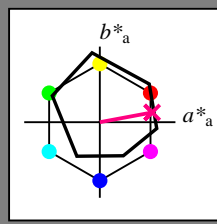


Input and Output: Offset Reflective System ORS18a for relative CIELAB hue $h_{ab,a,rel} = h_{ab}/360 = 10/360 = 0.02$

$H^*_- = B75R_-$

Data for any device (d) or elementary (e) colour:
 HIC^*_-

hue text for the colours of this page:
 $H^*_- = B75R_-$
triangle lightness T^*



ORS18a; adapted (a) CIELAB data

name	$L^*=L^*_a$	a^*_a	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
R _{-,Ma}	47.9	65.3	50.5	82.6	37
Y _{-,Ma}	90.3	-10.2	91.7	92.3	96
G _{-,Ma}	50.9	-62.8	34.9	71.9	150
C _{-,Ma}	58.6	-30.3	-45.0	54.2	236
B _{-,Ma}	25.7	31.0	-44.4	54.2	305
M _{-,Ma}	48.1	75.2	-8.3	75.7	353
N _{-,Ma}	18.0	0.0	0.0	0.0	0
W _{-,Ma}	95.4	0.0	0.0	0.0	0
R _{-,CIE}	39.9	58.7	27.9	65.0	25
Y _{-,CIE}	81.2	-2.8	71.5	71.6	92
G _{-,CIE}	52.2	-42.4	13.6	44.5	162
B _{-,CIE}	30.5	1.4	-46.4	46.4	271

Data for maximum colour (Ma):

$LabCh^*_{-,Ma}$: 48 69 12 70 10

$HIC^*_{-,Ma}$: B75R_100_100_

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

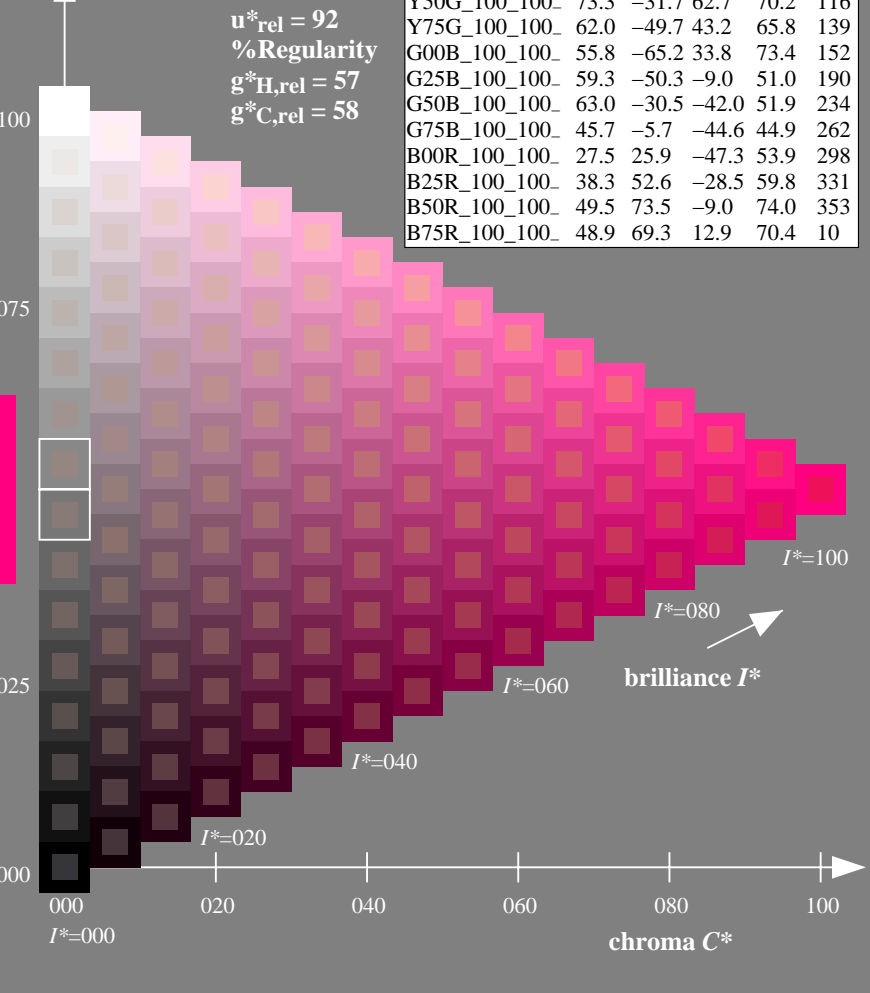
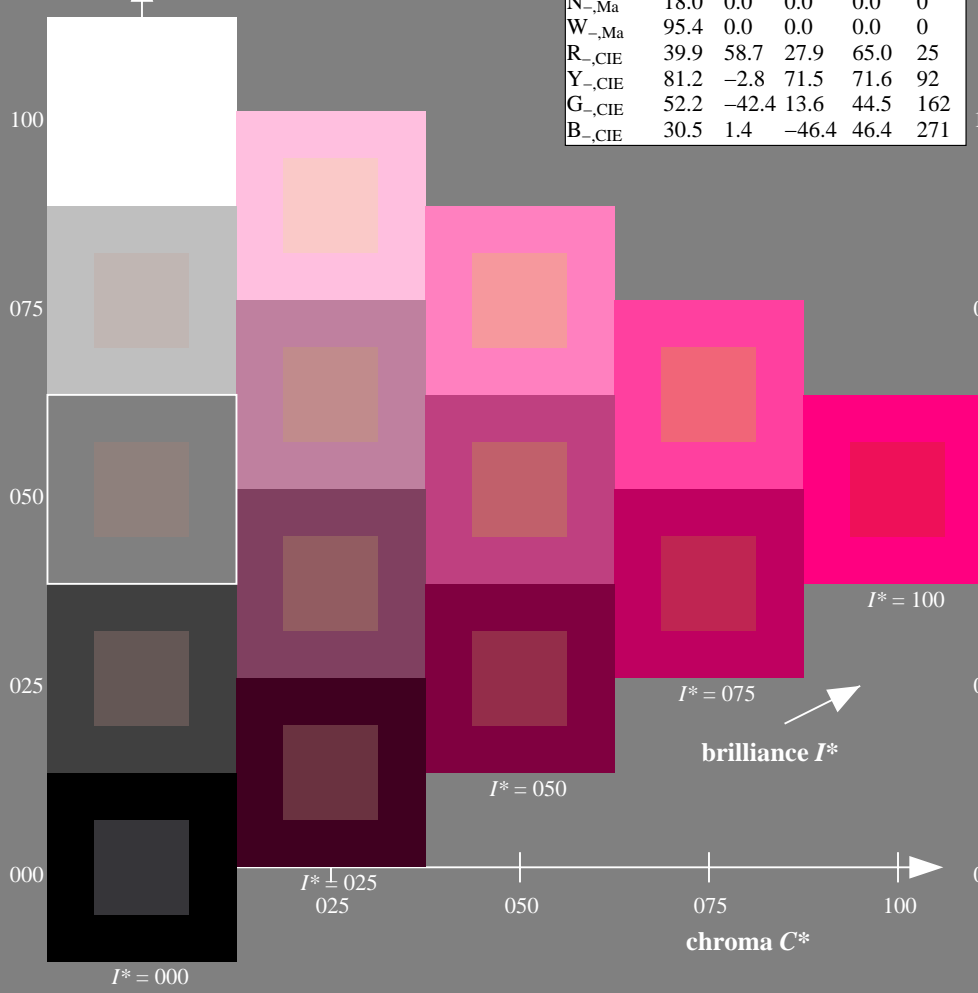
1.0 0.0 0.5 1.0 1.0

triangle lightness T^*

%Gamut
 $u^*_{rel} = 92$
%Regularity
 $g^*_{H,rel} = 57$
 $g^*_{C,rel} = 58$

ORS20a; adapted (a) CIELAB data

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	31
R25Y_100_100_	56.8	48.0	50.5	69.6	46
R50Y_100_100_	68.6	25.0	63.9	68.6	68
R75Y_100_100_	80.6	4.8	77.2	77.3	86
Y00G_100_100_	90.2	-9.6	88.2	88.7	96
Y25G_100_100_	83.2	-18.4	79.9	81.9	102
Y50G_100_100_	73.3	-31.7	62.7	70.2	116
Y75G_100_100_	62.0	-49.7	43.2	65.8	139
G00B_100_100_	55.8	-65.2	33.8	73.4	152
G25B_100_100_	59.3	-50.3	-9.0	51.0	190
G50B_100_100_	63.0	-30.5	-42.0	51.9	234
G75B_100_100_	45.7	-5.7	-44.6	44.9	262
B00R_100_100_	27.5	25.9	-47.3	53.9	298
B25R_100_100_	38.3	52.6	-28.5	59.8	331
B50R_100_100_	49.5	73.5	-9.0	74.0	353
B75R_100_100_	48.9	69.3	12.9	70.4	10



see similar files: <http://130.149.60.45/~farbmetrik/RE42/RE42.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20130201-RE42/RE42L0FA.TXT /PS
application for measurement of display output

TUB material: code=rh4ta

Input and Output: Television Luminous System TLS00a for relative CIELAB hue $h_{ab,a,rel} = h_{ab}/360 = 352/360 = 0.97$

$H^*_e = B75R_e$

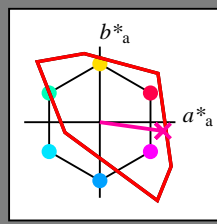
Data for any device (d) or elementary (e) colour:

HIC^*_e

hue text for the colours of this page:

$H^*_e = B75R_e$

triangle lightness T^*



TLS00a; adapted (a) CIELAB data

name	$L^*=L^*_a$	a^*_a	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
Re,Ma	50.9	78.3	37.3	86.7	25
Ye,Ma	83.7	-3.4	84.5	84.5	92
Ge,Ma	85.1	-64.6	20.7	67.9	162
Ce,Ma	79.0	-34.2	-25.7	42.8	216
Be,Ma	59.2	1.7	-56.6	56.6	271
Me,Ma	57.1	94.1	-57.4	110.3	328
Ne,Ma	0.0	0.0	0.0	0.0	0
We,Ma	95.4	0.0	0.0	0.0	0
Re,CIE	39.9	58.7	27.9	65.0	25
Ye,CIE	81.2	-2.8	71.5	71.6	92
Ge,CIE	52.2	-42.4	13.6	44.5	162
Be,CIE	30.5	1.4	-46.4	46.4	271

Data for maximum colour (Ma):

$LabCh^*_{e, Ma}: 52\ 83\ -11\ 84\ 352$

$HIC^*_{e, Ma}: B75R_100_100_e$

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

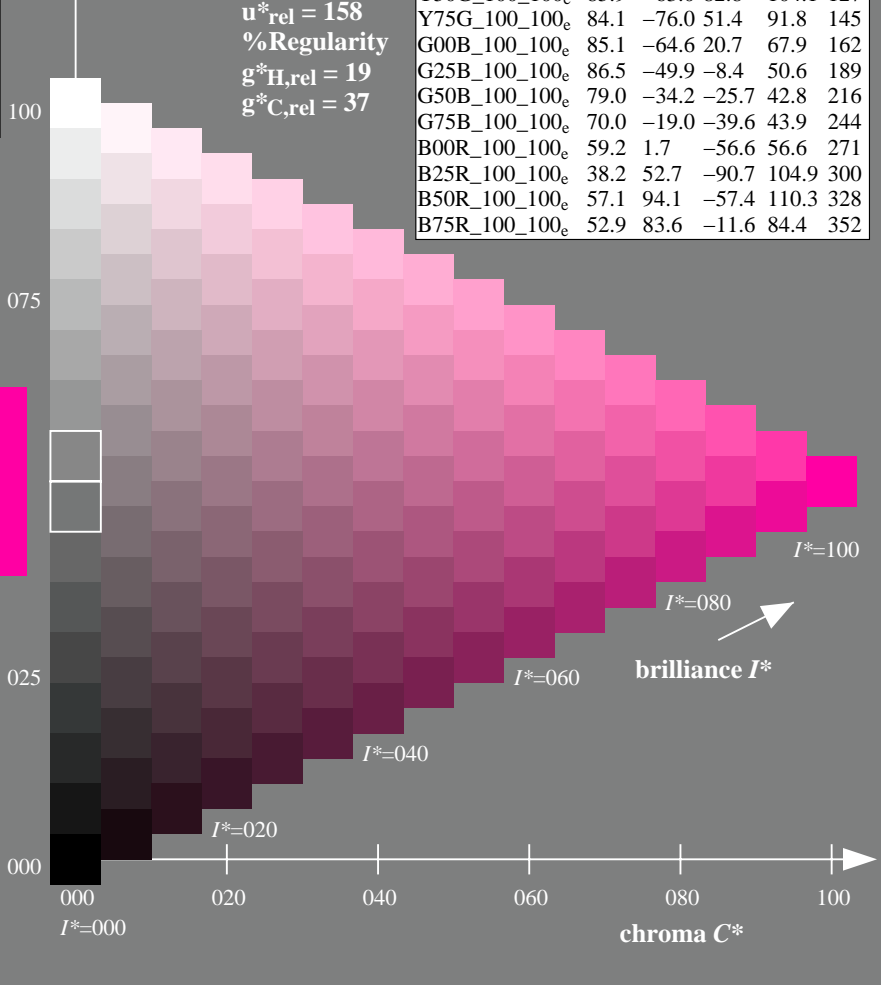
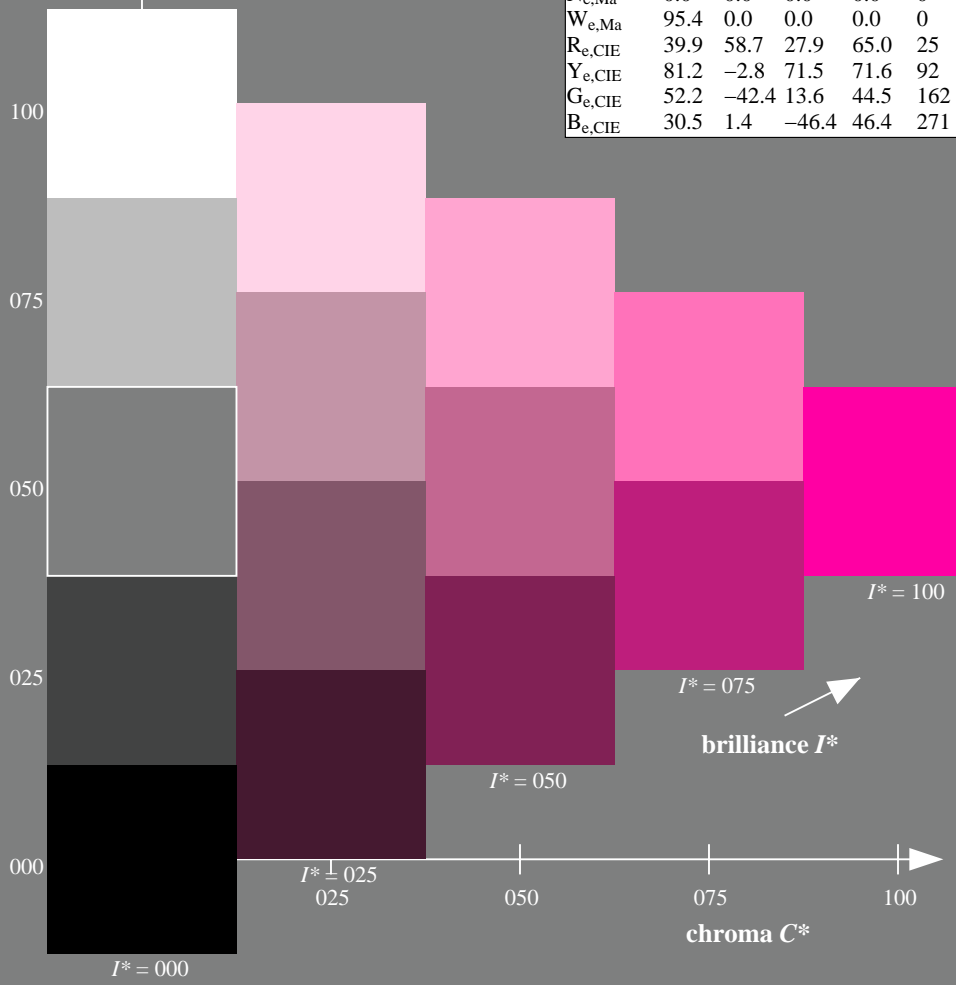
1.0 0.0 0.61 1.0 1.0

triangle lightness T^*

TLS00a; adapted (a) CIELAB data

H^*_e	$L^*=L^*_a$	a^*_a	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
R00Y_100_100_e	50.9	78.3	37.3	86.7	25
R25Y_100_100_e	51.3	74.4	64.8	98.7	41
R50Y_100_100_e	63.1	42.7	70.8	82.7	58
R75Y_100_100_e	73.5	18.3	77.7	79.8	76
Y00G_100_100_e	83.7	-3.4	84.5	84.5	92
Y25G_100_100_e	91.0	-29.9	88.9	93.8	108
Y50G_100_100_e	85.9	-63.0	82.8	104.1	127
Y75G_100_100_e	84.1	-76.0	51.4	91.8	145
G00B_100_100_e	85.1	-64.6	20.7	67.9	162
G25B_100_100_e	86.5	-49.9	-8.4	50.6	189
G50B_100_100_e	79.0	-34.2	-25.7	42.8	216
G75B_100_100_e	70.0	-19.0	-39.6	43.9	244
B00R_100_100_e	59.2	1.7	-56.6	56.6	271
B25R_100_100_e	38.2	52.7	-90.7	104.9	300
B50R_100_100_e	57.1	94.1	-57.4	110.3	328
B75R_100_100_e	52.9	83.6	-11.6	84.4	352

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

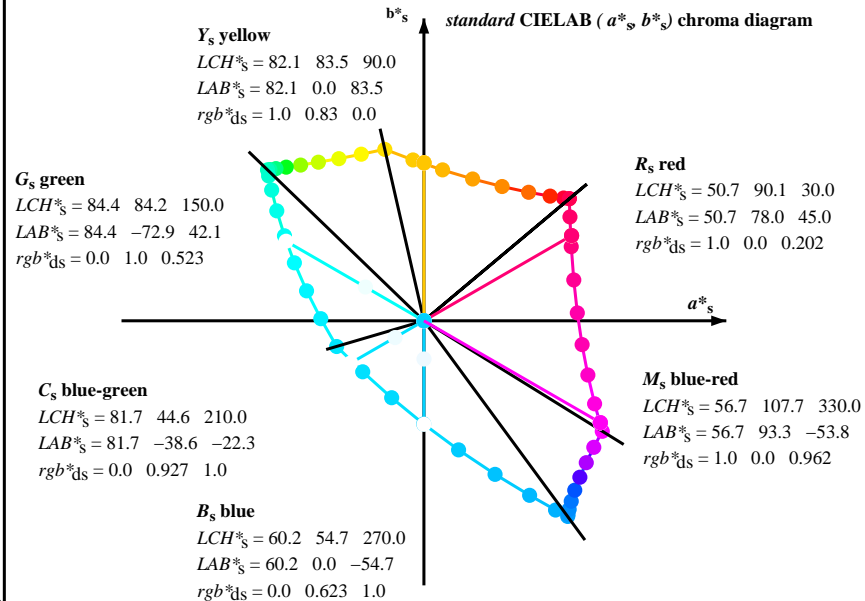
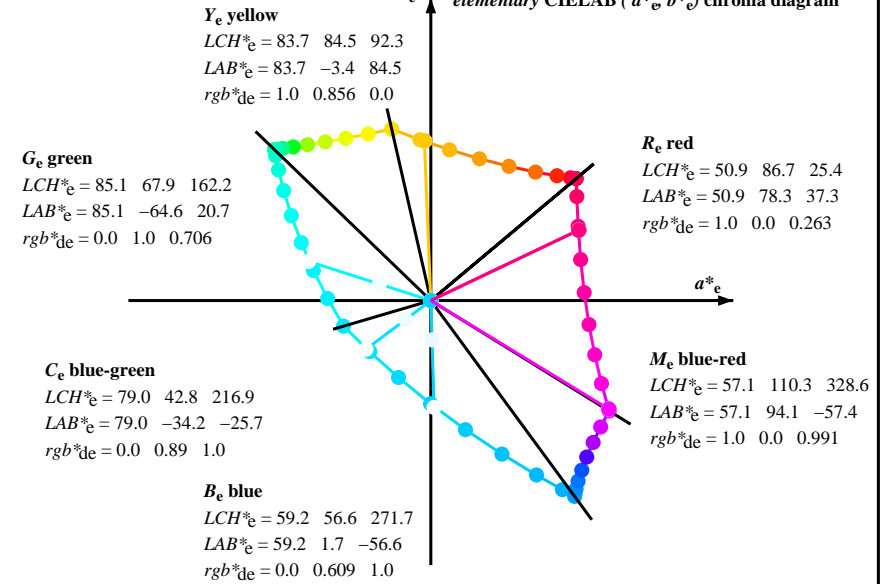
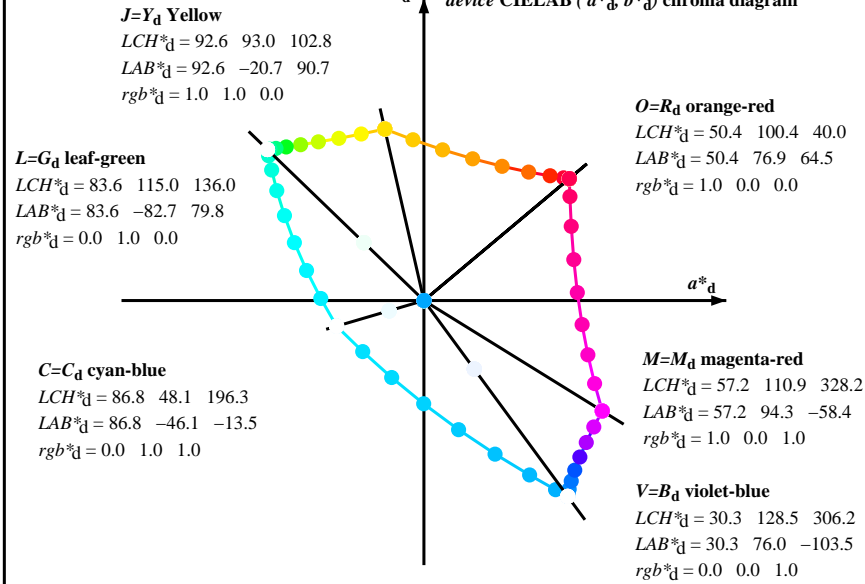


see similar files: http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT /PS
technical information: http://www.ps.bam.de or http://130.149.60.45/~farbmetrik

TUB registration: 20130201-RE42/RE42L0FA.TXT /PS
application for measurement of display output, no separation

TUB material: 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 $RYGCBM_e$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$



Notes to the CIELAB chroma diagrams (a^*_d, b^*_d), (a^*_s, b^*_s), (a^*_e, b^*_e)

- For the rgb^*_e -input values the CIELAB data LCH^*_e and LAB^*_e have been calculated.
- For the calculation of the standard hue angle $h_{ab,s}$ use for any device values rgb^*_d the equation:

$$h_{ab,s} = atan [r^*_d \cos(30) + g^*_d \cos(150)] / [r^*_d \sin(30) + g^*_d \sin(150) + b^*_d \sin(270)] \quad (1)$$
- For the 48 or 360 equally spaced standard hue angles $h_{ab,s}$ of the colours of maximum chroma use the seven hue angles of the 60 degree colours s : $h_{ab,s} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0, 390.0$ ($i=0,6$) and the equations for a 48 and 360 step hue circle:

$$h_{48ab,sij} = h_{ab,si} + j [h_{ab,si+1} - h_{ab,si}] / 8 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 7) \quad (2)$$

$$h_{360ab,sij} = h_{ab,si} + j [h_{ab,si+1} - h_{ab,si}] / 60 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 59) \quad (3)$$
- For the 48 or 360 elementary hue angles $h_{ab,e}$ of the colours of maximum chroma use the seven hue angles of the elementary colours e : $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6, 385.5$ ($i=0,6$) and the equations for a 48 and 360 step elementary hue circle:

$$h_{48ab,eij} = h_{ab,ei} + j [h_{ab,ei+1} - h_{ab,ei}] / 8 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 7) \quad (4)$$

$$h_{360ab,eij} = h_{ab,ei} + j [h_{ab,ei+1} - h_{ab,ei}] / 60 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 59) \quad (5)$$
- For any elementary hue angle $h_{ab,e}$ there is a well defined device hue angle $h_{ab,d}$ see the following tables, columns 1 to 5 or 1 to 4.
- The values rgb^*_e produce the output of the device-independent elementary hues

see similar files: <http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT>
 technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20130201-RE42/RE42L0FA.TXT /PS
 application for measurement of display output, no separation

TUB material: 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* 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 1.0 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 1.0 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 1.0 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	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	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	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	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	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	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	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	50.9 78.3 37.3 86.7 385



see similar files: <http://130.149.60.45/~farbmetrik/RE42/RE42LOFA.TXT>
 technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20130201-RE42/RE42LOFA.TXT /PS
 application for measurement of display output, no separation

TUB material: 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 columns for device colors (LAB*, dsx361Mi), elementary colors (LAB*, dex361Mi), and RGB values (rgb*, ds361Mi). Rows represent various color patches from 128 to 139.

see similar files: http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT
technical information: http://www.ps.bam.de or http://130.149.60.45/~farbmetrik

TUB registration: 20130201-RE42/RE42L0FA.TXT /.PS
application for measurement of display output, no separation
TUB material: code=rha4ta

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

$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	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}
139	165	175	0.0	1.0	0.25	83.8	-80.5	69.1	106.1	139	0.0	1.0	0.25	83.8
139	166	176	0.0	1.0	0.266	83.8	-80.2	67.6	104.9	139	0.0	1.0	0.267	83.8
140	167	177	0.0	1.0	0.283	83.8	-79.9	66.1	103.7	140	0.0	1.0	0.283	83.8
140	168	178	0.0	1.0	0.3	83.8	-79.6	64.6	102.5	140	0.0	1.0	0.3	83.8
141	169	179	0.0	1.0	0.316	83.9	-79.2	63.1	101.3	141	0.0	1.0	0.317	83.9
141	170	180	0.0	1.0	0.333	83.9	-78.8	61.7	100.1	141	0.0	1.0	0.333	83.9
142	171	181	0.0	1.0	0.35	83.9	-78.4	60.2	98.9	142	0.0	1.0	0.35	83.9
142	172	182	0.0	1.0	0.366	84.0	-78.0	58.8	97.7	142	0.0	1.0	0.367	84.0
143	173	183	0.0	1.0	0.383	84.0	-77.6	57.2	96.4	143	0.0	1.0	0.383	84.0
144	174	184	0.0	1.0	0.4	84.0	-77.1	55.4	94.9	144	0.0	1.0	0.4	84.0
145	175	185	0.0	1.0	0.416	84.1	-76.6	53.6	93.5	145	0.0	1.0	0.417	84.1
145	176	185	0.0	1.0	0.433	84.1	-76.1	51.8	92.1	145	0.0	1.0	0.433	84.1
146	177	186	0.0	1.0	0.45	84.2	-75.6	50.0	90.6	146	0.0	1.0	0.45	84.2
147	178	187	0.0	1.0	0.466	84.2	-75.0	48.3	89.2	147	0.0	1.0	0.467	84.2
147	179	188	0.0	1.0	0.483	84.3	-74.4	46.6	87.8	147	0.0	1.0	0.483	84.3
148	180	189	0.0	1.0	0.5	84.3	-73.7	44.9	86.4	148	0.0	1.0	0.5	84.3
149	181	190	0.0	1.0	0.516	84.4	-73.2	42.9	84.8	149	0.0	1.0	0.517	84.4
150	182	191	0.0	1.0	0.533	84.4	-72.6	40.9	83.3	150	0.0	1.0	0.533	84.4
151	183	192	0.0	1.0	0.55	84.5	-71.9	39.0	81.8	151	0.0	1.0	0.55	84.5
152	184	193	0.0	1.0	0.566	84.5	-71.2	37.0	80.3	152	0.0	1.0	0.567	84.5
153	185	194	0.0	1.0	0.583	84.6	-70.5	35.2	78.8	153	0.0	1.0	0.583	84.6
154	186	195	0.0	1.0	0.6	84.6	-69.7	33.3	77.3	154	0.0	1.0	0.6	84.6
155	187	195	0.0	1.0	0.616	84.7	-68.9	31.5	75.8	155	0.0	1.0	0.617	84.7
156	188	196	0.0	1.0	0.633	84.8	-68.1	29.5	74.3	156	0.0	1.0	0.633	84.8
157	189	197	0.0	1.0	0.65	84.8	-67.4	27.4	72.8	157	0.0	1.0	0.65	84.8
159	190	198	0.0	1.0	0.666	84.9	-66.7	25.4	71.3	159	0.0	1.0	0.667	84.9
160	191	199	0.0	1.0	0.683	85.0	-65.8	23.4	69.9	160	0.0	1.0	0.683	85.0
161	192	200	0.0	1.0	0.7	85.1	-65.0	21.4	68.4	161	0.0	1.0	0.7	85.1
163	193	201	0.0	1.0	0.716	85.2	-64.0	19.5	67.0	163	0.0	1.0	0.717	85.2
164	194	202	0.0	1.0	0.733	85.2	-63.1	17.6	65.5	164	0.0	1.0	0.733	85.2
165	195	203	0.0	1.0	0.75	85.3	-62.0	15.9	64.0	165	0.0	1.0	0.75	85.3
167	196	204	0.0	1.0	0.766	85.4	-61.2	13.7	62.8	167	0.0	1.0	0.767	85.4
169	197	205	0.0	1.0	0.783	85.5	-60.4	11.5	61.5	169	0.0	1.0	0.783	85.5
170	198	206	0.0	1.0	0.8	85.6	-59.5	9.5	60.2	170	0.0	1.0	0.8	85.6
172	199	206	0.0	1.0	0.816	85.7	-58.5	7.5	59.0	172	0.0	1.0	0.817	85.7
174	200	207	0.0	1.0	0.833	85.8	-57.4	5.5	57.7	174	0.0	1.0	0.833	85.8
176	201	208	0.0	1.0	0.85	85.9	-56.3	3.7	56.4	176	0.0	1.0	0.85	85.9
177	202	209	0.0	1.0	0.866	86.0	-55.1	1.9	55.2	177	0.0	1.0	0.867	86.0
180	203	210	0.0	1.0	0.883	86.1	-54.1	0.0	54.1	180	0.0	1.0	0.883	86.1
182	204	211	0.0	1.0	0.9	86.2	-53.2	-2.1	53.2	182	0.0	1.0	0.9	86.2
184	205	212	0.0	1.0	0.916	86.3	-52.2	-4.2	52.4	184	0.0	1.0	0.917	86.3
187	206	213	0.0	1.0	0.933	86.4	-51.1	-6.3	51.5	187	0.0	1.0	0.933	86.4
189	207	214	0.0	1.0	0.95	86.5	-50.0	-8.2	50.7	189	0.0	1.0	0.95	86.5
191	208	215	0.0	1.0	0.966	86.6	-48.8	-10.1	49.8	191	0.0	1.0	0.967	86.6
194	209	216	0.0	1.0	0.983	86.7	-47.5	-11.8	48.9	194	0.0	1.0	0.983	86.7
196	210	216	0.0	1.0	1.0	86.8	-46.1	-13.5	48.1	196	0.0	1.0	1.0	86.8
C _d			0.0	0.927	1.0	81.7	-38.6	-22.2	44.7	210	C _s	0.0	1.0	1.0
			0.0	0.89	1.0	79.1	-34.2	-25.7	42.9	216	C _e	0.0	1.0	1.0

1-113830-L0 RE420-73 LAB*la0, YN=0%, XYZnw=0.0, 0.0, 0.0, 84.2, 88.6, 96.5, LAB*nw=0.0, 0.0, 0.0, 95.4, 0.0, 0.0

Output: sRGB standard device; no separation, D65, page 9/29

TUB-test chart RE42; hue code: $H^*_e=B75R_e$
 48 step hue circles; $rgb-LabCh^*$ tables

input: $rgb/cmyk \rightarrow rgb_{de}$
 output: 3D-linearization to rgb^*_{de}

1-113830-F0 C M Y O L V

see similar files: <http://130.149.60.45/~farbmetrik/RE42/RE42LOFA.TXT>
<http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20130201-RE42/RE42LOFA.TXT /.PS
 application for measurement of display output, no separation
 TUB material: 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 $RYGCBM_e$; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_d	$dd361M$	LAB^*_d	$dsx361Mi$ (x=LabCh)	C_d	rgb^*_s	$ds361Mi$	LAB^*_s	$dsx361Mi$ (x=LabCh)	$210C_s$	0.0	1.0	1.0	rgb^*_e	$de361Mi$	LAB^*_e	$dex361Mi$ (x=LabCh)	$216C_e$	0.0	1.0	1.0	rgb^*_d	rgb^*_s	rgb^*_e						
196	210	216	0.0	1.0	1.0	86.8	-46.1	-13.5	48.1	196	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
199	211	217	0.0	0.983	1.0	85.6	-44.6	-15.8	47.3	199	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
202	212	218	0.0	0.966	1.0	84.5	-42.9	-17.9	46.5	202	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
205	213	219	0.0	0.95	1.0	83.3	-41.1	-19.8	45.7	205	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
208	214	220	0.0	0.933	1.0	82.1	-39.3	-21.7	44.9	208	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
212	215	221	0.0	0.916	1.0	80.9	-37.4	-23.4	44.1	212	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
215	216	222	0.0	0.9	1.0	79.7	-35.4	-24.9	43.3	215	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
218	217	223	0.0	0.883	1.0	78.5	-33.4	-26.3	42.5	218	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
221	218	224	0.0	0.866	1.0	77.4	-31.5	-28.1	42.2	221	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
225	219	225	0.0	0.85	1.0	76.2	-29.9	-30.2	42.5	225	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
228	220	226	0.0	0.833	1.0	75.0	-28.1	-32.3	42.8	228	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
232	221	227	0.0	0.816	1.0	73.8	-26.1	-34.2	43.1	232	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
236	222	227	0.0	0.8	1.0	72.6	-24.0	-36.0	43.3	236	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
239	223	228	0.0	0.783	1.0	71.4	-21.8	-37.7	43.6	239	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
243	224	229	0.0	0.766	1.0	70.2	-19.5	-39.3	43.9	243	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
247	225	230	0.0	0.75	1.0	69.1	-17.0	-40.7	44.1	247	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
250	226	231	0.0	0.733	1.0	67.9	-15.3	-42.9	45.5	250	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
253	227	232	0.0	0.716	1.0	66.7	-13.5	-44.9	46.9	253	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
256	228	233	0.0	0.7	1.0	65.5	-11.4	-46.9	48.3	256	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
259	229	234	0.0	0.683	1.0	64.4	-9.2	-48.8	49.7	259	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
262	230	235	0.0	0.666	1.0	63.2	-6.8	-50.6	51.1	262	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
265	231	236	0.0	0.65	1.0	62.0	-4.2	-52.3	52.5	265	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
268	232	237	0.0	0.633	1.0	60.9	-1.5	-53.9	53.9	268	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
270	233	237	0.0	0.616	1.0	59.7	0.8	-55.6	55.7	270	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
272	234	238	0.0	0.6	1.0	58.6	2.9	-57.7	57.8	272	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
274	235	239	0.0	0.583	1.0	57.4	5.1	-59.7	59.9	274	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
276	236	240	0.0	0.566	1.0	56.3	7.4	-61.6	62.1	276	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
278	237	241	0.0	0.55	1.0	55.2	10.0	-63.5	64.2	278	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
280	238	242	0.0	0.533	1.0	54.0	12.6	-65.2	66.4	280	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
283	239	243	0.0	0.516	1.0	52.9	15.4	-66.8	68.5	283	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
285	240	244	0.0	0.5	1.0	51.7	18.3	-68.3	70.7	285	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
286	241	245	0.0	0.483	1.0	50.7	20.6	-70.2	73.2	286	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
287	242	246	0.0	0.466	1.0	49.6	22.9	-72.1	75.7	287	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
288	243	247	0.0	0.45	1.0	48.6	25.4	-74.0	78.2	288	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
290	244	248	0.0	0.433	1.0	47.5	28.0	-75.7	80.7	290	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
291	245	248	0.0	0.416	1.0	46.5	30.6	-77.4	83.2	291	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
292	246	249	0.0	0.4	1.0	45.4	33.3	-79.0	85.7	292	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
294	247	250	0.0	0.383	1.0	44.3	36.2	-80.5	88.2	294	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
295	248	251	0.0	0.366	1.0	43.4	38.7	-82.0	90.7	295	0.0	0.74	1.0	68.4	-16.0	-41.9	45.0	249	0.0	0.35	1.0	0.0	0.721	1.0	67.0	-13.9	-44.4	46.6	252	0.0	0.35	1.0
296	249	252	0.0	0.35	1.0	42.5	41.0	-83.6	93.2	296	0.0	0.735	1.0	68.0	-15.4	-42.6	45.5	250	0.0	0.333	1.0	0.0	0.716	1.0	66.7	-13.3	-45.0	47.1	253	0.0		

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* ddx361Mi (x=LabCh)	rgb* ds361Mi	LAB* dsx361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)	rgb* dd361Mi	LAB* dex361Mi (x=LabCh)
301	255	258	0.0 0.25 1.0	37.1 55.9 -92.3 107.9 301	0.0 0.707 1.0 66.1 -12.3 -46.0 47.8 255	0.0 0.25 1.0	0.0 0.69 1.0 64.9 -10.1 -48.0 49.2 258	0.0 0.25 1.0	0.0 0.69 1.0 64.9 -10.1 -48.0 49.2 258	
301	256	258	0.0 0.233 1.0	36.5 57.6 -93.4 109.7 301	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	0.0 0.685 1.0 64.6 -9.4 -48.6 49.6 258	
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	0.0 0.68 1.0 64.2 -8.7 -49.1 50.0 259	
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	0.0 0.675 1.0 63.8 -8.0 -49.7 50.4 260	
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	0.0 0.67 1.0 63.5 -7.2 -50.2 50.9 261	
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	0.0 0.665 1.0 63.1 -6.5 -50.8 51.3 262	
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	0.0 0.66 1.0 62.8 -5.7 -51.3 51.7 263	
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	0.0 0.655 1.0 62.4 -5.0 -51.8 52.1 264	
304	263	265	0.0 0.116 1.0	32.3 70.0 -100.3 123.3 304	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	0.0 0.65 1.0 62.1 -4.2 -52.3 52.5 265	
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	0.0 0.645 1.0 61.7 -3.4 -52.8 53.0 266	
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	0.0 0.64 1.0 61.4 -2.5 -53.2 53.4 267	
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	0.0 0.635 1.0 61.0 -1.7 -53.7 53.8 268	
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	0.0 0.63 1.0 60.6 -0.8 -54.1 54.2 269	
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	0.0 0.624 1.0 60.3 0.0 -54.6 54.7 269	
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	0.0 0.617 1.0 59.8 0.8 -55.6 55.7 270	
306	270	271	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306	0.0 0.624 1.0 60.2 0.0 -54.7 54.8 270	0.0 0.0 1.0	0.0 0.609 1.0 59.3 1.7 -56.5 56.6 271	0.0 0.0 1.0	0.0 0.609 1.0 59.3 1.7 -56.5 56.6 271	
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	0.0 0.602 1.0 58.7 2.7 -57.5 57.6 272	
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	0.0 0.594 1.0 58.2 3.7 -58.4 58.6 273	
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	0.0 0.586 1.0 57.7 4.8 -59.4 59.7 274	
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	0.0 0.578 1.0 57.1 5.8 -60.3 60.7 275	
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	0.0 0.57 1.0 56.6 7.0 -61.2 61.7 276	
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	0.0 0.563 1.0 56.1 8.1 -62.0 62.7 277	
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	0.0 0.555 1.0 55.5 9.3 -62.9 63.7 278	
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	0.0 0.547 1.0 55.0 10.5 -63.7 64.7 279	
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	0.0 0.539 1.0 54.5 11.7 -64.5 65.7 280	
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	0.0 0.531 1.0 53.9 13.0 -65.3 66.7 281	
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	0.0 0.524 1.0 53.4 14.3 -66.1 67.7 282	
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	0.0 0.516 1.0 52.9 15.6 -66.8 68.7 283	
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	0.0 0.508 1.0 52.3 16.9 -67.5 69.7 284	
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	0.0 0.5 1.0 51.8 18.3 -68.2 70.7 285	
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	0.0 0.488 1.0 51.0 19.9 -69.6 72.5 285	
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	0.0 0.476 1.0 50.3 21.6 -71.0 74.3 286	
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	0.0 0.464 1.0 49.5 23.3 -72.4 76.1 287	
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	0.0 0.452 1.0 48.8 25.1 -73.7 77.9 288	
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	0.0 0.44 1.0 48.0 26.9 -75.0 79.8 289	
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 -76.2 81.6 290	0.333 0.0 1.0	0.0 0.428 1.0 47.2 28.8 -76.2 81.6 290	
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	0.0 0.416 1.0 46.5 30.7 -77.4 83.4 291	
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	0.0 0.404 1.0 45.7 32.7 -78.5 85.2 292	
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	0.0 0.392 1.0 44.9 34.7 -79.7 87.0 293	
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	0.0 0.38 1.0 44.2 36.8 -80.7 88.8 294	
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	0.417 0.0 1.0	0.0 0.364 1.0 43.3 39.2 -82.2 91.2 295	
310	296	296	0.433 0.0 1.0	36.7 78.9 -92.7 121.8 310	0.0 0.353 1.0 42.7 40.7 -83.3 92.8 296	0.433 0.0 1.0	0.0 0.345 1.0 42.3 41.7 -84.0 93.9 296	0.433 0.0 1.0	0.0 0.345 1.0 42.3 41.7 -84.0 93.9 296	
310	297	297	0.45 0.0 1.0	37.2 79.1 -92.0 121.3 310	0.0 0.333 1.0 41.6 43.5 -85.2 95.7 297	0.45 0.0 1.0	0.0 0.327 1.0 41.3 44.4 -85.8 96.7 297	0.45 0.0 1.0	0.0 0.327 1.0 41.3 44.4 -85.8 96.7 297	
311	298	298	0.466 0.0 1.0	37.6 79.3 -91.2 120.9 311	0.0 0.313 1.0 40.5 46.3 -87.0 98.6 298	0.467 0.0 1.0	0.0 0.308 1.0 40.3 47.1 -87.5 99.4 298	0.467 0.0 1.0	0.0 0.308 1.0 40.3 47.1 -87.5 99.4 298	
311	299	299	0.483 0.0 1.0	38.1 79.6 -90.4 120.5 311	0.0 0.293 1.0 39.5 49.2 -88.7 101.5 299	0.483 0.0 1.0	0.0 0.289 1.0 39.2 49.9 -89.1 102.2 299	0.483 0.0 1.0	0.0 0.289 1.0 39.2 49.9 -89.1 102.2 299	
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	0.0 0.27 1.0 38.2 52.8 -90.6 105.0 300	

1-1131030-L0 RE420-73 LAB*la0, YN=0%, XYZnw=0.0, 0.0, 0.0, 84.2, 88.6, 96.5, LAB*nw=0.0, 0.0, 0.0, 95.4, 0.0, 0.0

Output: sRGB standard device; no separation, D65, page 11/29

TUB-test chart RE42; hue code: H*_e=B75R_e
48 step hue circles; rgb-LabCh*tables

input: rgb/cmyk -> rgb_{de}
output: 3D-linearization to rgb*_{de}

see similar files: http://130.149.60.45/~farbmetrik/RE42/RE42LOFA.TXT /PS
technical information: http://www.ps.bam.de or http://130.149.60.45/~farbmetrik

TUB registration: 20130201-RE42/RE42LOFA.TXT /PS
application for measurement of display output, no separation

TUB material: code=rha4ta

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 $RYGCBM_e$; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_d	$dd361M$	LAB^*_d	$dsx361Mi$ (x=LabCh)	rgb^*_s	$ds361Mi$	LAB^*_s	$dsx361Mi$ (x=LabCh)	rgb^*_e	$dd361M$	LAB^*_e	$dex361Mi$ (x=LabCh)	rgb^*_e	$dd361M$																		
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	38.5	79.8	-89.7	120.0	311									
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	303	0.567	0.0	1.0			
313	305	304	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	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	55.1	89.2	-37.5	96.8	337	1.0	0.0	0.85		
336	340	338	1.0	0.0	0.833	55.1	89.4	-38.6	97.4	336	1.0	0.0	0.778	54.5	87																			

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

<i>h_{ab,d}</i>	<i>h_{ab,s}</i>	<i>h_{ab,e}</i>	<i>rgb[*]_{dd361M}</i>	<i>LAB[*]_{ddx361Mi (x=LabCh)}</i>	<i>rgb[*]_{ds361Mi}</i>	<i>LAB[*]_{dsx361Mi (x=LabCh)}</i>	<i>rgb[*]_{dd361Mi}</i>	<i>rgb[*]_{de361Mi}</i>	<i>LAB[*]_{dex361Mi (x=LabCh)}</i>	<i>rgb[*]_{dd361Mi}</i>	<i>rgb[*]_{dd}</i>	<i>rgb[*]_{ds}</i>	<i>rgb[*]_{de}</i>
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

1-1131230-L0 RE420-73 LAB*la0, YN=0%, XYZnw=0.0, 0.0, 0.0, 84.2, 88.6, 96.5, LAB*nw=0.0, 0.0, 0.0, 95.4, 0.0, 0.0

Output: sRGB standard device; no separation, D65, page 13/29

TUB-test chart RE42; hue code: H*e=B75R_e
48 step hue circles; *rgb-LabCh**tables

input: *rgb/cmyk* -> *rgb_{de}*
output: 3D-linearization to *rgb*_{de}*

see similar files: http://130.149.60.45/~farbmetrik/RE42/RE42LOFA.TXT /PS
technical information: http://www.ps.bam.de or http://130.149.60.45/~farbmetrik

TUB registration: 20130201-RE42/RE42LOFA.TXT /.PS
application for measurement of display output, no separation
TUB material: code=rha4ta

http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT /.PS; 3D-linearization F: 3D-linearization RE42/RE42LE30FA.DAT in file (F), page 15/29



Table with columns: r/g, g/b, b/r, i/r, r/g, g/b, b/r, h/s, r/g, g/b, b/r, LabCH*Fde, LabCH*Fde, r/g, g/b, b/r, DF*Fde, h/s, LabCH*Fde, LabCH*Fde, r/g, g/b, b/r. It contains a large grid of numerical data for color calibration.

Mean color difference of this page: delta E* = 0.8

input: rgb/cmyk -> rgbde output: 3D-linearization to rgb*de

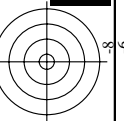
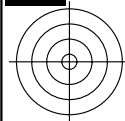
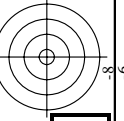
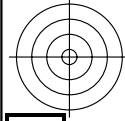
http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT /.PS; 3D-linearization F: 3D-linearization RE42/RE42LE30FA.DAT in file (F), page 16/29

Table with 80 columns (n#) and 80 rows (m#). Columns include: n#, H#*C*E, rgb*E, iet*E, hsa*E, rgb*E, LabC*H*E, LabCH*E, rgb*E, LabCH*E, DP*E, hsa*E, rgb*E, LabCH*E, rgb*E. Each cell contains numerical data representing color differences and linearization parameters.

Mean color difference of this page: delta E** = 0.6

TUB-test chart RE42; hue code: H#e=B75Re colors and differences, AE**

input: rbg/cmyk -> rgbde output: 3D-linearization to rbg*de



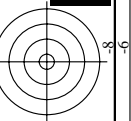
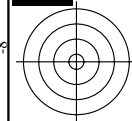
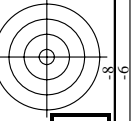
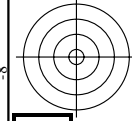
http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT /.PS; 3D-linearization F: 3D-linearization RE42/RE42LE30FA.DAT in file (F), page 17/29

Table with 16 columns: n, HHC*File, rpb*File, iet*File, hsa*File, rpb*File, LabCh*File, LabCh*File, rpb*File, rpb*File, hsa*File, DF*File, hsa*File, rpb*File, LabCh*File, LabCh*File. Rows 81-161.

Mean color difference of this page: delta E** = 0.6

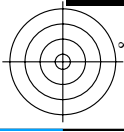
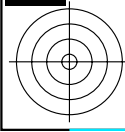
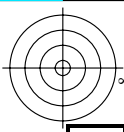
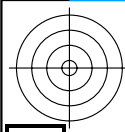
TUB-test chart RE42; hue code: H*e=B75Re colors and differences, AE**

input: rgb/cmyk -> rgbd output: 3D-linearization to rpb*de



http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT /.PS; 3D-linearization
F: 3D-linearization RE42/RE42L30FA.DAT in file (F), page 18/29

n	HC* ^{File}	rgb* ^{File}	icc* ^{File}	hsa* ^{File}	rgb* ^{File}	LabCH* ^{File}	rgb* ^{File}	LabCH* ^{File}	DE* ^{File}	rgb* ^{File}	LabCH* ^{File}	DE* ^{File}	rgb* ^{File}	LabCH* ^{File}	DE* ^{File}				
162	ROY5_025_025	0.25	0.0	0.25	0.0	0.065	0.248	0.077	0.016	12.1	20.4	10.6	23.2	27.4	1.6	375	78.3	86.7	25.4
163	ROY5_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
164	B5R3_037_037	0.25	0.0	0.375	0.187	0.311	0.241	0.086	0.233	13.1	30.7	-15.3	28.9	37.9	1.4	330	94.1	-57.4	110.3
165	B5R3_037_037	0.25	0.0	0.375	0.187	0.311	0.187	0.086	0.233	13.1	30.7	-15.3	28.9	37.9	1.4	330	94.1	-57.4	110.3
166	B5R3_037_037	0.25	0.0	0.375	0.187	0.311	0.131	0.148	0.474	18.9	26.6	-46.0	54.1	30.0	0.7	254	52.7	-90.7	104.9
167	B5R3_037_037	0.25	0.0	0.375	0.187	0.311	0.131	0.148	0.474	18.9	26.6	-46.0	54.1	30.0	0.7	254	52.7	-90.7	104.9
168	B1K3_075_075	0.25	0.0	0.625	0.262	0.262	0.131	0.248	0.597	28.9	21.5	-49.8	54.1	24.3	0.2	247	44.9	-79.7	86.9
169	B1K3_075_075	0.25	0.0	0.625	0.262	0.262	0.078	0.333	0.728	35.7	19.6	-56.4	54.1	24.3	0.2	247	44.9	-79.7	86.9
170	B1R1_100_100	0.25	0.0	1.0	0.5	0.5	0.043	0.417	0.862	44.0	18.4	-62.1	64.8	28.6	0.5	241	50.2	-71.1	74.3
171	ROY5_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
172	ROY5_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
173	B5R3_037_037	0.25	0.0	0.375	0.187	0.311	0.241	0.086	0.233	13.1	30.7	-15.3	28.9	37.9	1.4	330	94.1	-57.4	110.3
174	B5R3_037_037	0.25	0.0	0.375	0.187	0.311	0.241	0.086	0.233	13.1	30.7	-15.3	28.9	37.9	1.4	330	94.1	-57.4	110.3
175	B1K3_075_075	0.25	0.0	0.625	0.262	0.262	0.078	0.333	0.728	35.7	19.6	-56.4	54.1	24.3	0.2	247	44.9	-79.7	86.9
176	B1K3_075_075	0.25	0.0	0.625	0.262	0.262	0.078	0.333	0.728	35.7	19.6	-56.4	54.1	24.3	0.2	247	44.9	-79.7	86.9
177	B0R1_087_087	0.25	0.0	0.875	0.437	0.437	0.247	0.163	0.163	18.8	11.6	-7.6	13.8	32.6	0.5	330	94.1	-57.4	110.3
178	B0R1_087_087	0.25	0.0	0.875	0.437	0.437	0.247	0.163	0.163	18.8	11.6	-7.6	13.8	32.6	0.5	330	94.1	-57.4	110.3
179	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
180	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
181	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
182	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
183	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
184	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
185	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
186	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
187	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
188	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
189	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
190	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
191	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
192	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
193	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
194	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
195	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
196	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
197	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
198	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
199	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
200	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
201	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
202	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
203	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
204	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
205	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
206	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
207	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
208	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
209	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
210	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
211	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
212	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
213	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
214	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
215	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
216	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
217	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
218	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
219	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
220	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
221	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
222	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	352	83.6	-11.6	84.4
223	Y0G6_025_025	0.25	0.0	0.25	0.0	0.154	0.241	0.082	0.072	12.6	21.8	-4.0	22.0	23.0	1.6	35			



http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT /.PS; 3D-linearization F: 3D-linearization RE42/RE42LE30FA.DAT in file (F), page 20/29

Table with columns: n, HHC*F0e, rgb*F0e, iet*F0e, ihs*F0e, rgb*F0e, LabCH*F0e, LabCH*F0e, rgb*F0e, DP*F0e, rgh*F0e, LabCH*F0e, LabCH*F0e, rgb*F0e. Rows contain numerical data for various color channels and measurements.

Mean color difference of this page: delta E** = 0.4

TUB-test chart RE42; hue code: H*e=B75Re colors and differences, AE**

input: rgb/cmlyk -> rghbde output: 3D-linearization to rgb*de

http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT /.PS; 3D-linearization F: 3D-linearization RE42/RE42LE30FA.DAT in file (F), page 21/29

Table with 10 columns: n, HHC*F0, rgb*F0, iet*F0, Hsa*F0, rgb*F0, LabCh*F0, LabCh*F0, LabCh*F0, LabCh*F0. Rows 405-485.

Mean color difference of this page: delta E* = 0.4

TUB-test chart RE42; hue code: H*e=B75Re colors and differences, AE*
input: rgb/cmyk -> rgbde output: 3D-linearization to rgb*de

http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT /.PS; 3D-linearization F: 3D-linearization RE42/RE42LE30FA.DAT in file (F), page 23/29

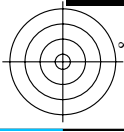
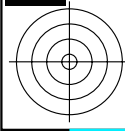
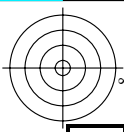
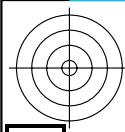
Table with 10 columns: n, HHC*F0, rgb*F0, iZt*F0, Hsa*F0, rgb*F0, LabCh*F0, LabCh*F0, LabCh*F0, LabCh*F0. Rows 567-647.

delta E** = 0.3

Mean color difference of this page:

TUB-test chart RE42; hue code: H*e=B75Re colors and differences, ΔE**

input: rgb*cmYk -> rgbde output: 3D-linearization to rgb*de



http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT /.PS; 3D-linearization
F: 3D-linearization RE42/RE42LE30FA.DAT in file (F), page 24/29

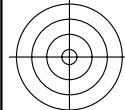
Table with 28 columns: n, HHC*F0, rpb*F0, iet*F0, Hsa*F0, rpb*F0, LabCH*F0, LabCH*F0, rpb*F0, DF*F0, rpb*F0, LabCH*F0, LabCH*F0, rpb*F0, DF*F0, rpb*F0, LabCH*F0, LabCH*F0, rpb*F0, DF*F0, rpb*F0, LabCH*F0, LabCH*F0, rpb*F0, DF*F0, rpb*F0, LabCH*F0, LabCH*F0, rpb*F0, DF*F0. Each row contains numerical data for a specific color patch.

Mean color difference of this page: delta E** = 2.5

RE420-TN; Page 24/29-F

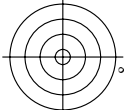
TUB-test chart RE42; hue code: H*e=B75Re
colors and differences, ΔE**

input: rgb*cmlyk -> rbg*de
output: 3D-linearization to rbg*de



TUB registration: 20130201-RE42/RE42LOFA.TXT /.PS
 application for measurement of display output, no separation

TUB material: code=rha4ta



n	HC*File	rgb*File	ief*File	hsa*File	rgb*File	LabCh*File	LabCh*File	rgb*File	DP*File	hsa*File	rgb*File	LabCh*File	LabCh*File	DP*File	hsa*File	rgb*File	LabCh*File	LabCh*File	DP*File
729	NW_1000c	0.875	1.0	1.0	0.875	0.986	1.0	1.0	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
730	G50B_100.012c	0.875	1.0	1.0	0.875	0.986	1.0	1.0	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
731	G50B_100.025c	0.75	1.0	1.0	0.75	0.972	1.0	1.0	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
732	G50B_100.050c	0.625	1.0	1.0	0.625	0.958	1.0	1.0	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
733	G50B_100.100c	0.5	1.0	1.0	0.5	0.945	1.0	1.0	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
734	G50B_100.0625c	0.375	1.0	1.0	0.375	0.931	1.0	1.0	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
735	G50B_100.075c	0.25	1.0	1.0	0.25	0.917	1.0	1.0	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
736	G50B_100.0875c	0.125	1.0	1.0	0.125	0.903	1.0	1.0	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
737	G50B_100.1012c	0.0	1.0	1.0	0.0	0.89	1.0	1.0	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
738	ROXY_100.012c	0.875	0.875	1.0	0.875	0.907	0.875	0.875	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
739	NW_0875c	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
740	G50B_087.012c	0.75	0.875	0.875	0.75	0.861	0.875	0.875	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
741	G50B_087.025c	0.625	0.875	0.875	0.625	0.847	0.875	0.875	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
742	G50B_087.050c	0.5	0.875	0.875	0.5	0.833	0.875	0.875	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
743	G50B_087.100c	0.375	0.875	0.875	0.375	0.819	0.875	0.875	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
744	G50B_087.0625c	0.25	0.875	0.875	0.25	0.806	0.875	0.875	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
745	G50B_087.075c	0.125	0.875	0.875	0.125	0.792	0.875	0.875	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
746	G50B_087.0875c	0.0	0.875	0.875	0.0	0.778	0.875	0.875	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
747	ROXY_100.025c	0.875	0.75	0.75	0.875	0.75	0.882	0.75	0.875	0.875	0.75	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
748	NW_0575c	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
749	G50B_075.012c	0.625	0.75	0.75	0.625	0.736	0.75	0.75	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
750	G50B_075.025c	0.5	0.75	0.75	0.5	0.722	0.75	0.75	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
751	G50B_075.050c	0.375	0.75	0.75	0.375	0.708	0.75	0.75	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
752	G50B_075.100c	0.25	0.75	0.75	0.25	0.695	0.75	0.75	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
753	G50B_075.0625c	0.125	0.75	0.75	0.125	0.681	0.75	0.75	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
754	G50B_075.075c	0.0	0.75	0.75	0.0	0.667	0.75	0.75	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
755	ROXY_100.0375c	0.875	0.625	1.0	0.625	0.625	0.625	0.625	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
756	ROXY_087.025c	0.875	0.625	0.625	0.875	0.25	0.812	0.390	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
757	ROXY_087.050c	0.875	0.625	0.625	0.875	0.25	0.797	0.375	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
758	NW_0625c	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
759	G50B_062.012c	0.5	0.625	0.625	0.5	0.611	0.625	0.55	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
760	G50B_062.025c	0.375	0.625	0.625	0.375	0.597	0.625	0.55	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
761	G50B_062.050c	0.25	0.625	0.625	0.25	0.583	0.625	0.55	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
762	G50B_062.100c	0.125	0.625	0.625	0.125	0.569	0.625	0.55	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
763	G50B_062.0625c	0.0	0.625	0.625	0.0	0.556	0.625	0.55	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
764	ROXY_100.050c	1.0	0.5	0.5	1.0	0.5	0.631	3.1	18.6	43.3	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
765	ROXY_087.0575c	0.875	0.5	0.5	0.875	0.375	0.687	3.90	18.6	43.3	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
766	ROXY_075.025c	0.875	0.5	0.5	0.875	0.375	0.687	3.90	18.6	43.3	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
767	ROXY_075.050c	0.875	0.5	0.5	0.875	0.375	0.687	3.90	18.6	43.3	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
768	NW_0500c	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
769	G50B_050.012c	0.375	0.5	0.5	0.375	0.486	0.5	45.6	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
770	G50B_050.025c	0.25	0.5	0.5	0.25	0.472	0.5	43.6	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
771	G50B_050.050c	0.125	0.5	0.5	0.125	0.458	0.5	41.5	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
772	G50B_050.100c	0.0	0.5	0.5	0.0	0.445	0.5	39.5	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
773	ROXY_100.0625c	1.0	0.375	0.375	1.0	0.375	0.509	61.2	39.1	18.6	43.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
774	ROXY_100.075c	0.875	0.375	0.375	0.875	0.375	0.509	61.2	39.1	18.6	43.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
775	ROXY_087.0575c	0.875	0.375	0.375	0.875	0.375	0.509	61.2	39.1	18.6	43.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
776	ROXY_075.0575c	0.875	0.375	0.375	0.875	0.375	0.509	61.2	39.1	18.6	43.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
777	ROXY_062.025c	0.625	0.375	0.375	0.625	0.375	0.44	48.5	19.5	9.3	21.6	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0
778	NW_0375c	0.375	0.375	0.375	0.375	0.375	0.407	42.1	9.7	4.6	10.8	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0
779	G50B_037.012c	0.25	0.375	0.375	0.25	0.375	0.407	35.7	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
780	G50B_037.025c	0.125	0.375	0.375	0.125	0.363	0.407	33.6	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
781	G50B_037.050c	0.0	0.375	0.375	0.0	0.333	0.407	29.6	-12.8	-9.6	16.0	21.6	25.4	95.4	0.0	0.0	0.0	0.0	0.0
782	ROXY_100.1075c	1.0	0.25	0.25	1.0	0.25	0.447	62.6	58.7	27.9	65.0	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0
783	ROXY_087.1025c	0.875	0.25	0.25	0.875	0.25	0.447	62.6	58.7	27.9	65.0	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0
784	G50B_075.109c	0.75	0.25	0.25	0.75	0.25	0.381	49.3	18.6	43.3	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
785	G50B_075.1075c	0.625	0.25	0.25	0.625	0.25	0.363	47.1	9.7	4.6	10.8	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0
786	ROXY_050.012c	0.5	0.25	0.25	0.5	0.249	0.315	36.5	19.5	9.3	21.6	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0
787	ROXY_050.025c	0.375	0.25	0.25	0.375	0.249	0.315	30.2	9.7	4.6	10.8	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0
788	NW_0250c	0.25	0.25	0.25	0.25	0.25	0.25	23.8	0.0	0.0	0.0	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
789	G50B_025.012c	0.125	0.25	0.25	0.125	0.236	0.25	21.8	-4.2	-3.2	5.3	95.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
790	G50B_025.025c	0.0	0.25	0.25	0.0	0.222	0.25	19.7	-8.8	-6.4	10.8	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0
791	G50B_025.050c	1.0	0.125	0.125	1.0	0.125	0.355	16.4	68.5	32.6	75.8	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0
792	ROXY_087.075c	0.875	0.125	0.125	0.875	0.125	0.320	14.1	58.7	27.9	65.0	25.4	95.4	0.0	0.0	0.0	0.0	0.0	0.0
793	ROXY_075.0625c	0.75	0.125	0.125	0.75	0.125	0.289	12.7	48.9	23.3	54.2	25.4	95.4	0.0	0.				

TUB registration: 20130201-RE42/RE42L0FA.TXT /.PS
application for measurement of display output, no separation

TUB material: code=rha4ta

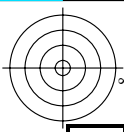
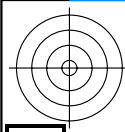
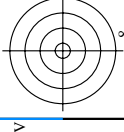
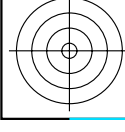


Table with 30 columns (n, HH, rgb, iZ, iRs, rGb, LabCh, LabCh*, rGb*, LabCh, LabCh*, rGb*, DP, rGb*, LabCh, LabCh*, rGb*) and 890 rows of numerical data.

Mean color difference of this page: delta E** = 0.6



see similar files: <http://130.149.60.45/~farbmetrik/RE42/RE42.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB-test chart RE42; hue code: H*e=B75Re
colors and differences, ΔE**

input: rgb/cmyk -> rgbde
output: 3D-linearization to rGb*de

RE42-70N; Page 26/29-F

L-1132530-F0

L-1132530-F0

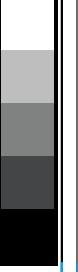
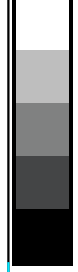
Table with 16 columns: n, HC*File, rgb*File, iet*File, ihs*File, rgb*File, LabCH*File, LabCH*File, LabCH*File, DP*File, ihs*File, rgb*File, LabCH*File, LabCH*File, LabCH*File, LabCH*File. Rows contain numerical data for various color patches.

Mean color difference of this page: $\Delta E^*_{ab} = 0.3$

input: rgb/cmyk -> rgbd
output: 3D-linearization to rgb*de

TUB registration: 20130201-RE42/RE42L0FA.TXT /.PS
 application for measurement of display output, no separation

TUB material: code=rha4ta



http://130.149.60.45/~farbmetrik/RE42/RE42L0FA.TXT /.PS; 3D-linearization
 F: 3D-linearization RE42/RE42LE30FA.DAT in file (F), page 29/29

n	HC*File	rgb*File	iet*File	hsa*File	rgb*File	LabCH*File	hsa*File	LabCH*File	rgb*File	DF*File	hsa*File	rgb*File	LabCH*File	hsa*File	LabCH*File	rgb*File	DF*File	hsa*File	LabCH*File	rgb*File
1053	NW_086de	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866
1054	NW_093de	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933
1055	NW_100de	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1056	NW_006de	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1057	NW_006de	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066
1058	NW_013de	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133
1059	NW_020de	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2
1060	NW_026de	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266
1061	NW_033de	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333
1062	NW_040de	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4
1063	NW_046de	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466
1064	NW_053de	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533
1065	NW_060de	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6
1066	NW_066de	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666
1067	NW_073de	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734
1068	NW_080de	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8
1069	NW_086de	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866
1070	NW_093de	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933
1071	NW_100de	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1072	NW_006de	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1073	NW_006de	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066
1074	ROY_100_100de	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1075	GS0B_100_100de	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1076	Y06C_100_100de	0.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1077	B00C_100_100de	0.0	0.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1078	B08C_100_100de	0.0	0.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1079	B50R_100_100de	0.0	0.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0

Mean color difference of this page: $\Delta E^*_{ab} = 0.3$

see similar files: <http://130.149.60.45/~farbmetrik/RE42/RE42.HTM>
 technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

input: *rgb/cmyk* -> *rgbde*
 output: 3D-linearization to *rgb*_de*

TUB-test chart RE42; hue code: H*_e=B75R_e
 colors and differences, ΔE^*_{ab}