.6	360	1.0 1.0 1.0	95.4 0.0 0.0	
1061	NW_033a	0.333 0.333 0.333	0.333 0.0	0.333 360	0.333 0.333 0.333	31.7 0.0 0.0	0.333 0.333 0.333	34.0 0.0 0.0	325.3 2.2	360	1.0 1.0 1.0	95.4 0.0 0.0	
1062	NW_040a	0.4 0.4 0.4	0.4 0.0	0.4 360	0.4 0.4 0.4	38.1 0.0 0.0	0.4 0.4 0.4	40.8 0.0 0.0	325.3 2.6	360	1.0 1.0 1.0	95.4 0.0 0.0	
1063	NW_046a	0.466 0.466 0.466	0.466 0.0	0.466 360	0.466 0.466 0.466	44.4 0.0 0.0	0.466 0.466 0.466	47.3 0.0 0.0	325.4 2.8	360	1.0 1.0 1.0	95.4 0.0 0.0	
1064	NW_053a	0.533 0.533 0.533	0.533 0.0	0.533 360	0.533 0.533 0.533	50.8 0.0 0.0	0.533 0.533 0.533	53.7 0.0 0.0	325.3 2.9	360	1.0 1.0 1.0	95.4 0.0 0.0	
1065	NW_060a	0.6 0.6 0.6	0.6 0.0	0.6 360	0.6 0.6 0.6	57.2 0.0 0.0	0.6 0.6 0.6	60.0 0.0 0.0	325.3 2.8	360	1.0 1.0 1.0	95.4 0.0 0.0	
1066	NW_066a	0.666 0.666 0.666	0.666 0.0	0.666 360	0.666 0.666 0.666	63.5 0.0 0.0	0.666 0.666 0.666	66.1 0.0 0.0	325.2 2.6	360	1.0 1.0 1.0	95.4 0.0 0.0	
1067	NW_073a	0.734 0.734 0.734	0.734 0.0	0.734 360	0.734 0.734 0.734	70.0 0.0 0.0	0.734 0.734 0.734	72.3 0.0 0.0	325.2 2.2	360	1.0 1.0 1.0	95.4 0.0 0.0	
1068	NW_080a	0.8 0.8 0.8	0.8 0.0	0.8 360	0.8 0.8 0.8	76.3 0.0 0.0	0.8 0.8 0.8	78.1 0.0 0.0	325.2 1.8	360	1.0 1.0 1.0	95.4 0.0 0.0	
1069	NW_086a	0.866 0.866 0.866	0.866 0.0	0.866 360	0.866 0.866 0.866	82.6 0.0 0.0	0.866 0.866 0.866	83.9 0.0 0.0	325.2 1.3	360	1.0 1.0 1.0	95.4 0.0 0.0	
1070	NW_093a	0.933 0.933 0.933	0.933 0.0	0.933 360	0.933 0.933 0.933	89.0 0.0 0.0	0.933 0.933 0.933	89.7 0.0 0.0	325.2 0.6	360	1.0 1.0 1.0	95.4 0.0 0.0	
1071	NW_100a	1.0 1.0 1.0	1.0 0.0	1.0 360	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	325.2 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0	
1072	NW_000a	0.0 0.0 0.0	0.0 0.0	0.0 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0	
1073	NW_100a	1.0 1.0 1.0	1.0 0.0	1.0 360	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	325.2 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0	
1074	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5	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
1075	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	86.8 -46.1 -13.5	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
1076	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	92.6 -20.7 90.7	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
1077	B00R_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	30.3 76.0 -103.5	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
1078	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	83.6 -82.7 79.8	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
1079	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	57.2 94.3 -58.4	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

delta E* = 1.0

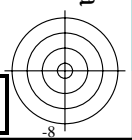
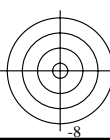


grafico TUB-QI41; codice di tinte: H*d=Y25Gd
colori e la differenza, ΔE*

immettere: rgb/cmyk -> rgbd
uscita: trasferire a rgbd