

input: *rgb/cmyk* $\rightarrow$ *rgb/cmyk*
output: no change

Input and Output: Offset Reflective System ORS18a for relative CIELAB hue $h_{ab,a,rel} = h_{ab}/360 = 328/360 = 0.91$

$H^*_e = B50R_e$

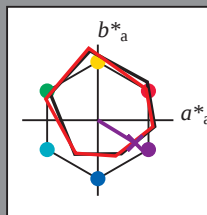
Data for any device (d) or elementary (e) colour:

HIC^*_e

hue text for the colours of this page:

$H^*_e = B50R_e$

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
Y _e ,Ma	82.9	-3.5	87.8	87.9
G _e ,Ma	52.4	-67.1	21.5	70.5
C _e ,Ma	56.6	-39.7	-29.9	49.8
B _e ,Ma	37.9	1.3	-45.4	45.4
M _e ,Ma	34.8	49.2	-30.0	57.7
N _e ,Ma	17.7	0.0	0.0	0
W _e ,Ma	95.4	0.0	0.0	0
R _e ,CIE	39.9	58.7	27.9	65.0
Y _e ,CIE	81.2	-2.8	71.5	71.6
G _e ,CIE	52.2	-42.4	13.6	44.5
B _e ,CIE	30.5	1.4	-46.4	46.4

Data for maximum colour (Ma):

$LabCh^*_e, Ma: 34\ 49\ -30\ 57\ 328$

$HIC^*_e, Ma: B50R_100_100_e$

$rgbic^*_e, Ma:$

0.4 0.0 1.0 1.0 1.0

triangle lightness T^*

% Gamut

$u^*_{rel} = 92$

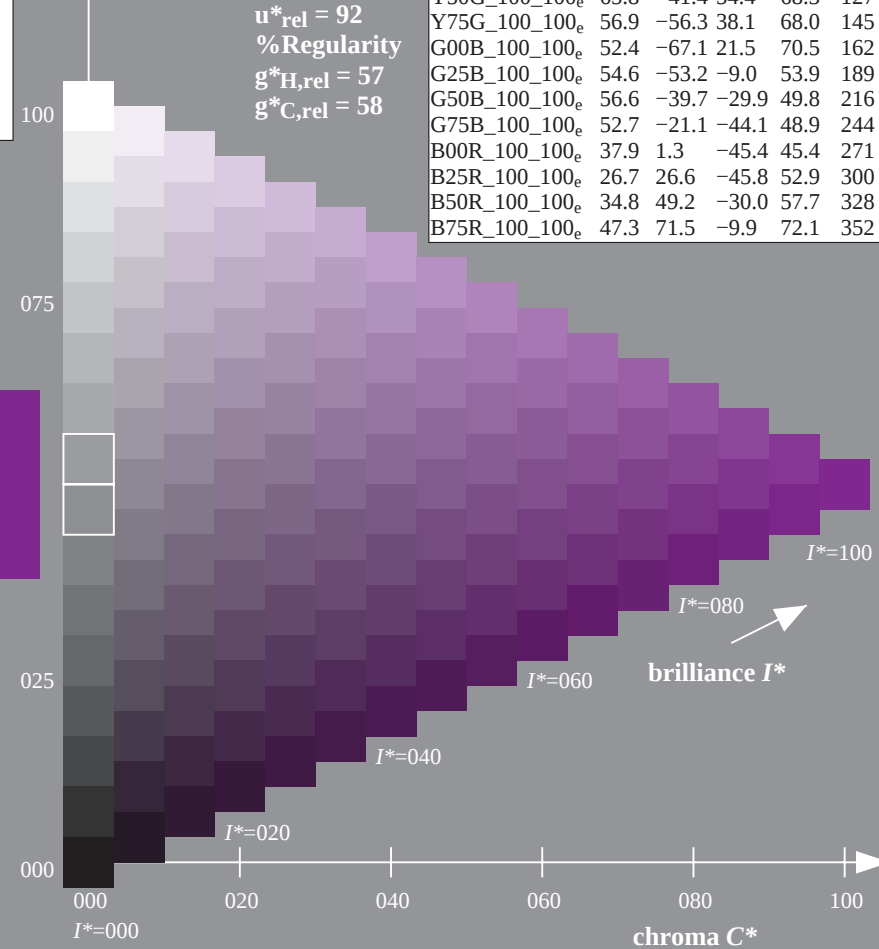
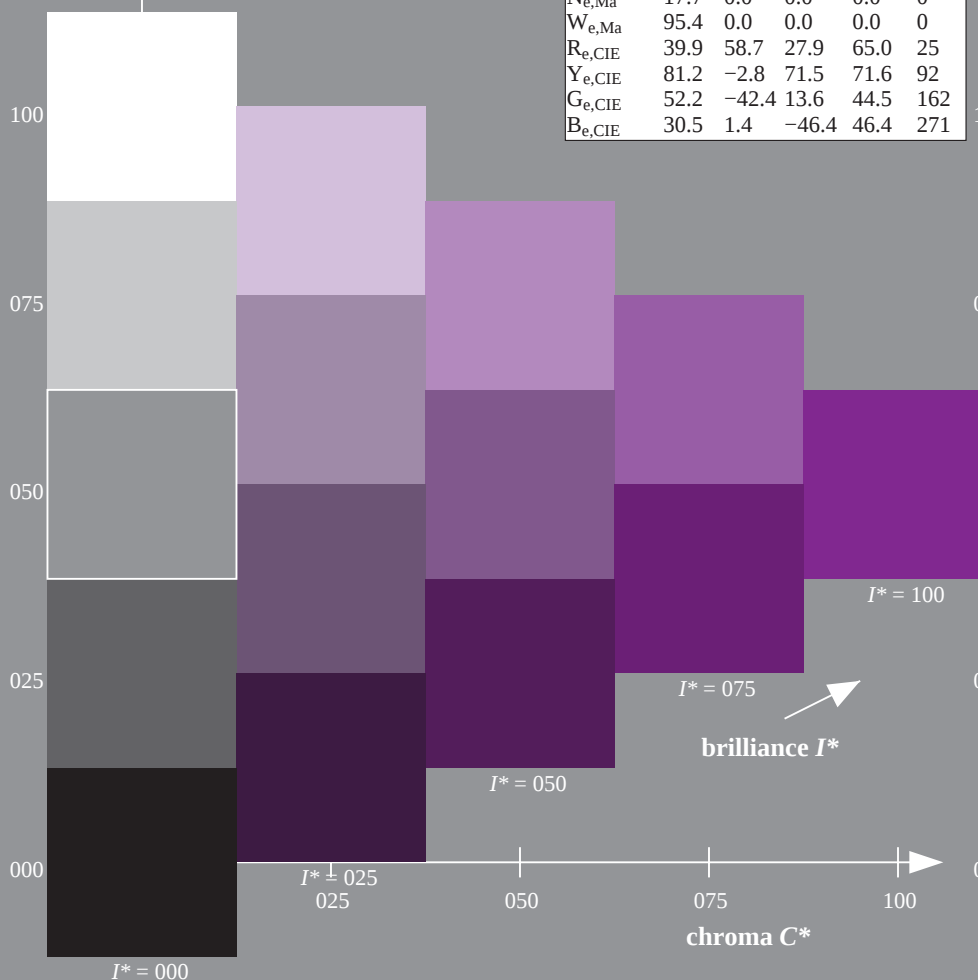
% Regularity

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

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

ORS20a; adapted (a) CIELAB data

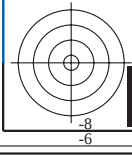
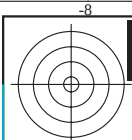
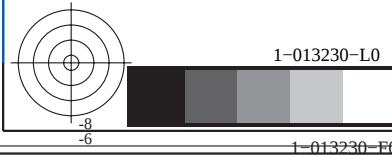
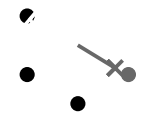
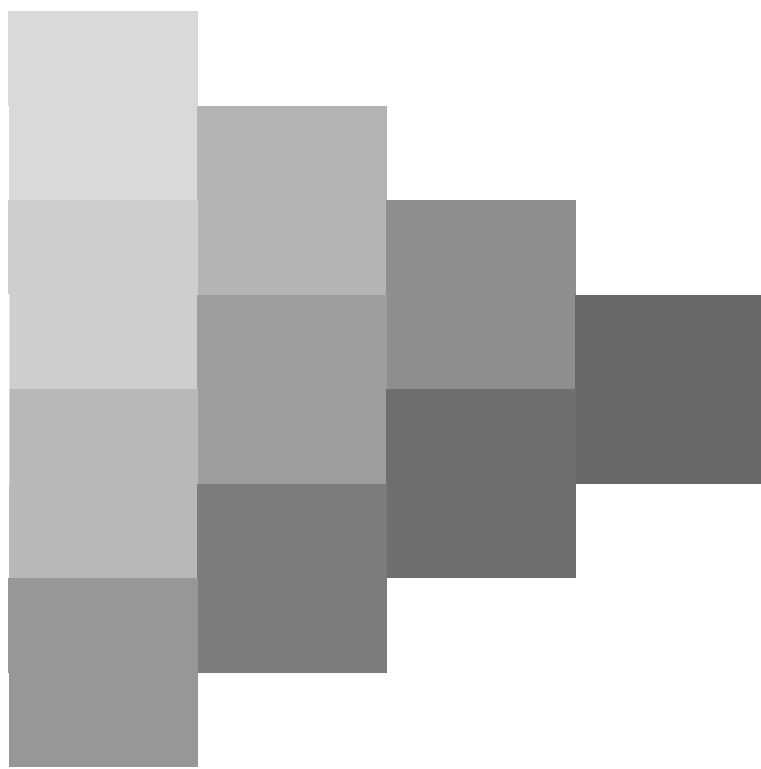
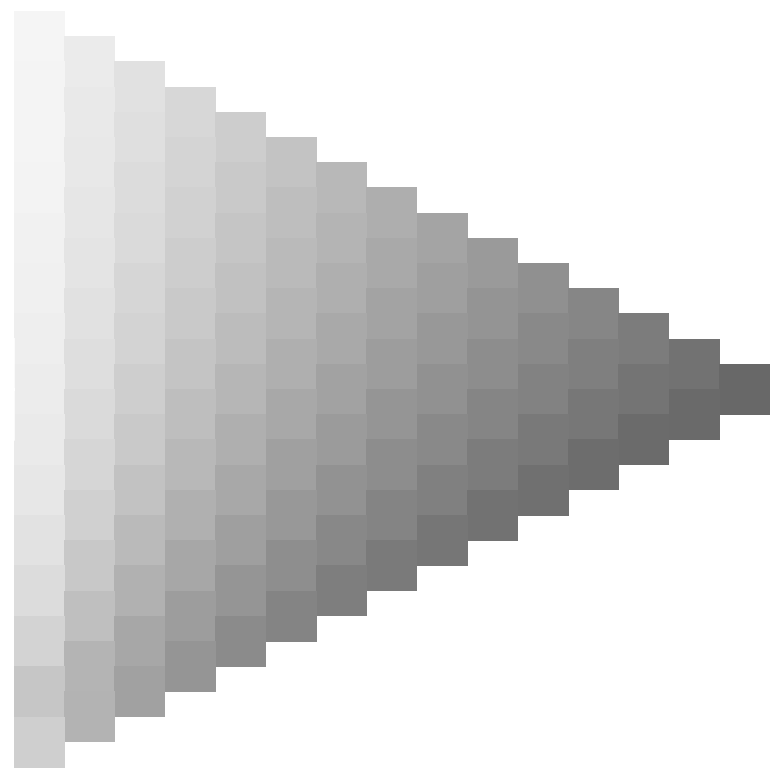
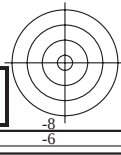
H^*_e	$L^*=L^*_a a^*_a$	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
R00Y_100_100 _e	47.6	64.9	30.9	71.9
R25Y_100_100 _e	51.5	54.2	47.2	71.9
R50Y_100_100 _e	60.3	35.6	59.0	68.9
R75Y_100_100 _e	70.4	17.0	72.2	74.1
Y00G_100_100 _e	82.9	-3.5	87.8	87.9
Y25G_100_100 _e	76.9	-25.5	75.9	80.1
Y50G_100_100 _e	65.8	-41.4	54.4	68.3
Y75G_100_100 _e	56.9	-56.3	38.1	68.0
G00B_100_100 _e	52.4	-67.1	21.5	70.5
G25B_100_100 _e	54.6	-53.2	-9.0	53.9
G50B_100_100 _e	56.6	-39.7	-29.9	49.8
G75B_100_100 _e	52.7	-21.1	-44.1	48.9
B00R_100_100 _e	37.9	1.3	-45.4	45.4
B25R_100_100 _e	26.7	26.6	-45.8	52.9
B50R_100_100 _e	34.8	49.2	-30.0	57.7
B75R_100_100 _e	47.3	71.5	-9.9	72.1



1-013130-L0 RE350-71

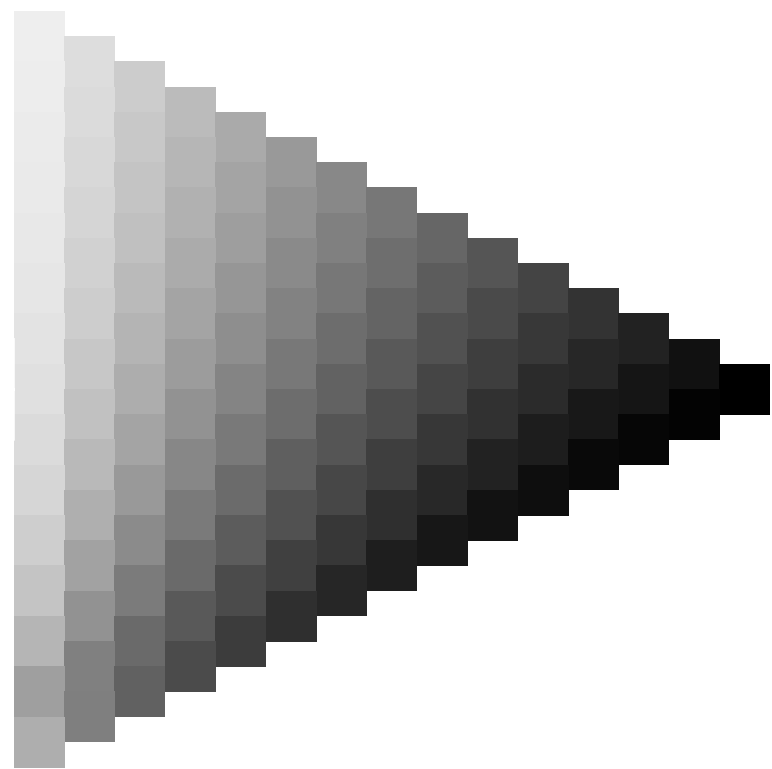
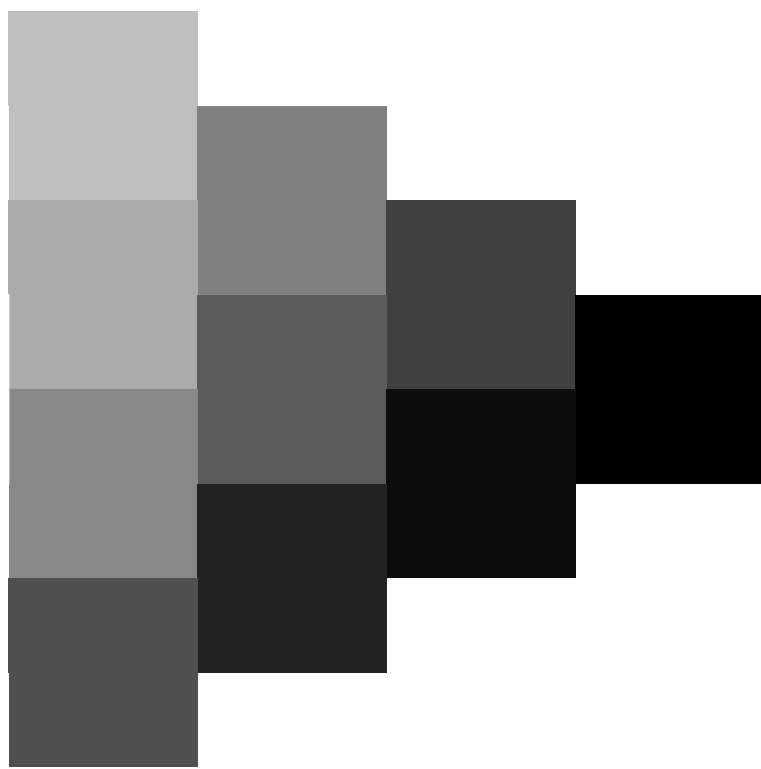
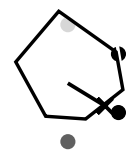
TUB-test chart RE35; hue code: $H^*_e = B50R_e$
Test chart according to DIN 33872, 3D=0, de=1, cmyk

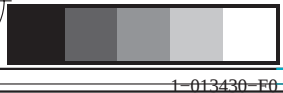
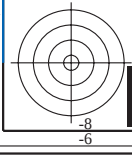
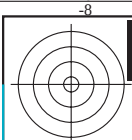
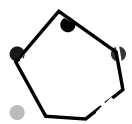
input: $rgb/cmyk \rightarrow rgb_e$
output: transfer to $cmyk_e$





see similar files: <http://130.149.60.45/~farbmetrik/RE35/RE35.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>





Input and Output: Offset Reflective System ORS18a for relative CIELAB hue $h_{ab,a,rel} = h_{ab}/360 = 328/360 = 0.91$

$H^*_e = B50R_e$

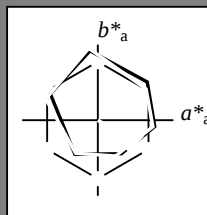
Data for any device (d) or elementary (e) colour:

HIC^*_e

hue text for the colours of this page:

$H^*_e = B50R_e$

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):

$LabCh^*_e, Ma: 34\ 49\ -30\ 57\ 328$

$HIC^*_e, Ma: B50R_100_100_e$

$rgbic^*_e, Ma:$

0.4 0.0 1.0 1.0 1.0

triangle lightness T^*

% Gamut

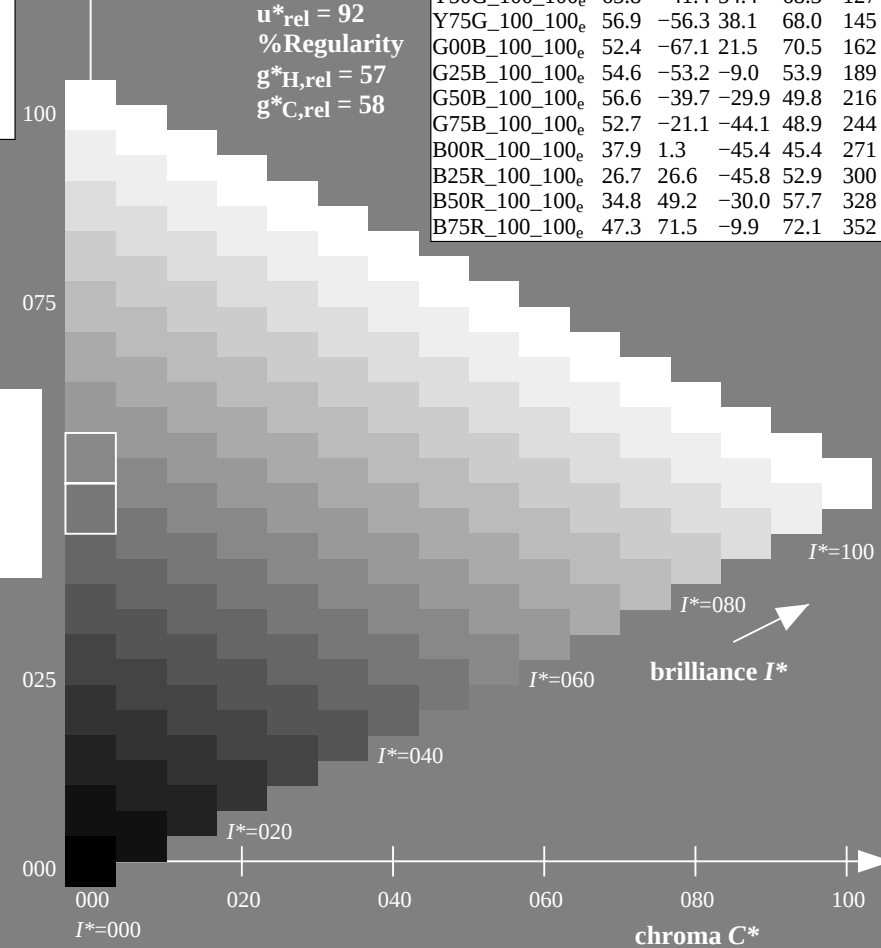
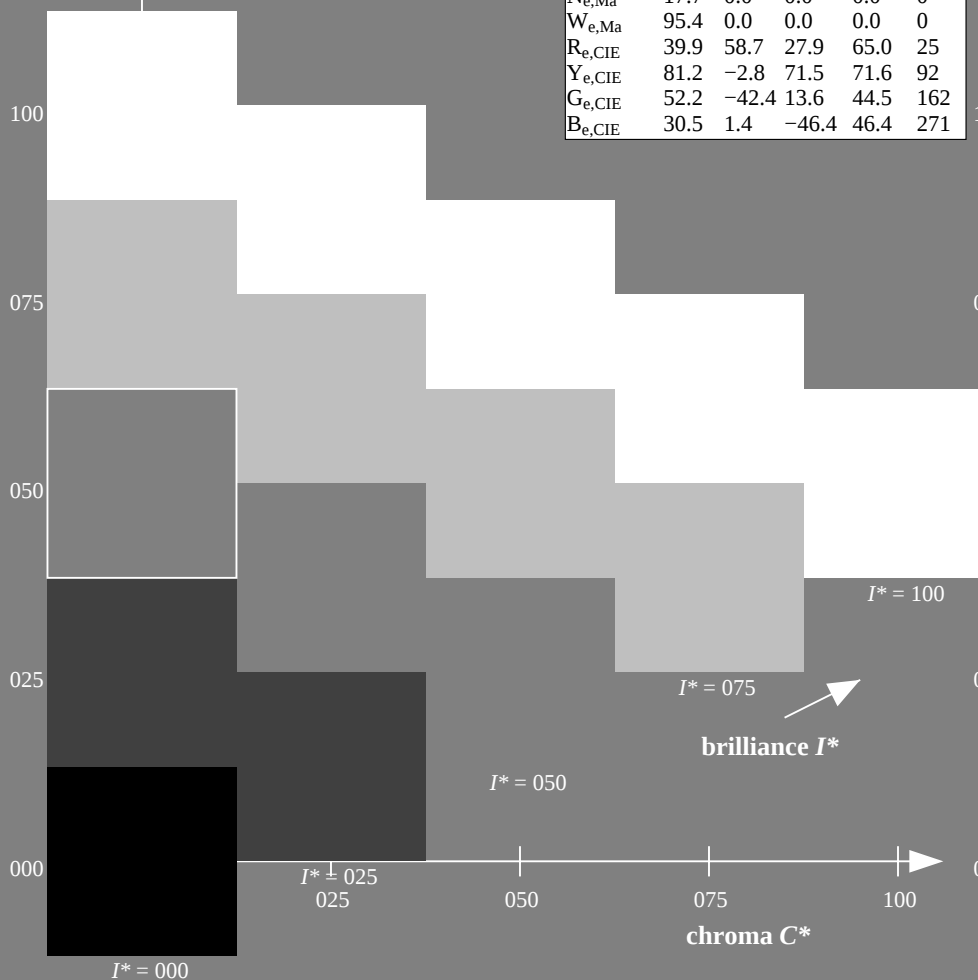
$u^*_{rel} = 92$

% Regularity

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

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

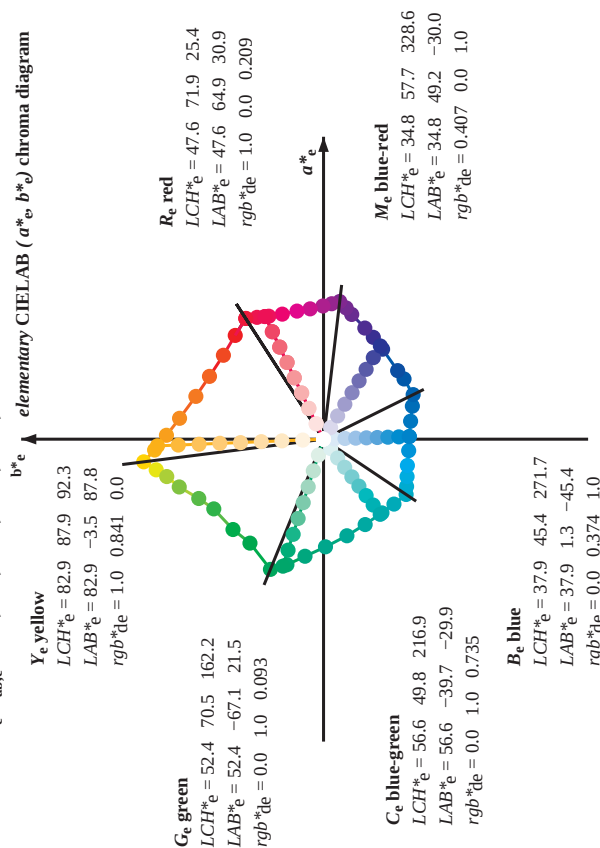
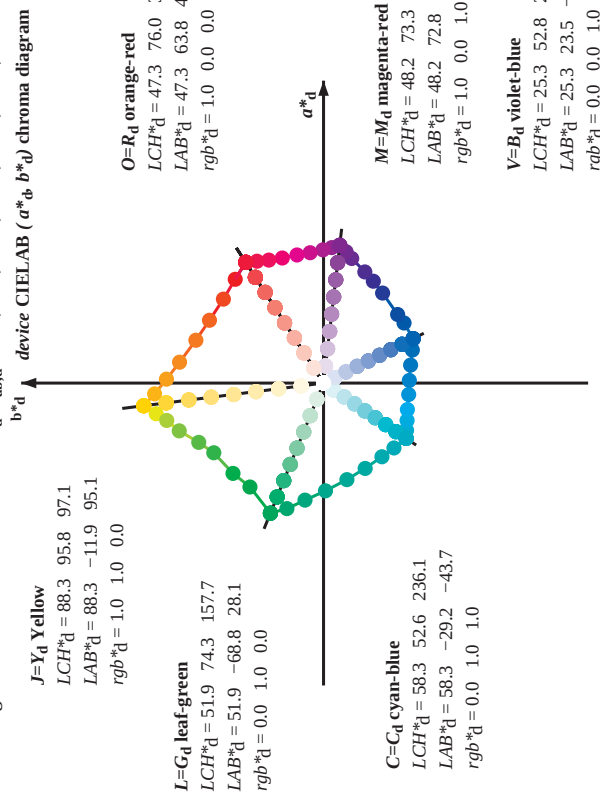
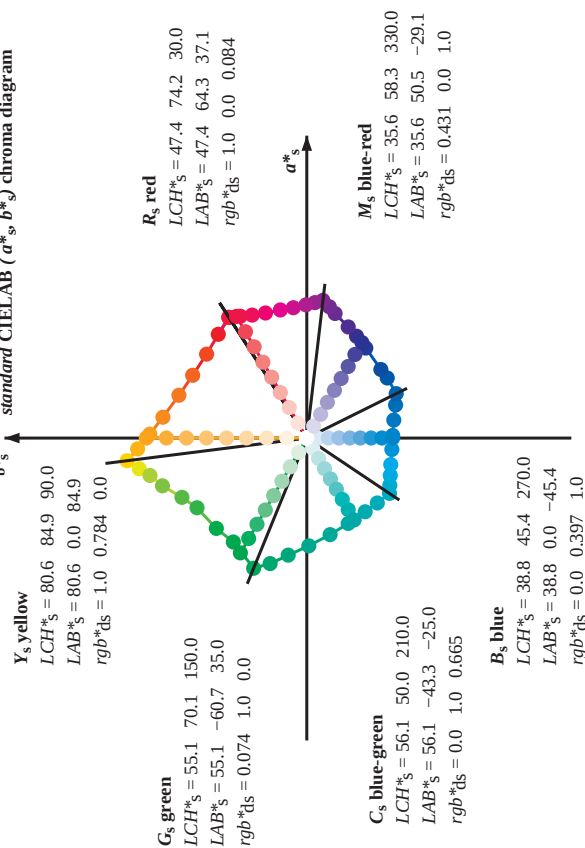
ORS20a; adapted (a) CIELAB data					
H^*_e	$L^*=L^*_a$	a^*_a	b^*_a	$C^*_{ab,a}$	$h^*_{ab,a}$
$R00Y_100_100_e$	47.6	64.9	30.9	71.9	25
$R25Y_100_100_e$	51.5	54.2	47.2	71.9	41
$R50Y_100_100_e$	60.3	35.6	59.0	68.9	58
$R75Y_100_100_e$	70.4	17.0	72.2	74.1	76
$Y00G_100_100_e$	82.9	-3.5	87.8	87.9	92
$Y25G_100_100_e$	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 RE35; hue code: $H^*_e = B50R_e$
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_e; $h_{ab,d} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours RYGBCM_d; $h_{ab,d} = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3$; Six hue angles of the elementary colours RYGBCM_e; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

**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:
$$h_{48ab,sij} = h_{ab,si} + j \frac{h_{ab,si+1} - h_{ab,si}}{1/8} \quad (i=0,1,...,5; j=0,1,...,7) \quad (2)$$
$$h_{360ab,sij} = h_{ab,si} + j \frac{h_{ab,si+1} - h_{ab,si}}{1/60} \quad (i=0,1,...,5; j=0,1,...,59) \quad (3)$$
- 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:
$$h_{48ab,eij} = h_{ab,ei} + j \frac{h_{ab,ei+1} - h_{ab,ei}}{1/8} \quad (i=0,1,...,5; j=0,1,...,7) \quad (4)$$
$$h_{360ab,eij} = h_{ab,ei} + j \frac{h_{ab,ei+1} - h_{ab,ei}}{1/60} \quad (i=0,1,...,5; j=0,1,...,59) \quad (5)$$
- 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

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

input: $rgb/cmyk \rightarrow rgb_e$
output: transfer to $cmyk_e$ TUB-test chart RE35; hue code: $H^*_e=B50R_e$
48 step hue circles; $rgb-LabCh$ -tables

<http://130.149.60.45/~farbmatrik/RE35/RE35LONP.PDF> /PS; transfer output N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 10/33

Data of Maximum color M in colorimetric system Offset standard print; separation cmv66*, D65 for input or output; Six hue angles of the 60 degree standard colours RYGCBS; $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;

Six hue angles of the device colours RYGBM: $h_{\text{ab,d}} = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3$; Six hue angles of the elementary colours RYGBM: $h_{\text{ab,e}} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

see similar files: <http://130.149.60.45/~farbmatrik/RE35/RE35.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

RE350-71	LAB* _{a0} , YN=0%, XY _{Znw} =2.4, 2.5, 2.6, 85.1, 88.8, 104.3, LAB* _{nw} =17.7, 0.0, 0.0, 95.5, 0.0, 0.0	input: <i>rgb/cmyk</i> → <i>rgbe</i> output: transfer to <i>cmyk_e</i>
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TUB-test chart RE35; hue code: H*_e=B50R_e
48 step hue circles: *rgb-LabCh**tables

1-013030-I 0	RF350-71	LAB*a0	YN=0%	XYZ _{nw} =24.2	5.26	85.1888	104.3	LAB*nw=177.0	0.0	95.5	0.0	0.0
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Output: Offset standard print: separation cmv6*: D65, page 10/33

Output: Offset standard print: separation cmv6*: D65, page 10/33

technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

<http://130.149.60.45/~farbmatrik/RE35/RE35LONP.PDF> /PS; transfer output N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 12/33

<http://130.149.60.45/~farbmatrik/RE35/RE35LONP.PDF> /PS; transfer output N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 12/33

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 RYGCMB*: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;

Six hue angles of the device colours $RYGBM_d$: $h_{ab,d} = 32.8, 97.2, 157.8, 236.2, 296.4, 353.3$; Six hue angles of the elementary colours $RYGBM_e$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$										$rgb^*_{d361Mi}(x^*, y^*, z^*)$																						
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_{d361M}	$LAB^*_{d361Mi}(x^*, y^*, z^*)$	rgb^*_{s361Mi}	$LAB^*_{s361Mi}(x^*, y^*, z^*)$	rgb^*_{d361Mi}	$LAB^*_{d361Mi}(x^*, y^*, z^*)$	rgb^*_{e361Mi}	$LAB^*_{e361Mi}(x^*, y^*, z^*)$	rgb^*_{d361Mi}	$LAB^*_{d361Mi}(x^*, y^*, z^*)$																				
127	115	120	0.5	1.0	0.0	72.7	-31.3	66.0	73.1	115	0.418	1.0	0.0	70.7	-35.1	60.9	70.7	121	0.5	1.0	0.0	0.327	1.0	0.0	65.8	-41.3	54.4	68.4	127	0.5	1.0	0.0
128	116	121	0.483	1.0	0.0	72.2	-32.1	65.0	72.5	116	0.4	1.0	0.0	69.7	-35.8	59.8	69.7	121	0.483	1.0	0.0	0.315	1.0	0.0	65.1	-42.3	53.5	68.3	128	0.483	1.0	0.0
129	117	122	0.466	1.0	0.0	71.7	-32.9	63.9	71.9	117	0.383	1.0	0.0	69.2	-36.5	58.6	69.1	122	0.467	1.0	0.0	0.303	1.0	0.0	64.3	-43.3	52.5	68.2	129	0.467	1.0	0.0
130	118	123	0.45	1.0	0.0	71.2	-33.7	62.9	71.4	118	0.369	1.0	0.0	68.5	-37.4	57.7	68.8	123	0.45	1.0	0.0	0.292	1.0	0.0	63.6	-44.3	51.5	68.1	130	0.45	1.0	0.0
131	119	124	0.433	1.0	0.0	70.7	-34.5	61.8	70.8	119	0.359	1.0	0.0	67.9	-38.3	56.9	68.7	124	0.433	1.0	0.0	0.28	1.0	0.0	62.8	-45.3	50.6	67.9	131	0.433	1.0	0.0
133	120	125	0.416	1.0	0.0	70.2	-35.2	60.8	70.2	120	0.349	1.0	0.0	67.3	-39.2	56.2	68.6	125	0.417	1.0	0.0	0.269	1.0	0.0	62.1	-46.2	49.5	67.8	133	0.417	1.0	0.0
134	121	126	0.4	1.0	0.0	69.6	-35.9	59.7	69.6	121	0.339	1.0	0.0	66.6	-40.2	55.4	68.5	126	0.4	1.0	0.0	0.257	1.0	0.0	61.3	-47.2	48.5	67.7	134	0.4	1.0	0.0
135	121	127	0.383	1.0	0.0	69.1	-36.5	58.6	69.1	121	0.329	1.0	0.0	66.0	-41.1	54.6	68.4	127	0.383	1.0	0.0	0.244	1.0	0.0	60.7	-48.1	47.5	67.6	135	0.383	1.0	0.0
136	122	128	0.366	1.0	0.0	68.3	-37.7	57.4	68.7	123	0.319	1.0	0.0	65.3	-42.0	53.8	68.3	128	0.367	1.0	0.0	0.229	1.0	0.0	60.3	-49.0	46.5	67.6	136	0.367	1.0	0.0
137	123	129	0.35	1.0	0.0	67.3	-39.2	56.2	68.6	124	0.309	1.0	0.0	64.7	-42.8	53.0	68.2	129	0.35	1.0	0.0	0.214	1.0	0.0	59.9	-49.9	45.4	67.6	137	0.35	1.0	0.0
138	126	130	0.333	1.0	0.0	66.2	-40.8	54.9	68.4	126	0.299	1.0	0.0	64.1	-43.7	52.2	68.1	130	0.333	1.0	0.0	0.199	1.0	0.0	59.5	-50.8	44.4	67.5	138	0.333	1.0	0.0
139	128	131	0.316	1.0	0.0	65.1	-42.3	53.6	68.2	128	0.289	1.0	0.0	63.4	-44.5	51.3	68.0	131	0.317	1.0	0.0	0.184	1.0	0.0	59.1	-51.7	43.3	67.5	140	0.317	1.0	0.0
140	129	132	0.3	1.0	0.0	64.0	-43.7	52.2	68.1	129	0.28	1.0	0.0	62.8	-45.4	50.5	67.9	132	0.3	1.0	0											

1-0131130-1.0	RE350-71	LAB* _{a0}	YN=0%	XYZ _{nw} =2.4, 2.5, 2.6	85.1, 88.8, 104.3	LAB* _{nw} =17.7, 0.0, 0.0	95.5, 0.0, 0.0
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input: *rgb/cmyk* - $>$ *rgbe*
output: transfer to *cmyke*
48 step hue circles: *rab-LabCh**tables

Output: Offset standard print; separation cmyn6*, D65, page 12/33

Output: Offset standard print; separation cmyn6*, D65, page 12/33

TUB-test chart RE35; hue code: H*e=B50Re										input: rgb/cmyk -> rgbe														
48 step hue circles; rgb-LabCh*tables										output: transfer to cmyke														
h _{ab,s}	h _{ab,d}	h _{ab,e}	rgb*d361M	LAB*d361M (x=LabCh)	C _d	rgb*ds361M	LAB*ds361M (x=LabCh)	rgb*d361M	LAB*ds361M (x=LabCh)	rgb*de361M	LAB*de361M (x=LabCh)	rgb*dd361M	LAB*dd361M	rgb*dd361M	LAB*dd361M									
216	236	216	0.0	1.0	58.3	-29.2	-43.7	52.6	236	C _d	0.0	1.0	0.666	56.1	-43.2	-24.9	50.0	210	C _s	0.0	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	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	1.0	1.0

Output: Offset standard print; separation cmykn6*, D65

input: rgb/cmyk -> rgbe

output: transfer to cmyke

1-0131330-L0 RE350-71 LAB* h_{ab} , YN=0%, XY Z $_{nw}$ =2.4, 2.5, 2.6, 85.1, 88.8, 104.3, LAB* nw =17.7, 0.0, 95.5, 0.0, 0.0

TUB-test chart RE35; hue code: H* $_{e}$ =B50R $_{e}$
48 step hue circles; $rgb-LabCh$ *tables

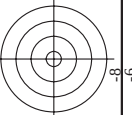
input: $rgb/cmyk -> rgb_e$
output: transfer to $cmyk_e$

Output: Offset standard print; separation cmyk6*, D65, page 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 RYGBM; $h_{ab,d} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;										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; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$																																																																																																																																																																																																																																																																																																																				
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_{ds361M}	$LAB^*_{ds361MI} (x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI} (x=LabCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI} (x=LabCh)$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rgb^*_{ds361MI}$	$rg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nj	HC*Fe	rgb_Fe	icr_Fe	hsa_Fe	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	0.125	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
2/666	R25Y_100_100e	0.25	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
3/675	R38Y_100_100e	0.375	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
4/684	R50Y_100_100e	0.5	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
5/693	R63Y_100_100e	0.625	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
6/702	R75Y_100_100e	0.75	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
7/711	R88Y_100_100e	0.875	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
8/720	Y00C_100_100e	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	0.0	0.0
9/639	Y13C_100_100e	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	0.0	0.0
10/558	Y25C_100_100e	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	0.0	0.0
11/477	Y38C_100_100e	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	0.0	0.0
12/396	Y50C_100_100e	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	0.0	0.0
13/315	Y63C_100_100e	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	0.0	0.0
14/234	Y75C_100_100e	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	0.0	0.0
15/153	Y88C_100_100e	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	0.0	0.0
16/72	G00C_100_100e	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	0.0	0.0
17/73	G13C_100_100e	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	0.0	0.0
18/74	G25C_100_100e	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	0.0	0.0
19/75	G38C_100_100e	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	0.0	0.0
20/76	G50C_100_100e	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	0.0	0.0
21/77	G63C_100_100e	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	0.0	0.0
22/78	G75C_100_100e	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	0.0	0.0
23/79	G88C_100_100e	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	0.0	0.0
24/80	C00B_100_100e	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	0.0
25/71	C13B_100_100e	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	0.0
26/62	C25B_100_100e	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	0.0
27/63	C38B_100_100e	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	0.0
28/44	C50B_100_100e	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	0.0
29/35	C63B_100_100e	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	0.0
30/26	C75B_100_100e	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	0.0
31/17	C88B_100_100e	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	0.0
32/8	B00M_100_100e	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	0.0
33/89	B13M_100_100e	0.125	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	0.0
34/170	B25M_100_100e	0.25	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	0.0
35/251	B38M_100_100e	0.375	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	0.0
36/332	B50M_100_100e	0.5	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	0.0
37/413	B63M_100_100e	0.625	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	0.0
38/494	B75M_100_100e	0.75	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	0.0
39/575	B88M_100_100e	0.875	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	0.0
40/656	M00R_100_100e	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	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	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

Mean color difference of this page: $\Delta E^* = 17.3$

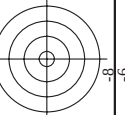


TUB material: code=rha4ta
nyn6 (CMYK)

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

[illegible]

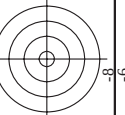
Mean color difference of this page: $\Delta E^* = 12.3$



TUB material: code=rha4ta
nyn6 (CMYK)

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

[illegible]

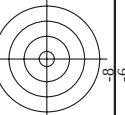


TUB material: code=rha4ta
nyn6 (CMYK)

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

colors and differences, ΔE^* ,

[illegible]



TUB material: code=rha4ta
nyn6 (CMYK)

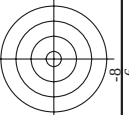
input: *rgb/cmyk* → *rgb*_e
output: transfer to *cmyk*_e

[illegible]

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

TUB-test chart RE35; hue code: $H_e^*=B50R_e$
 colors and differences ΔE^* ,

see similar files: <http://130.149.60.45/~farbmatrik/RE35/RE35.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>



TUB material: code=rha4ta
nyn6 (CMYK)

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



RE350-7N, Page 29/33-F

n	HC ^{Fe}	rgb^{Fe}	kr^{Fe}	hsa^{Fe}	$LabCh^{Fe}$	rgb^{Fe}	$LabCh^{Fe}$	DE^{Fe}	hsa^{Fe}	rgb^{Fe}	$LabCh^{Fe}$	$delta F^{Fe} = 2.3$
729	NW_100 ₀	1.0	1.0	1.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0
730	G50B_100_012 ₀	0.875	1.0	1.0	0.125	0.937	210	0.0	0.0	0.0	0.0	0.0
731	G50B_100_025 ₀	0.75	1.0	1.0	0.25	0.875	210	0.0	0.0	0.0	0.0	0.0
732	G50B_100_037 ₀	0.625	1.0	1.0	0.375	0.812	210	0.0	0.0	0.0	0.0	0.0
733	G50B_100_050 ₀	0.5	1.0	1.0	0.5	0.75	210	0.0	0.0	0.0	0.0	0.0
734	G50B_100_062 ₀	0.375	1.0	1.0	0.625	0.687	210	0.0	0.0	0.0	0.0	0.0
735	G50B_100_075 ₀	0.25	1.0	1.0	0.75	0.625	210	0.0	0.0	0.0	0.0	0.0
736	G50B_100_087 ₀	0.125	1.0	1.0	0.875	0.562	210	0.0	0.0	0.0	0.0	0.0
737	G50B_100_100 ₀	0.0	1.0	1.0	1.0	0.5	210	0.0	0.0	0.0	0.0	0.0
738	ROXY_100_012 ₀	1.0	0.875	0.875	1.0	0.125	937	330	0.0	0.0	0.0	0.0
739	ROXY_100_025 ₀	0.875	0.875	0.875	1.0	0.25	937	330	0.0	0.0	0.0	0.0
740	G50B_087_012 ₀	0.625	0.875	0.875	1.0	0.375	937	330	0.0	0.0	0.0	0.0
741	G50B_087_025 ₀	0.5	0.875	0.875	1.0	0.5	937	330	0.0	0.0	0.0	0.0
742	G50B_087_037 ₀	0.375	0.875	0.875	1.0	0.625	937	330	0.0	0.0	0.0	0.0
743	G50B_087_050 ₀	0.25	0.875	0.875	1.0	0.75	937	330	0.0	0.0	0.0	0.0
744	G50B_087_062 ₀	0.125	0.875	0.875	1.0	0.875	937	330	0.0	0.0	0.0	0.0
745	G50B_087_075 ₀	0.0	0.875	0.875	1.0	1.0	937	330	0.0	0.0	0.0	0.0
746	G50B_087_087 ₀	1.0	0.875	0.875	1.0	0.875	937	330	0.0	0.0	0.0	0.0
747	ROXY_100_025 ₀	0.875	0.75	0.75	1.0	0.25	9812	390	0.0	0.0	0.0	0.0
748	ROXY_100_037 ₀	0.75	0.75	0.75	1.0	0.375	9812	390	0.0	0.0	0.0	0.0
749	ROXY_100_050 ₀	0.625	0.75	0.75	1.0	0.5	9812	390	0.0	0.0	0.0	0.0
750	G50B_075_012 ₀	0.5	0.75	0.75	1.0	0.625	9812	390	0.0	0.0	0.0	0.0
751	G50B_075_025 ₀	0.375	0.75	0.75	1.0	0.75	9812	390	0.0	0.0	0.0	0.0
752	G50B_075_037 ₀	0.25	0.75	0.75	1.0	0.875	9812	390	0.0	0.0	0.0	0.0
753	G50B_075_050 ₀	0.125	0.75	0.75	1.0	1.0	9812	390	0.0	0.0	0.0	0.0
754	G50B_075_062 ₀	0.0	0.75	0.75	1.0	0.875	9812	390	0.0	0.0	0.0	0.0
755	ROXY_100_037 ₀	0.875	0.75	0.75	1.0	0.375	9812	390	0.0	0.0	0.0	0.0
756	ROXY_100_050 ₀	0.75	0.75	0.75	1.0	0.5	9812	390	0.0	0.0	0.0	0.0
757	ROXY_100_075 ₀	0.625	0.75	0.75	1.0	0.625	9812	390	0.0	0.0	0.0	0.0
758	NW_050 ₀	0.625	0.625	0.625	0.625	0.625	360	0.0	0.0	0.0	0.0	0.0
759	G50B_062_012 ₀	0.5	0.625	0.625	0.625	0.5	360	0.0	0.0	0.0	0.0	0.0

n	HIC*Fe	rgb_Fe	icr_Fe	hsa_Fe	rgb_Fe	LabCh*Fe	LabCh*Fe	rgb_Fe	LabCh*Fe	DF*Fe	hsa_Fe	rgb_Fe	LabCh*Fe
810	NW_100a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
811	BOOR_100.012a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
812	BOOR_100.025a	0.75 0.75 1.0	1.0 1.0 1.0	270	0.75 0.843 1.0	81.0 0.3 1.0	82.2 0.1 1.0	0.0 0.0 0.0	297.6 3.1	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
813	BOOR_100.037a	0.625 0.625 1.0	1.0 1.0 1.0	270	0.625 0.765 1.0	73.8 0.5 1.0	78.1 0.6 1.0	0.0 0.0 0.0	303.7 7.9	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
814	BOOR_100.050a	0.5 0.5 1.0	1.0 1.0 1.0	270	0.5 0.687 1.0	66.7 0.6 1.0	69.3 0.5 1.0	0.0 0.0 0.0	302.6 11.4	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
815	BOOR_100.062a	0.375 0.375 1.0	1.0 1.0 1.0	270	0.375 0.609 1.0	59.5 0.8 1.0	62.8 0.6 1.0	0.0 0.0 0.0	304.0 22.4	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
816	BOOR_100.075a	0.25 0.25 1.0	1.0 1.0 1.0	270	0.25 0.531 1.0	52.3 1.0 1.0	55.8 0.8 1.0	0.0 0.0 0.0	302.8 25.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
817	BOOR_100.087a	0.125 0.125 1.0	1.0 1.0 1.0	270	0.125 0.452 1.0	45.1 1.2 1.0	48.1 0.8 1.0	0.0 0.0 0.0	303.1 29.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
818	BOOR_100.100a	0.0 0.0 1.0	1.0 1.0 1.0	270	0.0 0.374 1.0	37.9 1.3 1.0	40.4 0.8 1.0	0.0 0.0 0.0	302.8 32.6	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
819	YOCG_100.012a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
820	NW_087a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
821	BOOR_087.012a	0.75 0.75 1.0	1.0 1.0 1.0	270	0.75 0.843 1.0	81.0 0.3 1.0	82.2 0.1 1.0	0.0 0.0 0.0	297.6 3.1	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
822	BOOR_087.025a	0.625 0.625 1.0	1.0 1.0 1.0	270	0.625 0.765 1.0	73.8 0.5 1.0	78.1 0.6 1.0	0.0 0.0 0.0	303.7 7.9	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
823	BOOR_087.037a	0.5 0.5 1.0	1.0 1.0 1.0	270	0.5 0.687 1.0	66.7 0.6 1.0	69.3 0.5 1.0	0.0 0.0 0.0	302.6 11.4	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
824	BOOR_087.050a	0.375 0.375 1.0	1.0 1.0 1.0	270	0.375 0.609 1.0	59.5 0.8 1.0	62.8 0.6 1.0	0.0 0.0 0.0	304.0 22.4	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
825	BOOR_087.062a	0.25 0.25 1.0	1.0 1.0 1.0	270	0.25 0.531 1.0	52.3 1.0 1.0	55.8 0.8 1.0	0.0 0.0 0.0	302.8 25.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
826	BOOR_087.075a	0.125 0.125 1.0	1.0 1.0 1.0	270	0.125 0.452 1.0	45.1 1.2 1.0	48.1 0.8 1.0	0.0 0.0 0.0	303.1 29.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
827	BOOR_087.087a	0.0 0.0 1.0	1.0 1.0 1.0	270	0.0 0.374 1.0	37.9 1.3 1.0	40.4 0.8 1.0	0.0 0.0 0.0	302.8 32.6	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
828	YOCG_100.012a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
829	YOCG_100.025a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
830	NW_075a	0.75 0.75 1.0	1.0 1.0 1.0	360	0.75 0.843 1.0	81.0 0.3 1.0	82.2 0.1 1.0	0.0 0.0 0.0	297.6 3.1	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
831	BOOR_075.012a	0.625 0.625 1.0	1.0 1.0 1.0	270	0.625 0.765 1.0	73.8 0.5 1.0	78.1 0.6 1.0	0.0 0.0 0.0	303.7 7.9	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
832	BOOR_075.025a	0.5 0.5 1.0	1.0 1.0 1.0	270	0.5 0.687 1.0	66.7 0.6 1.0	69.3 0.5 1.0	0.0 0.0 0.0	302.6 11.4	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
833	BOOR_075.037a	0.375 0.375 1.0	1.0 1.0 1.0	270	0.375 0.609 1.0	59.5 0.8 1.0	62.8 0.6 1.0	0.0 0.0 0.0	304.0 22.4	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
834	BOOR_075.050a	0.25 0.25 1.0	1.0 1.0 1.0	270	0.25 0.531 1.0	52.3 1.0 1.0	55.8 0.8 1.0	0.0 0.0 0.0	302.8 25.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
835	BOOR_075.062a	0.125 0.125 1.0	1.0 1.0 1.0	270	0.125 0.452 1.0	45.1 1.2 1.0	48.1 0.8 1.0	0.0 0.0 0.0	303.1 29.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
836	BOOR_075.075a	0.0 0.0 1.0	1.0 1.0 1.0	270	0.0 0.374 1.0	37.9 1.3 1.0	40.4 0.8 1.0	0.0 0.0 0.0	302.8 32.6	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
837	YOCG_075.012a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
838	YOCG_075.025a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
839	YOCG_075.037a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
840	NW_050a	0.375 0.375 1.0	1.0 1.0 1.0	270	0.375 0.609 1.0	59.5 0.8 1.0	62.8 0.6 1.0	0.0 0.0 0.0	302.8 25.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
841	BOOR_050.012a	0.375 0.375 1.0	1.0 1.0 1.0	270	0.375 0.609 1.0	59.5 0.8 1.0	62.8 0.6 1.0	0.0 0.0 0.0	302.8 25.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
842	BOOR_050.025a	0.25 0.25 1.0	1.0 1.0 1.0	270	0.25 0.531 1.0	52.3 1.0 1.0	55.8 0.8 1.0	0.0 0.0 0.0	303.1 29.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
843	BOOR_050.037a	0.125 0.125 1.0	1.0 1.0 1.0	270	0.125 0.452 1.0	45.1 1.2 1.0	48.1 0.8 1.0	0.0 0.0 0.0	303.1 29.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
844	BOOR_050.050a	0.0 0.0 1.0	1.0 1.0 1.0	270	0.0 0.374 1.0	37.9 1.3 1.0	40.4 0.8 1.0	0.0 0.0 0.0	302.8 32.6	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
845	YOCG_100.050a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
846	YOCG_075.037a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0 0.0 0.0	95.4 1.0 1.0	95.4 1.0 1.0
847	YOCG_075.050a	0.75 0.75 1.0	1.0 1.0 1.0	360	0.75 0.843 1.0	81.0 0.3 1.0	82.2 0.1 1.0	0.0 0.0 0.0	297.6 3.1	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
848	YOCG_075.062a	0.625 0.625 1.0	1.0 1.0 1.0	270	0.625 0.765 1.0	73.8 0.5 1.0	78.1 0.6 1.0	0.0 0.0 0.0	303.7 7.9	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
849	NW_050a	0.375 0.375 1.0	1.0 1.0 1.0	270	0.375 0.609 1.0	59.5 0.8 1.0	62.8 0.6 1.0	0.0 0.0 0.0	302.8 25.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
850	BOOR_050.012a	0.375 0.375 1.0	1.0 1.0 1.0	270	0.375 0.609 1.0	59.5 0.8 1.0	62.8 0.6 1.0	0.0 0.0 0.0	302.8 25.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
851	BOOR_050.025a	0.25 0.25 1.0	1.0 1.0 1.0	270	0.25 0.531 1.0	52.3 1.0 1.0	55.8 0.8 1.0	0.0 0.0 0.0	303.1 29.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
852	BOOR_050.037a	0.125 0.125 1.0	1.0 1.0 1.0	270	0.125 0.452 1.0	45.1 1.2 1.0	48.1 0.8 1.0	0.0 0.0 0.0	303.1 29.2	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
853	BOOR_050.050a	0.0 0.0 1.0	1.0 1.0 1.0	270	0.0 0.374 1.0	37.9 1.3 1.0	40.4 0.8 1.0	0.0 0.0 0.0	302.8 32.6	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
854	BOOR_050.062a	0.0 0.0 1.0	1.0 1.0 1.0	270	0.0 0.374 1.0	37.9 1.3 1.0	40.4 0.8 1.0	0.0 0.0 0.0	302.8 32.6	248 0.0 0.0	0.0 0.0 0.0	379 1.3 1.0	379 1.3 1.0
855	YOCG_100.062a	0.875 0.875 1.0	1.0 1.0 1.0	360	0.875 0.921 1.0	95.4 1.0 1.0	95.5 1.0 1.0	0.0 0.0 0.0	103.6 6.7	360 1.0 1.0	0.0		

http://130.149.60.45/~farbmatrik/RE35/RE35L0NP.PDF /.PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 31/33

TUB-test chart RE35; hue code: $H_e^*=B50R_e$
 colors and differences ΔE^* ,

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

n	HIC ^{Fe}	rgb_{Fe}	lrc_{Fe}	$h_{\alpha_{Fe}}$	rgb^{Fe}	$LabCh^{Fe}$	$LabCh^{Fe}$	rgb^{Me}	DE^{Fe}	h_{α}^{Me}	rgb^{Me}	$LabCh^{Me}$	n
891	NW_100k	1.0	1.0	1.0	0.0	95.4	1.0	0.0	0.0	1.0	1.0	95.4	891
892	B50R_100_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	892
893	B50R_100_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	893
894	B50R_100_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	894
895	B50R_100_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	895
896	B50R_100_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	896
897	B50R_100_075e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	897
898	B50R_100_087e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	898
899	B50R_100_100e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	899
900	NW_067e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	900
901	B50R_087_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	901
902	B50R_087_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	902
903	B50R_087_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	903
904	B50R_087_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	904
905	B50R_087_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	905
906	B50R_087_075e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	906
907	B50R_087_087e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	907
908	B50R_087_100e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	908
909	NW_075e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	909
910	B50R_075_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	910
911	B50R_075_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	911
912	B50R_075_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	912
913	B50R_075_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	913
914	B50R_075_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	914
915	B50R_075_075e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	915
916	B50R_075_087e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	916
917	B50R_075_100e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	917
918	NW_087e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	918
919	B50R_087_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	919
920	B50R_087_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	920
921	NW_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	921
922	B50R_062_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	922
923	B50R_062_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	923
924	B50R_062_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	924
925	B50R_062_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	925
926	B50R_062_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	926
927	NW_050_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	927
928	NW_050_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	928
929	NW_050_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	929
930	NW_050_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	930
931	NW_050_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	931
932	B50R_050_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	932
933	B50R_050_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	933
934	B50R_050_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	934
935	B50R_050_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	935
936	B50R_050_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	936
937	NW_037_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	937
938	NW_037_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	938
939	NW_037_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	939
940	NW_037_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	940
941	NW_037_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	941
942	B50R_037_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	942
943	B50R_037_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	943
944	B50R_037_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	944
945	B50R_037_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	945
946	B50R_037_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	946
947	NW_025_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	947
948	NW_025_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	948
949	NW_025_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	949
950	NW_025_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	950
951	NW_025_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	951
952	B50R_025_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	952
953	B50R_025_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	953
954	B50R_025_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	954
955	B50R_025_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	955
956	B50R_025_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	956
957	NW_012_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	957
958	NW_012_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	958
959	NW_012_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	959
960	NW_012_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	960
961	NW_012_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	961
962	B50R_012_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	962
963	B50R_012_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	963
964	B50R_012_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	964
965	B50R_012_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	965
966	B50R_012_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	966
967	NW_007_012e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	967
968	NW_007_025e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	968
969	NW_007_037e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	969
970	NW_007_050e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	970
971	NW_007_062e	1.0	0.875	1.0	0.0	87.9	6.1	-3.7	7.2	328.6	0.0	0.0	971

see similar files: <http://130.149.60.45/~farbmatrik/RE35/RE35.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

n	HC*Fe	rgb*Fe	icT*Fe	hsa*Fe	rgb*Fe	LabCh*Fe	LabCh*Fe	DF*Fe	haM,e	rgb*Me	LabCh*Me	0.0	0.0	0.0
1053	NW_086e	0.866 0.866 0.866	0.866 0.866 0.866	360 360 360	0.866 0.866 0.866	85.0 0.0 0.0	89.4 -0.1 0.0	204.5 4.4 0.1	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1054	NW_093e	0.933 0.933 0.933	0.933 0.933 0.933	360 360 360	0.933 0.933 0.933	90.2 0.0 0.0	92.2 0.0 0.0	177.8 1.9 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1055	NW_100e	1.0 1.0 1.0	1.0 1.0 1.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	95.4 0.0 0.0	61.5 0.0 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1056	NW_000e	0.0 0.0 0.0	0.0 0.0 0.0	360 360 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	96.3 0.0 0.1	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1057	NW_006e	0.066 0.066 0.066	0.066 0.066 0.066	360 360 360	0.066 0.066 0.066	22.8 0.0 0.0	22.3 -0.1 0.0	151.6 0.5 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1058	NW_013e	0.133 0.133 0.133	0.133 0.133 0.133	360 360 360	0.133 0.133 0.133	30.4 0.0 0.0	-0.2 -0.5 0.6	242.3 2.4 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1059	NW_020e	0.2 0.2 0.2	0.2 0.2 0.2	360 360 360	0.2 0.2 0.2	33.2 0.0 0.0	-0.4 -0.8 0.9	243.3 5.7 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1060	NW_026e	0.266 0.266 0.266	0.266 0.266 0.266	360 360 360	0.266 0.266 0.266	38.3 0.0 0.0	-0.4 -0.7 0.8	240.2 7.2 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1061	NW_033e	0.333 0.333 0.333	0.333 0.333 0.333	360 360 360	0.333 0.333 0.333	43.6 0.0 0.0	-0.4 -0.6 0.8	235.4 8.6 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1062	NW_040e	0.4 0.4 0.4	0.4 0.4 0.4	360 360 360	0.4 0.4 0.4	48.8 0.0 0.0	-0.4 -0.6 0.7	234.3 8.6 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1063	NW_046e	0.466 0.466 0.466	0.466 0.466 0.466	360 360 360	0.466 0.466 0.466	53.9 0.0 0.0	-0.5 0.6 0.7	235.2 7.8 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1064	NW_053e	0.533 0.533 0.533	0.533 0.533 0.533	360 360 360	0.533 0.533 0.533	59.1 0.0 0.0	-0.5 0.6 0.7	231.6 7.7 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1065	NW_060e	0.6 0.6 0.6	0.6 0.6 0.6	360 360 360	0.6 0.6 0.6	64.3 0.0 0.0	-0.4 0.5 0.6	233.5 7.3 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1066	NW_066e	0.666 0.666 0.666	0.666 0.666 0.666	360 360 360	0.666 0.666 0.666	69.5 0.0 0.0	-0.2 0.3 0.2	225.3 6.1 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1067	NW_073e	0.734 0.734 0.734	0.734 0.734 0.734	360 360 360	0.734 0.734 0.734	74.7 0.0 0.0	-0.2 0.3 0.2	221.2 4.9 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1068	NW_080e	0.8 0.8 0.8	0.8 0.8 0.8	360 360 360	0.8 0.8 0.8	79.9 0.0 0.0	-0.1 0.1 0.1	220.3 4.3 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1069	NW_086e	0.866 0.866 0.866	0.866 0.866 0.866	360 360 360	0.866 0.866 0.866	85.0 0.0 0.0	0.0 0.0 0.0	125.8 2.0 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1070	NW_093e	0.933 0.933 0.933	0.933 0.933 0.933	360 360 360	0.933 0.933 0.933	90.2 0.0 0.0	0.0 0.0 0.0	92.4 0.0 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1071	NW_100e	1.0 1.0 1.0	1.0 1.0 1.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	78.4 2.3 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1072	NW_000e	0.0 0.0 0.0	0.0 0.0 0.0	360 360 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	275.2 0.1 0.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0	0.0	0.0
1073	RO0Y_100_100e	1.0 1.0 1.0	1.0 1.0 1.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	-0.1 0.1 0.1	53.6 10.5 0.0	378 378 378	1.0 0.0 0.0	0.0 0.0 0.0	0.0	0.0	0.0
1074	RO0Y_100_100e	1.0 1.0 1.0	1.0 1.0 1.0	360 360 360	1.0 1.0 1.0	95.4 0.0 0.0	-0.1 0.1 0.1	53.6 10.5 0.0	378 378 378	1.0 0.0 0.0	0.0 0.0 0.0	0.0	0.0	0.0
1075	Y00B_100_100e	0.0 1.0 0.0	0.0 1.0 0.0	360 360 360	0.0 1.0 0.0	56.6 -39.7 29.9	-28.4 -45.4 53.6	237.9 19.1 19.5	360 360 360	1.0 0.0 0.0	0.0 0.0 0.0	0.0	0.0	0.0
1076	Y00B_100_100e	0.0 1.0 0.0	0.0 1.0 0.0	360 360 360	0.0 1.0 0.0	56.6 -39.7 29.9	-28.4 -45.4 53.6	237.9 19.1 19.5	360 360 360	1.0 0.0 0.0	0.0 0.0 0.0	0.0	0.0	0.0
1077	B00R_100_100e	0.0 0.0 1.0	0.0 0.0 1.0	360 360 360	0.0 0.0 1.0	37.9 21.5 4.4	25.3 25.1 24.6	299.0 25.4 24.8	360 360 360	1.0 0.0 0.0	0.0 0.0 0.0	0.0	0.0	0.0
1078	B00R_100_100e	0.0 0.0 1.0	0.0 0.0 1.0	360 360 360	0.0 0.0 1.0	37.9 21.5 4.4	25.3 25.1 24.6	299.0 25.4 24.8	360 360 360	1.0 0.0 0.0	0.0 0.0 0.0	0.0	0.0	0.0
1079	B50R_100_100e	1.0 0.0 1.0	1.0 0.0 1.0	360 360 360	1.0 0.0 1.0	34.8 49.2 -36.0	-3.2 75.3 75.3	357.5 38.7 29.3	293 293 293	0.407 0.0 0.0	0.0 0.0 0.0	0.0	0.0	0.0

Mean color difference of this page: delta E** = 7.6

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

TUB-test chart RE35; hue code: H*e=B50Re
colors and differences, ΔE**