

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

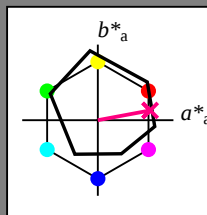
Data for any device (d) or elementary (e) colour:

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

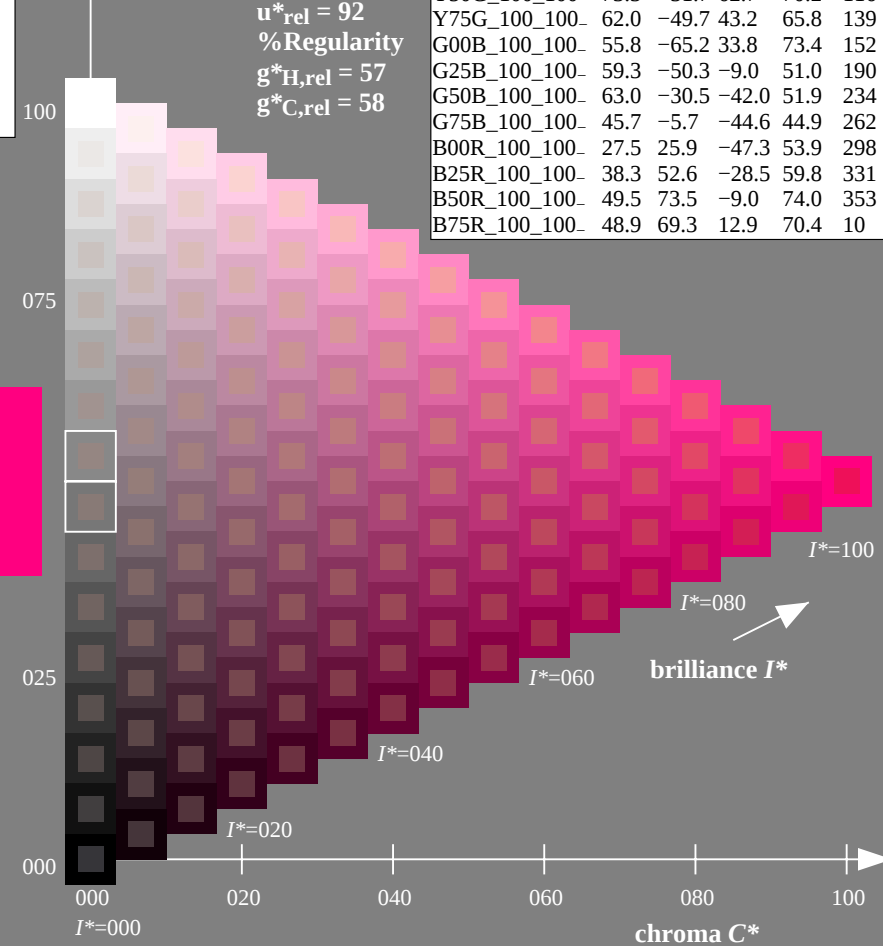
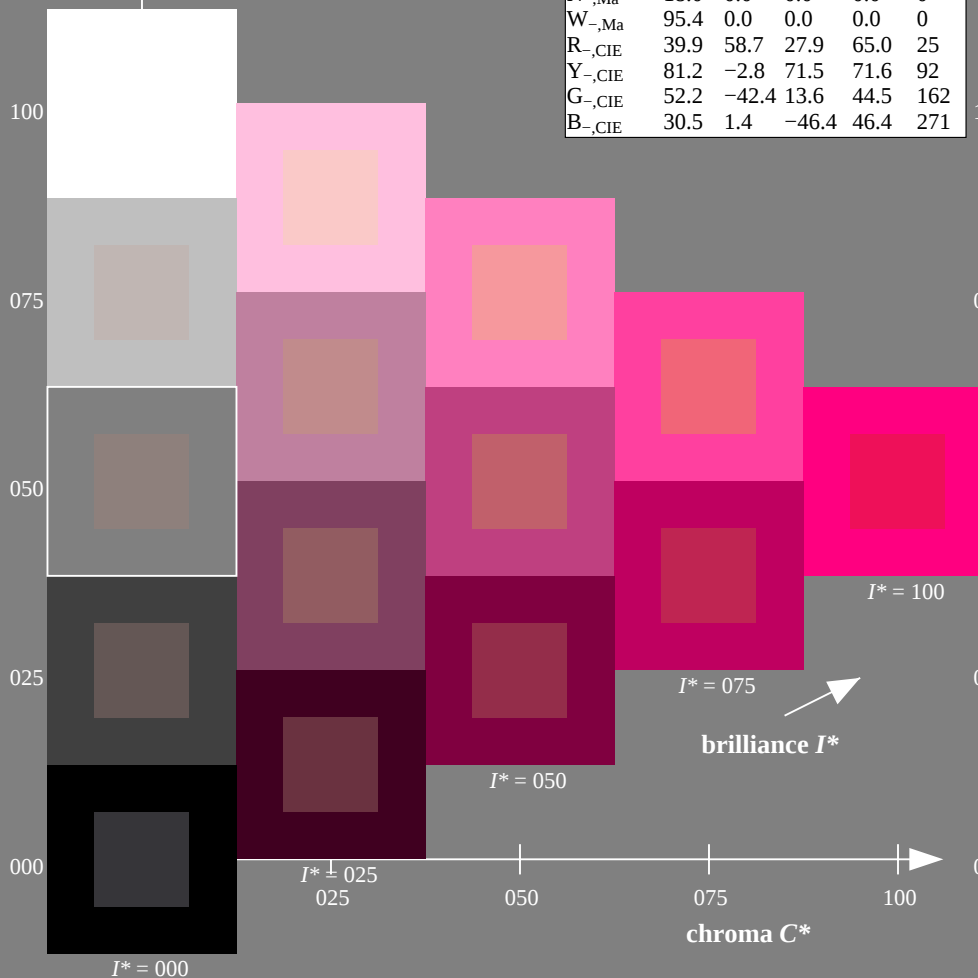
$u^*_{rel} = 92$

% 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



http://130.149.60.45/~farbmatrik/RE45/RE45L0NA.TXT /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 2/33

Input and Output: Offset Reflective System ORS18a for relative CIELAB hue $h_{ab,a,rel} = h_{ab}/360 = 352/360 = 0.97$

$H^*_e = B75R_e$

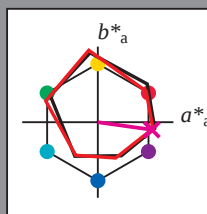
Data for any device (d) or elementary (e) colour:

HIC^*_e

hue text for the colours of this page:

$H^*_e = B75R_e$

triangle lightness T^*



name	$L^*=L^*_a$	a^*_a	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
R_e, Ma	47.6	64.9	30.9	71.9	25
Y_e, Ma	82.9	-3.5	87.8	87.9	92
G_e, Ma	52.4	-67.1	21.5	70.5	162
C_e, Ma	56.6	-39.7	-29.9	49.8	216
B_e, Ma	37.9	1.3	-45.4	45.4	271
M_e, Ma	34.8	49.2	-30.0	57.7	328
N_e, Ma	17.7	0.0	0.0	0.0	0
W_e, Ma	95.4	0.0	0.0	0.0	0
R_e, CIE	39.9	58.7	27.9	65.0	25
Y_e, CIE	81.2	-2.8	71.5	71.6	92
G_e, CIE	52.2	-42.4	13.6	44.5	162
B_e, CIE	30.5	1.4	-46.4	46.4	271

Data for maximum colour (Ma):

$LabCh^*_e, Ma$: 47 71 -9 72 352

HIC^*_e, Ma : B75R_100_100_e

$rgbic^*_e, Ma$:

0.94 0.0 1.0 1.0 1.0

triangle lightness T^*

% Gamut

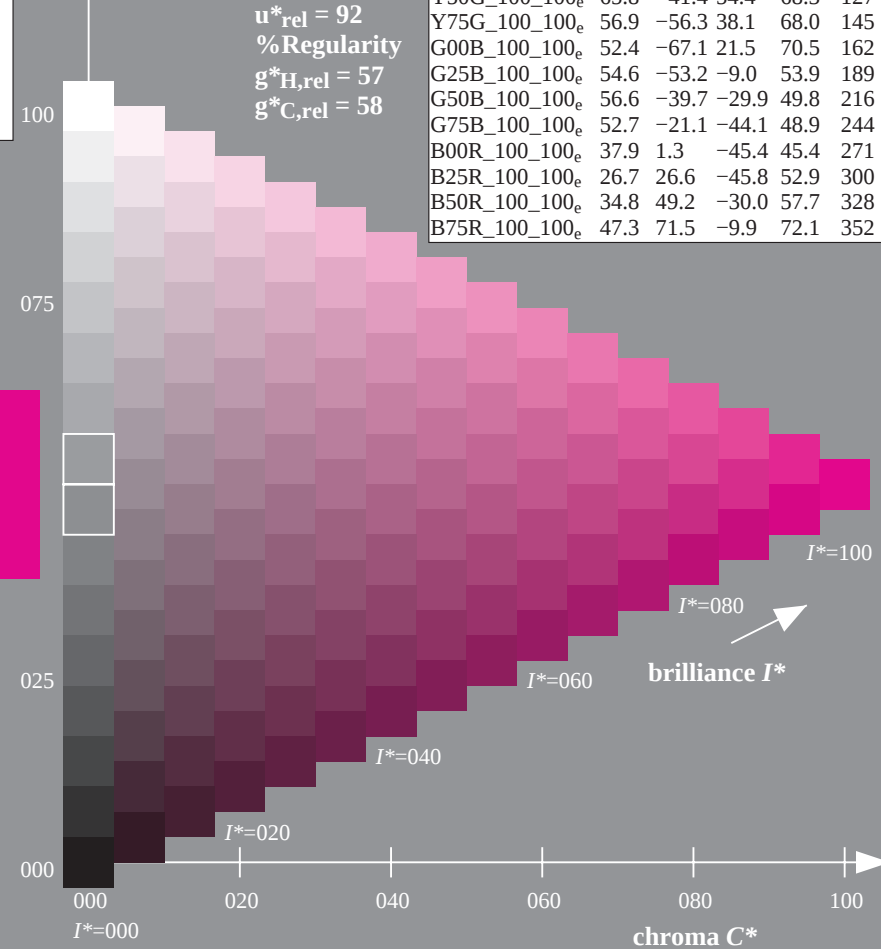
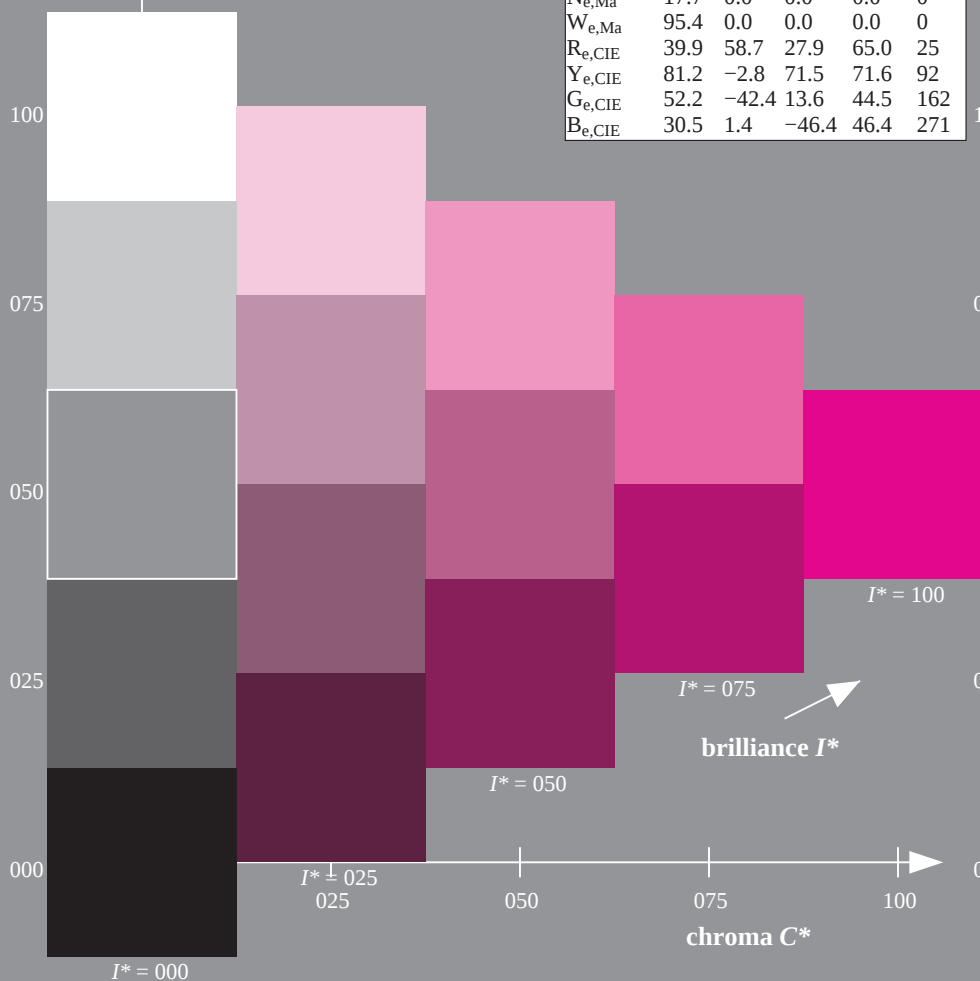
$u^*_{rel} = 92$

% Regularity

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

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

H^*_e	$L^*=L^*_a$	a^*_a	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
$R00Y_100_100_e$	47.6	64.9	30.9	71.9	25
$R25Y_100_100_e$	51.5	54.2	47.2	71.9	41
$R50Y_100_100_e$	60.3	35.6	59.0	68.9	58
$R75Y_100_100_e$	70.4	17.0	72.2	74.1	76
$Y00G_100_100_e$	82.9	-3.5	87.8	87.9	92
$Y25G_100_100_e$	76.9	-25.5	75.9	80.1	108
$Y50G_100_100_e$	65.8	-41.4	54.4	68.3	127
$Y75G_100_100_e$	56.9	-56.3	38.1	68.0	145
$G00B_100_100_e$	52.4	-67.1	21.5	70.5	162
$G25B_100_100_e$	54.6	-53.2	-9.0	53.9	189
$G50B_100_100_e$	56.6	-39.7	-29.9	49.8	216
$G75B_100_100_e$	52.7	-21.1	-44.1	48.9	244
$B00R_100_100_e$	37.9	1.3	-45.4	45.4	271
$B25R_100_100_e$	26.7	26.6	-45.8	52.9	300
$B50R_100_100_e$	34.8	49.2	-30.0	57.7	328
$B75R_100_100_e$	47.3	71.5	-9.9	72.1	352



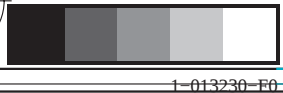
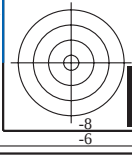
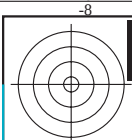
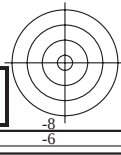
1-013130-L0 RE450-71

TUB-test chart RE45; hue code: $H^*_e = B75R_e$
Test chart according to DIN 33872, 3D=0, de=1, cmyk

input: $rgb/cmyk \rightarrow rgb_e$
output: transfer to $cmyk_e$

TUB registration: 20150701-RE45/RE45L0NA.TXT /PS
application for measurement of offset print output, separation cmyk6 (CMYK)

TUB material: code=rha4ta



see similar files: <http://130.149.60.45/~farbmetrik/RE45/RE45.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

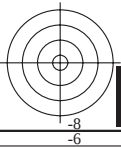
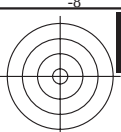
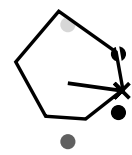
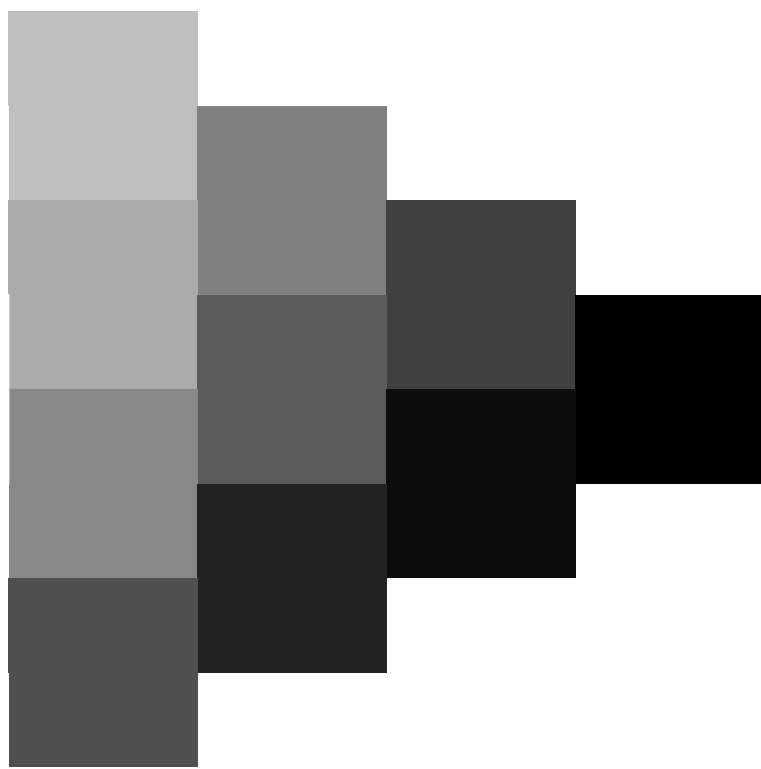
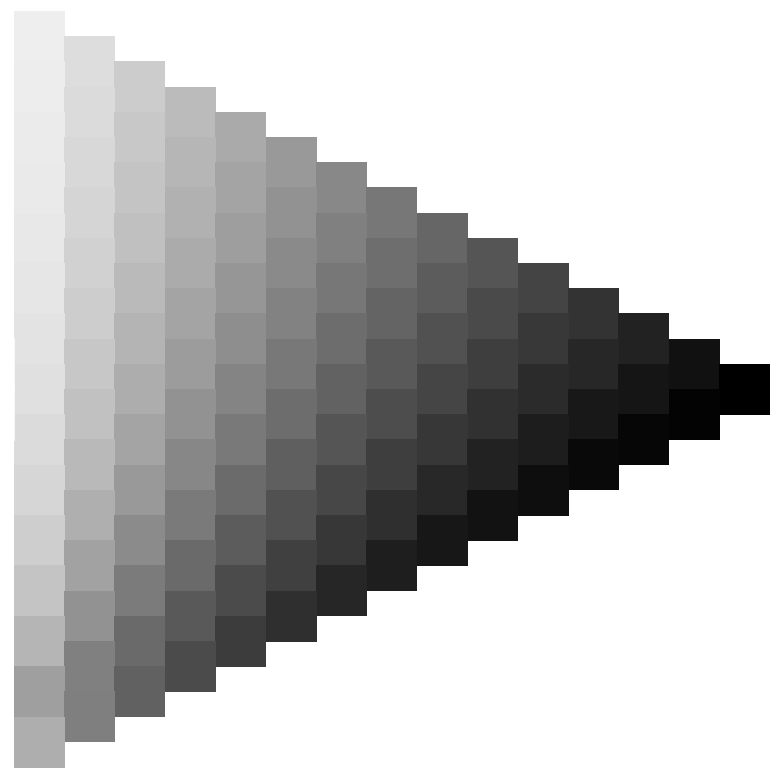
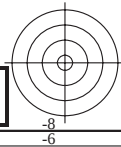
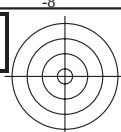


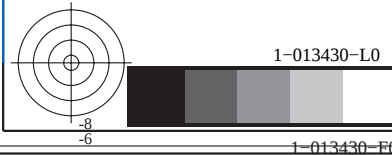
1-013230-L0 RE450-71

TUB-test chart RE45; hue code: $H^*_e=B75R_e$
Test chart according to DIN 33872, 3D=0, de=1, cmyk

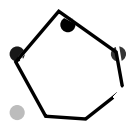
input: $rgb/cmyk \rightarrow rgb_e$
output: transfer to $cmyk_e$

1-013230-E0





see similar files: <http://130.149.60.45/~farbmetrik/RE45/RE45.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



1-013430-L0 RE450-71

TUB-test chart RE45; hue code: $H^*_e=B75R_e$
Test chart according to DIN 33872, 3D=0, de=1, cmyk

input: $rgb/cmyk \rightarrow rgb_e$
output: transfer to $cmyk_e$

1-013430-E0

Input and Output: Offset Reflective System ORS18a for relative CIELAB hue $h_{ab,a,rel} = h_{ab}/360 = 352/360 = 0.97$

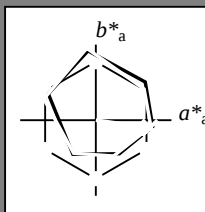
Data for any device (d) or elementary (e) colour:

HIC_e^*

hue text for the colours of this page:

$H_e^* = B75R_e$

triangle lightness T^*



ORS20a; adapted (a) CIELAB data					
name	$L^*=L_a^*$	a_a^*	b_a^*	$C_{ab,a}^*$	$h_{ab,a}^*$
$R_{e,Ma}$	47.6	64.9	30.9	71.9	25
$Y_{e,Ma}$	82.9	-3.5	87.8	87.9	92
$G_{e,Ma}$	52.4	-67.1	21.5	70.5	162
$C_{e,Ma}$	56.6	-39.7	-29.9	49.8	216
$B_{e,Ma}$	37.9	1.3	-45.4	45.4	271
$M_{e,Ma}$	34.8	49.2	-30.0	57.7	328
$N_{e,Ma}$	17.7	0.0	0.0	0.0	0
$W_{e,Ma}$	95.4	0.0	0.0	0.0	0
$R_{e,CIE}$	39.9	58.7	27.9	65.0	25
$Y_{e,CIE}$	81.2	-2.8	71.5	71.6	92
$G_{e,CIE}$	52.2	-42.4	13.6	44.5	162
$B_{e,CIE}$	30.5	1.4	-46.4	46.4	271

Data for maximum colour (Ma):

$LabCh_e^*, Ma: 47\ 71\ -9\ 72\ 352$

$HIC_e^*, Ma: B75R_100_100_e$

$rgbic_e^*, Ma:$

0.94 0.0 1.0 1.0 1.0

triangle lightness T^*

% Gamut

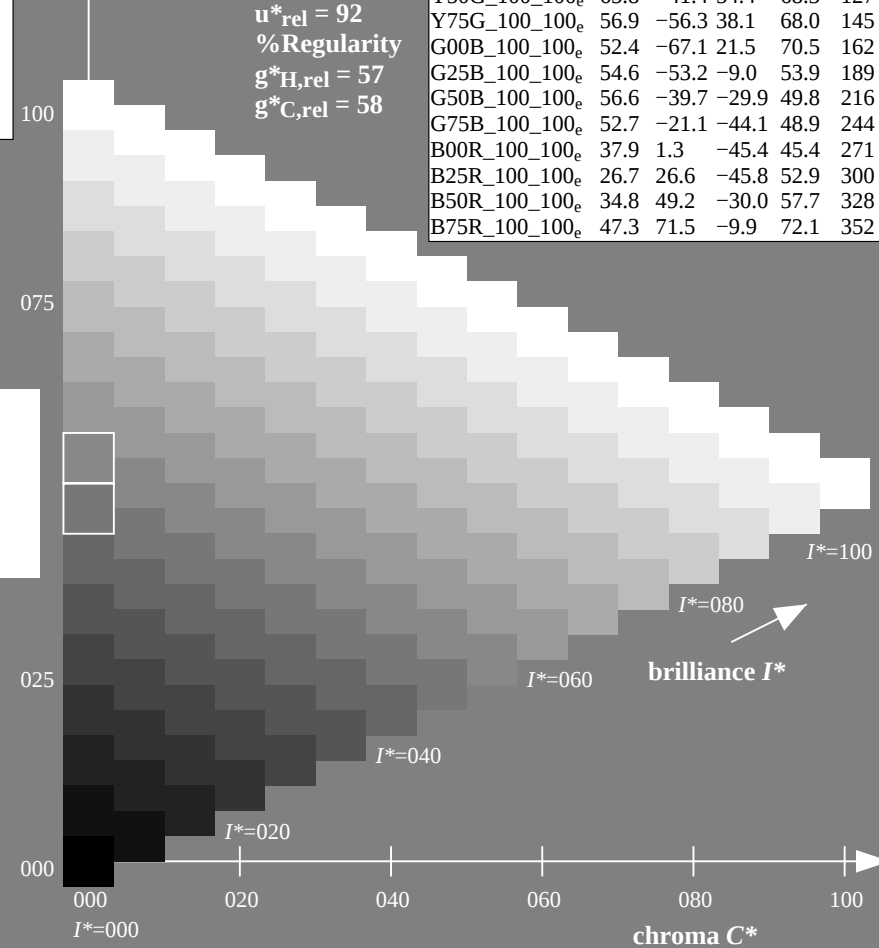
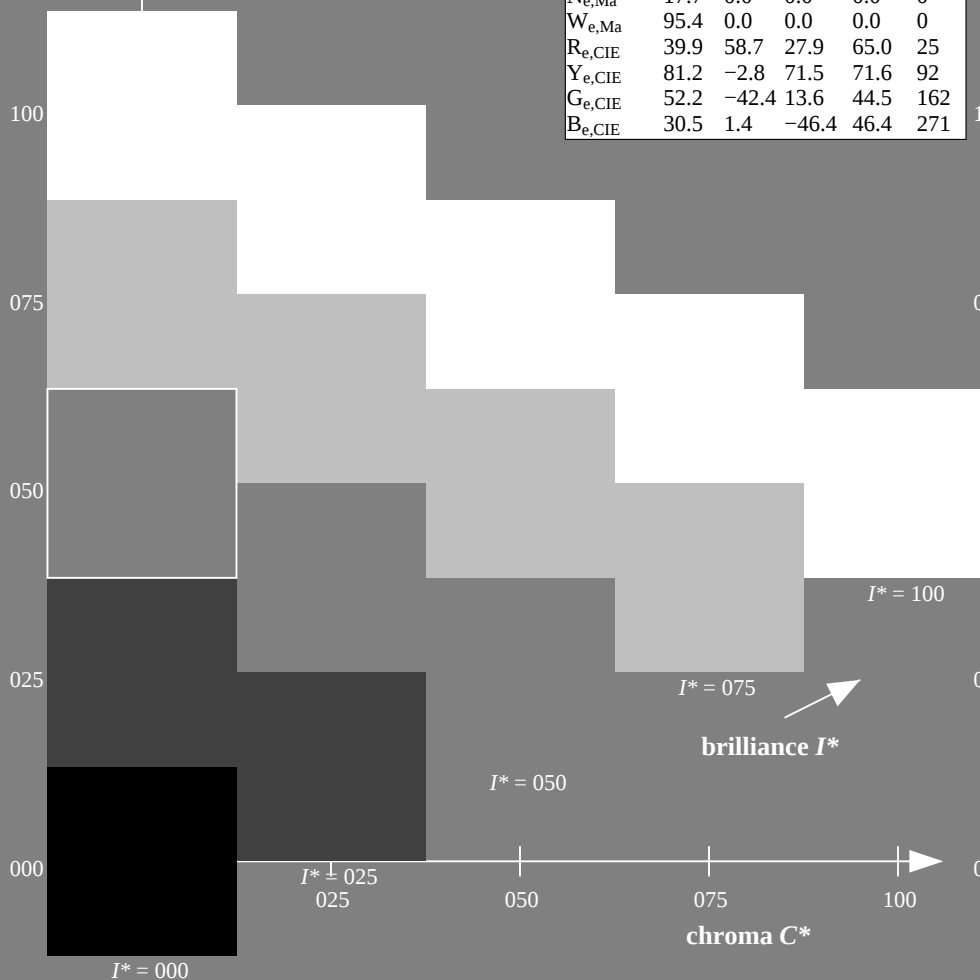
$u_{rel}^* = 92$

% Regularity

$g_{H,rel}^* = 57$

$g_{C,rel}^* = 58$

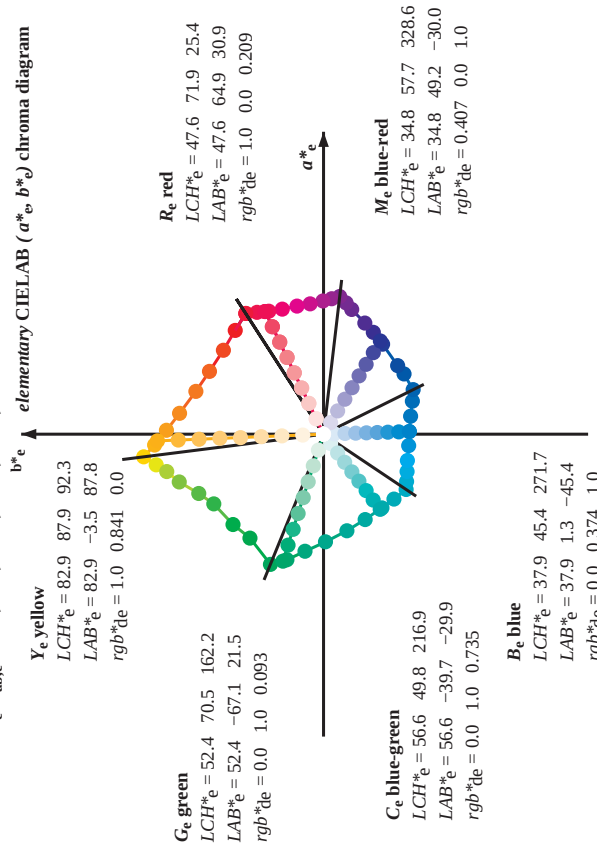
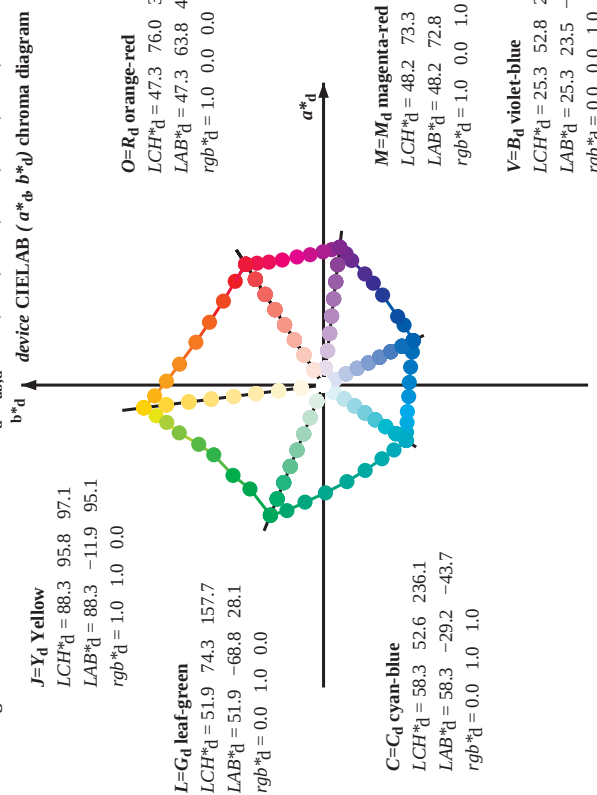
ORS20a; adapted (a) CIELAB data					
H_e^*	$L^*=L_a^*$	a_a^*	b_a^*	$C_{ab,a}^*$	$h_{ab,a}^*$
$R00Y_100_100_e$	47.6	64.9	30.9	71.9	25
$R25Y_100_100_e$	51.5	54.2	47.2	71.9	41
$R50Y_100_100_e$	60.3	35.6	59.0	68.9	58
$R75Y_100_100_e$	70.4	17.0	72.2	74.1	76
$Y00G_100_100_e$	82.9	-3.5	87.8	87.9	92
$Y25G_100_100_e$	76.9	-25.5	75.9	80.1	108
$Y50G_100_100_e$	65.8	-41.4	54.4	68.3	127
$Y75G_100_100_e$	56.9	-56.3	38.1	68.0	145
$G00B_100_100_e$	52.4	-67.1	21.5	70.5	162
$G25B_100_100_e$	54.6	-53.2	-9.0	53.9	189
$G50B_100_100_e$	56.6	-39.7	-29.9	49.8	216
$G75B_100_100_e$	52.7	-21.1	-44.1	48.9	244
$B00R_100_100_e$	37.9	1.3	-45.4	45.4	271
$B25R_100_100_e$	26.7	26.6	-45.8	52.9	300
$B50R_100_100_e$	34.8	49.2	-30.0	57.7	328
$B75R_100_100_e$	47.3	71.5	-9.9	72.1	352



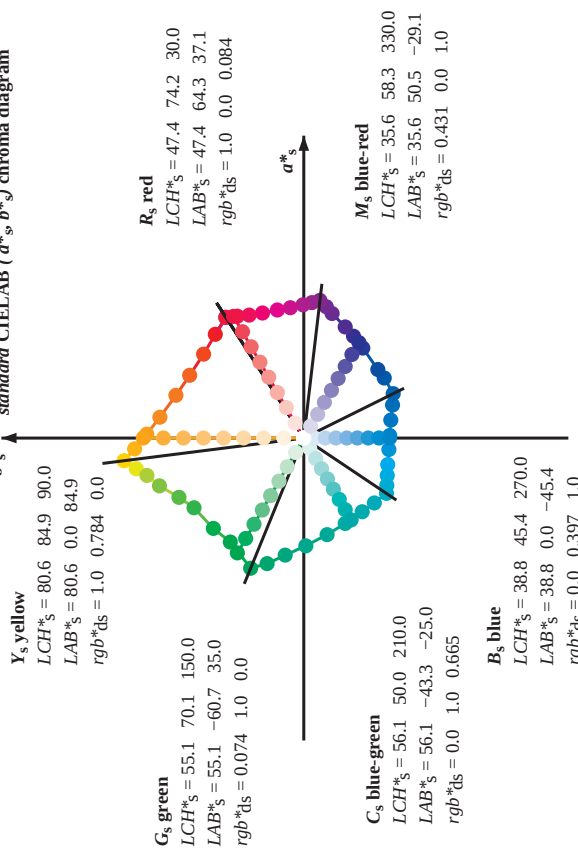
TUB-test chart RE45; hue code: $H_e^* = B75R_e$
Test chart according to DIN 33872, 3D=0, de=1, cmyk

input: $rgb/cmyk \rightarrow rgb_e$
output: transfer to $cmyk_e$

Data of Maximum color M in colorimetric system Offset standard print; separation cmyk6*, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBCM; $h_{ab,d} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours RYGBCM; $h_{ab,d} = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3$; Six hue angles of the elementary colours RYGBCM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$



standard CIELAB (a^*_s, b^*_s) chroma diagram



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} use for any device values rgb^*_e the equation:

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

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

$$h_{360ab,sij} = h_{ab,si} + j \left[\frac{h_{ab,si+1} - h_{ab,si}}{1/60} \right] \quad (i=0,1,...,5; j=0,1,...,59) \quad (3)$$
- For the 48 or 360 elementary hue angles h_{ab} 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:

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

$$h_{360ab,eij} = h_{ab,ei} + j \left[\frac{h_{ab,ei+1} - h_{ab,ei}}{1/60} \right] \quad (i=0,1,...,5; j=0,1,...,59) \quad (5)$$
- For any elementary hue angle h_{ab} 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

1-013630-L0 RE450-71 LAB*_{ab,0} YN=0%, XY Z_{nm}=2.4, 2.5, 2.6, 85.1, 88.8, 104.3, LAB*_{nm}=17.7, 0.0, 0.0, 95.5, 0.0, 0.0

TUB-test chart RE45; hue code: H*_e=B75R_e
48 step hue circles; $rgb-LabCh$ -tables

input: $rgb/cmyk -> rgb_e$
output: transfer to $cmyk_e$

Output: Offset standard print; separation cmyk6*, D65, page 7/33

http://130.149.60.45/~farbmatrik/RE45/RE45L0NA.TXT /PS; transfer output N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 9/33

Six hue angles of the 60 degree standard colours RYGBCM; h _{ab,d} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;										Six hue angles of the elementary colours RYGBCM; h _{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6									
Data of Maximum color M in colorimetric system Offset standard print; separation cmyk6; D65 for input or output;										Six hue angles of the elementary colours RYGBCM; h _{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6									
h _{ab,d} h _{ab,s} h _{ab,e} r _g b* d _{64M} LAB* d _{64M} (x=LabCh)										r _g b* d _{64M} LAB* d _{64M} (x=LabCh)									
32.8	30.0	25.4	1.0	0.0	0.0	47.3	63.8	41.2	76.0	32.8	1.0	0.0	0.209	47.6	64.9	30.9	71.9	25	330.0
40.4	37.5	33.8	1.0	0.125	0.0	51.2	54.9	46.7	72.1	40.4	1.0	0.007	0.0	47.6	63.4	41.6	75.8	33	30.0
50.0	45.0	42.1	1.0	0.25	0.0	56.0	44.4	53.0	69.1	50.0	1.0	0.148	0.0	52.1	53.0	48.1	71.6	42	330.0
61.1	52.5	50.5	1.0	0.375	0.0	61.4	33.2	60.3	68.8	61.1	1.0	0.25	0.0	56.0	44.5	53.0	69.2	49	30.0
71.4	60.0	58.8	1.0	0.5	0.0	67.2	22.6	67.6	71.2	71.4	1.0	0.375	0.0	60.3	35.6	59.0	69.0	58	330.0
81.7	67.5	67.2	1.0	0.625	0.0	73.6	11.0	76.1	76.9	81.7	1.0	0.442	0.0	64.5	27.8	64.5	70.2	66	30.0
88.5	75.0	75.6	1.0	0.75	0.0	79.2	2.0	83.0	83.1	88.5	1.0	0.55	0.0	69.8	18.3	71.3	73.6	75	330.0
93.6	82.5	83.9	1.0	0.875	0.0	84.2	-5.7	89.4	89.6	93.6	1.0	0.655	0.0	75.0	9.0	77.9	78.5	83	30.0
97.1	90.0	92.3	1.0	1.0	0.0	88.3	-11.9	95.1	95.8	97.1	1.0	0.871	0.0	83.0	-3.4	87.8	87.9	92	330.0
100.3	97.5	101.0	1.0	0.875	1.0	85.8	-16.2	88.6	90.0	100.3	1.0	0.842	0.0	85.8	-16.2	88.4	89.9	100	30.0
103.3	105.0	109.7	1.0	0.75	1.0	82.9	-19.7	83.0	85.3	103.3	1.0	0.599	1.0	76.2	-26.6	74.3	78.9	109	330.0
108.3	112.5	118.5	1.0	0.625	1.0	77.0	-25.2	76.3	80.4	108.3	1.0	0.455	1.0	71.4	-33.4	63.2	71.6	117	30.0
115.3	120.0	127.2	1.0	0.5	1.0	72.7	-31.3	66.0	73.1	115.3	1.0	0.327	1.0	65.8	-41.3	54.4	68.4	127	330.0
122.4	127.5	136.0	1.0	0.375	1.0	68.9	-36.9	58.1	68.8	122.4	1.0	0.244	1.0	60.7	-48.1	47.5	67.6	135	30.0
134.9	135.0	144.7	1.0	0.25	1.0	60.8	-47.8	47.8	67.6	134.9	1.0	0.124	1.0	57.4	-54.9	38.9	67.4	144	330.0
144.6	142.5	153.4	1.0	0.125	1.0	57.4	-54.9	38.9	67.3	144.6	1.0	0.047	1.0	54.0	-63.8	32.7	71.7	152	30.0
157.7	150.0	162.2	1.0	0.0	1.0	51.9	-68.8	28.1	74.3	157.7	1.0	0.093	1.0	52.4	-67.0	21.5	70.5	162	330.0
163.7	157.5	169.0	1.0	0.125	0.0	52.5	-66.4	19.3	69.1	163.7	1.0	0.209	53.1	-63.5	12.8	64.9	168	175	30.0
170.9	165.0	175.9	1.0	0.25	0.0	53.2	-61.9	9.8	62.7	170.9	1.0	0.311	53.7	-59.7	4.3	59.9	175	182	330.0
181.0	172.5	182.7	1.0	0.375	0.0	54.1	-56.9	-1.0	56.9	181.0	1.0	0.387	54.2	-56.4	-2.2	56.5	182	189	30.0
193.5	180.0	189.6	1.0	0.5	0.0	54.8	-51.0	-12.3	52.5	193.5	1.0	0.46	54.6	-53.1	-8.9	54.0	189	203	330.0
205.9	187.5	196.4	1.0	0.625	0.0	55.8	-45.1	-21.9	50.1	205.9	1.0	0.598	55.6	-46.5	-19.9	50.7	203	216	30.0
218.4	190.0	203.2	1.0	0.75	0.0	56.7	-38.9	-30.9	49.7	218.4	1.0	0.662	56.1	-43.4	-24.7	50.1	209	223	330.0
227.3	202.5	210.1	1.0	0.875	0.0	57.5	-34.3	-37.2	50.6	227.3	1.0	0.736	56.7	-39.7	-29.9	49.8	216	230	30.0
236.1	210.0	216.9	1.0	1.0	0.0	58.3	-29.2	-43.7	52.6	236.1	1.0	0.819	57.2	-36.4	-34.4	50.3	223	237	330.0
240.3	217.5	223.8	1.0	0.875	1.0	55.2	-25.0	-43.9	50.5	240.3	1.0	0.922	57.9	-32.5	-39.7	51.4	230	244	30.0
245.8	225.0	230.6	1.0	0.75	1.0	51.7	-19.7	-44.1	48.3	245.8	1.0	0.974	1.0	57.7	-28.3	-43.7	52.2	237	330.0
252.5	232.5	237.5	1.0	0.625	1.0	47.7	-13.9	-44.4	46.5	252.5	1.0	0.785	1.0	52.7	-21.1	-44.1	49.0	244	30.0
262.3	240.0	244.3	1.0	0.5	1.0	42.7	-6.0	-45.0	45.4	262.3	1.0	0.659	1.0	48.9	-15.4	-44.3	47.1	250	330.0
271.7	247.5	251.2	1.0	0.375	1.0	37.9	1.3	-45.4	45.4	271.7	1.0	0.555	1.0	45.0	-9.4	-44.8	45.9	258	30.0
281.6	255.0	258.0	1.0	0.25	1.0	33.3	9.4	-46.0	47.0	281.6	1.0	0.472	1.0	41.7	-4.3	-45.1	45.4	264	330.0
290.3	262.5	264.8	1.0	0.125	1.0	28.6	17.4	-46.9	50.1	290.3	1.0	0.375	1.0	37.9	1.4	-45.3	45.5	271	30.0
296.4	270.0	271.7	1.0	0.0	1.0	25.3	23.5	-47.3	52.8	296.4	1.0	0.291	1.0	34.9	6.8	-45.9	46.5	278	330.0
306.7	277.5	278.8	1.0	0.125	0.0	29.3	31.8	-42.6	53.1	306.7	1.0	0.188	1.0	31.0	13.3	-46.6	48.5	285	30.0
312.7	285.0	285.9	1.0	0.25	0.0	31.5	36.2	-39.2	53.4	312.7	1.0	0.079	1.0	27.4	19.6	-47.1	51.1	292	330.0
326.7	292.5	293.0	1.0	0.375	0.0	33.8	47.6	-31.2	56.9	326.7	1.0	0.046	0.0	26.8	26.6	-45.7	53.0	300	30.0
333.9	300.0	300.1	1.0	0.5	0.0	37.8	53.8	-26.3	59.9	333.9	1.0	0.126	0.0	29.4	31.9	-42.5	53.2	306	330.0
339.6	307.5	307.2	1.0	0.625	0.0	40.9	58.8	-21.8	62.7	339.6	1.0	0.265	0.0	31.8	37.7	-38.4	53.8	314	30.0
347.2	315.0	314.3	1.0	0.75	0.0	43.1	65.9	-14.9	67.6	347.2	1.0	0.324	0.0	32.9	43.2	-34.8	55.5	321	330.0
350.2	322.5	321.4	1.0	0.875	0.0	45.9	69.4	-11.9	70.5	350.2	1.0	0.407	0.0	34.9	49.3	-30.0	57.7	328	30.0
353.3	330.0	328.6	1.0	1.0	0.0	48.2	72.8	-8.5	73.3	353.3	1.0	0.529	0.0	38.6	55.0	-25.3	60.6	335	330.0
356.5	337.5	335.7	1.0	0.875	1.0	46.2	71.6	-4.3	71.7	356.5	1.0	0.678	0.0	41.9	61.9	-19.0	64.8	342	30.0
360.3	345.0	342.8	1.0	0.75	1.0	47.5	48.1	70.4	0.3	360.3	1.0	0.842	0.0	45.2	68.6	-12.7	69.8	349	330.0
365.8	352.5	349.9	1.0	0.625	1.0	46.2	48.0	68.9	7.1	365.8	1.0	0.949	0.0	47.3	71.5	-9.9	72.2	352	30.0
371.6	360.0	357.0	1.0	0.5	1.0	47.7	67.7	14.0	69.1	371.6	1.0	0.765	48.2	70.6	-0.1	70.6	359	368	330.0
378.2	367.5	364.1	1.0	0.375	1.0	47.7	66.1	21.8	69.6	378.2	1.0	0.563	47.9	68.4	10.6	69.2	368	376	30.0
383.9	375.0	371.2	1.0	0.25	1.0	47.7	65.0	28.9	71.2	383.9	1.0	0.408	47.8	66.7	19.8	69.6	376	385	330.0
388.6	382.5	378.3	1.0	0.125	1.0	47.3	63.8	41.2	76.0	388.6	1.0	0.209	47.6	64.9	30.9	71.9	385	392.8	30.0
392.8	390.0	385.4	1.0	0.0	1.0	47.3	63.8	41.2	76.0	392.8	1.0	0.209	47.6	64.9	30.9	71.9	385	392.8	30.0

Six hue angles of the device colours RYGBM; $h_{ab,d} = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3$; Six hue angles of the elementary colours RYGBM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$										Six hue angles of the device colours RYGBM; $h_{ab,d} = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3$; Six hue angles of the elementary colours RYGBM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$																								
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$		
88	75	75	1.0	0.75	0.0	79.2	2.0	83.0	83.1	88	1.0	0.543	0.0	69.4	19.0	70.7	73.2	75	1.0	0.75	0.0	1.0	0.55	0.0	69.8	18.3	71.3	73.6	75	1.0	0.75	0.0		
89	76	76	1.0	0.766	0.0	79.9	1.0	83.9	83.9	89	1.0	0.555	0.0	70.0	17.9	71.6	73.8	76	1.0	0.767	0.0	1.0	0.564	0.0	70.5	17.0	72.2	74.2	76	1.0	0.767	0.0		
89	77	77	1.0	0.783	0.0	80.6	0.0	84.8	84.8	89	1.0	0.567	0.0	70.7	16.7	72.4	74.3	77	1.0	0.783	0.0	1.0	0.577	0.0	71.2	15.8	73.1	74.8	77	1.0	0.783	0.0		
90	78	78	1.0	0.8	0.0	81.2	-0.9	85.7	85.7	90	1.0	0.579	0.0	71.3	15.6	73.3	74.9	78	1.0	0.8	0.0	1.0	0.591	0.0	71.9	14.5	74.0	75.4	78	1.0	0.8	0.0		
91	79	80	1.0	0.816	0.0	81.9	-1.9	86.5	86.5	91	1.0	0.591	0.0	71.9	14.4	74.1	75.5	79	1.0	0.817	0.0	1.0	0.604	0.0	72.6	13.1	74.9	76.0	80	1.0	0.817	0.0		
91	80	81	1.0	0.833	0.0	82.6	-3.0	87.4	87.4	91	1.0	0.604	0.0	72.5	13.2	74.9	76.6	80	1.0	0.833	0.0	1.0	0.618	0.0	73.3	11.8	75.8	76.7	81	1.0	0.833	0.0		
92	81	82	1.0	0.85	0.0	83.2	-4.0	88.2	88.3	92	1.0	0.616	0.0	73.2	12.0	75.6	76.6	81	1.0	0.85	0.0	1.0	0.635	0.0	74.1	10.4	76.8	77.5	82	1.0	0.85	0.0		
93	82	83	1.0	0.866	0.0	83.9	-5.1	89.0	89.2	93	1.0	0.629	0.0	73.8	10.7	76.5	77.2	82	1.0	0.867	0.0	1.0	0.655	0.0	75.0	9.0	77.9	78.5	83	1.0	0.867	0.0		
93	83	84	1.0	0.883	0.0	84.5	-6.1	89.8	90.0	93	1.0	0.648	0.0	74.7	9.5	77.5	78.1	83	1.0	0.883	0.0	1.0	0.675	0.0	75.9	7.6	79.1	79.5	84	1.0	0.883	0.0		
94	84	85	1.0	0.9	0.0	85.1	-6.9	90.6	90.8	94	1.0	0.666	0.0	75.5	8.3	78.6	79.0	84	1.0	0.9	0.0	1.0	0.696	0.0	76.8	6.1	80.2	80.5	85	1.0	0.9	0.0		
94	85	86	1.0	0.916	0.0	85.6	-7.7	91.3	91.7	94	1.0	0.684	0.0	76.3	7.0	79.6	79.9	85	1.0	0.917	0.0	1.0	0.716	0.0	77.8	4.6	81.3	81.5	86	1.0	0.917	0.0		
95	86	87	1.0	0.933	0.0	86.1	-8.5	92.1	92.5	95	1.0	0.703	0.0	77.1	5.6	80.6	80.8	86	1.0	0.933	0.0	1.0	0.736	0.0	78.7	3.1	82.4	82.5	87	1.0	0.933	0.0		
95	87	88	1.0	0.95	0.0	86.7	-9.3	92.9	93.3	95	1.0	0.721	0.0	78.0	4.3	81.6	81.7	87	1.0	0.95	0.0	1.0	0.759	0.0	79.7	1.5	83.6	83.6	88	1.0	0.95	0.0		
96	88	90	1.0	0.966	0.0	87.2	-10.2	93.6	94.2	96	1.0	0.739	0.0	78.8	2.9	82.5	82.6	88	1.0	0.967	0.0	1.0	0.787	0.0	80.8	0.0	85.0	85.0	90	1.0	0.967	0.0		
96	89	91	1.0	0.983	0.0	87.8	-11.1	94.3	95.0	96	1.0	0.76	0.0	79.7	1.5	83.6	83.6	89	1.0	0.983	0.0	1.0	0.814	0.0	81.9	-1.7	86.5	86.5	91	1.0	0.983	0.0		
97	90	92	1.0	1.0	0.0	88.3	-11.9	95.1	95.8	97	1.0	0.785	0.0	80.7	0.0	84.9	84.9	90	1.0	1.0	0.0	1.0	0.842	0.0	83.0	-3.4	87.8	87.9	92	1.0	1.0	0.0		
97	91	93	0.983	1.0	0.0	88.0	-12.5	94.2	95.1	97	1.0	0.809	0.0	81.7	-1.4	86.2	86.2	91	0.983	1.0	0.0	1.0	0.871	0.0	84.1	-5.3	89.2	89.4	93	0.983	1.0	0.0		
98	92	94	0.966	1.0	0.0	87.7	-13.1	93.4	94.3	98	1.0	0.834	0.0	82.7	-3.0	87.5	87.5	92	0.967	1.0	0.0	1.0	0.91	0.0	85.4	-7.3	91.1	91.4	94	0.967	1.0	0.0		
98	93	95	0.95	1.0	0.0	87.3	-13.7	92.5	93.5	98	1.0	0.859	0.0	83.6	-4.5	88.7	88.8	93	0.95	1.0	0.0	1.0	0.951	0.0	86.8	-9.4	93.0	93.4	95	0.95	1.0	0.0		
98	94	96	0.933	1.0	0.0	87.0	-14.3	91.6	92.7	98	1.0	0.887	0.0	84.7	-6.2	90.0	90.3	94	0.933	1.0	0.0	1.0	0.993	0.0	88.1	-11.5	94.8	95.5	96	0.933	1.0	0.0		
99	95	98	0.916	1.0	0.0	86.6	-14.8	90.8	92.0	99	1.0	0.923	0.0	85.8	-7.9	91.7	92.0	95	0.917	1.0	0.0	1.0	0.963	1.0	0.0	87.6	-13.2	93.2	94.1	98	0.917	1.0	0.0	
99	96	99	0.9	1.0	0.0	86.3	-15.4	89.9	91.2	99	1.0	0.958	0.0	87.0	-9.7	93.3	93.8	96	0.9	1.0	0.0	1.0	0.917	1.0	0.0	86.7	-14.8	90.8	92.0	99	0.9	1.0	0.0	
100	97	100	0.883	1.0	0.0	86.0	-15.9	89.0	90.4	100	1.0	0.994	0.0	88.2	-11.5	94.8	95.6	97	0.883	1.0	0.0	1.0	0.871	1.0	0.0	85.8	-16.2	88.4	89.9	100	0.883	1.0	0.0	
100	98	101	0.866	1.0	0.0	85.6	-16.4	88.2	89.7	100	1.0	0.968	1.0	0.0	87.7	-13.0	93.5	94.4	98	0.867	1.0	0.0	1.0	0.823	1.0	0.0	84.7	-17.7	86.3	88.1	101	0.867	1.0	0.0
100	99	102	0.85	1.0	0.0	85.2	-16.9	87.4	89.1	100	1.0	0.929	1.0	0.0	86.9	-14.4	91.4	92.6	99	0.85	1.0	0.0	1.0	0.774	1.0	0.0	83.5	-19.0	84.1	86.2	102	0.85	1.0	0.0
101	100	103	0.833	1.0	0.0	84.8	-17.4	86.7	88.4	101	0.89	1.0	0.0	86.2	-15.7	89.4	90.8	100	0.833	1.0	0.0	1.0	0.735	1.0	0.0	82.3	-20.3	82.2	84.7	103	0.833	1.0	0.0	
101	101	105	0.816	1.0	0.0	84.5	-17.9	86.0	87.8	101	0.849	1.0	0.0	85.3	-16.9	87.5	89.1	101	0.817	1.0	0.0	1.0	0.706	1.0	0.0	80.9	-21.7	80.7	83.6	105	0.817	1.0	0.0	
102	102	106	0.8	1.0	0.0	84.1	-18.3	85.2	87.2	102	0.807	1.0	0.0	84.3	-18.1	85.6	87.5	102	0.8	1.0	0.0	1.0	0.676	1.0	0.0	79.5	-23.0	79.1	82.4	106	0.8	1.0	0.0	
102	103	107	0.783	1.0	0.0	83.7	-18.8	84.5	86.5	102	0.765	1.0	0.0	83.3	-19.2	83.7	85.9	103	0.783	1.0	0.0	1.0	0.647	1.0	0.0	78.1	-24.3	77.5	81.3	107	0.783	1.0	0.0	
102	104	108	0.766	1.0	0.0	83.3	-19.2	83.7	85.9	102	0.734	1.0	0.0	82.2	-20.4	82.2	84.7	104	0.767	1.0	0.0	1.0	0.62	1.0	0.0	76.9	-25.5	75.9	80.1	108	0.767	1.0	0.0	
103	105	109	0.75	1.0	0.0	82.9	-19.7	83.0	85.3	103	0.709	1.0	0.0	81.0	-21.6	80.9	83.7	105	0.75	1.0	0.0	1.0	0.599	1.0	0.0	76.2	-26.6	74.3	78.9	109	0.75	1.0	0.0	
104	106	110	0.733	1.0	0.0	82.2	-20.5	82.1	84.6	104	0.684	1.0	0.0	79.9	-22.7	79.5	82.7	106	0.733	1.0	0.0	1.0	0.578	1.0	0.0	75.5	-27.7	72.6	77.7	110	0.733			

Data of Maximum color M in colorimetric system Offset standard print; separation cmyk6* D65 for input or output; Six hue angles of the 60 degree standard colours RYGCBM; $h_{ab,d} = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3$; Six hue angles of the elementary colours RYGCBM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$										Six hue angles of the device colours RYGCBM; $h_{ab,d} = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3$; Six hue angles of the elementary colours RYGCBM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$																																																																																																																																																																																																																																																																																																																																										
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	rgb^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	rgb^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab^*_{d361M}	Lab

Data of Maximum color M in colorimetric system										Six hue angles of the device colours RYGBM _d : $h_{ab,d} = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3$; 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^*_{d361M}	LAB^*_{d361M}	LAB^*_{d361M}	rgb^*_{d361M}	LAB^*_{d361M}	LAB^*_{d361M}	C_d	$rgb^*_{ds361Mi}$	$LAB^*_{ds361Mi}$	$LAB^*_{ds361Mi}$	rgb^*_{d361Mi}	$rgb^*_{de361Mi}$	$LAB^*_{de361Mi}$	$rgb^*_{dd361Mi}$	$rgb^*_{dd361Mi}$	$rgb^*_{dd361Mi}$				
236	210	216	0.0	1.0	1.0	58.3	-29.2	-43.7	52.6	236	0.0	1.0	0.666	56.1	-43.2	-24.9	50.0	210	0.0	1.0	1.0	
236	211	217	0.0	0.983	1.0	57.9	-28.7	-43.7	52.3	236	0.0	1.0	0.676	56.2	-42.8	-25.7	50.0	211	0.0	0.983	1.0	
237	212	218	0.0	0.966	1.0	57.5	-28.1	-43.8	52.0	237	0.0	1.0	0.686	56.3	-42.3	-26.4	50.0	212	0.0	0.967	1.0	
237	213	219	0.0	0.95	1.0	57.1	-27.5	-43.8	51.8	237	0.0	1.0	0.696	56.4	-41.8	-27.1	49.9	213	0.0	0.95	1.0	
238	214	220	0.0	0.933	1.0	56.7	-26.9	-43.9	51.5	238	0.0	1.0	0.706	56.4	-41.3	-27.8	49.9	214	0.0	0.933	1.0	
238	215	221	0.0	0.916	1.0	56.2	-26.4	-43.9	51.2	238	0.0	1.0	0.716	56.5	-40.8	-28.5	49.9	215	0.0	0.917	1.0	
239	216	222	0.0	0.9	1.0	55.8	-25.8	-43.9	50.9	239	0.0	1.0	0.726	56.6	-40.2	-29.2	49.8	216	0.0	0.9	1.0	
240	217	223	0.0	0.883	1.0	55.4	-25.2	-43.9	50.7	240	0.0	1.0	0.736	56.7	-39.7	-29.9	49.8	217	0.0	0.883	1.0	
240	218	224	0.0	0.866	1.0	55.0	-24.6	-43.9	50.4	240	0.0	1.0	0.746	56.7	-39.1	-30.5	49.8	218	0.0	0.867	1.0	
241	219	225	0.0	0.85	1.0	54.5	-23.9	-44.0	50.1	241	0.0	1.0	0.758	56.8	-38.6	-31.2	49.8	219	0.0	0.85	1.0	
242	220	226	0.0	0.833	1.0	54.1	-23.2	-44.0	49.8	242	0.0	1.0	0.772	56.9	-38.1	-32.0	49.9	220	0.0	0.833	1.0	
242	221	227	0.0	0.816	1.0	53.6	-22.5	-44.1	49.5	242	0.0	1.0	0.786	57.0	-37.7	-32.7	50.0	221	0.0	0.817	1.0	
243	222	227	0.0	0.8	1.0	53.1	-21.8	-44.1	49.2	243	0.0	1.0	0.8	57.1	-37.2	-33.4	50.1	222	0.0	0.8	1.0	
244	223	228	0.0	0.783	1.0	52.7	-21.1	-44.1	48.9	244	0.0	1.0	0.814	57.2	-36.6	-34.2	50.2	223	0.0	0.783	1.0	
245	224	229	0.0	0.766	1.0	52.2	-20.4	-44.1	48.6	245	0.0	1.0	0.828	57.3	-36.1	-34.9	50.3	224	0.0	0.767	1.0	
245	225	230	0.0	0.75	1.0	51.7	-19.7	-44.1	48.3	245	0.0	1.0	0.842	57.4	-35.6	-35.6	50.4	225	0.0	0.75	1.0	
246	226	231	0.0	0.733	1.0	51.2	-18.9	-44.2	48.1	246	0.0	1.0	0.856	57.5	-35.0	-36.3	50.5	226	0.0	0.733	1.0	
247	227	232	0.0	0.716	1.0	50.7	-18.1	-44.3	47.8	247	0.0	1.0	0.87	57.5	-34.4	-36.9	50.7	227	0.0	0.717	1.0	
248	228	233	0.0	0.7	1.0	50.1	-17.4	-44.3	47.6	248	0.0	1.0	0.884	57.6	-33.9	-37.7	50.8	228	0.0	0.7	1.0	
249	229	234	0.0	0.683	1.0	49.6	-16.6	-44.3	47.4	249	0.0	1.0	0.899	57.7	-33.4	-38.4	51.1	229	0.0	0.683	1.0	
250	230	235	0.0	0.666	1.0	49.1	-15.8	-44.4	47.1	250	0.0	1.0	0.913	57.8	-32.9	-39.2	51.3	230	0.0	0.667	1.0	
251	231	236	0.0	0.65	1.0	48.5	-15.0	-44.4	46.9	251	0.0	1.0	0.927	57.9	-32.3	-39.9	51.5	231	0.0	0.65	1.0	
252	232	237	0.0	0.633	1.0	48.0	-14.3	-44.4	46.6	252	0.0	1.0	0.941	58.0	-31.7	-40.7	51.7	232	0.0	0.633	1.0	
253	233	237	0.0	0.616	1.0	47.4	-13.4	-44.5	46.4	253	0.0	1.0	0.955	58.1	-31.2	-41.4	51.9	233	0.0	0.617	1.0	
254	234	238	0.0	0.6	1.0	46.7	-12.3	-44.6	46.3	254	0.0	1.0	0.969	58.2	-30.6	-42.1	52.2	234	0.0	0.6	1.0	
255	235	239	0.0	0.583	1.0	46.1	-11.3	-44.7	46.1	255	0.0	1.0	0.983	58.2	-29.9	-42.8	52.4	235	0.0	0.583	1.0	
257	236	240	0.0	0.566	1.0	45.4	-10.2	-44.8	46.0	257	0.0	1.0	0.997	58.3	-29.3	-43.5	52.6	236	0.0	0.567	1.0	
258	237	241	0.0	0.55	1.0	44.7	-9.1	-44.9	45.8	258	0.0	1.0	0.976	1.0	57.7	-28.4	-43.7	52.2	237	0.0	0.55	1.0
259	238	242	0.0	0.533	1.0	44.1	-8.1	-45.0	45.7	259	0.0	1.0	0.946	1.0	57.0	-27.3	-43.8	51.7	238	0.0	0.533	1.0
261	239	243	0.0	0.516	1.0	43.4	-7.0	-45.0	45.5	261	0.0	1.0	0.916	1.0	56.3	-26.3	-43.8	51.2	239	0.0	0.517	1.0
262	240	244	0.0	0.5	1.0	42.7	-6.0	-45.0	45.4	262	0.0	1.0	0.886	1.0	55.5	-25.3	-43.8	50.7	240	0.0	0.5	1.0
263	241	245	0.0	0.483	1.0	42.1	-5.0	-45.1	45.4	263	0.0	1.0	0.861	1.0	54.9	-24.3	-43.9	50.3	241	0.0	0.483	1.0
264	242	246	0.0	0.466	1.0	41.4	-4.0	-45.2	45.4	264	0.0	1.0	0.838	1.0	54.2	-23.3	-44.0	49.9	242	0.0	0.467	1.0
266	243	247	0.0	0.45	1.0	40.8	-3.0	-45.3	45.4	266	0.0	1.0	0.815	1.0	53.6	-22.4	-44.0	49.5	243	0.0	0.45	1.0
267	244	248	0.0	0.433	1.0	40.2	-2.1	-45.3	45.4	267	0.0	1.0	0.793	1.0	53.0	-21.4	-44.1	49.1	244	0.0	0.433	1.0
268	245	248	0.0	0.416	1.0	39.5	-1.1	-45.4	45.4	268	0.0	1.0	0.77	1.0	52.3	-20.5	-44.1	48.7	245	0.0	0.417	1.0
269	246	249	0.0	0.4	1.0	38.9	-0.1	-45.4	45.4	269	0.0	1.0	0.748	1.0	51.7	-19.6	-44.1	48.4	246	0.0	0.4	1.0
271	247	250	0.0	0.383	1.0	38.2	0.8	-45.4	45.4	271	0.0	1.0	0.729	1.0	51.1	-18.7	-44.2	48.1	247	0.0	0.383	1.0
272	248	251	0.0	0.366	1.0	37.6	1.8	-45.5	45.5	272	0.0	1.0	0.711	1.0	50.5	-17.8	-44.2	47.8	248	0.0	0.367	1.0
273	249	252	0.0	0.35	1.0	37.0	2.9	-45.6	45.7	273	0.0	1.0	0.692	1.0	49.9	-16.9	-44.3	47.5	249	0.0	0.35	1.0
275	250	253	0.0	0.333	1.0	36.4	4.0	-45.7	45.9	275	0.0	1.0	0.673	1.0	49.3	-16.1	-44.3	47.3	250	0.0	0.333	1.0
276	251	254	0.0	0.316	1.0	35.7	5.1	-45.8	46.1	276	0.0	1.0	0.654	1.0	48.7	-15.2	-44.3	47.0	251	0.0	0.317	1.0
277	252	255	0.0	0.3	1.0	35.1	6.1	-45.9	46.3	277	0.0	1.0	0.636	1.0	48.1	-14.3	-44.3	46.7	252	0.0	0.3	1.0
279	253	256	0.0	0.283	1.0	34.5	7.2	-46.0	46.5	279	0.0	1.0	0.62	1.0	47.6	-13.5	-44.4	46.5	253	0.0	0.283	1.0
280	254	257	0.0	0.266	1.0	33.9	8.3	-46.0	46.7	280	0.0	1.0	0.607	1.0	47.1	-12.7	-44.5	46.4	254	0.0	0.267	1.0
281	255	258	0.0	0.25	1.0	33.3	9.4	-46.0	47.0	281	0.0	1.0	0.594	1.0	46.5	-11.9	-44.6	46.3	255	0.0	0.25	1.0

1-0131330-1.0

RE450-71

LAB*a0, YN=0%, XYZnw=2.4, 2.5, 2.6, 85.1, 88.8, 104.3, LAB*rw=17.7, 0.0, 0.0, 95.5, 0.0, 0.0

input: $rgb/cmyk -> rgb_e$
output: transfer to $cmyk_e$

Output: Offset standard print: separation cmykn6*, D65

1-0131330-L0 RE450-71 LAB*lab, YN=0%, XY Znw=2.4, 2.5, 2.6, 85.1, 88.8, 104.3, LAB*rw=17.7, 0.0, 95.5, 0.0, 0.0

TUB-test chart RE45; hue code: H*_e=B75R_e
48 step hue circles; rgb-LabCh*tables

input: rgb/cmyk -> rgbe
output: transfer to cmyke

Output: Offset standard print; separation cmyk6*, D65, page 14/38

Data of Maximum color M in colorimetric system Offset standard print; separation cmyk6* D65 for input or output; Six hue angles of the 60 degree standard colours RYGBCM; $h_{ab,d} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;										
Six hue angles of the device colours RYGBCM; $h_{ab,d} = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3$; Six hue angles of the elementary colours RYGBCM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$										
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=LabCh)$	$rgb^*_{ds361MI}$	
281	255	258	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	281
282	256	258	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	282
283	257	259	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	283
284	258	260	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	284
285	259	261	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	285
286	260	262	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	286
287	261	263	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	287
288	262	264	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	288
289	263	265	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	289
290	264	266	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	290
291	265	267	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	291
292	266	268	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	292
293	267	269	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	293
294	268	270	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	294
295	269	271	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	295
296	270	272	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	296
297	271	273	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	297
298	272	274	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	298
299	273	275	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	299
300	274	276	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	300
301	275	277	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	301
302	276	278	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	302
303	277	279	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	303
304	278	280	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	304
305	279	281	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	305
306	280	282	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	306
307	281	283	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	307
308	282	284	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	308
309	283	285	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	309
310	284	286	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	310
311	285	287	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	311
312	286	288	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	312
313	287	289	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	313
314	288	290	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	314
315	289	291	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	315
316	290	292	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	316
317	291	293	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	317
318	292	294	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	318
319	293	295	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	319
320	294	296	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	320
321	295	297	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	321
322	296	298	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	322
323	297	299	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	323
324	298	300	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	324
325	299	301	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	325
326	300	302	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	326
327	301	303	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	327
328	302	304	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	328
329	303	305	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	329
330	304	306	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	330
331	305	307	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	331
332	306	308	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	332
333	307	309	0.0	2.5	1.0	33.3	9.4	-46.0	47.0	333

input: $rgb/cmyk -> rgb_e$
output: transfer to $cmyk_e$

TUB-test chart RE45; hue code: $H^*_e=B75R_e$
48 step hue circles; $rgb-LabCh$ *tables

see similar files: <http://130.149.60.45/~farbmatrik/RE45/RE45.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

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http://130.149.60.45/~farbmetrik/RE45/RE45L0NA.TXT /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 17/33

Data of Maximum color M in colorimetric system Offset standard print; separation cmyk6*, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBCM; $h_{ab,d}$ = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0;									
Six hue angles of the device colours RYGBCM; $h_{ab,d}$ = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3; Six hue angles of the elementary colours RYGBCM; $h_{ab,e}$ = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6									
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	$rgb^*_{ab,d}$	$LAB^*_{ab,d}$	$LAB^*_{ab,s}$	$rgb^*_{ab,s}$	$rgb^*_{ab,e}$	$LAB^*_{ab,e}$	$rgb^*_{ab,e}$
360	345	342	1.0	0.0	0.75	48.1	70.4	0.3	70.4
361	346	343	1.0	0.0	0.733	48.1	70.3	1.3	70.3
362	347	344	1.0	0.0	0.716	48.1	70.1	2.2	70.1
363	348	345	1.0	0.0	0.7	48.1	69.9	3.1	70.0
364	349	346	1.0	0.0	0.683	48.1	69.7	4.0	69.8
365	350	347	1.0	0.0	0.666	48.0	69.5	4.9	69.7
366	351	348	1.0	0.0	0.65	48.0	69.3	5.7	69.5
367	352	349	1.0	0.0	0.633	48.0	69.0	6.6	69.3
368	353	350	1.0	0.0	0.616	48.0	68.8	7.5	69.2
369	354	351	1.0	0.0	0.6	47.9	68.7	8.5	69.2
370	355	352	1.0	0.0	0.583	47.9	68.6	9.4	69.2
371	356	353	1.0	0.0	0.566	47.9	68.4	10.3	69.2
372	357	354	1.0	0.0	0.55	47.8	68.2	11.2	69.2
373	358	355	1.0	0.0	0.533	47.8	68.1	12.1	69.1
374	359	356	1.0	0.0	0.516	47.7	67.9	13.1	69.1
375	360	357	1.0	0.0	0.5	47.7	67.7	14.0	69.1
376	361	358	1.0	0.0	0.483	47.7	67.5	15.0	69.2
377	362	359	1.0	0.0	0.466	47.7	67.3	16.1	69.2
378	363	360	1.0	0.0	0.45	47.7	67.2	17.1	69.3
379	364	361	1.0	0.0	0.433	47.7	67.0	18.2	69.4
380	365	362	1.0	0.0	0.416	47.7	66.7	19.2	69.5
381	366	363	1.0	0.0	0.4	47.7	66.5	20.3	69.5
382	367	364	1.0	0.0	0.383	47.7	66.3	21.3	69.6
383	368	365	1.0	0.0	0.366	47.7	66.1	22.3	69.7
384	369	366	1.0	0.0	0.35	47.7	66.0	23.2	69.9
385	370	367	1.0	0.0	0.333	47.7	65.8	24.2	70.2
386	371	368	1.0	0.0	0.316	47.7	65.7	25.1	70.4
387	372	369	1.0	0.0	0.3	47.7	65.6	26.0	70.6
388	373	370	1.0	0.0	0.283	47.7	65.4	27.0	70.8
389	374	371	1.0	0.0	0.266	47.7	65.2	27.9	71.0
390	375	372	1.0	0.0	0.25	47.7	65.0	28.9	71.2
391	376	373	1.0	0.0	0.233	47.6	65.0	29.7	71.5
392	377	374	1.0	0.0	0.216	47.6	64.9	30.5	71.8
393	378	375	1.0	0.0	0.2	47.6	64.9	31.4	72.1
394	379	376	1.0	0.0	0.183	47.5	64.8	32.2	72.4
395	380	377	1.0	0.0	0.166	47.5	64.7	33.0	72.7
396	381	378	1.0	0.0	0.15	47.5	64.6	33.9	72.9
397	382	379	1.0	0.0	0.133	47.4	64.5	34.7	73.2
398	383	380	1.0	0.0	0.116	47.4	64.4	35.5	73.6
399	384	381	1.0	0.0	0.1	47.4	64.3	36.3	73.9
400	385	382	1.0	0.0	0.083	47.4	64.3	37.1	74.2
401	386	383	1.0	0.0	0.066	47.4	64.2	37.9	74.6
402	387	384	1.0	0.0	0.049	47.4	64.1	38.7	74.9
403	388	385	1.0	0.0	0.033	47.3	64.0	39.5	75.3
404	389	386	1.0	0.0	0.016	47.3	63.9	40.3	75.6
405	390	387	1.0	0.0	0.0	47.3	63.8	41.2	76.0
406	391	388	1.0	0.0	0.0	47.3	63.8	42.1	76.3
407	392	389	1.0	0.0	0.0	47.3	63.8	43.0	76.6
408	393	390	1.0	0.0	0.0	47.3	63.8	43.9	76.9
409	394	391	1.0	0.0	0.0	47.3	63.8	44.8	77.2
410	395	392	1.0	0.0	0.0	47.3	63.8	45.7	77.5
411	396	393	1.0	0.0	0.0	47.3	63.8	46.6	77.8
412	397	394	1.0	0.0	0.0	47.3	63.8	47.5	78.1
413	398	395	1.0	0.0	0.0	47.3	63.8	48.4	78.4
414	399	396	1.0	0.0	0.0	47.3	63.8	49.3	78.7
415	400	397	1.0	0.0	0.0	47.3	63.8	50.2	79.0
416	401	398	1.0	0.0	0.0	47.3	63.8	51.1	79.3
417	402	399	1.0	0.0	0.0	47.3	63.8	52.0	79.6
418	403	400	1.0	0.0	0.0	47.3	63.8	52.9	79.9
419	404	401	1.0	0.0	0.0	47.3	63.8	53.8	80.2
420	405	402	1.0	0.0	0.0	47.3	63.8	54.7	80.5
421	406	403	1.0	0.0	0.0	47.3	63.8	55.6	80.8
422	407	404	1.0	0.0	0.0	47.3	63.8	56.5	81.1
423	408	405	1.0	0.0	0.0	47.3	63.8	57.4	81.4
424	409	406	1.0	0.0	0.0	47.3	63.8	58.3	81.7
425	410	407	1.0	0.0	0.0	47.3	63.8	59.2	82.0
426	411	408	1.0	0.0	0.0	47.3	63.8	60.1	82.3
427	412	409	1.0	0.0	0.0	47.3	63.8	61.0	82.6
428	413	410	1.0	0.0	0.0	47.3	63.8	61.9	82.9
429	414	411	1.0	0.0	0.0	47.3	63.8	62.8	83.2
430	415	412	1.0	0.0	0.0	47.3	63.8	63.7	83.5
431	416	413	1.0	0.0	0.0	47.3	63.8	64.6	83.8
432	417	414	1.0	0.0	0.0	47.3	63.8	65.5	84.1
433	418	415	1.0	0.0	0.0	47.3	63.8	66.4	84.4
434	419	416	1.0	0.0	0.0	47.3	63.8	67.3	84.7
435	420	417	1.0	0.0	0.0	47.3	63.8	68.2	85.0
436	421	418	1.0	0.0	0.0	47.3	63.8	69.1	85.3
437	422	419	1.0	0.0	0.0	47.3	63.8	70.0	85.6
438	423	420	1.0	0.0	0.0	47.3	63.8	70.9	85.9
439	424	421	1.0	0.0	0.0	47.3	63.8	71.8	86.2
440	425	422	1.0	0.0	0.0	47.3	63.8	72.7	86.5
441	426	423	1.0	0.0	0.0	47.3	63.8	73.6	86.8
442	427	424	1.0	0.0	0.0	47.3	63.8	74.5	87.1
443	428	425	1.0	0.0	0.0	47.3	63.8	75.4	87.4
444	429	426	1.0	0.0	0.0	47.3	63.8	76.3	87.7
445	430	427	1.0	0.0	0.0	47.3	63.8	77.2	88.0
446	431	428	1.0	0.0	0.0	47.3	63.8	78.1	88.3
447	432	429	1.0	0.0	0.0	47.3	63.8	79.0	88.6
448	433	430	1.0	0.0	0.0	47.3	63.8	79.9	88.9
449	434	431	1.0	0.0	0.0	47.3	63.8	80.8	89.2
450	435	432	1.0	0.0	0.0	47.3	63.8	81.7	89.5
451	436	433	1.0	0.0	0.0	47.3	63.8	82.6	89.8
452	437	434	1.0	0.0	0.0	47.3	63.8	83.5	90.1
453	438	435	1.0	0.0	0.0	47.3	63.8	84.4	90.4
454	439	436	1.0	0.0	0.0	47.3	63.8	85.3	90.7
455	440	437	1.0	0.0	0.0	47.3	63.8	86.2	91.0
456	441	438	1.0	0.0	0.0	47.3	63.8	87.1	91.3
457	442	439	1.0	0.0	0.0	47.3	63.8	88.0	91.6
458	443	440	1.0	0.0	0.0	47.3	63.8	88.9	91.9
459	444	441	1.0	0.0	0.0	47.3	63.8	89.8	92.2
460	445	442	1.0	0.0	0.0	47.3	63.8	90.7	92.5
461	446	443	1.0	0.0	0.0	47.3	63.8	91.6	92.8
462	447	444	1.0	0.0	0.0	47.3	63.8	92.5	93.1
463	448	445	1.0	0.0	0.0	47.3	63.8	93.4	93.4
464	449	446	1.0	0.0	0.0	47.3	63.8	94.3	93.7
465	450	447	1.0	0.0	0.0	47.3	63.8	95.2	94.0
466	451	448	1.0	0.0	0.0	47.3	63.8	96.1	94.3
467	452	449	1.0	0.0	0.0	47.3	63.8	97.0	94.6
468	453	450	1.0	0.0	0.0	47.3	63.8	97.9	94.9
469	454	451	1.0	0.0	0.0	47.3	63.8	98.8	95.2
470	455	452	1.0	0.0	0.0	47.3	63.8	99.7	95.5

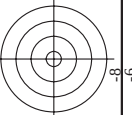
<http://130.149.60.45/~farbmatrik/RE45/RE45L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 18/33

nj	HC*Fe	rgb_Fe	icr_Fe	hsa_Fa	rgb_Fe	LabCh*Fe	rgb_Fe	LabCh*Fe	DF*Fe	hsa_Fe	rgb_Fe	LabCh*Fe	DF*Fe	hsa_Fe	rgb_Fe	LabCh*Fe	DF*Fe	hsa_Fe
0/648	R00Y_100_100e	1.0	0.0	0.0	1.0	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
1/657	R13Y_100_100e	1.0	0.125	0.0	1.0	0.0	0.007	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
2/666	R25Y_100_100e	1.0	0.25	0.0	1.0	0.0	0.133	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
3/675	R38Y_100_100e	1.0	0.375	0.0	1.0	0.0	0.249	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
4/684	R50Y_100_100e	1.0	0.5	0.0	1.0	0.0	0.349	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
5/693	R63Y_100_100e	1.0	0.625	0.0	1.0	0.0	0.455	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
6/702	R75Y_100_100e	1.0	0.75	0.0	1.0	0.0	0.563	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
7/711	R88Y_100_100e	1.0	0.875	0.0	1.0	0.0	0.675	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
8/720	Y00C_100_100e	1.0	1.0	0.0	1.0	0.0	0.841	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
9/639	Y13C_100_100e	0.875	1.0	0.0	1.0	0.0	0.875	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
10/558	Y25C_100_100e	0.75	1.0	0.0	1.0	0.0	0.875	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
11/477	Y38C_100_100e	0.625	1.0	0.0	1.0	0.0	0.875	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
12/396	Y50C_100_100e	0.5	1.0	0.0	1.0	0.0	0.875	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
13/315	Y63C_100_100e	0.375	1.0	0.0	1.0	0.0	0.875	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
14/234	Y75C_100_100e	0.25	1.0	0.0	1.0	0.0	0.875	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
15/153	Y88C_100_100e	0.125	1.0	0.0	1.0	0.0	0.875	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
16/72	G00C_100_100e	0.0	1.0	0.0	1.0	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
17/73	G13C_100_100e	0.0	1.0	0.125	1.0	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
18/74	G25C_100_100e	0.0	1.0	0.25	1.0	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
19/75	G38C_100_100e	0.0	1.0	0.375	1.0	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
20/76	G50C_100_100e	0.0	1.0	0.5	1.0	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
21/77	G63C_100_100e	0.0	1.0	0.625	1.0	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
22/78	G75C_100_100e	0.0	1.0	0.75	1.0	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
23/79	G88C_100_100e	0.0	1.0	0.875	1.0	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
24/80	C00B_100_100e	0.0	1.0	1.0	1.0	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
25/71	C13B_100_100e	0.0	0.875	1.0	1.0	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
26/62	C25B_100_100e	0.0	0.75	1.0	1.0	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
27/53	C38B_100_100e	0.0	0.625	1.0	1.0	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
28/44	C50B_100_100e	0.0	0.5	1.0	1.0	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
29/35	C63B_100_100e	0.0	0.375	1.0	1.0	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
30/26	C75B_100_100e	0.0	0.25	1.0	1.0	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
31/17	C88B_100_100e	0.0	0.125	1.0	1.0	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
32/8	B00M_100_100e	0.0	0.0	1.0	1.0	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
33/89	B13M_100_100e	0.125	0.0	1.0	1.0	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
34/170	B25M_100_100e	0.25	0.0	1.0	1.0	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
35/251	B38M_100_100e	0.375	0.0	1.0	1.0	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
36/332	B50M_100_100e	0.5	0.0	1.0	1.0	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
37/413	B63M_100_100e	0.625	0.0	1.0	1.0	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
38/494	B75M_100_100e	0.75	0.0	1.0	1.0	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
39/575	B88M_100_100e	0.875	0.0	1.0	1.0	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
40/656	M00R_100_100e	1.0	0.0	1.0	1.0	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
41/655	M13R_100_100e	1.0	0.0	0.875	1.0	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
42/654	M25R_100_100e	1.0	0.0	0.75	1.0	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
43/653	M38R_100_100e	1.0	0.0	0.625	1.0	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
44/652	M50R_100_100e	1.0	0.0	0.5	1.0	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
45/651	M63R_100_100e	1.0	0.0	0.375	1.0	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
46/650	M75R_100_100e	1.0	0.0	0.25	1.0	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
47/649	M88R_100_100e	1.0	0.0	0.125	1.0	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
48/648	R00Y_100_100e	1.0	0.0	0.0	1.0	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
49/0	NW_00e	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
50/91	NW_01e	0.125	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
51/182	NW_02e	0.25	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
52/273	NW_03e	0.375	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
53/364	NW_04e	0.5	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
54/455	NW_05e	0.625	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
55/546	NW_06e	0.75	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
56/637	NW_07e	0.875	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
57/728	NW_10e	1.0	1.0	1.0	1.0	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

Mean color difference of this page:
delta E* = 17.3

TUB-test chart RE45; hue code: H*_e=B75R_e
colors and differences, ΔE*

input: rgb/cmyk -> rgbe
output: transfer to cmyke

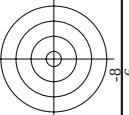


TUB material: code=rha4ta
nyn6 (CMYK)

input: *rgb/cmyk* -> *rgbe*
output: transfer to *cmyke*

UB-test chart RE45, hue code: $H_e^* = B/5R_e$
colors and differences, ΔE^*

[illegible]



TUB material: code=rha4ta
myn6 (CMYK)

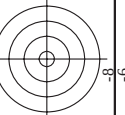
input: *rgb/cmyk* $\rightarrow$ *rgb*_e
output: transfer to *cmyk*_e

TUB-test chart RE45; hue code: $H_e^* = B75R_e$
colors and differences. ΔE^* ,

DE450-7N Page 20/23-F

Mean color difference of this range:

 $\Delta E^* = 11.0$



TUB material: code=rha4ta
nyn6 (CMYK)

input: *rgb/cmyk* -> *rgb*
output: transfer to *cmyk*

RE450-7N, Page 21/33-F

[illegible]



TUB material: code=rha4ta
nyn6 (CMYK)

input: *rgb/cmyk* → *rgb*
output: transfer to *cmyk*

colors and differences, ΔE^* ,

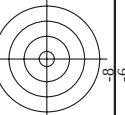
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E450-7N, Page 22/33-F

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TUB material: code=rha4ta
nyn6 (CMYK)

input: *rgb/cmyk* → *rgb*_e
output: transfer to *cmyk*_e

1-0132230-F0

[illegible]

see similar files: <http://130.149.60.45/~farbmatrik/RE45/RE45.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

TUB-test chart RE45; hue code: $H_e^*=B75R_e$

input: *rgb/cmyk* -> *rgb*_e
output: transfer to *cmyk*_e

	n	HC*Fe	rgb^*Fe	ict^*Fe	hs_a^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	$LabCh^*Fe$	DE^*Fe	hs_{a1}^*Fe	rgb^*Fe	$LabCh^*Fe$
3224	R0Y0_050_050a	0.5	0.0	0.5	0.5	0.5	390	32.4	15.4	35.9	25.4	34.6	35.9	34.6
3225	R0Y0_050_050a	0.5	0.125	0.5	0.5	0.5	376	32.7	34.0	35.9	25.4	34.6	35.9	34.6
3226	R0Y0_050_050a	0.5	0.25	0.5	0.5	0.5	360	32.9	34.0	35.9	25.4	34.6	35.9	34.6
3227	B61R_050_050a	0.5	0.375	0.5	0.5	0.5	344	33.3	0.0	0.5	0.0	0.125	34.5	35.7
3228	B0R0_050_050a	0.5	0.5	0.5	0.5	0.5	328	0.0	0.5	0.0	0.0	0.25	34.6	38.0
3229	B40R_062_062a	0.5	0.0	0.625	0.625	0.312	319	0.186	0.0	0.625	0.312	319	0.186	0.0
3230	B34R_057_075a	0.5	0.0	0.75	0.75	0.375	311	0.153	0.0	0.75	0.375	311	0.153	0.0
3231	B29R_087_087a	0.5	0.0	0.875	0.875	0.437	305	0.089	0.0	0.875	0.437	305	0.089	0.0
3232	R23Y_050_100a	0.5	0.0	1.0	1.0	0.5	300	0.045	0.0	1.0	0.5	300	0.045	0.0
3233	R23Y_050_100a	0.5	0.125	0.0	0.5	0.25	44	0.0	0.066	0.0	0.346	27.1	23.6	35.9
3234	R0Y0_050_037a	0.5	0.125	0.125	0.5	0.375	332	0.370	0.5	0.124	0.372	38.6	26.0	1.9
3235	R0Y0_050_037a	0.5	0.125	0.25	0.5	0.375	331	0.400	0.72	0.124	0.372	38.6	26.0	1.9
3236	B65R_050_037a	0.5	0.125	0.375	0.336	0.375	330	0.427	0.124	0.5	0.338	18.4	-11.2	21.6
3237	B65R_050_037a	0.5	0.125	0.5	0.5	0.375	316	0.261	0.125	0.625	0.345	19.2	-19.0	27.0
3238	B30R_062_050a	0.5	0.125	0.625	0.625	0.5	316	0.203	0.125	0.625	0.345	19.2	-19.0	27.0
3239	B30R_062_050a	0.5	0.125	0.75	0.75	0.625	0.437	0.307	0.203	0.125	0.75	0.625	0.345	19.2
3240	B25R_087_075a	0.5	0.125	0.875	0.875	0.75	0.5	0.0	0.159	0.125	0.875	0.342	19.9	-33.7
3241	B20R_100_087a	0.5	0.125	1.0	1.0	0.875	0.562	0.295	0.0	0.125	1.0	0.345	19.7	-41.4
3242	R50Y_050_037a	0.5	0.25	0.0	0.5	0.25	44	0.0	0.066	0.0	0.346	27.1	23.6	35.9
3243	R31Y_050_037a	0.5	0.25	0.125	0.5	0.375	332	0.370	0.5	0.124	0.372	38.6	26.0	1.9
3244	R0Y0_050_025a	0.5	0.25	0.25	0.5	0.25	376	0.329	0.340	0.375	25.4	34.6	35.9	34.6
3245	R0Y0_050_025a	0.5	0.25	0.375	0.360	0.345	360	0.487	0.249	0.302	44.6	16.7	17.9	25.4
3246	B30R_062_037a	0.5	0.25	0.5	0.5	0.375	330	0.351	0.249	0.5	44.5	17.8	-2.4	18.0
3247	B30R_062_037a	0.5	0.25	0.625	0.625	0.5	311	0.326	0.25	0.625	42.0	13.0	-15.1	19.9
3248	B19R_057_050a	0.5	0.25	0.75	0.75	0.5	300	0.272	0.25	0.75	41.6	13.8	-22.9	28.4
3249	B19R_057_050a	0.5	0.25	0.875	0.875	0.625	293	0.287	0.25	0.875	42.8	12.8	-22.9	28.4
3250	B13K_100_075a	0.5	0.25	1.0	1.0	0.625	289	0.25	0.35	1.0	43.5	12.6	-35.2	37.4
3251	R61Y_050_050a	0.5	0.375	0.5	0.5	0.375	276	0.351	0.249	0.302	44.6	16.7	17.9	25.4
3252	R61Y_050_050a	0.5	0.375	0.625	0.625	0.5	276	0.351	0.249	0.302	44.6	16.7	17.9	25.4
3253	R0Y0_050_012a	0.5	0.375	0.625	0.625	0.5	276	0.351	0.249	0.302	44.6	16.7	17.9	25.4
3254	R0Y0_050_012a	0.5	0.375	0.75	0.75	0.625	289	0.375	0.475	0.875	53.7	62.2	-23.2	24.1
3255	B50R_062_050a	0.5	0.375	0.75	0.75	0.625	289	0.375	0.475	0.875	53.7	62.2	-23.2	24.1
3256	B50R_062_050a	0.5	0.375	0.875	0.875	0.75	281	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3257	B11R_087_050a	0.5	0.375	1.0	1.0	0.625	281	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3258	B0R0_050_062a	0.5	0.5	0.5	0.5	0.5	280	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3259	B0R0_050_062a	0.5	0.625	0.625	0.625	0.312	270	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3260	Y0C0_050_037a	0.5	0.625	0.625	0.625	0.312	270	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3261	Y0C0_050_037a	0.5	0.625	0.75	0.75	0.625	289	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3262	Y0C0_050_037a	0.5	0.625	0.875	0.875	0.75	281	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3263	Y0C0_050_037a	0.5	0.625	1.0	1.0	0.875	281	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3264	NW_050a	0.5	0.5	0.5	0.5	0.5	360	0.5	0.5	0.5	56.5	0.0	0.0	0.0
3265	B0R0_062_012a	0.5	0.5	0.625	0.625	0.312	270	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3266	B0R0_062_012a	0.5	0.5	0.75	0.75	0.625	270	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3267	B0R0_062_012a	0.5	0.5	0.875	0.875	0.75	270	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3268	B0R0_100_050a	0.5	0.5	1.0	1.0	0.5	270	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3269	Y18C_062_062a	0.5	0.625	0.625	0.625	0.312	270	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3270	Y23C_062_050a	0.5	0.625	0.625	0.625	0.312	270	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3271	Y31C_062_037a	0.5	0.625	0.625	0.625	0.312	270	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3272	Y31C_062_037a	0.5	0.625	0.75	0.75	0.625	289	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3273	Y31C_062_037a	0.5	0.625	0.875	0.875	0.75	289	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3274	G0B0_062_012a	0.5	0.625	0.625	0.625	0.312	270	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3275	G0B0_062_012a	0.5	0.625	0.75	0.75	0.625	289	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3276	G0B0_062_012a	0.5	0.625	0.875	0.875	0.75	289	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3277	G0B0_062_012a	0.5	0.625	1.0	1.0	0.875	289	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3278	G0B0_062_012a	0.5	0.625	1.0	1.0	0.875	289	0.375	0.526	1.0	56.4	62.2	-23.2	24.1
3279	Y31C_075_075a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3280	Y30C_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3281	Y0C0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3282	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3283	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3284	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3285	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3286	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3287	Y41C_087_087a	0.5	0.875	0.875	0.875	0.437	115	0.343	0.875	0.0	62.9	31.6	51.8	60.7
3288	Y41C_087_087a	0.5	0.875	0.875	0.875	0.437	115	0.343	0.875	0.0	62.9	31.6	51.8	60.7
3289	Y60C_087_062a	0.5	0.875	0.875	0.875	0.625	127	0.401	0.875	0.125	63.5	31.0	48.0	51.2
3290	Y60C_087_062a	0.5	0.875	0.875	0.875	0.625	127	0.401	0.875	0.125	63.5	31.0	48.0	51.2
3291	G0B0_087_037a	0.5	0.875	0.875	0.875	0.375	136	0.401	0.875	0.375	66.4	28.1	29.6	34.0
3292	G0B0_087_037a	0.5	0.875	0.875	0.875	0.375	136	0.401	0.875	0.375	66.4	28.1	29.6	34.0
3293	G0B0_087_037a	0.5	0.875	0.875	0.875	0.375	136	0.401	0.875	0.375	66.4	28.1	29.6	34.0
3294	G0B0_087_037a	0.5	0.875	0.875	0.875	0.375	136	0.401	0.875	0.375	66.4	28.1	29.6	34.0
3295	G0B0_087_037a	0.5	0.875	0.875	0.875	0.375	136	0.401	0.875	0.375	66.4	28.1	29.6	34.0
3296	G0B0_087_037a	0.5	0.875	0.875	0.875	0.375	136	0.401	0.875	0.375	66.4	28.1	29.6	34.0
3297	Y31C_075_075a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3298	Y30C_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3299	Y0C0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3300	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3301	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3302	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3303	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3304	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3305	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3306	G0B0_075_050a	0.5	0.75	0.75	0.75	0.75	256	0.5	0.771	1.0	69.9	4.3	-22.4	22.9
3307	G0													

see similar files: <http://130.149.60.45/~farbmatrik/RE45/RE45.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

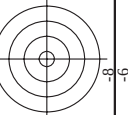
TUB-test chart RE45; hue code: $H_e^* = B75R_e$
colors and differences. ΔE^* ,

RE450-7N, Page 25/33-F

Mean color difference of this page:

input: *rgb/cmyk* → *rgbe*
output: transfer to *cmyke*

n	HIC ^{Fe}	rgb^{Fe}	lcr^{Fe}	h_{α}^{Fe}	rgb^{*Fe}	$LabCh^{*Fe}$	$LabCh^{Fe}$	rgb^{*Fe}	DE^{*Fe}	h_{α}^{*Fe}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$	$LabCh^{Me}$	rgb^{*Me}	DE^{*Me}	h_{α}^{*Me}	rgb^{*Me}	$LabCh^{*Me}$
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TUB material: code=rha4ta

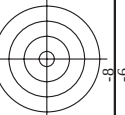
A target diagram with a score of 86. The score is written as 86, with a vertical line through the 6, indicating a score of 86.

input: *rgb/cmyk* → *rgb*
output: transfer to *cmyk*

TUB-
colors

see similar files: <http://130.149.60.45/~farbmatrik/RE45/RE45.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

[illegible]



TUB material: code=rha4ta
nyn6 (CMYK)

input: *rgb/cmyk* → *rgb*_e
output: transfer to *cmyk*_e

TUB-test chart RE45; hue colors and differences, ΔE^* ,[illegible]



TUB material: code=rha4ta
nyn6 (CMYK)

input: *rgb/cmyk* → *rgb*_e
output: transfer to *cmyk*_e

1

see similar files: <http://130.149.60.45/~farbmatrik/RE45/RE45.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

http://130.149.60.45/~farbmatrik/RE45/RE45L0NA.TXT /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 30/33

TUUB-test chart RE45; hue code: $H^*_e=B75R_e$
 colors and differences. ΔE^* ,

input: *rgb/cmyk* $\rightarrow$ *rgbe*
output: transfer to *cmvke*

[illegible]

<http://130.149.60.45/~farbmatrik/RE45/RE45L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 31/33

n	HIC*Fe	rgb-Fe	icL-Fe	hsL-Fe	LabCh*Fe	rgb*Fe	LabCh*Fe	rgb*Fe	LabCh*Fe	rgb*Me	LabCh*Me	0.0	0.0	0.0
891	NW_100e	1.0	1.0	1.0	95.4	1.0	95.4	1.0	139.6	0.0	1.0	95.4	0.0	0.0
892	B50R_100.012e	1.0	0.875	1.0	87.9	1.0	87.9	1.0	342.7	3.4	0.97	87.9	0.0	0.0
893	B50R_100.025e	1.0	0.75	1.0	80.3	1.0	80.3	1.0	345.3	6.1	0.97	80.3	0.0	0.0
894	B50R_100.037e	1.0	0.625	1.0	72.7	1.0	72.7	1.0	348.6	9.4	0.97	72.7	0.0	0.0
895	B50R_100.050e	1.0	0.5	1.0	65.1	1.0	65.1	1.0	348.6	13.0	0.97	65.1	0.0	0.0
896	B50R_100.062e	1.0	0.375	1.0	57.5	1.0	57.5	1.0	348.6	16.6	0.97	57.5	0.0	0.0
897	B50R_100.075e	1.0	0.25	1.0	50.0	1.0	50.0	1.0	348.6	20.2	0.97	50.0	0.0	0.0
898	B50R_100.087e	1.0	0.125	1.0	42.4	1.0	42.4	1.0	348.6	23.8	0.97	42.4	0.0	0.0
899	B50R_100.100e	1.0	0.0	1.0	34.8	1.0	34.8	1.0	348.6	27.4	0.97	34.8	0.0	0.0
900	GOB_100.012e	0.875	1.0	0.875	95.4	0.875	95.4	0.875	139.6	0.0	0.875	95.4	0.0	0.0
901	NW_087e	0.875	0.875	0.875	87.9	0.875	87.9	0.875	342.7	3.4	0.875	87.9	0.0	0.0
902	B50R_087.012e	0.875	0.75	0.875	80.3	0.875	80.3	0.875	345.3	6.1	0.875	80.3	0.0	0.0
903	B50R_087.025e	0.875	0.625	0.875	72.7	0.875	72.7	0.875	348.6	9.4	0.875	72.7	0.0	0.0
904	B50R_087.037e	0.875	0.5	0.875	65.1	0.875	65.1	0.875	348.6	13.0	0.875	65.1	0.0	0.0
905	B50R_087.050e	0.875	0.375	0.875	57.5	0.875	57.5	0.875	348.6	16.6	0.875	57.5	0.0	0.0
906	B50R_087.062e	0.875	0.25	0.875	50.0	0.875	50.0	0.875	348.6	20.2	0.875	50.0	0.0	0.0
907	B50R_087.075e	0.875	0.125	0.875	42.4	0.875	42.4	0.875	348.6	23.8	0.875	42.4	0.0	0.0
908	B50R_087.087e	0.875	0.0	0.875	34.8	0.875	34.8	0.875	348.6	27.4	0.875	34.8	0.0	0.0
909	GOB_100.025e	0.75	1.0	0.75	87.9	0.75	87.9	0.75	139.6	0.0	0.75	87.9	0.0	0.0
910	GOB_100.037e	0.75	0.875	0.75	80.3	0.75	80.3	0.75	342.7	3.4	0.75	80.3	0.0	0.0
911	NW_075e	0.75	0.75	0.75	72.7	0.75	72.7	0.75	345.3	6.1	0.75	72.7	0.0	0.0
912	B50R_075.012e	0.75	0.625	0.75	65.1	0.75	65.1	0.75	348.6	9.4	0.75	65.1	0.0	0.0
913	B50R_075.025e	0.75	0.5	0.75	57.5	0.75	57.5	0.75	348.6	13.0	0.75	57.5	0.0	0.0
914	B50R_075.037e	0.75	0.375	0.75	50.0	0.75	50.0	0.75	348.6	16.6	0.75	50.0	0.0	0.0
915	B50R_075.050e	0.75	0.25	0.75	42.4	0.75	42.4	0.75	348.6	20.2	0.75	42.4	0.0	0.0
916	B50R_075.062e	0.75	0.125	0.75	34.8	0.75	34.8	0.75	348.6	23.8	0.75	34.8	0.0	0.0
917	GOB_100.037e	0.75	0.875	0.75	80.3	0.75	80.3	0.75	342.7	3.4	0.75	80.3	0.0	0.0
918	GOB_100.050e	0.625	1.0	0.625	95.4	0.625	95.4	0.625	139.6	0.0	0.625	95.4	0.0	0.0
919	GOB_100.062e	0.625	0.875	0.625	87.9	0.625	87.9	0.625	342.7	3.4	0.625	87.9	0.0	0.0
920	GOB_100.075e	0.625	0.75	0.625	80.3	0.625	80.3	0.625	345.3	6.1	0.625	80.3	0.0	0.0
921	NW_062e	0.625	0.625	0.625	72.7	0.625	72.7	0.625	348.6	9.4	0.625	72.7	0.0	0.0
922	B50R_062.012e	0.625	0.5	0.625	65.1	0.625	65.1	0.625	348.6	13.0	0.625	65.1	0.0	0.0
923	B50R_062.025e	0.625	0.375	0.625	57.5	0.625	57.5	0.625	348.6	16.6	0.625	57.5	0.0	0.0
924	B50R_062.037e	0.625	0.25	0.625	50.0	0.625	50.0	0.625	348.6	20.2	0.625	50.0	0.0	0.0
925	B50R_062.050e	0.625	0.125	0.625	42.4	0.625	42.4	0.625	348.6	23.8	0.625	42.4	0.0	0.0
926	B50R_062.062e	0.625	0.0	0.625	34.8	0.625	34.8	0.625	348.6	27.4	0.625	34.8	0.0	0.0
927	GOB_100.050e	0.5	1.0	0.5	95.4	0.5	95.4	0.5	139.6	0.0	0.5	95.4	0.0	0.0
928	GOB_087.037e	0.5	0.875	0.5	87.9	0.5	87.9	0.5	342.7	3.4	0.5	87.9	0.0	0.0
929	GOB_087.050e	0.5	0.75	0.5	80.3	0.5	80.3	0.5	345.3	6.1	0.5	80.3	0.0	0.0
930	GOB_087.062e	0.5	0.625	0.5	72.7	0.5	72.7	0.5	348.6	9.4	0.5	72.7	0.0	0.0
931	NW_050e	0.5	0.5	0.5	65.1	0.5	65.1	0.5	348.6	13.0	0.5	65.1	0.0	0.0
932	B50R_050.012e	0.5	0.375	0.5	50.0	0.5	50.0	0.5	348.6	16.6	0.5	50.0	0.0	0.0
933	B50R_050.025e	0.5	0.25	0.5	42.4	0.5	42.4	0.5	348.6	20.2	0.5	42.4	0.0	0.0
934	B50R_050.037e	0.5	0.125	0.5	34.8	0.5	34.8	0.5	348.6	23.8	0.5	34.8	0.0	0.0
935	B50R_050.050e	0.5	0.0	0.5	27.4	0.5	27.4	0.5	348.6	27.4	0.5	27.4	0.0	0.0
936	GOB_100.062e	0.375	1.0	0.375	95.4	0.375	95.4	0.375	139.6	0.0	0.375	95.4	0.0	0.0
937	GOB_087.050e	0.375	0.875	0.375	87.9	0.375	87.9	0.375	342.7	3.4	0.375	87.9	0.0	0.0
938	GOB_087.062e	0.375	0.75	0.375	80.3	0.375	80.3	0.375	345.3	6.1	0.375	80.3	0.0	0.0
939	GOB_087.075e	0.375	0.625	0.375	72.7	0.375	72.7	0.375	348.6	9.4	0.375	72.7	0.0	0.0
940	GOB_087.087e	0.375	0.5	0.375	65.1	0.375	65.1	0.375	348.6	13.0	0.375	65.1	0.0	0.0
941	NW_037e	0.375	0.375	0.375	57.5	0.375	57.5	0.375	348.6	16.6	0.375	57.5	0.0	0.0
942	B50R_037.012e	0.375	0.25	0.375	50.0	0.375	50.0	0.375	348.6	20.2	0.375	50.0	0.0	0.0
943	B50R_037.025e	0.375	0.125	0.375	42.4	0.375	42.4	0.375	348.6	23.8	0.375	42.4	0.0	0.0
944	B50R_037.037e	0.375	0.0	0.375	34.8	0.375	34.8	0.375	348.6	27.4	0.375	34.8	0.0	0.0
945	GOB_100.075e	0.25	1.0	0.25	95.4	0.25	95.4	0.25	139.6	0.0	0.25	95.4	0.0	0.0
946	GOB_087.062e	0.25	0.875	0.25	87.9	0.25	87.9	0.25	342.7	3.4	0.25	87.9	0.0	0.0
947	GOB_087.075e	0.25	0.75	0.25	80.3	0.25	80.3	0.25	345.3	6.1	0.25	80.3	0.0	0.0
948	GOB_087.087e	0.25	0.625	0.25	72.7	0.25	72.7	0.25	348.6	9.4	0.25	72.7	0.0	0.0
949	GOB_087.090e	0.25	0.5	0.25	65.1	0.25	65.1	0.25	348.6	13.0	0.25	65.1	0.0	0.0
950	GOB_087.102e	0.25	0.375	0.25	57.5	0.25	57.5	0.25	348.6	16.6	0.25	57.5	0.0	0.0
951	NW_025e	0.25	0.25	0.25	50.0	0.25	50.0	0.25	348.6	20.2	0.25	50.0	0.0	0.0
952	B50R_025.012e	0.25	0.125	0.25	42.4	0.25	42.4	0.25	348.6	23.8	0.25	42.4	0.0	0.0
953	B50R_025.025e	0.25	0.0	0.25	34.8	0.25	34.8	0.25	348.6	27.4	0.25	34.8	0.0	0.0
954	GOB_100.087e	0.125	1.0	0.125	95.4	0.125	95.4	0.125	139.6	0.0	0.125	95.4	0.0	0.0
955	GOB_087.075e	0.125	0.875	0.125	87.9	0.125	87.9	0.125	342.7	3.4	0.125	87.9	0.0	0.0
956	GOB_087.087e	0.125	0.75	0.125	80.3	0.125	80.3	0.125	345.3	6.1	0.125	80.3	0.0	0.0
957	GOB_087.090e	0.125	0.625	0.125	72.7	0.125	72.7	0.125	348.6	9.4	0.125	72.7	0.0	0.0
958	GOB_087.102e	0.125	0.5	0.125	65.1	0.125	65.1	0.125	348.6	13.0	0.125	65.1	0.0	0.0
959	GOB_087.037e	0.125	0.375	0.125	57.5	0.125	57.5	0.125	348.6	16.6	0.125	57.5	0.0	0.0
960	GOB													

n	H*Ch*Fe	rgb*Fe	icL*Fe	hsa*Fe	rgb*Fe	LabCh*Fe	LabCh*Fe	rgb*Fe	DF*Fe	hsa*Fe	rgb*Fe	LabCh*Fe	LabCh*Fe
972	NW 000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
973	NW 012e	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
974	NW 025e	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
975	NW 037e	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
976	NW 050e	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
977	NW 062e	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
978	NW 075e	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
979	NW 087e	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
980	NW 100e	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
981	NW 000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
982	NW 012e	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
983	NW 025e	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
984	NW 037e	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
985	NW 050e	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
986	NW 062e	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
987	NW 075e	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
988	NW 087e	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
989	NW 100e	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
990	NW 000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
991	NW 012e	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
992	NW 025e	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
993	NW 037e	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
994	NW 050e	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
995	NW 062e	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
996	NW 075e	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
997	NW 087e	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
998	NW 100e	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
999	NW 000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1000	NW 012e	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
1001	NW 025e	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
1002	NW 037e	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
1003	NW 050e	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
1004	NW 062e	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
1005	NW 075e	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
1006	NW 087e	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
1007	NW 100e	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1008	NW 000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1009	NW 012e	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
1010	NW 025e	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
1011	NW 037e	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
1012	NW 050e	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
1013	NW 062e	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
1014	NW 075e	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
1015	NW 087e	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
1016	NW 100e	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1017	NW 000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1018	NW 012e	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
1019	NW 025e	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
1020	NW 037e	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
1021	NW 050e	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
1022	NW 062e	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
1023	NW 075e	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
1024	NW 087e	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
1025	NW 100e	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1026	NW 000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1027	NW 012e	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
1028	NW 025e	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
1029	NW 037e	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
1030	NW 050e	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
1031	NW 062e	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
1032	NW 075e	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
1033	NW 087e	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
1034	NW 100e	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1035	NW 000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1036	NW 012e	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
1037	NW 025e	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
1038	NW 037e	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
1039	NW 050e	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
1040	NW 062e	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
1041	NW 075e	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
1042	NW 087e	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
1043	NW 100e	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1044	NW 000e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1045	NW 012e	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125	0.125
1046	NW 025e	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
1047	NW 037e	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375	0.375
1048	NW 050e	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.5
1049	NW 062e	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625	0.625
1050	NW 075e	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75	0.75
1051	NW 087e	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875	0.875
1052	NW 100e	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0

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TUB-test chart RE45; hue code: H*e=B75Re
colors and differences, ΔE*

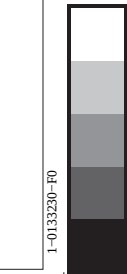
input: rgb/cmyk -> rgbe
output: transfer to cmyke

Mean color difference of this page:

delta E* = 5.5



TUB material: code=rha4ta
nyn6 (CMYK)



input: *rgb/cmyk* → *rgbe*
output: transfer to *cmyk_e*

TUB-test chart RE45; hue code: $H_e^* = B75R_e$
colors and differences, ΔE^* ,

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1-0133230-F0

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