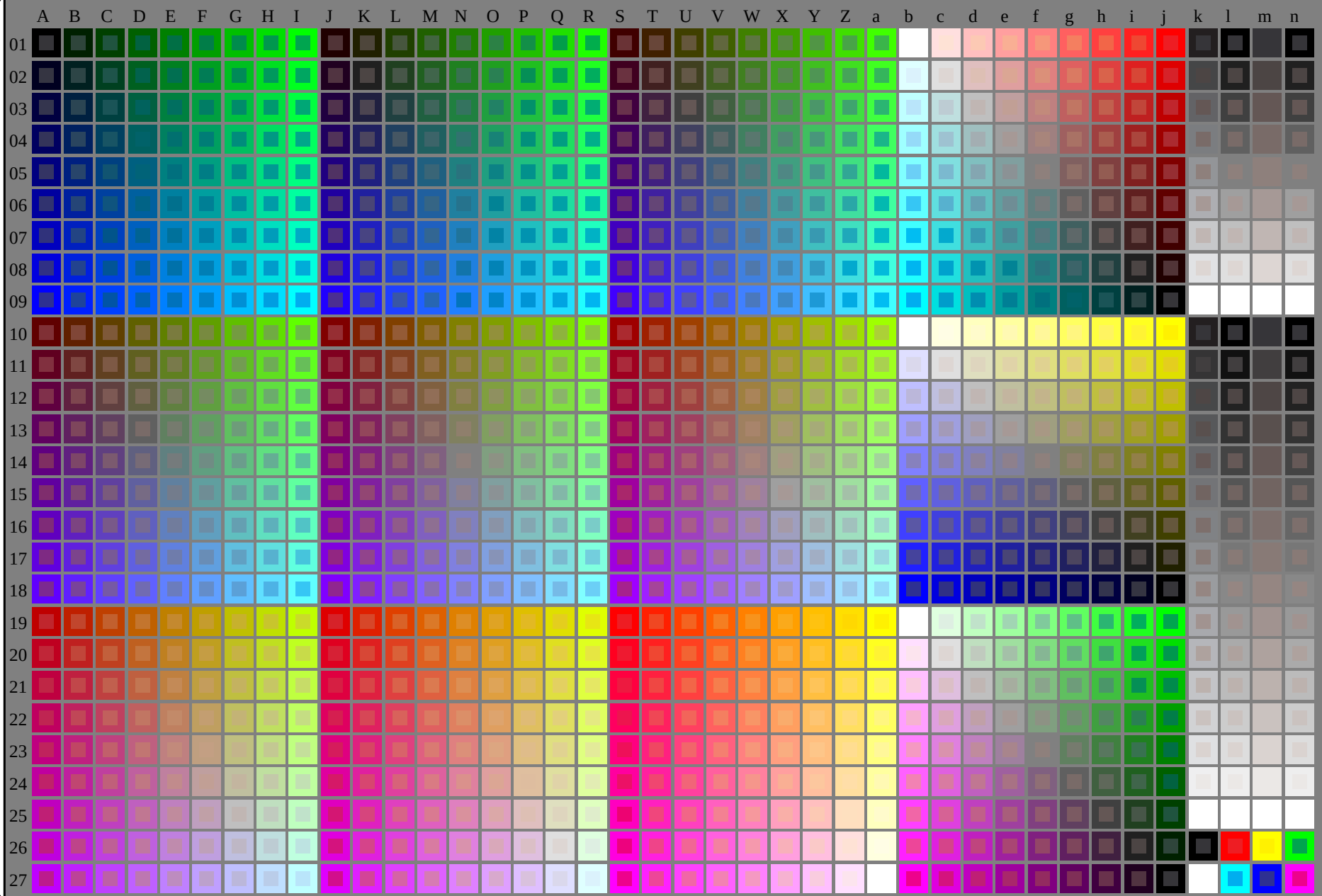


see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /.PS
application for measurement of laser printer output

TUB material: code=rha4ta



RE710-7N_RGB 1-003034-L0

TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872

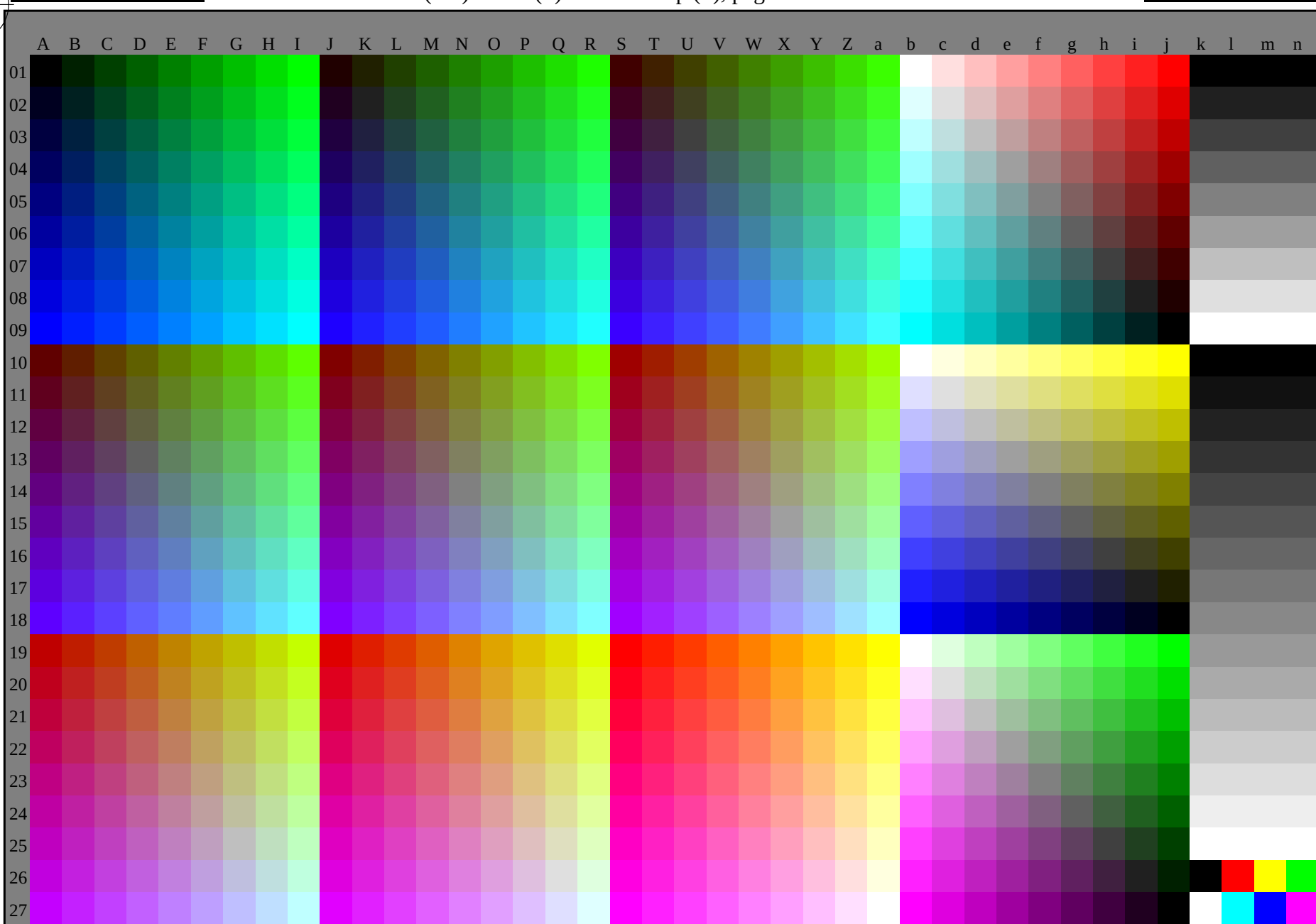
Test chart G with 40x27=1080 colours; equidistant 9 or 16 step colour scales; Colour data in column (A-n): **rgb** (A_j + k26_n27), 000n (k), w (l), nnn0 (m), www (n), 3D = 0

input: *rgb/cmyk* -> *rgb/cmyk*
output: no change

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /PS
application for measurement of laser printer output, no separation rgb (RGB)

TUB material: code=rha4ta



RE710-70 1-003134-L0

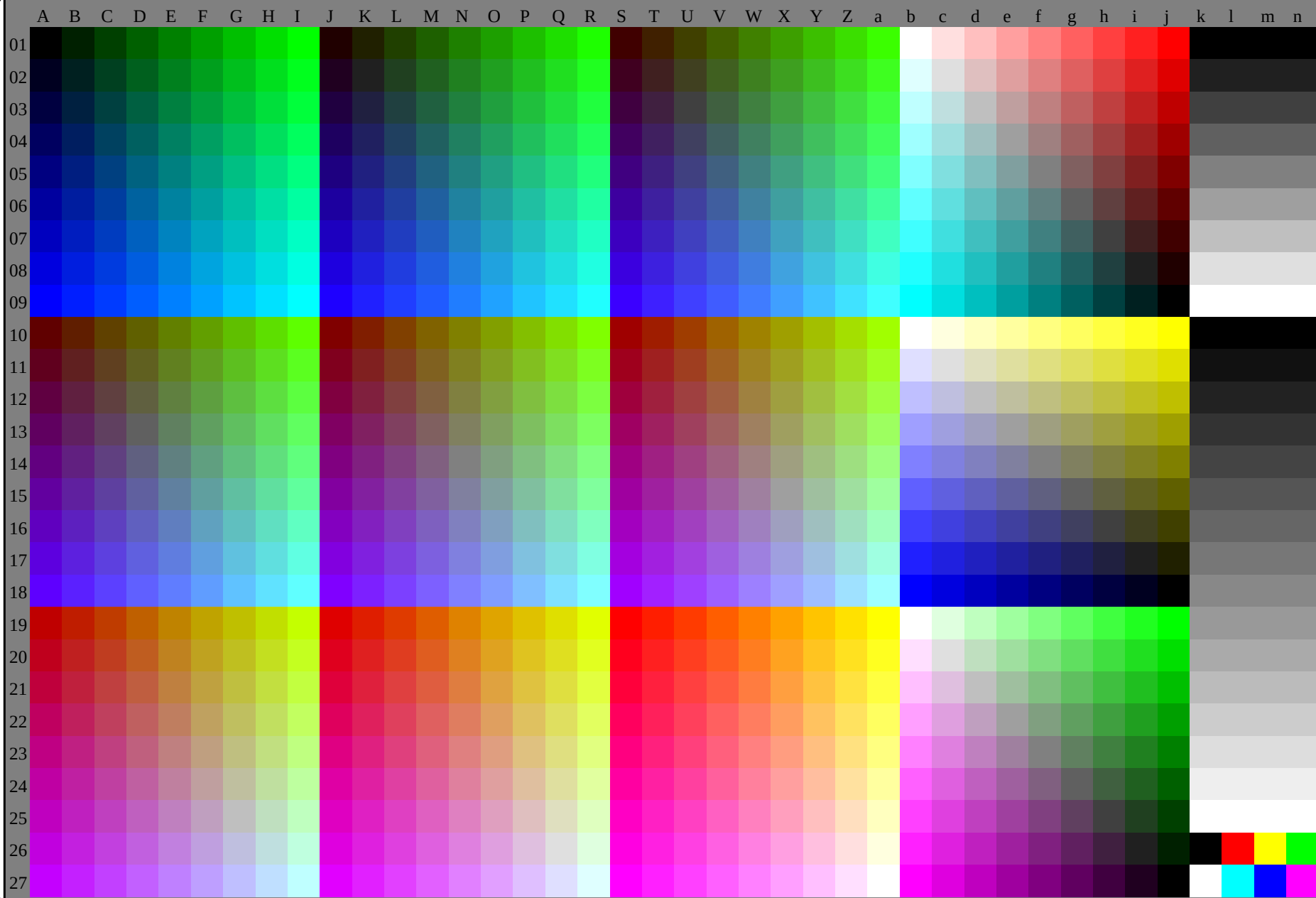
TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872, 3D=0, $de=0$, rgb

input: $rgb/cmyk \rightarrow rgb_d$
output: transfer to rgb_d

1-003134-F0

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /.PS
application for measurement of laser printer output, no separation rgb (RGB)
TUB material: code=rha4ta



RE710-70 1-003234-L0

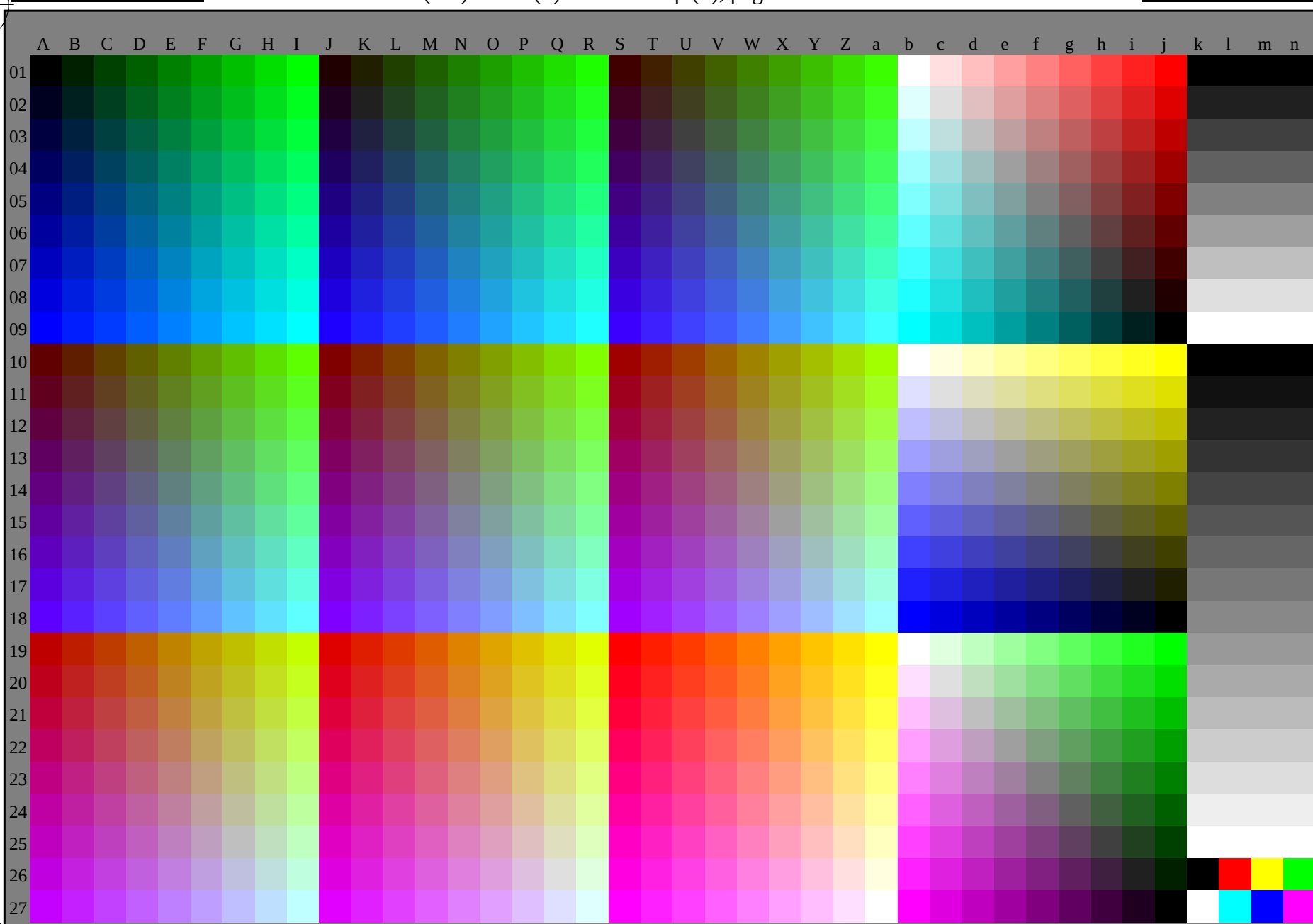
TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872

input: $rgb/cmyk \rightarrow rgb_d$
output: transfer to rgb_d

1-003234-F0

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /.PS
application for measurement of laser printer output, no separation rgb (RGB)
TUB material: code=rha4ta



RE710-70 1-003334-L0

TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872

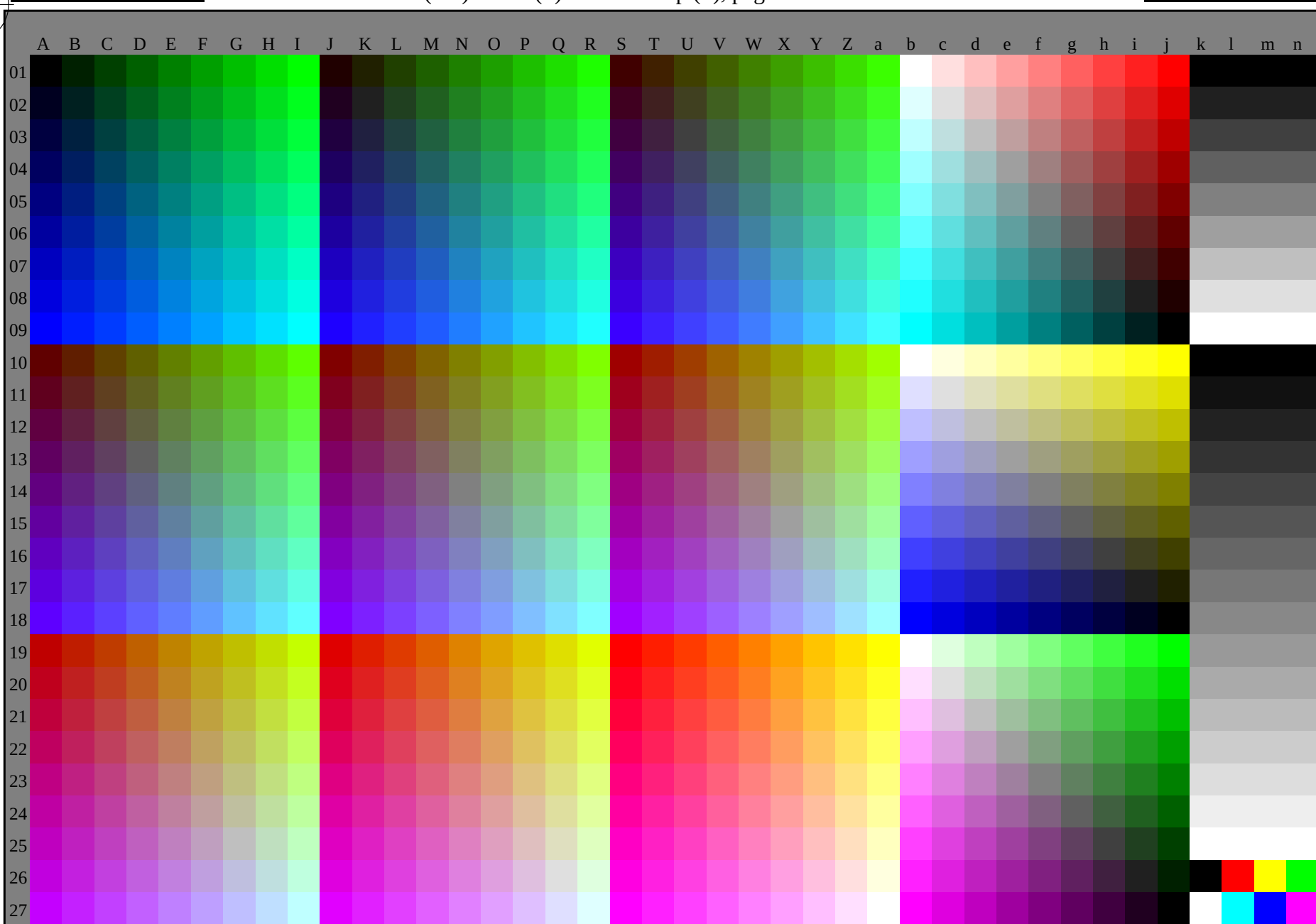
Test chart G with 40x27=1080 colours; equidistant 9 or 16 step colour scales; Colour data in column (A-n);, 3D = 0

input: $rgb/cmyk \rightarrow rgb_d$
output: transfer to rgb_d

1-003334-F0

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /.PS TUB material: code=rha4ta
application for measurement of laser printer output, no separation rgb (RGB)



RE710-70 1-003434-L0

TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872

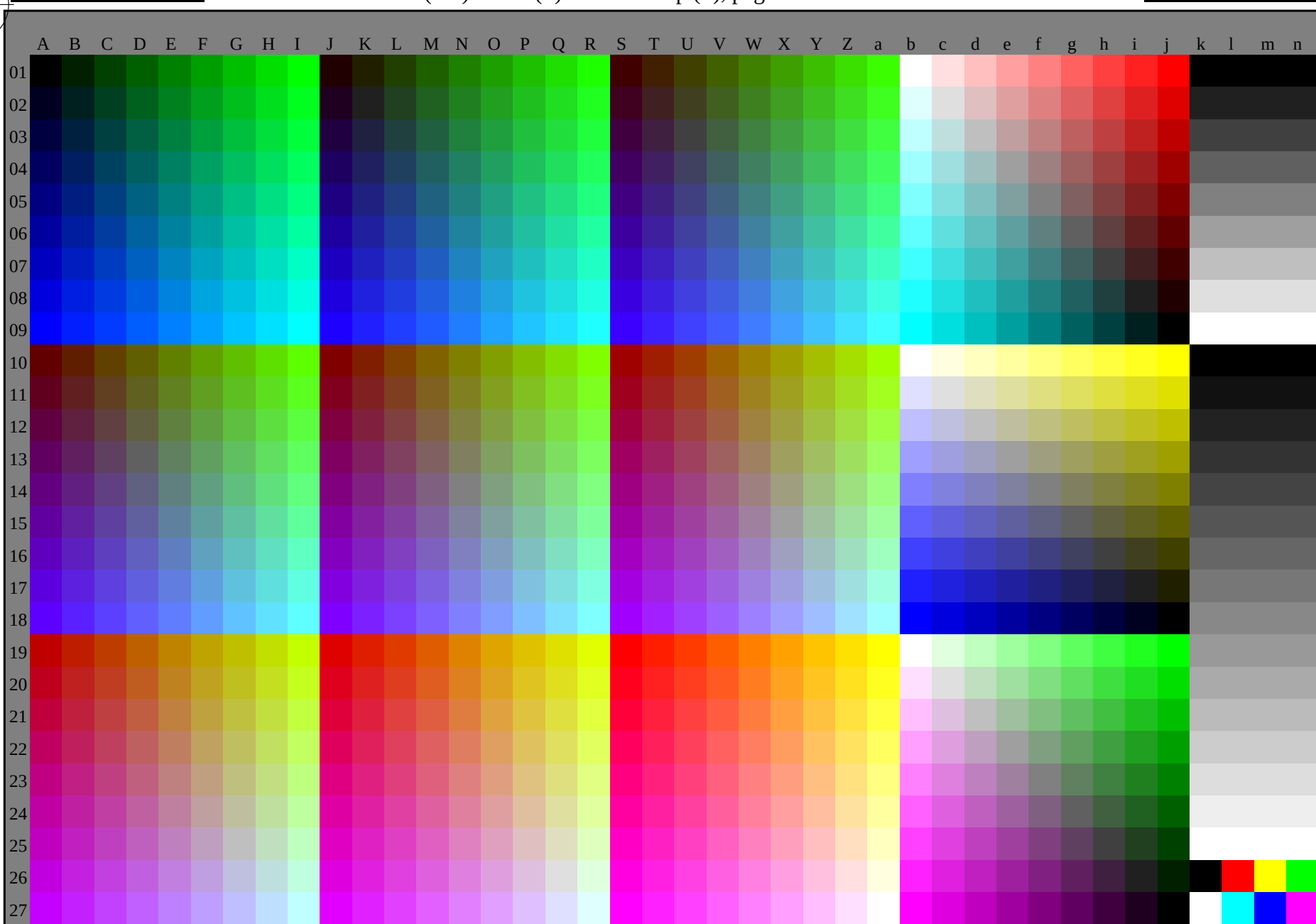
Test chart G with 40x27=1080 colours; equidistant 9 or 16 step colour scales; Colour data in column (A-n);, 3D = 0

input: $rgb/cmyk \rightarrow rgb_d$
output: transfer to rgb_d

1-003434-F0

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /.PS TUB material: code=rha4ta
application for measurement of laser printer output, no separation rgb (RGB)



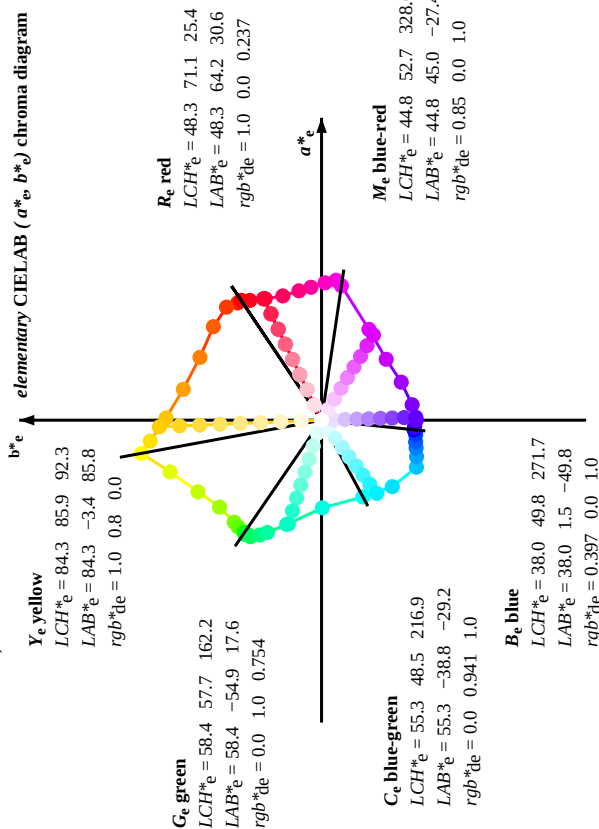
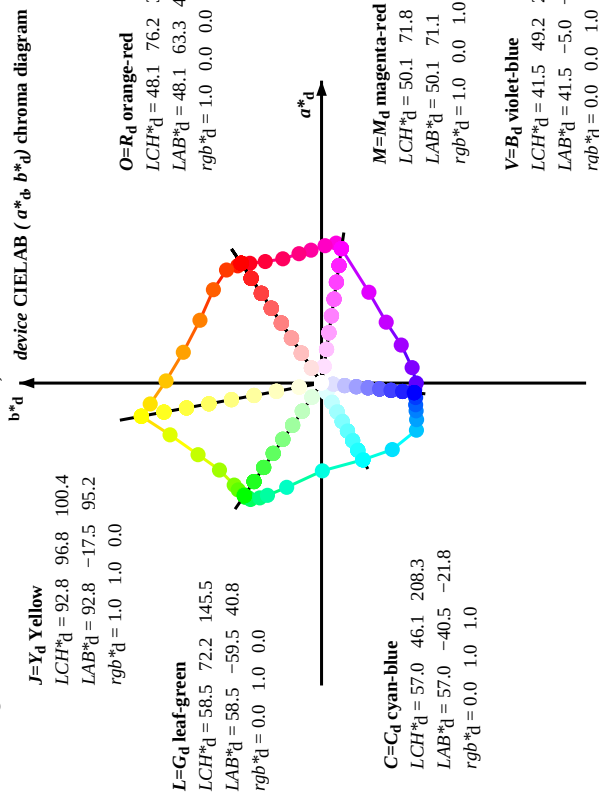
RE710-70 1-003534-L0

TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872

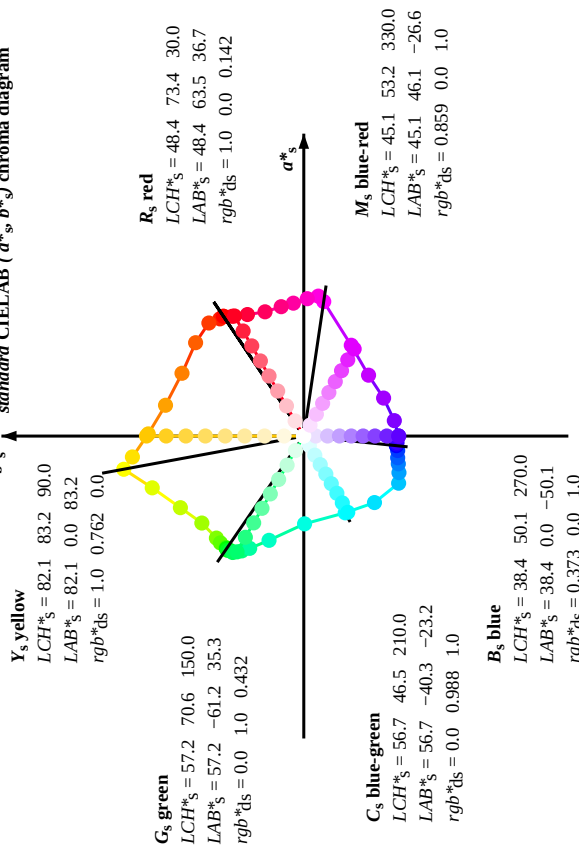
input: $rgb/cmyk \rightarrow rgb_d$
output: transfer to rgb_d

1-003534-F0

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 RYGCMB₆; $h_{ab,6} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours RYGCMB₆; $h_{ab,6} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; Six hue angles of the elementary colours RYGCMB₆; $h_{ab,6} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$



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:

$$h_{ab,s} = \arctan \left[\frac{r^*_d \cos(30) + g^*_d \sin(150)}{1 / r^*_d \sin(30) + g^*_d \sin(150)} + b^*_d \sin(270) \right] \quad (1)$$
- 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: $h_{ab,6} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0, 390.0$ ($i=0,6$) and the equations for a 48 and 360 step hue circle:

$$h_{48ab,6j} = h_{ab,6i} + j [h_{ab,6i+1} - h_{ab,6i}] / 8 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 7) \quad (2)$$

$$h_{360ab,6j} = h_{ab,6i} + j [h_{ab,6i+1} - h_{ab,6i}] / 60 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 59) \quad (3)$$
- 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 : $h_{ab,6} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6, 385.5$ ($i=0,6$) and the equations for a 48 and 360 step elementary hue circle:

$$h_{48ab,6j} = h_{ab,6i} + j [h_{ab,6i+1} - h_{ab,6i}] / 8 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 7) \quad (4)$$

$$h_{360ab,6j} = h_{ab,6i} + j [h_{ab,6i+1} - h_{ab,6i}] / 60 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 59) \quad (5)$$
- 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

RE710-70 1-003634-L0

LAB₆*a₀, YN=0%, XY Zmw=2.0, 2.1, 2.1, 85.9, 90.9, 95.1, LAB₆*mw=15.8, 0.0, 0.0, 96.4, 0.0, 0.0, not adapted=adapted

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

TUB-test chart RE71; 1080 standard colours, $cf=0.9$
48 step hue circles; $rgb-LabCh$ *tables

input: $rgb/cmyk \rightarrow rgb_d$
output: transfer to rgb_d

http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 8/33

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;										Data of Maximum color M in colorimetric system Offset standard print; separation cmyk6*; D65 for input or output;									
Six hue angles of the device colours RYGBCM; h _{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6;										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 _{gb} * d _{64M} r _{gb} * d _{36IM} LAB* d _{64M} (x=LabCh)										r _{gb} * d _{36IM} LAB* d _{36IM} (x=LabCh)									
33.8	30.0	25.4	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8	1.0	0.0	0.0	0.143	48.5	63.6	36.7	73.4	30
35.6	37.5	33.8	1.0	0.125	0.0	48.8	62.0	44.3	76.2	35.6	1.0	0.164	0.0	49.2	61.4	46.2	76.8	37	1.0
40.0	40.0	42.1	1.0	0.25	0.0	49.9	59.8	50.2	78.1	40.0	1.0	0.318	0.0	52.8	54.3	76.8	45	1.0	0.279
49.1	52.5	50.5	1.0	0.375	0.0	55.1	49.4	57.2	75.6	49.1	1.0	0.401	0.0	56.9	46.2	59.1	75.0	52	1.0
62.6	60.0	58.8	1.0	0.5	0.0	63.4	33.2	64.2	72.4	62.6	1.0	0.475	0.0	61.8	36.6	63.3	73.1	60	1.0
77.4	67.5	67.2	1.0	0.625	0.0	72.5	16.3	73.1	74.9	77.4	1.0	0.617	0.0	71.9	17.6	72.7	74.8	76	1.0
89.2	75.0	75.6	1.0	0.75	0.0	81.3	1.1	82.3	82.3	89.2	1.0	0.75	0.0	81.3	1.1	82.3	82.3	89	1.0
96.9	82.5	83.9	1.0	0.875	0.0	88.7	-11.0	90.6	91.3	96.9	1.0	0.867	0.0	88.3	-10.1	90.2	90.7	96	1.0
100.4	90.0	92.3	1.0	1.0	0.0	92.8	-17.5	95.2	96.8	100.4	1.0	1.0	0.0	92.9	-17.4	95.3	96.9	100	1.0
108.8	97.5	101.0	0.875	1.0	0.0	83.7	-27.3	80.1	84.7	108.8	0.875	1.0	0.0	84.4	-26.8	81.2	85.5	108	1.0
120.1	105.0	109.7	0.75	1.0	0.0	74.4	-37.9	65.2	75.5	120.1	0.75	1.0	0.0	74.4	-37.8	65.3	75.5	120	1.0
130.4	112.5	118.5	0.625	1.0	0.0	67.3	-45.9	53.9	70.9	130.4	0.625	1.0	0.0	67.8	-45.4	54.8	71.2	129	0.84
139.3	120.0	127.2	0.5	1.0	0.0	61.7	-53.9	46.2	71.0	139.3	0.5	1.0	0.0	61.8	-53.8	46.2	71.0	139	0.84
142.0	127.5	136.0	0.375	1.0	0.0	60.5	-56.5	44.0	71.6	142.0	0.375	1.0	0.0	60.6	-56.2	44.2	71.6	141	0.667
145.1	135.0	144.7	0.25	1.0	0.0	58.6	-59.0	41.1	71.9	145.1	0.25	1.0	0.0	58.7	-58.9	41.1	71.9	145	0.561
145.5	142.5	153.4	0.125	1.0	0.0	58.5	-59.5	40.8	72.2	145.5	0.125	1.0	0.0	58.5	-59.4	40.9	72.2	145	0.377
145.5	150.0	162.2	0.0	1.0	0.0	58.5	-59.5	40.8	72.2	145.5	0.0	1.0	0.0	58.5	-59.4	40.9	72.2	145	0.0
146.1	157.5	169.0	0.0	1.0	0.125	57.9	-60.4	40.4	72.7	146.1	0.0	1.0	0.117	58.0	-60.3	40.5	72.7	146	0.0
147.2	165.0	175.9	0.0	1.0	0.25	57.6	-60.6	38.9	72.0	147.2	0.0	1.0	0.25	57.6	-60.5	38.9	72.0	147	0.0
148.5	172.5	182.7	0.0	1.0	0.375	57.2	-61.5	37.6	72.1	148.5	0.0	1.0	0.367	57.3	-61.4	37.7	72.1	148	0.0
151.6	180.0	189.6	0.0	1.0	0.5	57.1	-60.7	37.2	68.9	151.6	0.0	1.0	0.5	57.1	-60.6	37.2	68.9	151	0.0
154.2	187.5	196.4	0.0	1.0	0.625	57.3	-59.4	28.6	65.9	154.2	0.0	1.0	0.617	57.3	-59.4	28.9	66.2	154	0.0
161.5	195.0	203.2	0.0	1.0	0.75	58.4	-55.1	18.4	58.1	161.5	0.0	1.0	0.75	58.4	-55.0	18.4	58.1	161	0.0
180.5	202.5	210.1	0.0	1.0	0.875	59.9	-46.4	-0.4	46.4	180.5	0.0	1.0	0.867	59.8	-47.1	0.6	47.2	179	0.0
208.3	210.0	216.9	0.0	1.0	1.0	57.0	-40.5	-21.8	46.1	208.3	0.0	1.0	1.0	57.1	-40.5	-21.8	46.1	208	0.0
226.7	217.5	223.8	0.0	0.875	1.0	53.3	-35.2	-37.3	51.3	226.7	0.0	0.875	1.0	53.6	-35.7	-36.3	51.0	225	0.0
243.5	225.0	230.6	0.0	0.75	1.0	52.6	-24.9	-50.1	56.0	243.5	0.0	0.75	1.0	52.7	-24.8	-50.1	56.0	243	0.0
248.9	232.5	237.5	0.0	0.625	1.0	49.4	-19.3	-50.3	53.8	248.9	0.0	0.625	1.0	49.6	-19.6	-50.2	54.0	248	0.0
253.6	240.0	244.3	0.0	0.5	1.0	47.1	-14.6	-50.0	52.1	253.6	0.0	0.5	1.0	47.1	-14.6	-50.0	52.2	253	0.0
256.9	247.5	251.2	0.0	0.375	1.0	45.3	-11.4	-49.7	51.0	256.9	0.0	0.375	1.0	45.4	-11.6	-49.7	51.1	256	0.0
261.2	255.0	258.0	0.0	0.25	1.0	42.9	-7.6	-49.7	50.3	261.2	0.0	0.25	1.0	43.0	-7.6	-49.6	50.3	261	0.0
264.0	262.5	264.8	0.0	0.125	1.0	41.5	-5.0	-49.0	49.2	264.0	0.0	0.125	1.0	41.7	-5.1	-49.0	49.4	263	0.0
264.0	270.0	271.7	0.0	0.0	1.0	41.5	-5.0	-49.0	49.2	264.0	0.0	0.0	1.0	41.6	-5.0	-48.9	49.3	264	0.377
265.1	277.5	278.8	0.0	0.125	0.0	40.9	-4.1	-49.0	49.2	265.1	0.0	0.125	0.0	41.0	-4.2	-49.0	49.3	265	0.0
266.0	285.0	285.9	0.0	0.25	0.0	40.3	-3.3	-49.3	49.4	266.0	0.0	0.25	0.0	40.4	-3.3	-49.2	49.4	266	0.0
270.0	292.5	293.0	0.0	0.375	0.0	38.3	0.0	-50.1	50.1	270.0	0.0	0.375	0.0	38.5	-0.1	-50.0	50.0	269	0.0
279.6	300.0	300.1	0.5	0.0	1.0	36.4	8.1	-47.9	48.5	279.6	0.5	0.0	1.0	36.5	8.1	-48.6	47.9	279	0.0
295.4	307.5	307.2	0.625	0.0	1.0	37.3	20.1	-42.2	46.7	295.4	0.625	0.0	1.0	37.3	19.9	-42.6	46.9	294	0.0
313.1	315.0	314.3	0.75	0.0	1.0	41.4	32.1	-34.2	46.9	313.1	0.75	0.0	1.0	41.4	32.1	-34.2	47.0	313	0.0
332.4	322.5	321.4	0.875	0.0	1.0	45.7	48.0	-25.0	54.1	332.4	0.875	0.0	1.0	45.8	47.0	-25.0	54.1	332	0.0
351.5	330.0	328.6	1.0	0.0	1.0	50.1	71.1	-10.5	71.8	351.5	1.0	0.0	1.0	50.2	71.1	-10.4	71.9	351	0.0
354.0	337.5	335.7	1.0	0.0	0.875	48.7	74.0	-7.7	74.4	354.0	1.0	0.0	0.883	48.7	73.9	-7.8	74.4	354	0.0
358.5	345.0	342.8	1.0	0.0	0.75	48.3	72.7	-1.8	72.7	358.5	1.0	0.0	0.75	48.3	72.7	-1.8	72.7	358	0.0
364.5	352.5	349.9	1.0	0.0	0.625	48.3	70.3	5.5	70.5	364.5	1.0	0.0	0.625	48.3	70.3	5.5	70.5	364	0.0
369.8	360.0	357.0	1.0	0.0	0.5	48.3	68.4	11.9	69.5	369.8	1.0	0.0	0.5	48.3	68.4	11.9	69.5	369	0.0
377.3	367.5	364.1	1.0	0.0	0.375	48.4	65.6	20.4	68.8	377.3	1.0	0.0	0.375	48.4	65.6	20.4	68.8	377	0.0
384.8	375.0	371.2	1.0	0.0	0.25	48.3	64.2	29.8	70.8	384.8	1.0	0.0	0.25	48.3	64.2	29.8	70.8	384	0.0
390.8	382.5	378.3	1.0	0.0	0.125	48.4	63.4	37.8	73.8	390.8	1.0	0.0	0.125	48.4	63.4	37.8	73.8	390	0.0
393.8	390.0	385.4	1.0	0.0	0.0	48.1	63.3	42.5	76.2	393.8	1.0	0.0	0.0	48.1	63.3	42.5	76.3	393	1.0
RE710-70	1-003734-10																		

LAB*ab,0, YN=0%, XY Zmw=2.0, 2.1, 2.1, 85.9, 90.9, 95.1, LAB*nmw=15.8, 0.0, 0.0, 96.4, 0.0, 0.0, not adapted=adapted
Output: Offset standard print; separation cmyk6*, D65, page 8/33

TUB-test chart RE71; 1080 standard colours, cf=0.9
48 step hue circles; rgb-LabCh*tables
input: rgb/cmyk -> rgbd
output: transfer to rgbd

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; $h_{ab,d65} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;										Six hue angles of the 60 degree standard colours RYGBCM; $h_{ab,s} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$									
Six hue angles of the device colours RYGBCM; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; Six hue angles of the elementary colours RYGBCM; $h_{ab,s} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$										Six hue angles of the elementary colours RYGBCM; $h_{ab,s} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$									
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,d}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$	$h_{ab,s}$
89	75	75	75	75	75	75	75	75	75	89	75	75	75	75	75	75	75	75	75
90	76	76	76	76	76	76	76	76	76	90	76	76	76	76	76	76	76	76	76
91	77	77	77	77	77	77	77	77	77	91	77	77	77	77	77	77	77	77	77
92	78	78	78	78	78	78	78	78	78	92	78	78	78	78	78	78	78	78	78
93	79	80	80	80	80	80	80	80	80	93	79	80	80	80	80	80	80	80	80
94	80	81	81	81	81	81	81	81	81	94	80	81	81	81	81	81	81	81	81
95	81	82	82	82	82	82	82	82	82	95	81	82	82	82	82	82	82	82	82
96	82	83	83	83	83	83	83	83	83	96	82	83	83	83	83	83	83	83	83
97	83	84	84	84	84	84	84	84	84	97	83	84	84	84	84	84	84	84	84
97	84	85	85	85	85	85	85	85	85	97	84	85	85	85	85	85	85	85	85
98	85	86	86	86	86	86	86	86	86	98	85	86	86	86	86	86	86	86	86
98	86	87	87	87	87	87	87	87	87	98	86	87	87	87	87	87	87	87	87
99	87	88	88	88	88	88	88	88	88	99	87	88	88	88	88	88	88	88	88
99	88	89	89	89	89	89	89	89	89	99	88	89	89	89	89	89	89	89	89
99	89	90	90	90	90	90	90	90	90	99	89	90	90	90	90	90	90	90	90
100	90	92	92	92	92	92	92	92	92	100	90	92	92	92	92	92	92	92	92
101	91	93	93	93	93	93	93	93	93	101	91	93	93	93	93	93	93	93	93
102	92	94	94	94	94	94	94	94	94	102	92	94	94	94	94	94	94	94	94
103	93	95	95	95	95	95	95	95	95	103	93	95	95	95	95	95	95	95	95
104	94	96	96	96	96	96	96	96	96	104	94	96	96	96	96	96	96	96	96
106	95	98	98	98	98	98	98	98	98	106	95	98	98	98	98	98	98	98	98
107	96	99	99	99	99	99	99	99	99	107	96	99	99	99	99	99	99	99	99
108	97	100	100	100	100	100	100	100	100	108	97	100	100	100	100	100	100	100	100
109	98	101	101	101	101	101	101	101	101	109	98	101	101	101	101	101	101	101	101
111	99	102	102	102	102	102	102	102	102	111	99	102	102	102	102	102	102	102	102
112	100	103	103	103	103	103	103	103	103	112	100	103	103	103	103	103	103	103	103
114	101	105	105	105	105	105	105	105	105	114	101	105	105	105	105	105	105	105	105
115	102	106	106	106	106	106	106	106	106	115	102	106	106	106	106	106	106	106	106
117	103	107	107	107	107	107	107	107	107	117	103	107	107	107	107	107	107	107	107
118	104	108	108	108	108	108	108	108	108	118	104	108	108	108	108	108	108	108	108
120	105	109	109	109	109	109	109	109	109	120	105	109	109	109	109	109	109	109	109
121	106	110	110	110	110	110	110	110	110	121	106	110	110	110	110	110	110	110	110
122	107	112	112	112	112	112	112	112	112	122	107	112	112	112	112	112	112	112	112
124	108	113	113	113	113	113	113	113	113	124	108	113	113	113	113	113	113	113	113
125	109	114	114	114	114	114	114	114	114	125	109	114	114	114	114	114	114	114	114
126	110	115	115	115	115	115	115	115	115	126	110	115	115	115	115	115	115	115	115
128	111	116	116	116	116	116	116	116	116	128	111	116	116	116	116	116	116	116	116
129	112	117	117	117	117	117	117	117	117	129	112	117	117	117	117	117	117	117	117
131	113	119	119	119	119	119	119	119	119	131	113	119	119	119	119	119	119	119	119
132	114	120	120	120	120	120	120	120	120	132	114	120	120	120	120	120	120	120	120
133	115	121	121	121	121	121	121	121	121	133	115	121	121	121	121	121	121	121	121
134	116	122	122	122	122	122	122	122	122	134	116	122	122	122	122	122	122	122	122
135	117	123	123	123	123	123	123	123	123	135	117	123	123	123	123	123	123	123	123
136	118	124	124	124	124	124	124	124	124	136	118	124	124	124	124	124	124	124	124
138	119	126	126	126	126	126	126	126	126	138	119	126	126	126	126	126	126	126	126
139	120	127	127	127	127	127	127	127	127	139	120	127	127	127	127	127	127	127	127

input: $rgb/cmyk \rightarrow rgbd$
output: transfer to $rgbd$

TUB-test chart RE71; 1080 standard colours, $cf=0.9$
48 step hue circles; $rgb-LabCh$ *tables

1-0031034-L0
1-0031034-F0

RE710-70

Output: Offset standard print; separation cmyk6*, D65, page 11/38

Six hue angles of the device colours RYGBCM; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; 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^*_d361M	lab^*_d361M	rgb^*_d361Mi	lab^*_d361Mi	rgb^*_d361Mi	lab^*_d361Mi	rgb^*_d361Mi
139	120	127	0.5	1.0	0.0	61.7	-53.9	46.2	71.0
139	121	128	0.0	61.5	-53.9	46.2	71.0	139	139
140	122	129	0.466	1.0	0.0	61.4	-54.6	45.6	71.2
140	123	130	0.45	1.0	0.0	61.2	-54.9	45.4	71.2
140	124	131	0.433	1.0	0.0	61.0	-55.3	45.1	71.3
141	125	133	0.416	1.0	0.0	60.9	-55.6	44.8	71.4
141	126	134	0.4	1.0	0.0	60.7	-56.0	44.5	71.5
141	127	135	0.383	1.0	0.0	60.5	-56.3	44.2	71.6
142	128	136	0.366	1.0	0.0	60.3	-56.6	43.9	71.6
142	129	137	0.35	1.0	0.0	60.1	-57.0	43.5	71.7
143	130	138	0.333	1.0	0.0	59.8	-57.3	43.1	71.7
143	131	140	0.316	1.0	0.0	59.6	-57.7	42.7	71.8
143	132	141	0.3	1.0	0.0	59.3	-58.0	42.3	71.8
144	133	142	0.283	1.0	0.0	59.1	-58.3	41.9	71.8
144	134	143	0.266	1.0	0.0	58.9	-58.6	41.5	71.9
145	135	144	0.25	1.0	0.0	58.6	-59.0	41.1	71.9
145	136	145	0.233	1.0	0.0	58.6	-59.0	41.0	71.9
145	137	147	0.216	1.0	0.0	58.6	-59.1	41.0	72.0
145	138	148	0.2	1.0	0.0	58.5	-59.2	41.0	72.0
145	139	149	0.183	1.0	0.0	58.5	-59.3	40.9	72.0
145	140	150	0.166	1.0	0.0	58.5	-59.3	40.9	72.1
145	141	151	0.15	1.0	0.0	58.5	-59.4	40.9	72.1
145	142	152	0.133	1.0	0.0	58.5	-59.5	40.8	72.2
145	143	154	0.116	1.0	0.0	58.5	-59.5	40.8	72.2
145	144	155	0.1	1.0	0.0	58.5	-59.5	40.8	72.2
145	145	156	0.083	1.0	0.0	58.5	-59.5	40.8	72.2
145	146	157	0.066	1.0	0.0	58.5	-59.5	40.8	72.2
145	147	158	0.049	1.0	0.0	58.5	-59.5	40.8	72.2
145	148	159	0.033	1.0	0.0	58.5	-59.5	40.8	72.2
145	149	161	0.016	1.0	0.0	58.5	-59.5	40.8	72.2
145	150	162	0.0	1.0	0.0	58.5	-59.5	40.8	72.2
145	151	163	0.0	1.0	0.016	58.4	-59.6	40.8	72.2
145	152	164	0.0	1.0	0.033	58.3	-59.7	40.7	72.3
145	153	164	0.0	1.0	0.05	58.2	-59.9	40.7	72.4
145	154	165	0.0	1.0	0.066	58.2	-60.0	40.6	72.4
145	155	166	0.0	1.0	0.083	58.1	-60.1	40.5	72.5
146	156	167	0.0	1.0	0.1	58.0	-60.2	40.5	72.6
146	157	168	0.0	1.0	0.116	58.0	-60.3	40.4	72.6
146	158	169	0.0	1.0	0.133	57.9	-60.4	40.3	72.6
146	159	170	0.0	1.0	0.15	57.9	-60.4	40.1	72.5
146	160	171	0.0	1.0	0.166	57.8	-60.4	39.9	72.4
146	161	172	0.0	1.0	0.183	57.8	-60.5	39.7	72.4
146	162	173	0.0	1.0	0.2	57.7	-60.5	39.5	72.3
146	163	174	0.0	1.0	0.216	57.7	-60.5	39.3	72.2
147	164	175	0.0	1.0	0.233	57.6	-60.5	39.1	72.1
147	165	175	0.0	1.0	0.25	57.6	-60.6	38.9	72.0
LAB* h_{ab} , YN=0%, XY Zmw=2.0, 2.1, 2.1, 85.9, 90.9, 95.1, LAB* m_w =15.8, 0.0, 96.4, 0.0, 0.0, not adapted=adapted									
1-0031134-L0									
RE710-70									
TUB-test chart RE71; 1080 standard colours, $cf=0.9$ input: $rgb/cmyk - > rgb$ 48 step hue circles: $rab-LabCh$ *tables output: transfer to rab									

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; $h_{ab,d65} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;										Six hue angles of the elementary colours RYGBCM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$									
Six hue angles of the device colours RYGBCM; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; Six hue angles of the elementary colours RYGBCM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$										Six hue angles of the elementary colours RYGBCM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$									
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	$rgb^*_{ab,e}$	$rgb^*_{ab,d}$	$rgb^*_{ab,s}$	$lab^*_{ab,e}$	$lab^*_{ab,d}$	$lab^*_{ab,s}$	C_d	$rgb^*_{ab,e}$	$rgb^*_{ab,d}$	$rgb^*_{ab,s}$	$lab^*_{ab,e}$	$lab^*_{ab,d}$	$lab^*_{ab,s}$	$rgb^*_{ab,e}$	$rgb^*_{ab,d}$	$rgb^*_{ab,s}$	$lab^*_{ab,e}$
208	210	216	0.0	1.0	1.0	57.0	-40.5	-21.8	46.1	208	0.0	0.989	1.0	56.8	-40.2	-23.2	46.6	210	0.0
210	211	217	0.0	0.983	1.0	56.5	-40.2	-23.9	46.8	210	0.0	0.982	1.0	56.6	-40.1	-24.0	46.9	211	0.0
213	212	218	0.0	0.966	1.0	56.0	-39.7	-26.0	47.5	213	0.0	0.975	1.0	56.3	-39.9	-24.9	47.2	212	0.0
215	213	219	0.0	0.95	1.0	55.5	-39.1	-28.1	48.2	215	0.0	0.968	1.0	56.1	-39.7	-25.7	47.4	213	0.0
218	214	220	0.0	0.933	1.0	55.0	-38.4	-30.2	48.9	218	0.0	0.962	1.0	55.9	-39.5	-26.6	47.7	214	0.0
220	215	221	0.0	0.916	1.0	54.5	-37.6	-32.2	49.6	220	0.0	0.955	1.0	55.7	-39.2	-27.4	48.0	215	0.0
223	216	222	0.0	0.9	1.0	54.0	-36.7	-34.3	50.3	223	0.0	0.948	1.0	55.5	-39.0	-28.3	48.3	216	0.0
225	217	223	0.0	0.883	1.0	53.5	-35.7	-36.3	51.0	225	0.0	0.941	1.0	55.3	-38.7	-29.1	48.6	217	0.0
227	218	224	0.0	0.866	1.0	53.2	-34.6	-38.3	51.6	227	0.0	0.934	1.0	55.1	-38.4	-30.0	48.9	218	0.0
230	219	225	0.0	0.85	1.0	53.1	-33.5	-40.1	52.2	230	0.0	0.928	1.0	54.9	-38.1	-30.8	49.2	219	0.0
232	220	226	0.0	0.833	1.0	53.1	-32.3	-41.9	52.9	232	0.0	0.921	1.0	54.7	-37.8	-31.7	49.4	220	0.0
234	221	227	0.0	0.816	1.0	53.0	-31.0	-43.6	53.5	234	0.0	0.914	1.0	54.5	-37.4	-32.5	49.7	221	0.0
236	222	227	0.0	0.8	1.0	52.9	-29.6	-45.3	54.1	236	0.0	0.907	1.0	54.3	-37.1	-33.4	50.0	222	0.0
239	223	228	0.0	0.783	1.0	52.8	-28.1	-47.0	54.7	239	0.0	0.9	1.0	54.1	-36.7	-34.2	50.3	223	0.0
241	224	229	0.0	0.766	1.0	52.7	-26.5	-48.6	55.4	241	0.0	0.894	1.0	53.9	-36.3	-35.0	50.6	224	0.0
243	225	230	0.0	0.75	1.0	52.6	-24.9	-50.1	56.0	243	0.0	0.887	1.0	53.7	-35.9	-35.9	50.9	225	0.0
244	226	231	0.0	0.733	1.0	52.2	-24.1	-50.2	55.7	244	0.0	0.88	1.0	53.5	-35.4	-36.7	51.2	226	0.0
245	227	232	0.0	0.716	1.0	51.8	-23.4	-50.2	55.4	245	0.0	0.873	1.0	53.3	-35.0	-37.5	51.4	227	0.0
245	228	233	0.0	0.7	1.0	51.3	-22.6	-50.3	55.1	245	0.0	0.866	1.0	53.3	-34.5	-38.3	51.7	228	0.0
246	229	234	0.0	0.683	1.0	50.9	-21.1	-50.3	54.8	246	0.0	0.858	1.0	53.2	-34.0	-39.1	52.0	229	0.0
247	230	235	0.0	0.666	1.0	50.4	-21.1	-50.3	54.6	247	0.0	0.851	1.0	53.2	-33.5	-39.9	52.3	230	0.0
247	231	236	0.0	0.65	1.0	50.0	-20.4	-50.3	54.3	247	0.0	0.843	1.0	53.2	-33.0	-40.7	52.5	231	0.0
248	232	237	0.0	0.633	1.0	49.6	-19.6	-50.3	54.0	248	0.0	0.836	1.0	53.1	-32.4	-41.5	52.8	232	0.0
249	233	237	0.0	0.616	1.0	49.2	-19.0	-50.2	53.7	249	0.0	0.829	1.0	53.1	-31.9	-42.3	53.1	233	0.0
249	234	238	0.0	0.6	1.0	48.9	-18.3	-50.2	53.5	249	0.0	0.821	1.0	53.0	-31.3	-43.1	53.4	234	0.0
250	235	239	0.0	0.583	1.0	48.6	-17.7	-50.2	53.3	250	0.0	0.814	1.0	53.0	-30.7	-43.9	53.7	235	0.0
251	236	240	0.0	0.566	1.0	48.3	-17.1	-50.2	53.0	251	0.0	0.806	1.0	53.0	-30.1	-44.6	53.9	236	0.0
251	237	241	0.0	0.55	1.0	48.0	-16.5	-50.2	52.8	251	0.0	0.799	1.0	52.9	-29.4	-45.4	54.2	237	0.0
252	238	242	0.0	0.533	1.0	47.7	-15.8	-50.1	52.6	252	0.0	0.791	1.0	52.9	-28.8	-46.1	54.5	238	0.0
253	239	243	0.0	0.516	1.0	47.4	-15.2	-50.1	52.3	253	0.0	0.784	1.0	52.9	-28.1	-46.8	54.8	239	0.0
253	240	244	0.0	0.5	1.0	47.1	-14.6	-50.0	52.1	253	0.0	0.777	1.0	52.8	-27.4	-47.6	55.0	240	0.0
254	241	245	0.0	0.483	1.0	46.8	-14.0	-50.0	52.0	254	0.0	0.769	1.0	52.8	-26.7	-48.3	55.3	241	0.0
254	242	246	0.0	0.466	1.0	46.6	-13.8	-49.9	51.8	254	0.0	0.762	1.0	52.7	-26.0	-49.0	55.6	242	0.0
254	243	247	0.0	0.45	1.0	46.4	-13.3	-49.9	51.7	254	0.0	0.754	1.0	52.7	-25.3	-49.7	55.9	243	0.0
255	244	248	0.0	0.433	1.0	46.1	-12.9	-49.9	51.5	255	0.0	0.741	1.0	52.4	-24.4	-50.1	55.9	244	0.0
255	245	248	0.0	0.416	1.0	45.9	-12.5	-49.8	51.4	255	0.0	0.717	1.0	51.8	-23.3	-50.2	55.5	245	0.0
256	246	249	0.0	0.4	1.0	45.6	-12.1	-49.8	51.2	256	0.0	0.694	1.0	51.2	-22.3	-50.2	55.1	246	0.0
256	247	250	0.0	0.383	1.0	45.4	-11.6	-49.7	51.1	256	0.0	0.671	1.0	50.6	-21.3	-50.2	54.7	247	0.0
257	248	251	0.0	0.366	1.0	45.1	-11.2	-49.7	50.9	257	0.0	0.648	1.0	50.0	-20.2	-50.2	54.3	248	0.0
257	249	252	0.0	0.35	1.0	44.8	-10.7	-49.7	50.8	257	0.0	0.624	1.0	49.4	-19.2	-50.2	53.9	249	0.0
258	250	253	0.0	0.333	1.0	44.5	-10.2	-49.7	50.8	258	0.0	0.598	1.0	48.9	-18.2	-50.2	53.5	250	0.0
258	251	254	0.0	0.316	1.0	44.2	-9.6	-49.7	50.7	258	0.0	0.571	1.0	48.4	-17.2	-50.1	53.1	251	0.0
259	252	255	0.0	0.3	1.0	43.9	-9.1	-49.7	50.6	259	0.0	0.544	1.0	47.9	-16.2	-50.1	52.8	252	0.0
260	253	256	0.0	0.283	1.0	43.5	-8.6	-49.7	50.5	260	0.0	0.518	1.0	47.5	-15.2	-50.0	52.4	253	0.0
260	254	257	0.0	0.266	1.0	43.2	-8.1	-49.7	50.4	260	0.0	0.487	1.0	47.0	-14.2	-49.9	52.0	254	0.0
261	255	258	0.0	0.25	1.0	42.9	-7.6	-49.7	50.3	261	0.0	0.45	1.0	46.4	-13.3	-49.8	51.7	255	0.0
261	255	258	0.0	0.25	1.0	42.9	-7.6	-49.7	50.3	261	0.0	0.45	1.0	46.4	-13.3	-49.8	51.7	255	0.0

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

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

TUB-test chart RE71; 1080 standard colours, $cf=0.9$
48 step hue circles; $rgb-LabCh$ *tables

Six hue angles of the device colours RYGBM; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; 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 elementary colours RYGBM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$																						
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	$rgb^*_{ab,d}$	$rgb^*_{ab,s}$	$rgb^*_{ab,e}$	$LAB^*_{ab,d}$	$LAB^*_{ab,s}$	$LAB^*_{ab,e}$	$rgb^*_{ab,d}$	$rgb^*_{ab,s}$	$rgb^*_{ab,e}$	$LAB^*_{ab,d}$	$LAB^*_{ab,s}$	$LAB^*_{ab,e}$	$rgb^*_{ab,d}$	$rgb^*_{ab,s}$	$rgb^*_{ab,e}$	$LAB^*_{ab,d}$	$LAB^*_{ab,s}$	$LAB^*_{ab,e}$	$rgb^*_{ab,d}$	$rgb^*_{ab,s}$	$rgb^*_{ab,e}$									
279	300	300	0.5	0.0	1.0	36.4	8.1	-47.9	48.5	279	0.657	0.0	1.0	38.4	23.4	-40.4	46.8	300	0.5	0.0	1.0	0.658	0.0	1.0	38.4	23.5	-40.4	46.8	300	0.5	0.0	1.0
281	301	301	0.516	0.0	1.0	36.5	9.8	-47.3	48.3	281	0.664	0.0	1.0	38.6	24.1	-40.0	46.8	301	0.517	0.0	1.0	0.665	0.0	1.0	38.6	24.2	-40.0	46.8	301	0.517	0.0	1.0
283	302	302	0.533	0.0	1.0	36.6	11.5	-46.7	48.1	283	0.671	0.0	1.0	38.8	24.8	-39.6	46.8	302	0.533	0.0	1.0	0.672	0.0	1.0	38.8	24.9	-39.6	46.8	302	0.533	0.0	1.0
285	303	303	0.55	0.0	1.0	36.8	13.1	-46.0	47.8	285	0.678	0.0	1.0	39.1	25.5	-39.2	46.9	303	0.55	0.0	1.0	0.678	0.0	1.0	39.1	25.5	-39.2	46.9	303	0.55	0.0	1.0
288	304	304	0.566	0.0	1.0	36.9	14.7	-45.2	47.6	288	0.685	0.0	1.0	39.3	26.2	-38.8	46.9	304	0.567	0.0	1.0	0.685	0.0	1.0	39.3	26.2	-38.8	46.9	304	0.567	0.0	1.0
290	305	304	0.583	0.0	1.0	37.0	16.3	-44.4	47.3	290	0.692	0.0	1.0	39.5	26.9	-38.3	46.9	304	0.583	0.0	1.0	0.692	0.0	1.0	39.5	26.8	-38.3	46.9	304	0.583	0.0	1.0
292	306	305	0.6	0.0	1.0	37.1	17.8	-43.6	47.1	292	0.699	0.0	1.0	39.8	27.6	-37.8	46.9	306	0.6	0.0	1.0	0.698	0.0	1.0	39.7	27.5	-37.9	46.9	305	0.6	0.0	1.0
294	307	306	0.616	0.0	1.0	37.2	19.3	-42.6	46.8	294	0.706	0.0	1.0	40.0	28.2	-37.4	46.9	307	0.617	0.0	1.0	0.705	0.0	1.0	39.9	28.1	-37.5	46.9	306	0.617	0.0	1.0
296	308	307	0.633	0.0	1.0	37.5	20.9	-41.8	46.7	296	0.713	0.0	1.0	40.2	28.9	-36.9	46.9	308	0.633	0.0	1.0	0.712	0.0	1.0	40.2	28.7	-37.0	46.9	307	0.633	0.0	1.0
299	309	308	0.65	0.0	1.0	38.1	22.6	-40.9	46.8	299	0.72	0.0	1.0	40.5	29.5	-36.4	46.9	309	0.65	0.0	1.0	0.718	0.0	1.0	40.4	29.3	-36.5	46.9	308	0.65	0.0	1.0
301	310	309	0.666	0.0	1.0	38.6	24.3	-39.9	46.8	301	0.728	0.0	1.0	40.7	30.2	-35.9	46.9	310	0.667	0.0	1.0	0.725	0.0	1.0	40.6	30.0	-36.0	46.9	309	0.667	0.0	1.0
303	311	310	0.683	0.0	1.0	39.2	26.0	-38.9	46.8	303	0.735	0.0	1.0	40.9	30.8	-35.3	47.0	311	0.683	0.0	1.0	0.732	0.0	1.0	40.8	30.6	-35.6	47.0	310	0.683	0.0	1.0
306	312	311	0.7	0.0	1.0	39.7	27.6	-37.8	46.8	306	0.742	0.0	1.0	41.2	31.4	-34.8	47.0	312	0.7	0.0	1.0	0.738	0.0	1.0	41.0	31.2	-35.1	47.0	311	0.7	0.0	1.0
308	313	312	0.716	0.0	1.0	40.3	29.1	-36.7	46.9	308	0.749	0.0	1.0	41.4	32.0	-34.3	47.0	313	0.717	0.0	1.0	0.745	0.0	1.0	41.3	31.7	-34.5	47.0	312	0.717	0.0	1.0
310	314	313	0.733	0.0	1.0	40.8	30.6	-35.5	46.9	310	0.755	0.0	1.0	41.6	32.9	-33.9	47.3	314	0.733	0.0	1.0	0.752	0.0	1.0	41.5	32.4	-34.1	47.1	313	0.733	0.0	1.0
313	315	314	0.75	0.0	1.0	41.4	32.1	-34.2	46.9	313	0.762	0.0	1.0	41.8	33.7	-33.6	47.7	315	0.75	0.0	1.0	0.758	0.0	1.0	41.7	33.2	-33.8	47.4	314	0.75	0.0	1.0
315	316	315	0.766	0.0	1.0	42.0	34.3	-33.4	47.9	315	0.768	0.0	1.0	42.1	34.6	-33.3	48.0	316	0.767	0.0	1.0	0.764	0.0	1.0	41.9	34.0	-33.5	47.8	315	0.767	0.0	1.0
318	317	316	0.783	0.0	1.0	42.5	36.5	-32.5	48.9	318	0.775	0.0	1.0	42.3	35.4	-32.9	48.4	317	0.783	0.0	1.0	0.77	0.0	1.0	42.1	34.8	-33.2	48.2	316	0.783	0.0	1.0
320	318	317	0.8	0.0	1.0	43.1	38.6	-31.4	49.8	320	0.781	0.0	1.0	42.5	36.3	-32.5	48.8	318	0.8	0.0	1.0	0.776	0.0	1.0	42.3	35.6	-32.8	48.5	317	0.8	0.0	1.0
323	319	318	0.816	0.0	1.0	43.7	40.8	-30.2	50.8	323	0.788	0.0	1.0	42.7	37.1	-32.2	49.2	319	0.817	0.0	1.0	0.782	0.0	1.0	42.5	36.4	-32.5	48.9	318	0.817	0.0	1.0
326	320	319	0.833	0.0	1.0	44.3	42.9	-28.9	51.7	326	0.794	0.0	1.0	43.0	37.9	-31.7	49.5	320	0.833	0.0	1.0	0.789	0.0	1.0	42.8	37.2	-32.1	49.2	319	0.833	0.0	1.0
328	321	320	0.85	0.0	1.0	44.8	45.0	-27.4	52.7	328	0.801	0.0	1.0	43.2	38.8	-31.3	49.9	321	0.85	0.0	1.0	0.795	0.0	1.0	43.0	38.0	-31.7	49.6	320	0.85	0.0	1.0
331	322	321	0.866	0.0	1.0	45.4	47.0	-25.9	53.7	331	0.807	0.0	1.0	43.4	39.6	-30.9	50.3	322	0.867	0.0	1.0	0.801	0.0	1.0	43.2	38.8	-31.3	49.9	321	0.867	0.0	1.0
333	323	322	0.883	0.0	1.0	46.0	49.6	-24.5	55.3	333	0.814	0.0	1.0	43.6	40.5	-30.4	50.7	323	0.883	0.0	1.0	0.807	0.0	1.0	43.4	39.6	-30.9	50.3	321	0.883	0.0	1.0
336	324	322	0.9	0.0	1.0	46.6	52.8	-23.2	57.7	336	0.82	0.0	1.0	43.8	41.3	-29.9	51.0	324	0.9	0.0	1.0	0.813	0.0	1.0	43.6	40.4	-30.4	50.6	322	0.9	0.0	1.0
338	325	323	0.916	0.0	1.0	47.2	56.0	-21.7	60.0	338	0.827	0.0	1.0	44.1	42.1	-29.4	51.4	325	0.917	0.0	1.0	0.819	0.0	1.0	43.8	41.2	-30.0	51.0	323	0.917	0.0	1.0
341	326	324	0.933	0.0	1.0	47.8	59.1	-19.9	62.4	341	0.833	0.0	1.0	44.3	42.9	-28.9	51.8	326	0.933	0.0	1.0	0.826	0.0	1.0	44.0	42.0	-29.5	51.3	324	0.933	0.0	1.0
343	327	325	0.95	0.0	1.0	48.4	62.2	-17.9	64.8	343	0.84	0.0	1.0	44.5	43.7	-28.3	52.2	327	0.95	0.0	1.0	0.832	0.0	1.0	44.2	42.7	-29.0	51.7	325	0.95	0.0	1.0
346	328	326	0.966	0.0	1.0	48.9	65.3	-15.7	67.1	346	0.846	0.0	1.0	44.7	44.5	-27.7	52.5	328	0.967	0.0	1.0	0.838	0.0	1.0	44.5	43.5	-28.5	52.0	326	0.967	0.0	1.0
349	329	327	0.983	0.0	1.0	49.5	68.2	-13.2	69.5	349	0.853	0.0	1.0	45.0	45.3	-27.1	52.9	329	0.983	0.0	1.0	0.844	0.0	1.0	44.7	44.3	-27.9	52.4	327	0.983	0.0	1.0
351	330	328	1.0	0.0	1.0	50.1	71.1	-10.5	71.8	351	0.859	0.0	1.0	45.2	46.1	-26.5	53.3	330	1.0	0.0	1.0	0.85	0.0	1.0	44.9	45.0	-27.4	52.8	328	1.0	0.0	1.0
351	331	329	1.0	0.0	0.983	49.9	71.5	-10.1	72.2	351	0.866	0.0	1.0	45.4	46.9	-25.9	53.7	331	1.0	0.0	0.983	0.856	0.0	1.0	45.1	45.8	-26.8	53.1	329	1.0	0.0	0.983
352	332	330	1.0	0.0	0.966	49.7	71.9	-9.8	72.5	352	0.872	0.0	1.0	45.6	47.7	-25.3	54.0	332	1.0	0.0	0.967	0.862	0.0	1.0	45.3	46.5	-26.2	53.5	330	1.0	0.0	0.967
352	333	331	1.0	0.0	0.95	49.6	72.3	-9.4	72.9	352	0.879	0.0	1.0	45.9	48.7	-24.7	54.7	333	1.0	0.0	0.95	0.869	0.0	1.0	45.5	47.3	-25.6	53.8	331	1.0	0.0	0.95
352	334	332	1.0	0.0	0.933	49.4	72.7	-9.0	73.2	352	0.885	0.0	1.0	46.1	50.0	-24.3	55.6	334	1.0	0.0	0.933	0.875	0.0	1.0	45.7	48.0	-25.0	54.2	332	1.0	0.0	0.933
353	335	333	1.0	0.0	0.916	49.2	73.1	-8.6	73.6	353	0.892	0.0	1.0	46.3	51.3	-23.8	56.6	335	1.0	0.0	0.917	0.881	0.0	1.0	46.0	49.2	-24.6	55.0	333	1.0	0.0	0.917
353	336	334	1.0	0.0	0.9	49.0	73.4	-8.2	73.9	353	0.898	0.0	1.0	46.6	52.5	-23.3	57.5	336	1.0	0.0	0.9	0.887	0.0	1.0	46.2	50.4	-24.1	55.9	334	1.0	0.0	0.9
353	337	335	1.0	0.0	0.883	48.8	73.8	-7.9	74.3	353	0.905	0.0	1.0	46.8	53.8	-22.7	58.4	337	1.0	0.0	0.883	0.893	0.0	1.0	46.4	51.6	-23.7	56.8	335	1.0	0.0	0.883
354	338	336	1.0	0.0	0.866	48.6	74.0	-7.3	74.3	354	0.911	0.0	1.0	47.0	55.0	-22.1	59.3	338	1.0	0.0	0.867	0.899	0.0	1.0	46.6	52.8	-23.2	57.7	336	1.0	0.0	0.867
354	339	337	1.0	0.0	0.85	48.6	73.8	-6.5	74.1	354	0.918	0.0	1.0	47.3	56.3	-21.5	60.3	339	1.0	0.0	0.85	0.906	0.0	1.0	46.8	53.9	-22.6	58.5	337	1.0	0.0	0.85
355	340	338	1.0	0.0	0.833	48.5	73.6	-5.7	73.9	355	0.924	0.0	1.0	47.5	57.5	-20.8	61.2	340	1.0	0.0	0.833	0.912	0.0	1.0	47.1	55.1	-22.1	59.4	338	1.0	0.0	0.833

TUB registration: 20150701-RE71/RE71L0NA.TXT /PS
application for measurement of laser printer output, no separation rgb (RGB)

TUB material: code=rha4ta

see similar files: <http://130.149.60.45/~farbmatrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

<http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 18/33

nf	HfC*Fd	rgb_Fd	ic_Fd	hs_Fd	rgb_Fd	LabCh*Fd	rgb_Fd	DF*Fd	hs_Md	rgb_Md	LabCh*Md				
0/648	R00Y_100_100d	1.0	0.0	1.0	0.5	390	1.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
1/657	R13Y_100_100d	1.0	0.125	1.0	0.5	370	1.0	0.116	0.0	0.0	48.1	63.3	42.5	76.2	33.8
2/666	R25Y_100_100d	1.0	0.25	1.0	0.5	340	1.0	0.233	0.0	0.0	48.1	63.3	42.5	76.2	33.8
3/675	R38Y_100_100d	1.0	0.375	1.0	0.5	310	1.0	0.366	0.0	0.0	48.1	63.3	42.5	76.2	33.8
4/684	R50Y_100_100d	1.0	0.5	1.0	0.5	280	1.0	0.5	0.0	0.0	48.1	63.3	42.5	76.2	33.8
5/693	R63Y_100_100d	1.0	0.625	1.0	0.5	250	1.0	0.633	0.0	0.0	48.1	63.3	42.5	76.2	33.8
6/702	R75Y_100_100d	1.0	0.75	1.0	0.5	220	1.0	0.766	0.0	0.0	48.1	63.3	42.5	76.2	33.8
7/711	R88Y_100_100d	1.0	0.875	1.0	0.5	86	1.0	0.875	0.0	0.0	48.1	63.3	42.5	76.2	33.8
8/720	Y00C_100_100d	1.0	1.0	1.0	0.5	90	1.0	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
9/639	Y13C_100_100d	0.875	1.0	1.0	0.5	97	0.883	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
10/558	Y25C_100_100d	0.625	1.0	1.0	0.5	104	0.766	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
11/477	Y38C_100_100d	0.375	1.0	1.0	0.5	112	0.633	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
12/396	Y50C_100_100d	0.25	1.0	1.0	0.5	120	0.5	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
13/315	Y63C_100_100d	0.125	1.0	1.0	0.5	128	0.366	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
14/234	Y75C_100_100d	0.0	1.0	1.0	0.5	136	0.233	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
15/153	Y88C_100_100d	0.125	1.0	1.0	0.5	143	0.116	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
16/72	G00C_100_100d	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
17/73	G13C_100_100d	0.0	1.0	1.0	0.5	157	0.0	1.0	0.116	0.0	48.1	63.3	42.5	76.2	33.8
18/74	G25C_100_100d	0.0	1.0	1.0	0.5	164	0.0	1.0	0.233	0.0	48.1	63.3	42.5	76.2	33.8
19/75	G38C_100_100d	0.0	1.0	1.0	0.5	172	0.0	1.0	0.366	0.0	48.1	63.3	42.5	76.2	33.8
20/76	G50C_100_100d	0.0	1.0	1.0	0.5	180	0.0	1.0	0.5	0.0	48.1	63.3	42.5	76.2	33.8
21/77	G63C_100_100d	0.0	1.0	1.0	0.5	188	0.0	1.0	0.633	0.0	48.1	63.3	42.5	76.2	33.8
22/78	G75C_100_100d	0.0	1.0	1.0	0.5	196	0.0	1.0	0.766	0.0	48.1	63.3	42.5	76.2	33.8
23/79	G88C_100_100d	0.0	1.0	1.0	0.5	203	0.0	1.0	0.883	0.0	48.1	63.3	42.5	76.2	33.8
24/80	C00B_100_100d	0.0	1.0	1.0	0.5	210	0.0	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
25/81	C13B_100_100d	0.0	0.875	1.0	1.0	217	0.0	0.883	1.0	0.0	48.1	63.3	42.5	76.2	33.8
26/62	C25B_100_100d	0.0	0.75	1.0	1.0	224	0.0	0.766	1.0	0.0	48.1	63.3	42.5	76.2	33.8
27/53	C38B_100_100d	0.0	0.625	1.0	1.0	232	0.0	0.633	1.0	0.0	48.1	63.3	42.5	76.2	33.8
28/44	C50B_100_100d	0.0	0.5	1.0	1.0	240	0.0	0.5	1.0	0.0	48.1	63.3	42.5	76.2	33.8
29/35	C63B_100_100d	0.0	0.375	1.0	1.0	248	0.0	0.366	1.0	0.0	48.1	63.3	42.5	76.2	33.8
30/26	C75B_100_100d	0.0	0.25	1.0	1.0	256	0.0	0.233	1.0	0.0	48.1	63.3	42.5	76.2	33.8
31/17	C88B_100_100d	0.0	0.125	1.0	1.0	263	0.0	0.116	1.0	0.0	48.1	63.3	42.5	76.2	33.8
32/8	B00M_100_100d	0.0	1.0	1.0	0.5	270	0.0	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
33/89	B13M_100_100d	0.125	1.0	1.0	0.5	277	0.116	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
34/170	B25M_100_100d	0.25	1.0	1.0	0.5	284	0.233	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
35/251	B38M_100_100d	0.375	1.0	1.0	0.5	292	0.366	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
36/332	B50M_100_100d	0.5	1.0	1.0	0.5	300	0.5	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
37/413	B63M_100_100d	0.625	1.0	1.0	0.5	308	0.633	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
38/494	B75M_100_100d	0.75	1.0	1.0	0.5	316	0.766	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
39/575	B88M_100_100d	0.875	1.0	1.0	0.5	323	0.883	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
40/656	M00R_100_100d	1.0	0.0	1.0	0.5	330	1.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
41/655	M13R_100_100d	1.0	0.0	1.0	0.5	337	1.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
42/654	M25R_100_100d	1.0	0.0	1.0	0.5	344	1.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
43/653	M38R_100_100d	1.0	0.0	1.0	0.5	352	1.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
44/652	M50R_100_100d	1.0	0.0	1.0	0.5	360	1.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
45/651	M63R_100_100d	1.0	0.0	1.0	0.5	368	1.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
46/650	M75R_100_100d	1.0	0.0	1.0	0.5	376	1.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
47/649	M88R_100_100d	1.0	0.0	1.0	0.5	383	1.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
48/648	R00Y_100_100d	1.0	0.0	1.0	0.5	390	1.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
49/0	NW_000d	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
50/91	NW_013d	0.125	0.0	0.0	0.0	360	0.125	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
51/182	NW_025d	0.25	0.0	0.0	0.0	360	0.25	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
52/273	NW_038d	0.375	0.0	0.0	0.0	360	0.375	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
53/364	NW_050d	0.5	0.0	0.0	0.0	360	0.5	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
54/455	NW_063d	0.625	0.0	0.0	0.0	360	0.625	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
55/546	NW_075d	0.75	0.0	0.0	0.0	360	0.75	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
56/637	NW_088d	0.875	0.0	0.0	0.0	360	0.875	0.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8
57/728	NW_100d	1.0	1.0	1.0	0.0	360	1.0	1.0	1.0	1.0	48.1	63.3	42.5	76.2	33.8

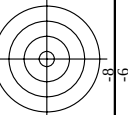
Mean color difference of this page:

delta E* = 1.2

RE710-7N, Page 18/33-F

TUB-test chart RE71; 1080 standard colours, cf=0.9
colors and differences, ΔE^*

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



TUB material: code=rha4ta

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

TUB-
colors

DF710-7N Date 20/33-F

see similar files: <http://130.149.60.45/~farbmatrik/RE71/RE71.HTM>
 technical information: <http://www.ps.uni-bam.de> or <http://130.149.60.45/~farbmatrik>

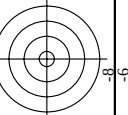
TUB registration: 20150701-RE71/RE71L0NA.TXT /PS
application for measurement of laser printer output, no separation rgb (RGB)

TUB material: code=rha4ta

see similar files: <http://130.149.60.45/~farbmatrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

<http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 21/33

n	H1C*Fd	rgb_Fd	icd_Fd	h3a_Fd	rgb**Fd	LabCH*Fd	LabCH**Fd	rgb**Nd	LabCH*Nd	DF**Fd	h3a**Fd	rgb**Nd	LabCH*Nd
81	B00Y_012_0124	0.125 0.0 0.125	0.125 0.0 0.125	390 330 330	5.3 9.5 33.5	7.9 19.8 8.8	17.2 1.4 0.5	0.125 0.0 0.125	17.2 1.4 0.5	19.6 8.5 389	1.0 0.0 0.0	48.1 63.3 42.5	76.2 33.8
82	B00R_012_0124	0.125 0.0 0.125	0.125 0.0 0.125	390 330 330	5.3 9.5 33.5	7.9 19.8 8.8	17.2 1.4 0.5	0.125 0.0 0.125	17.2 1.4 0.5	19.6 8.5 389	1.0 0.0 0.0	48.1 63.3 42.5	76.2 33.8
83	B25R_025_0254	0.125 0.0 0.125	0.125 0.0 0.125	209 20.8 20.8	-11.9 12.1 279.6	-0.5 20.5 24.6	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	30.0 19.5 300	0.5 0.0 0.0	36.4 8.1 -47.9	48.5 279.6
84	B15R_037_0374	0.125 0.0 0.125	0.125 0.0 0.125	209 20.8 20.8	-11.9 12.1 279.6	-0.5 20.5 24.6	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	30.0 19.5 300	0.5 0.0 0.0	36.4 8.1 -47.9	48.5 279.6
85	B11R_050_0504	0.125 0.0 0.125	0.125 0.0 0.125	209 20.8 20.8	-11.9 12.1 279.6	-0.5 20.5 24.6	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	30.0 19.5 300	0.5 0.0 0.0	36.4 8.1 -47.9	48.5 279.6
86	B09R_062_0624	0.125 0.0 0.125	0.125 0.0 0.125	281 281 281	-17.3 24.7 265.9	-31.3 23.0 36.8	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	37.3 27.3 174	0.282 0.183 0.0	40.4 -3.4 -49.2	49.2 265.9
87	B07R_075_0754	0.125 0.0 0.125	0.125 0.0 0.125	279 279 279	-34.5 34.5 34.5	-37.7 36.9 265.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
88	B06R_087_0874	0.125 0.0 0.125	0.125 0.0 0.125	279 279 279	-34.5 34.5 34.5	-37.7 36.9 265.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
89	B05R_100_1004	0.125 0.0 0.125	0.125 0.0 0.125	277 277 277	-49.0 49.2 100.4	-49.0 49.2 100.4	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
90	Y00C_012_0124	0.125 0.0 0.125	0.125 0.0 0.125	360 360 360	25.8 0.0 0.0	-2.1 19.0 12.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
91	NW_0124	0.125 0.0 0.125	0.125 0.0 0.125	360 360 360	25.8 0.0 0.0	-2.1 19.0 12.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
92	B00R_025_0124	0.125 0.0 0.125	0.125 0.0 0.125	360 360 360	25.8 0.0 0.0	-2.1 19.0 12.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
93	B00R_037_0254	0.125 0.0 0.125	0.125 0.0 0.125	360 360 360	25.8 0.0 0.0	-2.1 19.0 12.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
94	B00R_050_0374	0.125 0.0 0.125	0.125 0.0 0.125	360 360 360	25.8 0.0 0.0	-2.1 19.0 12.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
95	B00R_062_0504	0.125 0.0 0.125	0.125 0.0 0.125	360 360 360	25.8 0.0 0.0	-2.1 19.0 12.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
96	B00R_075_0504	0.125 0.0 0.125	0.125 0.0 0.125	360 360 360	25.8 0.0 0.0	-2.1 19.0 12.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
97	B00R_087_0754	0.125 0.0 0.125	0.125 0.0 0.125	360 360 360	25.8 0.0 0.0	-2.1 19.0 12.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
98	B00R_100_0874	0.125 0.0 0.125	0.125 0.0 0.125	360 360 360	25.8 0.0 0.0	-2.1 19.0 12.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
99	Y50C_025_0254	0.125 0.0 0.125	0.125 0.0 0.125	360 360 360	25.8 0.0 0.0	-2.1 19.0 12.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
100	G00B_025_0124	0.125 0.0 0.125	0.125 0.0 0.125	150 150 150	31.2 31.2 31.2	-7.4 5.1 9.0	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
101	G50B_025_0124	0.125 0.0 0.125	0.125 0.0 0.125	150 150 150	31.2 31.2 31.2	-7.4 5.1 9.0	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
102	G75B_037_0254	0.125 0.0 0.125	0.125 0.0 0.125	240 240 240	33.6 33.6 33.6	-12.5 13.0 208.3	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
103	G44B_062_0374	0.125 0.0 0.125	0.125 0.0 0.125	240 240 240	33.6 33.6 33.6	-12.5 13.0 208.3	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
104	G88B_050_0374	0.125 0.0 0.125	0.125 0.0 0.125	240 240 240	33.6 33.6 33.6	-12.5 13.0 208.3	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
105	G00B_075_0624	0.125 0.0 0.125	0.125 0.0 0.125	256 256 256	42.3 42.3 42.3	-3.9 36.8 31.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
106	G30B_100_0624	0.125 0.0 0.125	0.125 0.0 0.125	256 256 256	42.3 42.3 42.3	-3.9 36.8 31.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
107	G30B_100_0874	0.125 0.0 0.125	0.125 0.0 0.125	256 256 256	42.3 42.3 42.3	-3.9 36.8 31.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
108	Y68C_037_0374	0.125 0.0 0.125	0.125 0.0 0.125	131 131 131	32.2 32.2 32.2	-16.0 26.9 143.4	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
109	G00B_037_0254	0.125 0.0 0.125	0.125 0.0 0.125	150 150 150	31.2 31.2 31.2	-7.4 5.1 9.0	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
110	G50B_037_0254	0.125 0.0 0.125	0.125 0.0 0.125	150 150 150	31.2 31.2 31.2	-7.4 5.1 9.0	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
111	G68B_050_0374	0.125 0.0 0.125	0.125 0.0 0.125	240 240 240	33.6 33.6 33.6	-12.5 13.0 208.3	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
112	G75B_062_0374	0.125 0.0 0.125	0.125 0.0 0.125	240 240 240	33.6 33.6 33.6	-12.5 13.0 208.3	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
113	G44B_097_0624	0.125 0.0 0.125	0.125 0.0 0.125	240 240 240	33.6 33.6 33.6	-12.5 13.0 208.3	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
114	G00B_075_0624	0.125 0.0 0.125	0.125 0.0 0.125	256 256 256	42.3 42.3 42.3	-3.9 36.8 31.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
115	G44B_097_0504	0.125 0.0 0.125	0.125 0.0 0.125	240 240 240	33.6 33.6 33.6	-12.5 13.0 208.3	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
116	G68B_100_0874	0.125 0.0 0.125	0.125 0.0 0.125	256 256 256	42.3 42.3 42.3	-3.9 36.8 31.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
117	Y76C_050_0504	0.125 0.0 0.125	0.125 0.0 0.125	136 136 136	35.9 35.9 35.9	44.1 44.1 44.1	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
118	G00B_050_0374	0.125 0.0 0.125	0.125 0.0 0.125	150 150 150	31.2 31.2 31.2	-7.4 5.1 9.0	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
119	G15B_050_0374	0.125 0.0 0.125	0.125 0.0 0.125	169 169 169	41.6 41.6 41.6	-22.9 14.3 270.0	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
120	G34B_050_0374	0.125 0.0 0.125	0.125 0.0 0.125	191 191 191	48.8 48.8 48.8	-8.2 17.2 208.3	0.125 0.0 0.125	0.125 0.0 0.125	0.125 0.0 0.125	26.6 4.1 278	0.15 0.0 0.0	40.6 -3.8 -49.2	49.2 265.1
121	G50B_050_0374	0.125 0.0 0.125	0.125 0.0 0.125	210									



TUB material: code=rha4ta

RE710-7N, Page 23/33-F

[illegible]

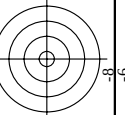
-see similar files: <http://130.149.60.45/~farbmatrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

<http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 25/33

n	H1C*Fd	rgb_Fd	icr_Fd	hsa_Fd	rgb*Fd	LabCH*Fd	rgb**Fd	LabCH**Fd	DF**Fd	hsa**d	rgb**d	LabCH**d
405	R00Y_062_062a	0.625 0.0 0.0	0.625 0.625 0.312	390	0.625 0.0 0.0	36.0 39.5	0.625 0.0 0.0	33.8 41.5	27.2	49.6	33.2	49.6
406	R00Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.114	36.1 39.9	0.625 0.0 0.114	33.8 41.5	27.2	49.6	33.2	49.6
407	R00Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.239	36.2 41.1	0.625 0.0 0.239	33.8 41.5	27.2	49.6	33.2	49.6
408	R00Y_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.364	36.3 42.3	0.625 0.0 0.364	33.8 41.5	27.2	49.6	33.2	49.6
409	R00Y_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.625 0.0 0.489	36.4 43.7	0.625 0.0 0.489	33.8 41.5	27.2	49.6	33.2	49.6
410	R00Y_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.614	36.5 45.1	0.625 0.0 0.614	33.8 41.5	27.2	49.6	33.2	49.6
411	R00Y_062_062a	0.625 0.0 0.75	0.625 0.625 0.312	318	0.625 0.0 0.739	36.6 46.5	0.625 0.0 0.739	33.8 41.5	27.2	49.6	33.2	49.6
412	R00Y_062_062a	0.625 0.0 0.875	0.625 0.625 0.312	306	0.625 0.0 0.864	36.7 47.9	0.625 0.0 0.864	33.8 41.5	27.2	49.6	33.2	49.6
413	R00Y_062_062a	0.625 0.1 0.0	0.625 0.625 0.312	294	0.625 0.114 0.0	36.8 49.3	0.625 0.114 0.0	33.8 41.5	27.2	49.6	33.2	49.6
414	R00Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	282	0.625 0.125 0.0	36.9 50.7	0.625 0.125 0.0	33.8 41.5	27.2	49.6	33.2	49.6
415	R00Y_062_062a	0.625 0.125 0.125	0.625 0.625 0.312	270	0.625 0.125 0.125	37.0 52.1	0.625 0.125 0.125	33.8 41.5	27.2	49.6	33.2	49.6
416	R00Y_062_062a	0.625 0.125 0.25	0.625 0.625 0.312	258	0.625 0.125 0.25	37.1 53.5	0.625 0.125 0.25	33.8 41.5	27.2	49.6	33.2	49.6
417	R00Y_062_062a	0.625 0.125 0.375	0.625 0.625 0.312	246	0.625 0.125 0.375	37.2 54.9	0.625 0.125 0.375	33.8 41.5	27.2	49.6	33.2	49.6
418	R00Y_062_062a	0.625 0.125 0.5	0.625 0.625 0.312	234	0.625 0.125 0.5	37.3 56.3	0.625 0.125 0.5	33.8 41.5	27.2	49.6	33.2	49.6
419	R00Y_062_062a	0.625 0.125 0.625	0.625 0.625 0.312	222	0.625 0.125 0.625	37.4 57.7	0.625 0.125 0.625	33.8 41.5	27.2	49.6	33.2	49.6
420	R00Y_062_062a	0.625 0.125 0.75	0.625 0.625 0.312	210	0.625 0.125 0.75	37.5 59.1	0.625 0.125 0.75	33.8 41.5	27.2	49.6	33.2	49.6
421	R00Y_062_062a	0.625 0.125 0.875	0.625 0.625 0.312	198	0.625 0.125 0.875	37.6 60.5	0.625 0.125 0.875	33.8 41.5	27.2	49.6	33.2	49.6
422	R00Y_062_062a	0.625 0.125 1.0	0.625 0.625 0.312	186	0.625 0.125 1.0	37.7 61.9	0.625 0.125 1.0	33.8 41.5	27.2	49.6	33.2	49.6
423	R00Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	174	0.625 0.25 0.0	37.8 63.3	0.625 0.25 0.0	33.8 41.5	27.2	49.6	33.2	49.6
424	R00Y_062_062a	0.625 0.25 0.125	0.625 0.625 0.312	162	0.625 0.25 0.125	37.9 64.7	0.625 0.25 0.125	33.8 41.5	27.2	49.6	33.2	49.6
425	R00Y_062_062a	0.625 0.25 0.25	0.625 0.625 0.312	150	0.625 0.25 0.25	38.0 66.1	0.625 0.25 0.25	33.8 41.5	27.2	49.6	33.2	49.6
426	R00Y_062_062a	0.625 0.25 0.375	0.625 0.625 0.312	138	0.625 0.25 0.375	38.1 67.5	0.625 0.25 0.375	33.8 41.5	27.2	49.6	33.2	49.6
427	R00Y_062_062a	0.625 0.25 0.5	0.625 0.625 0.312	126	0.625 0.25 0.5	38.2 68.9	0.625 0.25 0.5	33.8 41.5	27.2	49.6	33.2	49.6
428	R00Y_062_062a	0.625 0.25 0.625	0.625 0.625 0.312	114	0.625 0.25 0.625	38.3 70.3	0.625 0.25 0.625	33.8 41.5	27.2	49.6	33.2	49.6
429	R00Y_062_062a	0.625 0.25 0.75	0.625 0.625 0.312	102	0.625 0.25 0.75	38.4 71.7	0.625 0.25 0.75	33.8 41.5	27.2	49.6	33.2	49.6
430	R00Y_062_062a	0.625 0.25 0.875	0.625 0.625 0.312	90	0.625 0.25 0.875	38.5 73.1	0.625 0.25 0.875	33.8 41.5	27.2	49.6	33.2	49.6
431	R00Y_062_062a	0.625 0.25 1.0	0.625 0.625 0.312	78	0.625 0.25 1.0	38.6 74.5	0.625 0.25 1.0	33.8 41.5	27.2	49.6	33.2	49.6
432	R00Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	66	0.625 0.375 0.0	38.7 75.9	0.625 0.375 0.0	33.8 41.5	27.2	49.6	33.2	49.6
433	R00Y_062_062a	0.625 0.375 0.125	0.625 0.625 0.312	54	0.625 0.375 0.125	38.8 77.3	0.625 0.375 0.125	33.8 41.5	27.2	49.6	33.2	49.6
434	R00Y_062_062a	0.625 0.375 0.25	0.625 0.625 0.312	42	0.625 0.375 0.25	38.9 78.7	0.625 0.375 0.25	33.8 41.5	27.2	49.6	33.2	49.6
435	R00Y_062_062a	0.625 0.375 0.375	0.625 0.625 0.312	30	0.625 0.375 0.375	39.0 80.1	0.625 0.375 0.375	33.8 41.5	27.2	49.6	33.2	49.6
436	R00Y_062_062a	0.625 0.375 0.5	0.625 0.625 0.312	18	0.625 0.375 0.5	39.1 81.5	0.625 0.375 0.5	33.8 41.5	27.2	49.6	33.2	49.6
437	R00Y_062_062a	0.625 0.375 0.625	0.625 0.625 0.312	6	0.625 0.375 0.625	39.2 82.9	0.625 0.375 0.625	33.8 41.5	27.2	49.6	33.2	49.6
438	R00Y_062_062a	0.625 0.375 0.75	0.625 0.625 0.312	0	0.625 0.375 0.75	39.3 84.3	0.625 0.375 0.75	33.8 41.5	27.2	49.6	33.2	49.6
439	R00Y_062_062a	0.625 0.375 0.875	0.625 0.625 0.312	0	0.625 0.375 0.875	39.4 85.7	0.625 0.375 0.875	33.8 41.5	27.2	49.6	33.2	49.6
440	R00Y_062_062a	0.625 0.375 1.0	0.625 0.625 0.312	0	0.625 0.375 1.0	39.5 87.1	0.625 0.375 1.0	33.8 41.5	27.2	49.6	33.2	49.6
441	R00Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	76	0.625 0.5 0.0	39.6 88.5	0.625 0.5 0.0	33.8 41.5	27.2	49.6	33.2	49.6
442	R00Y_062_062a	0.625 0.5 0.125	0.625 0.625 0.312	64	0.625 0.5 0.125	39.7 89.9	0.625 0.5 0.125	33.8 41.5	27.2	49.6	33.2	49.6
443	R00Y_062_062a	0.625 0.5 0.25	0.625 0.625 0.312	52	0.625 0.5 0.25	39.8 91.3	0.625 0.5 0.25	33.8 41.5	27.2	49.6	33.2	49.6
444	R00Y_062_062a	0.625 0.5 0.375	0.625 0.625 0.312	40	0.625 0.5 0.375	39.9 92.7	0.625 0.5 0.375	33.8 41.5	27.2	49.6	33.2	49.6
445	R00Y_062_062a	0.625 0.5 0.5	0.625 0.625 0.312	28	0.625 0.5 0.5	40.0 94.1	0.625 0.5 0.5	33.8 41.5	27.2	49.6	33.2	49.6
446	R00Y_062_062a	0.625 0.5 0.625	0.625 0.625 0.312	16	0.625 0.5 0.625	40.1 95.5	0.625 0.5 0.625	33.8 41.5	27.2	49.6	33.2	49.6
447	R00Y_062_062a	0.625 0.5 0.75	0.625 0.625 0.312	4	0.625 0.5 0.75	40.2 96.9	0.625 0.5 0.75	33.8 41.5	27.2	49.6	33.2	49.6
448	R00Y_062_062a	0.625 0.5 0.875	0.625 0.625 0.312	0	0.625 0.5 0.875	40.3 98.3	0.625 0.5 0.875	33.8 41.5	27.2	49.6	33.2	49.6
449	R00Y_062_062a	0.625 0.5 1.0	0.625 0.625 0.312	0	0.625 0.5 1.0	40.4 99.7	0.625 0.5 1.0	33.8 41.5	27.2	49.6	33.2	49.6
450	R00Y_062_062a	0.625 0.625 0.0	0.625 0.625 0.312	90	0.625 0.625 0.0	40.5 101.1	0.625 0.625 0.0	33.8 41.5	27.2	49.6	33.2	49.6
451	R00Y_062_062a	0.625 0.625 0.125	0.625 0.625 0.312	78	0.625 0.625 0.125	40.6 102.5	0.625 0.625 0.125	33.8 41.5	27.2	49.6	33.2	49.6
452	R00Y_062_062a	0.625 0.625 0.25	0.625 0.625 0.312	66	0.625 0.625 0.25	40.7 103.9	0.625 0.625 0.25	33.8 41.5	27.2	49.6	33.2	49.6
453	R00Y_062_062a	0.625 0.625 0.375	0.625 0.625 0.312	54	0.625 0.625 0.375	40.8 105.3	0.625 0.625 0.375	33.8 41.5	27.2	49.6	33.2	49.6
454	R00Y_062_062a	0.625 0.625 0.5	0.625 0.625 0.312	42	0.625 0.625 0.5	40.9 106.7	0.625 0.625 0.5	33.8 41.5	27.2	49.6	33.2	49.6
455	R00Y_062_062a	0.625 0.625 0.625	0.625 0.625 0.312	30	0.625 0.625 0.625	41.0 108.1	0.625 0.625 0.625	33.8 41.5	27.2	49.6	33.2	49.6
456	R00Y_062_062a	0.625 0.625 0.75	0.625 0.625 0.312	18	0.625 0.625 0.75	41.1 109.5	0.625 0.625 0.75	33.8 41.5	27.2	49.6	33.2	49.6
457	R00Y_062_062a	0.625 0.625 0.875	0.625 0.625 0.312	6	0.625 0.625 0.875	41.2 110.9	0.625 0.625 0.875	33.8 41.5	27.2	49.6	33.2	49.6
458	R00Y_062_062a	0.625 0.625 1.0	0.625 0.625 0.312	0	0.625 0.625 1.0	41.3 112.3	0.625 0.625 1.0	33.8 41.5	27.2	49.6	33.2	49.6
459	R00Y_062_062a	0.625 0.75 0.0	0.625 0.625 0.312	90	0.625 0.75 0.0	41.4 113.7						

<http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 27/33

n	H1C*Fd	rgb_Fd	ic_Fd	h1a_Fd	rgb_Fd	LabCH*Fd	LabCH*Fd	rgb_Fd	LabCH*Fd	DE*Fd	h1a_Fd	rgb_Fd	LabCH*Fd
567	R00Y,087,087a	0.875, 0.0, 0.0	0.875, 0.875, 0.437	390	0.875, 0.0, 0.0	44.0, 55.4, 37.2	66.7, 66.7, 66.7	0.875, 0.0, 0.0	44.0, 55.4, 37.2	2.9, 389	389	0.0, 0.0, 0.0	48.1, 63.3, 42.5
568	R36Y,087,087a	0.875, 0.0, 0.125	0.875, 0.875, 0.437	382	0.875, 0.0, 0.116	44.3, 55.5, 37.2	66.7, 64.4, 30.4	0.875, 0.0, 0.125	43.3, 55.8, 34.5	35.2, 69.1	389	1.0, 0.0, 0.0	48.1, 63.3, 42.5
569	R20Y,087,087a	0.875, 0.0, 0.25	0.875, 0.875, 0.437	374	0.875, 0.0, 0.233	44.2, 56.4, 34.2	61.7, 23.8, 14.8	0.875, 0.0, 0.25	42.8, 56.3, 25.3	61.7, 24.1	375	1.0, 0.0, 0.0	48.1, 63.3, 42.5
570	R08Y,087,087a	0.875, 0.0, 0.375	0.875, 0.875, 0.437	365	0.875, 0.0, 0.364	44.3, 58.3, 15.4	60.4, 61.4, 6.3	0.875, 0.0, 0.375	40.9, 59.0, 12.1	60.2, 11.6	365	1.0, 0.0, 0.0	48.1, 63.3, 42.5
571	B70R,087,087a	0.875, 0.0, 0.5	0.875, 0.875, 0.437	355	0.875, 0.0, 0.511	44.3, 61.0, 6.7	61.4, 63.3, 359.3	0.875, 0.0, 0.5	43.3, 63.5, 359.9	60.2, 11.6	354	1.0, 0.0, 0.0	48.1, 63.3, 42.5
572	B63R,087,087a	0.875, 0.0, 0.625	0.875, 0.875, 0.437	346	0.875, 0.0, 0.641	44.3, 63.0, 6.7	63.3, 359.3, 68.6	0.875, 0.0, 0.625	43.3, 68.6, 68.6	60.2, 11.6	344	1.0, 0.0, 0.0	48.1, 63.3, 42.5
573	B56R,087,087a	0.875, 0.0, 0.75	0.875, 0.875, 0.437	338	0.875, 0.0, 0.758	44.3, 64.7, 6.3	65.0, 354.3, 68.6	0.875, 0.0, 0.75	43.3, 68.6, 68.6	60.2, 11.6	337	1.0, 0.0, 0.0	48.1, 63.3, 42.5
574	B50R,087,087a	0.875, 0.0, 0.875	0.875, 0.875, 0.437	330	0.875, 0.0, 0.875	44.3, 65.2, 6.2	62.9, 351.5, 68.6	0.875, 0.0, 0.875	43.3, 68.6, 68.6	60.2, 11.6	330	1.0, 0.0, 0.0	48.1, 63.3, 42.5
575	B44R,100,100a	0.875, 0.0, 1.0	0.875, 0.875, 0.437	323	0.875, 0.0, 1.0	44.6, 49.6, 6.2	55.3, 353.7, 68.6	0.875, 0.0, 1.0	43.3, 68.6, 68.6	60.2, 11.6	323	1.0, 0.0, 0.0	48.1, 63.3, 42.5
576	R13Y,087,087a	0.875, 0.125, 0.0	0.875, 0.875, 0.437	38	0.875, 0.116, 0.0	44.7, 54.1, 39.1	66.8, 35.8, 68.6	0.875, 0.125, 0.0	43.3, 68.6, 68.6	60.2, 11.6	37	1.0, 0.133, 0.0	48.1, 63.3, 42.5
577	R00Y,087,075a	0.875, 0.125, 0.125	0.875, 0.75, 0.5	390	0.875, 0.125, 0.125	50.3, 47.7, 31.8	57.2, 33.8, 68.6	0.875, 0.125, 0.125	43.3, 68.6, 68.6	60.2, 11.6	389	1.0, 0.0, 0.0	48.1, 63.3, 42.5
578	R35Y,087,075a	0.875, 0.125, 0.25	0.875, 0.75, 0.5	381	0.875, 0.125, 0.125	50.3, 47.7, 31.8	54.9, 29.6, 68.6	0.875, 0.125, 0.25	43.3, 68.6, 68.6	60.2, 11.6	382	1.0, 0.0, 0.15	48.1, 63.3, 42.5
579	R18Y,087,075a	0.875, 0.125, 0.375	0.875, 0.75, 0.5	371	0.875, 0.125, 0.362	50.3, 48.8, 18.6	52.3, 20.8, 68.6	0.875, 0.125, 0.375	43.3, 68.6, 68.6	60.2, 11.6	370	1.0, 0.0, 0.316	48.1, 63.3, 42.5
580	R00Y,087,075a	0.875, 0.125, 0.5	0.875, 0.75, 0.5	360	0.875, 0.125, 0.5	50.3, 51.3, 9.8	52.1, 9.8, 68.6	0.875, 0.125, 0.5	43.3, 68.6, 68.6	60.2, 11.6	360	1.0, 0.0, 0.5	48.1, 63.3, 42.5
581	B65R,087,075a	0.875, 0.125, 0.625	0.875, 0.75, 0.5	349	0.875, 0.125, 0.625	50.3, 53.6, 1.6	53.6, 1.7, 68.6	0.875, 0.125, 0.625	43.3, 68.6, 68.6	60.2, 11.6	348	1.0, 0.0, 0.683	48.1, 63.3, 42.5
582	B57R,087,075a	0.875, 0.125, 0.75	0.875, 0.75, 0.5	339	0.875, 0.125, 0.75	50.3, 55.3, 1.8	55.6, 354.9, 68.6	0.875, 0.125, 0.75	43.3, 68.6, 68.6	60.2, 11.6	337	1.0, 0.0, 0.85	48.1, 63.3, 42.5
583	B50R,087,075a	0.875, 0.125, 0.875	0.875, 0.75, 0.5	330	0.875, 0.125, 0.875	51.6, 53.3, 1.8	53.9, 331.1, 68.6	0.875, 0.125, 0.875	43.3, 68.6, 68.6	60.2, 11.6	330	1.0, 0.0, 1.0	48.1, 63.3, 42.5
584	B43R,100,087a	0.875, 0.125, 1.0	0.875, 0.75, 0.5	322	0.875, 0.125, 1.0	51.8, 41.1, 22.6	46.9, 331.1, 68.6	0.875, 0.125, 1.0	43.3, 68.6, 68.6	60.2, 11.6	322	1.0, 0.0, 1.0	48.1, 63.3, 42.5
585	R26Y,087,075a	0.875, 0.25, 0.0	0.875, 0.875, 0.437	46	0.875, 0.233, 0.0	46.2, 51.1, 44.8	68.0, 41.2, 68.6	0.875, 0.25, 0.0	43.3, 68.6, 68.6	60.2, 11.6	44	1.0, 0.266, 0.0	48.1, 63.3, 42.5
586	R15Y,087,075a	0.875, 0.25, 0.125	0.875, 0.75, 0.5	39	0.875, 0.233, 0.125	50.7, 46.2, 34.1	57.4, 36.4, 68.6	0.875, 0.25, 0.125	43.3, 68.6, 68.6	60.2, 11.6	39	1.0, 0.15, 0.0	48.1, 63.3, 42.5
587	R00Y,087,062a	0.875, 0.25, 0.25	0.875, 0.875, 0.437	490	0.875, 0.25, 0.25	56.1, 58.3, 38.3	68.6, 41.2, 68.6	0.875, 0.25, 0.25	43.3, 68.6, 68.6	60.2, 11.6	490	1.0, 0.0, 0.0	48.1, 63.3, 42.5
588	R10Y,087,062a	0.875, 0.25, 0.375	0.875, 0.75, 0.5	390	0.875, 0.25, 0.362	56.3, 61.3, 43.2	68.6, 41.2, 68.6	0.875, 0.25, 0.375	43.3, 68.6, 68.6	60.2, 11.6	390	1.0, 0.0, 0.183	48.1, 63.3, 42.5
589	R11Y,087,062a	0.875, 0.25, 0.5	0.875, 0.875, 0.437	367	0.875, 0.25, 0.488	56.3, 61.3, 43.2	68.6, 41.2, 68.6	0.875, 0.25, 0.5	43.3, 68.6, 68.6	60.2, 11.6	367	1.0, 0.0, 0.383	48.1, 63.3, 42.5
590	B68R,087,062a	0.875, 0.25, 0.625	0.875, 0.75, 0.5	353	0.875, 0.25, 0.636	56.3, 61.3, 43.2	68.6, 41.2, 68.6	0.875, 0.25, 0.625	43.3, 68.6, 68.6	60.2, 11.6	353	1.0, 0.0, 0.616	48.1, 63.3, 42.5
591	B57R,087,062a	0.875, 0.25, 0.75	0.875, 0.875, 0.437	341	0.875, 0.25, 0.75	56.3, 61.3, 43.2	68.6, 41.2, 68.6	0.875, 0.25, 0.75	43.3, 68.6, 68.6	60.2, 11.6	341	1.0, 0.0, 0.816	48.1, 63.3, 42.5
592	B43R,100,062a	0.875, 0.25, 0.875	0.875, 0.75, 0.5	330	0.875, 0.25, 0.875	57.4, 44.4, 20.6	45.9, 31.0, 68.6	0.875, 0.25, 0.875	43.3, 68.6, 68.6	60.2, 11.6	330	1.0, 0.0, 1.0	48.1, 63.3, 42.5
593	B42R,100,075a	0.875, 0.25, 1.0	0.875, 0.75, 0.5	321	0.875, 0.25, 1.0	57.7, 33.7, 20.6	39.5, 328.5, 68.6	0.875, 0.25, 1.0	43.3, 68.6, 68.6	60.2, 11.6	321	1.0, 0.0, 1.0	48.1, 63.3, 42.5
594	R13Y,087,075a	0.875, 0.375, 0.0	0.875, 0.875, 0.437	55	0.875, 0.364, 0.0	52.6, 58.6, 52.5	65.2, 53.6, 68.6	0.875, 0.375, 0.0	43.3, 68.6, 68.6	60.2, 11.6	54	1.0, 0.416, 0.0	48.1, 63.3, 42.5
595	R00Y,087,075a	0.875, 0.375, 0.125	0.875, 0.75, 0.5	49	0.875, 0.364, 0.125	53.5, 48.7, 50.5	54.9, 44.9, 68.6	0.875, 0.375, 0.125	43.3, 68.6, 68.6	60.2, 11.6	48	1.0, 0.316, 0.0	48.1, 63.3, 42.5
596	R31Y,087,075a	0.875, 0.375, 0.25	0.875, 0.75, 0.5	41	0.875, 0.364, 0.25	56.9, 59.1, 56.9	68.6, 41.2, 68.6	0.875, 0.375, 0.25	43.3, 68.6, 68.6	60.2, 11.6	39	1.0, 0.183, 0.0	48.1, 63.3, 42.5
597	R00Y,087,062a	0.875, 0.375, 0.375	0.875, 0.75, 0.5	390	0.875, 0.375, 0.375	62.1, 62.1, 62.1	68.6, 41.2, 68.6	0.875, 0.375, 0.375	43.3, 68.6, 68.6	60.2, 11.6	389	1.0, 0.0, 0.0	48.1, 63.3, 42.5
598	R26Y,087,059a	0.875, 0.375, 0.5	0.875, 0.75, 0.5	376	0.875, 0.375, 0.5	62.1, 62.1, 62.1	68.6, 41.2, 68.6	0.875, 0.375, 0.5	43.3, 68.6, 68.6	60.2, 11.6	376	1.0, 0.0, 0.233	48.1, 63.3, 42.5
599	R00Y,087,059a	0.875, 0.375, 0.625	0.875, 0.75, 0.5	360	0.875, 0.375, 0.625	62.1, 62.1, 62.1	68.6, 41.2, 68.6	0.875, 0.375, 0.625	43.3, 68.6, 68.6	60.2, 11.6	360	1.0, 0.0, 0.0	48.1, 63.3, 42.5
600	B61R,087,059a	0.875, 0.375, 0.75	0.875, 0.75, 0.5	344	0.875, 0.375, 0.758	62.1, 62.1, 62.1	68.6, 41.2, 68.6	0.875, 0.375, 0.75	43.3, 68.6, 68.6	60.2, 11.6	342	1.0, 0.0, 0.766	48.1, 63.3, 42.5
601	B61R,087,059a	0.875, 0.375, 0.875	0.875, 0.75, 0.5	335	0.875, 0.375, 0.875	62.1, 62.1, 62.1	68.6, 41.2, 68.6	0.875, 0.375, 0.875	43.3, 68.6, 68.6	60.2, 11.6	330	1.0, 0.0, 1.0	48.1, 63.3, 42.5
602	B40R,100,062a	0.875, 0.375, 1.0	0.875, 0.75, 0.5	334	0.875, 0.375, 1.0	63.4, 25.5, 18.9	61.7, 323.4, 68.6	0.875, 0.375, 1.0	43.3, 68.6, 68.6	60.2, 11.6	320	1.0, 0.0, 0.816	48.1, 63.3, 42.5
603	R58Y,087,075a	0.875, 0.5, 0.0	0.875, 0.875, 0.437	65	0.875, 0.5, 0.0	62.8, 64.8, 64.8	72.5, 68.6, 68.6	0.875, 0.5, 0.0	43.3, 68.6, 68.6	60.2, 11.6	65	1.0, 0.583, 0.0	48.1, 63.3, 42.5
604	R00Y,087,075a	0.875, 0.5, 0.125	0.875, 0.75, 0.5	60	0.875, 0.5, 0.125	61.6, 64.8, 64.8	68.6, 41.2, 68.6	0.875, 0.5, 0.125	43.3, 68.6, 68.6	60.2, 11.6	59	1.0, 0.5, 0.0	48.1, 63.3, 42.5
605	R38Y,087,062a	0.875, 0.5, 0.25	0.875, 0.875, 0.437	53	0.875, 0.489, 0.25	60.9, 60.9, 60.9	68.6, 41.2, 68.6	0.875, 0.5, 0.25	43.3, 68.6, 68.6	60.2, 11.6	52	1.0, 0.383, 0.0	48.1, 63.3, 42.5
606	R23Y,087,059a	0.875, 0.5, 0.375	0.875, 0.75, 0.5	44	0.875, 0.489, 0.25	60.9, 60.9, 60.9	68.6, 41.2, 68.6	0.875, 0.5, 0.375	43.3, 68.6, 68.6	60.2, 11			



TUB material: code=rha4ta

A target diagram with a bullseye in the center. The bullseye is marked with the number 85. The target is divided into concentric circles, and the bullseye is the innermost circle.

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

1: 1080 stan



n	HIC ^{F_d}	$h_{\text{sig}}^{\text{F}_d}$	rgb^{F_d}	$\text{LabCh}^{\text{F}_d}$	$\text{LabCh}^{\text{F}_d}$	rgb^{F_d}	$\text{LabCh}^{\text{F}_d}$	$\text{LabCh}^{\text{F}_d}$	DE^{F_d}	$\text{hansid}^{\text{F}_d}$	rgb^{Hsd}	$\text{LabCh}^{\text{Hsd}}$	$\text{LabCh}^{\text{Hsd}}$
648	R08Y_100_100d	1.0	0.5	390	62.2	33.8	42.5	76.2	73.8	30.8	0.3	383	42.5
649	R38Y_100_100d	1.0	0.5	383	48.1	63.3	48.1	48.1	63.3	30.8	0.3	383	42.5
650	R26Y_100_100d	1.0	0.5	376	64.2	29.8	70.8	24.8	1.0	0.0	0.116	48.1	48.1
651	R13Y_100_100d	1.0	0.5	368	65.6	21.1	68.9	17.8	1.0	0.0	0.233	48.1	48.1
652	R00Y_100_100d	1.0	0.5	360	68.4	11.9	69.5	9.8	1.0	0.0	0.366	48.1	48.1
653	R68R_100_100d	1.0	0.5	352	70.4	5.1	70.6	4.1	1.0	0.0	0.0	0.633	48.1
654	B61R_100_100d	1.0	0.5	344	72.9	7.2	72.9	357.9	1.0	0.0	0.0	0.766	48.1
655	B55R_100_100d	1.0	0.5	337	74.3	353.8	74.3	353.8	1.0	0.0	0.0	0.883	48.1
656	R11Y_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
657	R00Y_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
658	R36Y_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
659	R23Y_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
660	R23Y_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
661	R08Y_100_100d	1.0	0.5	337	74.3	353.8	74.3	353.8	1.0	0.0	0.0	0.0	50.1
662	B70R_100_100d	1.0	0.5	344	72.9	7.2	72.9	357.9	1.0	0.0	0.0	0.0	50.1
663	B68R_100_100d	1.0	0.5	352	70.4	5.1	70.6	4.1	1.0	0.0	0.0	0.0	50.1
664	B56R_100_100d	1.0	0.5	337	74.3	353.8	74.3	353.8	1.0	0.0	0.0	0.0	50.1
665	B50R_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
666	R23Y_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
667	R13Y_100_100d	1.0	0.5	366	65.6	21.1	68.9	17.8	1.0	0.0	0.0	0.0	50.1
668	R38Y_100_100d	1.0	0.5	383	48.1	63.3	48.1	48.1	63.3	30.8	0.3	383	42.5
669	R08Y_100_100d	1.0	0.5	376	64.2	29.8	70.8	24.8	1.0	0.0	0.0	0.0	50.1
670	R18Y_100_100d	1.0	0.5	360	68.4	11.9	69.5	9.8	1.0	0.0	0.0	0.0	50.1
671	R00Y_100_100d	1.0	0.5	352	70.4	5.1	70.6	4.1	1.0	0.0	0.0	0.0	50.1
672	B65R_100_100d	1.0	0.5	344	72.9	7.2	72.9	357.9	1.0	0.0	0.0	0.0	50.1
673	B57R_100_100d	1.0	0.5	337	74.3	353.8	74.3	353.8	1.0	0.0	0.0	0.0	50.1
674	B50R_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
675	R36Y_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
676	R26Y_100_100d	1.0	0.5	366	65.6	21.1	68.9	17.8	1.0	0.0	0.0	0.0	50.1
677	R13Y_100_100d	1.0	0.5	376	64.2	29.8	70.8	24.8	1.0	0.0	0.0	0.0	50.1
678	R08Y_100_100d	1.0	0.5	383	48.1	63.3	48.1	48.1	63.3	30.8	0.3	383	42.5
679	R00Y_100_100d	1.0	0.5	360	68.4	11.9	69.5	9.8	1.0	0.0	0.0	0.0	50.1
680	R11Y_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
681	B69R_100_100d	1.0	0.5	368	65.6	21.1	68.9	17.8	1.0	0.0	0.0	0.0	50.1
682	B69R_100_100d	1.0	0.5	368	65.6	21.1	68.9	17.8	1.0	0.0	0.0	0.0	50.1
683	B50R_100_100d	1.0	0.5	330	71.1	10.5	71.8	351.5	1.0	0.0	0.0	0.0	50.1
684	R50Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
685	R66Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
686	R31Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
687	R18Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
688	R00Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
689	R26Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
690	R00Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
691	B61R_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
692	B61R_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
693	R63Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
694	R58Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
695	R50Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
696	R38Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
697	R23Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
698	R00Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
699	R23Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
700	B65R_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
701	B65R_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
702	R76Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
703	R73Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
704	R68Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
705	R61Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
706	R50Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
707	R31Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
708	R00Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
709	R00Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
710	R50Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
711	R86Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
712	R86Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
713	R65Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
714	R61Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
715	R76Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
716	R68Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
717	R61Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
718	R60Y_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
719	B50R_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
720	Y00G_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
721	Y00G_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
722	Y00G_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
723	Y00G_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
724	Y00G_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
725	Y00G_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
726	Y00G_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
727	Y00G_100_100d	1.0	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
728	NW_100d	1.0	0.5	360	62.2	33.8	42.5	76.2	73.8	30.8	0.3	383	42.5

Mean color difference of this page:

$\Delta E^* = 11.2$

[illegible]

1

5

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

n	H1C*Fd	rgb_Fd	icr_Fd	hsa_Fd	rgb_Fd	LabCH*Fd	LabCH*Fd	rgb_Fd	DE*Fd	hsa_Fd	LabCH*Nd	rgb_Nd	LabCH*Nd
810	NW_100d	0.875 0.875 1.0	1.0 1.0 1.0	360 270	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	0.5 276.3	360 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
811	BOOR_100.012d	0.875 0.875 1.0	1.0 1.0 1.0	0.125 0.937	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
812	BOOR_100.025d	0.625 0.625 1.0	1.0 1.0 1.0	0.25 0.875	0.625 0.625 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	21.8 254.0	11.1 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
813	BOOR_100.037d	0.625 0.625 1.0	1.0 1.0 1.0	0.375 0.812	0.625 0.625 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	30.9 256.0	22.4 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
814	BOOR_100.050d	0.5 0.5 1.0	1.0 1.0 1.0	0.5 0.75	0.5 0.5 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	48.9 251.0	24.5 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
815	BOOR_100.062d	0.375 0.375 1.0	1.0 1.0 1.0	0.625 0.687	0.375 0.375 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	50.7 261.4	24.5 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
816	BOOR_100.075d	0.25 0.25 1.0	1.0 1.0 1.0	0.75 0.625	0.25 0.25 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	50.7 261.4	24.5 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
817	BOOR_100.087d	0.125 0.125 1.0	1.0 1.0 1.0	0.875 0.562	0.125 0.125 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	50.7 261.4	24.5 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
818	BOOR_100.100d	0.0 0.0 1.0	1.0 1.0 1.0	0.5 0.5	0.0 0.0 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	50.7 261.4	24.5 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
819	YOCOC_100.012d	0.875 0.875 1.0	1.0 1.0 1.0	0.125 0.937	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
820	NW_087d	0.875 0.875 1.0	1.0 1.0 1.0	0.875 0.875 1.0	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
821	BOOR_087.012d	0.875 0.875 1.0	1.0 1.0 1.0	0.875 0.875 1.0	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
822	BOOR_087.025d	0.625 0.625 1.0	1.0 1.0 1.0	0.875 0.875 1.0	0.625 0.625 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
823	BOOR_087.037d	0.5 0.5 1.0	1.0 1.0 1.0	0.875 0.875 1.0	0.5 0.5 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
824	BOOR_087.050d	0.375 0.375 1.0	1.0 1.0 1.0	0.875 0.875 1.0	0.375 0.375 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
825	BOOR_087.062d	0.25 0.25 1.0	1.0 1.0 1.0	0.875 0.875 1.0	0.25 0.25 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
826	BOOR_087.075d	0.125 0.125 1.0	1.0 1.0 1.0	0.875 0.875 1.0	0.125 0.125 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
827	BOOR_087.087d	0.0 0.0 1.0	1.0 1.0 1.0	0.875 0.875 1.0	0.0 0.0 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
828	YOCOC_100.012d	0.875 0.875 1.0	1.0 1.0 1.0	0.125 0.937	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
829	YOCOC_100.025d	0.875 0.875 1.0	1.0 1.0 1.0	0.25 0.875	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
830	YOCOC_100.037d	0.625 0.625 1.0	1.0 1.0 1.0	0.375 0.812	0.625 0.625 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
831	BOOR_075.012d	0.625 0.625 1.0	1.0 1.0 1.0	0.625 0.625 1.0	0.625 0.625 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
832	BOOR_075.025d	0.375 0.375 1.0	1.0 1.0 1.0	0.75 0.562	0.375 0.375 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
833	BOOR_075.037d	0.25 0.25 1.0	1.0 1.0 1.0	0.875 0.437	0.25 0.25 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
834	BOOR_075.050d	0.125 0.125 1.0	1.0 1.0 1.0	0.937 0.375	0.125 0.125 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
835	BOOR_075.062d	0.0 0.0 1.0	1.0 1.0 1.0	0.937 0.375	0.0 0.0 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
836	BOOR_075.075d	0.0 0.0 1.0	1.0 1.0 1.0	0.937 0.375	0.0 0.0 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
837	YOCOC_100.037d	0.875 0.875 1.0	1.0 1.0 1.0	0.375 0.812	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
838	YOCOC_100.050d	0.875 0.875 1.0	1.0 1.0 1.0	0.625 0.625 1.0	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
839	YOCOC_075.012d	0.625 0.625 1.0	1.0 1.0 1.0	0.625 0.625 1.0	0.625 0.625 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
840	NW_062d	0.625 0.625 1.0	1.0 1.0 1.0	0.625 0.625 1.0	0.625 0.625 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
841	BOOR_062.012d	0.625 0.625 1.0	1.0 1.0 1.0	0.625 0.625 1.0	0.625 0.625 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
842	BOOR_062.025d	0.375 0.375 1.0	1.0 1.0 1.0	0.75 0.562	0.375 0.375 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
843	BOOR_062.037d	0.25 0.25 1.0	1.0 1.0 1.0	0.875 0.437	0.25 0.25 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
844	BOOR_062.050d	0.125 0.125 1.0	1.0 1.0 1.0	0.937 0.375	0.125 0.125 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
845	YOCOC_100.062d	0.875 0.875 1.0	1.0 1.0 1.0	0.625 0.625 1.0	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
846	YOCOC_087.037d	0.875 0.875 1.0	1.0 1.0 1.0	0.375 0.812	0.875 0.875 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
847	YOCOC_075.025d	0.625 0.625 1.0	1.0 1.0 1.0	0.625 0.625 1.0	0.625 0.625 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
848	YOCOC_062.012d	0.625 0.625 1.0	1.0 1.0 1.0	0.625 0.625 1.0	0.625 0.625 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
849	NW_050d	0.375 0.375 1.0	1.0 1.0 1.0	0.5 0.5	0.375 0.375 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
850	BOOR_050.012d	0.375 0.375 1.0	1.0 1.0 1.0	0.5 0.5	0.375 0.375 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
851	BOOR_050.025d	0.25 0.25 1.0	1.0 1.0 1.0	0.625 0.562	0.25 0.25 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
852	BOOR_050.037d	0.125 0.125 1.0	1.0 1.0 1.0	0.75 0.437	0.125 0.125 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
853	BOOR_050.050d	0.0 0.0 1.0	1.0 1.0 1.0	0.875 0.375	0.0 0.0 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0
854	BOOR_050.062d	0.0 0.0 1.0	1.0 1.0 1.0	0.875 0.375	0.0 0.0 1.0	96.3 0.0 0.0	96.3 0.0 0.0	1.0 1.0 1.0	8.9 257.9	2.9 270	96.3 0.0 0.0	1.0 1.0 1.0	96.3 0.0 0.0

n	H1C*Fd	rgb_Fd	icr_Fd	hsa_Fd	rgb_Fd	LabCH*Fd	rgb*Fd	LabCH*Fd	DF*Fd	hsa_Fd	rgb*Nd	LabCH*Nd	0.0	0.0	0.0
891	NW_100d	1.0	1.0	1.0	1.0	96.3	1.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
892	B50R_100.012d	1.0	0.875	1.0	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
893	B50R_100.025d	1.0	0.75	1.0	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
894	B50R_100.037d	1.0	0.625	1.0	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
895	B50R_100.050d	1.0	0.5	1.0	0.5	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
896	B50R_100.062d	1.0	0.375	1.0	0.375	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
897	B50R_100.075d	1.0	0.25	1.0	0.25	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
898	B50R_100.087d	1.0	0.125	1.0	0.125	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
899	B50R_100.100d	1.0	0.0	1.0	0.0	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
900	COB_100.012d	0.875	1.0	0.875	1.0	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
901	NW_087d	0.875	0.875	0.875	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
902	B50R_087.012d	0.875	0.75	0.875	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
903	B50R_087.025d	0.875	0.625	0.875	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
904	B50R_087.037d	0.875	0.5	0.875	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
905	B50R_087.050d	0.875	0.375	0.875	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
906	B50R_087.062d	0.875	0.25	0.875	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
907	B50R_087.075d	0.875	0.125	0.875	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
908	B50R_087.087d	0.875	0.0	0.875	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
909	COB_100.025d	0.75	1.0	0.75	1.0	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
910	COB_100.037d	0.75	0.875	0.75	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
911	NW_075d	0.75	0.75	0.75	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
912	B50R_075.012d	0.75	0.625	0.75	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
913	B50R_075.025d	0.75	0.5	0.75	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
914	B50R_075.037d	0.75	0.375	0.75	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
915	B50R_075.050d	0.75	0.25	0.75	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
916	B50R_075.062d	0.75	0.125	0.75	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
917	B50R_075.075d	0.75	0.0	0.75	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
918	COB_100.037d	0.625	1.0	0.625	1.0	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
919	COB_100.050d	0.625	0.875	0.625	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
920	COB_100.062d	0.625	0.75	0.625	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
921	COB_100.075d	0.625	0.625	0.625	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
922	B50R_062.012d	0.625	0.5	0.625	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
923	B50R_062.025d	0.625	0.375	0.625	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
924	B50R_062.037d	0.625	0.25	0.625	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
925	B50R_062.050d	0.625	0.125	0.625	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
926	B50R_062.062d	0.625	0.0	0.625	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
927	COB_100.050d	0.5	1.0	0.5	1.0	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
928	COB_087.037d	0.5	0.875	0.5	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
929	COB_087.050d	0.5	0.75	0.5	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
930	COB_087.062d	0.5	0.625	0.5	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
931	NW_050d	0.5	0.5	0.5	0.5	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
932	B50R_050.012d	0.5	0.375	0.5	0.375	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
933	B50R_050.025d	0.5	0.25	0.5	0.25	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
934	B50R_050.037d	0.5	0.125	0.5	0.125	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
935	B50R_050.050d	0.5	0.0	0.5	0.5	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
936	COB_100.062d	0.375	1.0	0.375	1.0	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
937	COB_087.050d	0.375	0.875	0.375	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
938	COB_087.062d	0.375	0.75	0.375	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
939	COB_087.075d	0.375	0.625	0.375	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
940	NW_037d	0.375	0.5	0.375	0.5	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
941	B50R_037.012d	0.375	0.375	0.375	0.375	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
942	B50R_037.025d	0.375	0.25	0.375	0.375	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
943	B50R_037.037d	0.375	0.125	0.375	0.375	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
944	B50R_037.050d	0.375	0.0	0.375	0.375	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
945	COB_100.075d	0.25	1.0	0.25	1.0	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
946	COB_087.062d	0.25	0.875	0.25	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
947	COB_087.075d	0.25	0.75	0.25	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
948	COB_087.087d	0.25	0.625	0.25	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
949	COB_087.100d	0.25	0.5	0.25	0.5	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
950	COB_087.112d	0.25	0.375	0.25	0.375	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
951	NW_025d	0.25	0.25	0.25	0.25	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
952	B50R_025.012d	0.25	0.125	0.25	0.25	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
953	B50R_025.025d	0.25	0.0	0.25	0.25	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
954	COB_100.087d	0.125	1.0	0.125	1.0	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
955	COB_087.062d	0.125	0.875	0.125	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
956	COB_087.075d	0.125	0.75	0.125	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
957	COB_087.087d	0.125	0.625	0.125	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
958	COB_087.100d	0.125	0.5	0.125	0.5	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
959	COB_087.112d	0.125	0.375	0.125	0.375	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
960	COB_087.125d	0.125	0.25	0.125	0.25	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
961	NW_012d	0.125	0.125	0.125	0.125	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
962	B50R_012.012d	0.125	0.0	0.125	0.125	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
963	COB_100.100d	0.0	1.0	0.0	1.0	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
964	COB_087.087d	0.0	0.875	0.0	0.875	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
965	COB_087.100d	0.0	0.75	0.0	0.75	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
966	COB_087.112d	0.0	0.625	0.0	0.625	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
967	COB_087.125d	0.0	0.5	0.0	0.5	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
968	COB_087.137d	0.0	0.375	0.0	0.375	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
969	COB_087.150d	0.0	0.25	0.0	0.25	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
970	COB_087.162d	0.0	0.125	0.0	0.125	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0
971	NW_000d	0.0	0.0	0.0	0.0	96.3	0.0	96.3	0.0	360	1.0	96.3	0.0	0.0	0.0

Mean color difference of this page: $\Delta E^* = 19.8$

<http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 32/33

n	H1C*Fd	rgb_Fd	ic1_Fd	h1a_Fd	rgb_Fd	LabCH*Fd	LabCH*Fd	rgb_Fd	DE**Fd	h1a_Fd	LabCH*Fd	rgb_Fd	LabCH*Fd
972	NW_0004	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_0124	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_0254	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_0374	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_0504	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_0624	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_0754	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_0874	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_1004	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_0004	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_0124	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_0254	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25	0.25
992	NW_0374	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_0504	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_0624	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_0754	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_0874	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_1004	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_0004	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_0124	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_0254	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_0374	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_0504	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_0624	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_0754	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_0874	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_1004	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_0004	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_0124	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_0254	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_0374	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_0504	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_0624	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_0754	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_0874	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_1004	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_0004	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_0124	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_0254	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_0374	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_0504	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_0624	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_0754	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_0874	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_1004	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_0004	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_0124	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_0254	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_0374	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_0504	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_0624	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_0754	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_0874	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_1004	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_0004	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_0124	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_0254	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_0374	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_0504	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_0624	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_0754	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_0874	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_1004	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_0004	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_0124	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_0254	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_0374	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_0504	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_0624	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_0754	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_0874	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_1004	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0

Mean color difference of this page: $\Delta E^{*} = 6.3$

TUB-test chart RE71; 1080 standard colours, $cf=0.9$
input: *rgb/cmyk* -> *rgbd*
output: transfer to *rgbd*
colors and differences, ΔE^{*}

n	HC*Fd	rgb_Fd	ic*_Fd	h*_Fd	rgb*_Fd	LabCh*_Fd	h*_Fd	rgb*_Fd	LabCh*_Fd	DE*Fd	h*_d	rgb*_d	LabCh*_d	DE*Fd	h*_d	rgb*_d	LabCh*_d
1053	NW_086d	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866
1054	NW_093d	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933
1055	NW_100d	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1056	NW_006d	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066
1057	NW_013d	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133
1058	NW_020d	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2
1059	NW_026d	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266
1060	NW_033d	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333
1061	NW_040d	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4
1062	NW_046d	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466
1063	NW_053d	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533
1064	NW_060d	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6
1065	NW_066d	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666
1066	NW_073d	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734
1067	NW_080d	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8
1068	NW_086d	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866
1069	NW_093d	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933
1070	NW_100d	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1071	NW_006d	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066
1072	NW_013d	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133
1073	NW_020d	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2
1074	NW_026d	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266
1075	NW_033d	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333
1076	NW_040d	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4
1077	NW_046d	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466
1078	NW_053d	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533
1079	NW_060d	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6

Mean color difference of this page:

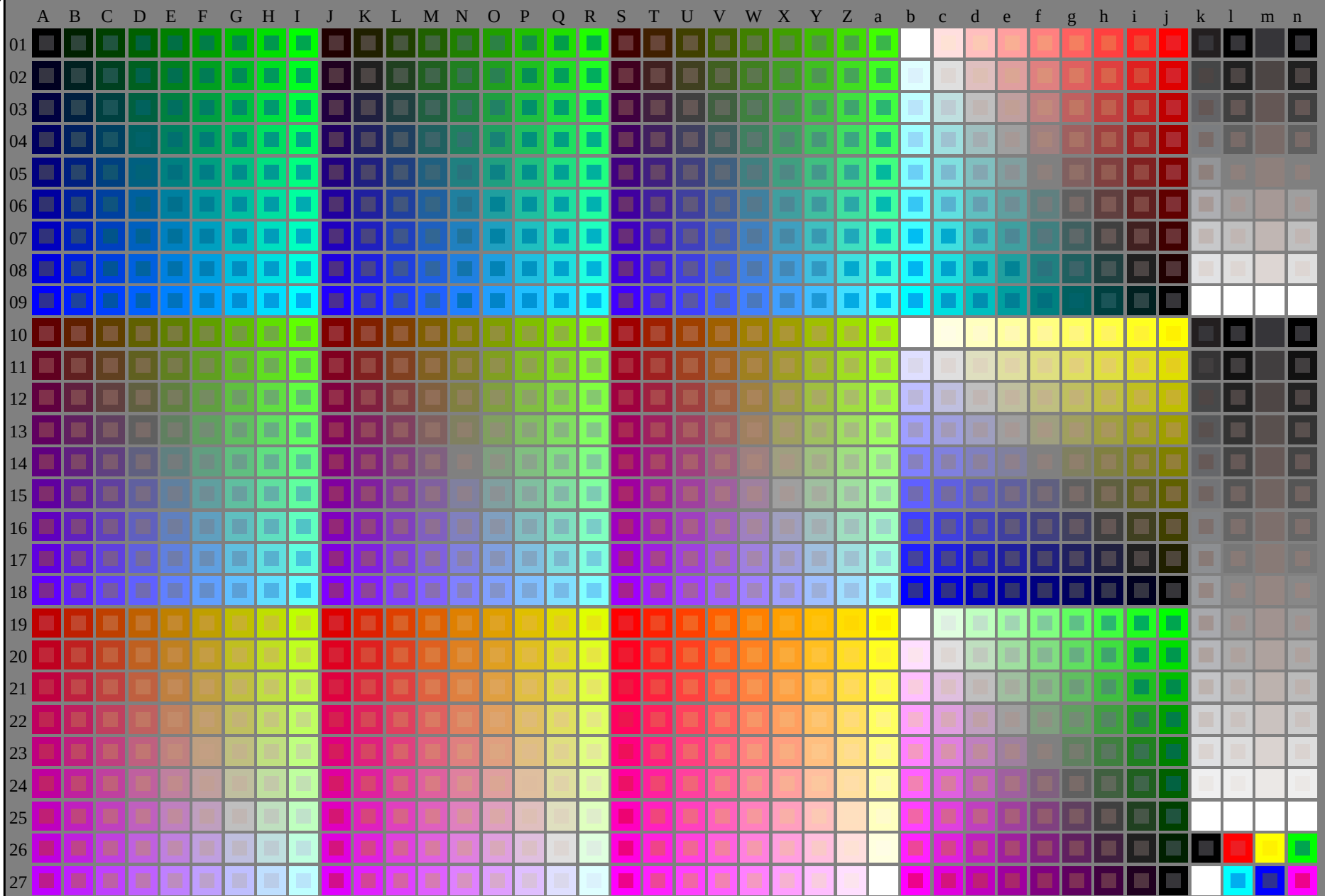
delta E* = 4.4

http://130.149.60.45/~farbmetrik/RE71/RE71L0NA.TXT /.PS; start output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 1/33

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /.PS
application for measurement of laser printer output

TUB material: code=rha4ta



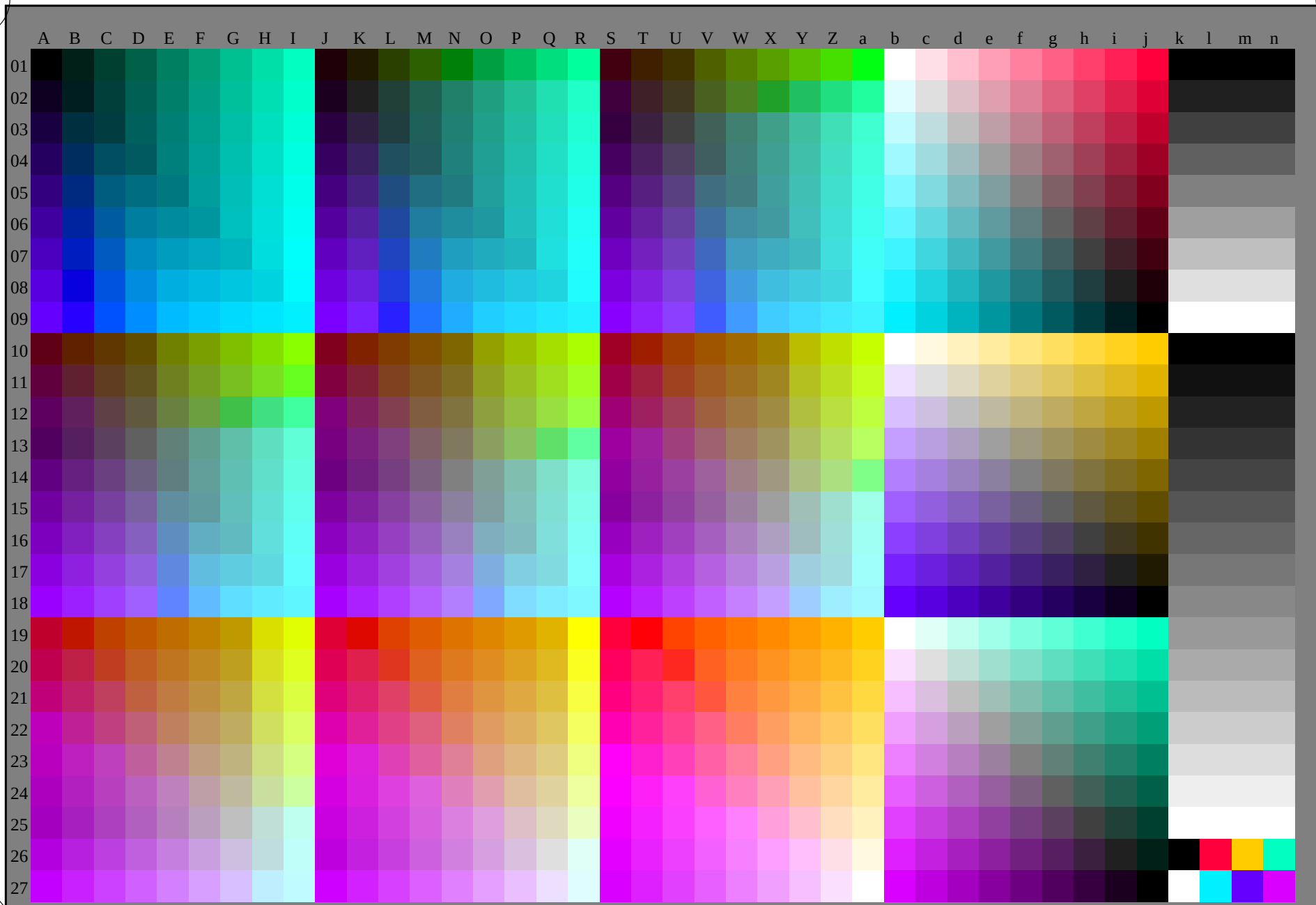
RE710-7N_RGB 1-013034-L0

TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872

input: *rgb/cmyk* -> *rgb/cmyk*
output: no change

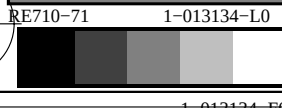
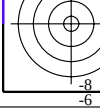
Test chart G with 40x27=1080 colours; equidistant 9 or 16 step colour scales; Colour data in column (A-n): *rgb* (A_j + k26_n27), 000n (k), w (l), nnn0 (m), www (n), 3D = 0

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



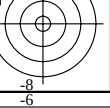
TUB registration: 20150701-RE71/RE71L0NA.TXT /.PS
application for measurement of laser printer output, no separation rgb (RGB)

TUB material: code=rha4ta



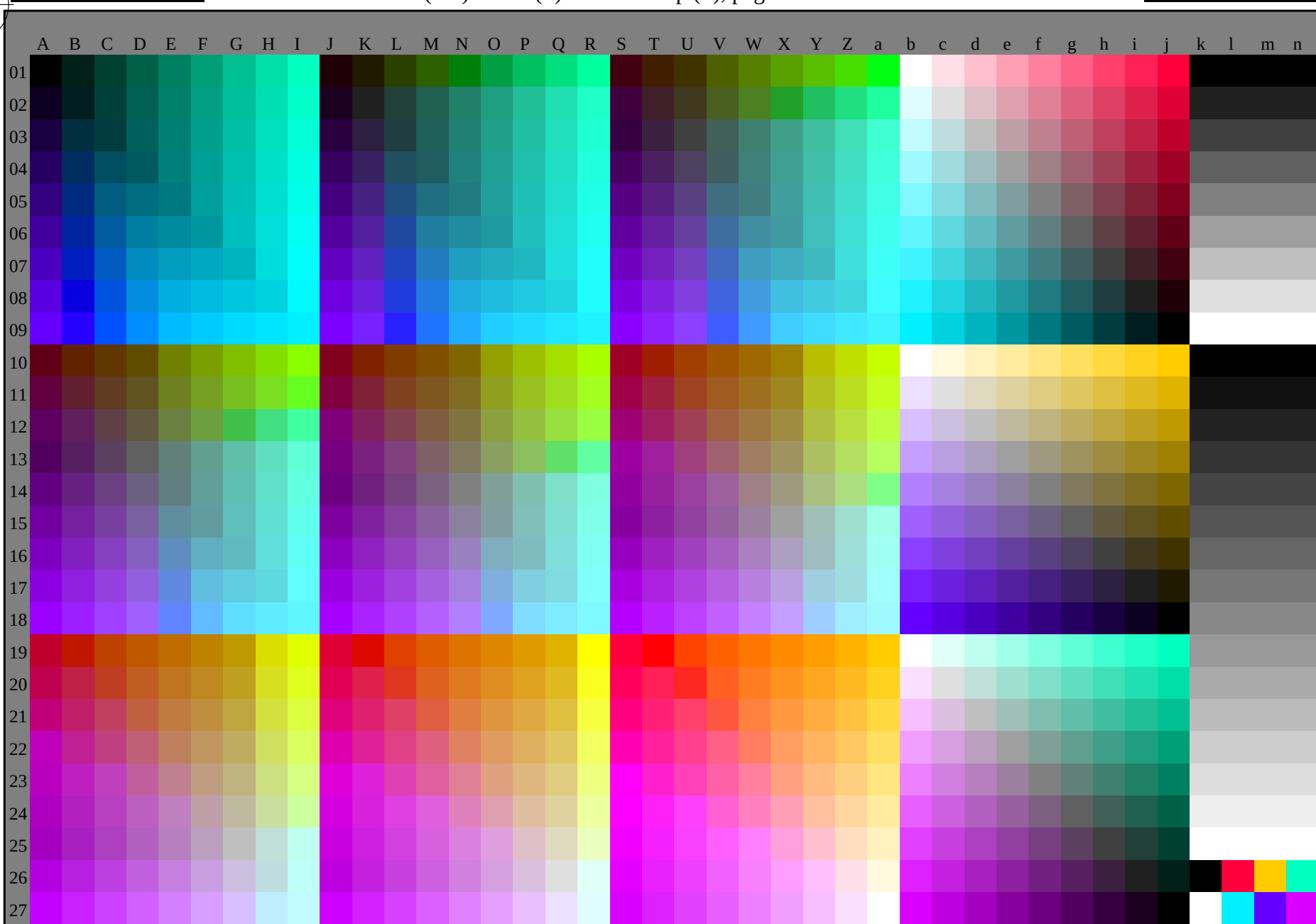
TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872, 3D=0, $de=1$, *rgb*

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



see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /PS
application for measurement of laser printer output, no separation rgb (RGB)
TUB material: code=rha4ta



RE710-71 1-013234-L0

TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872

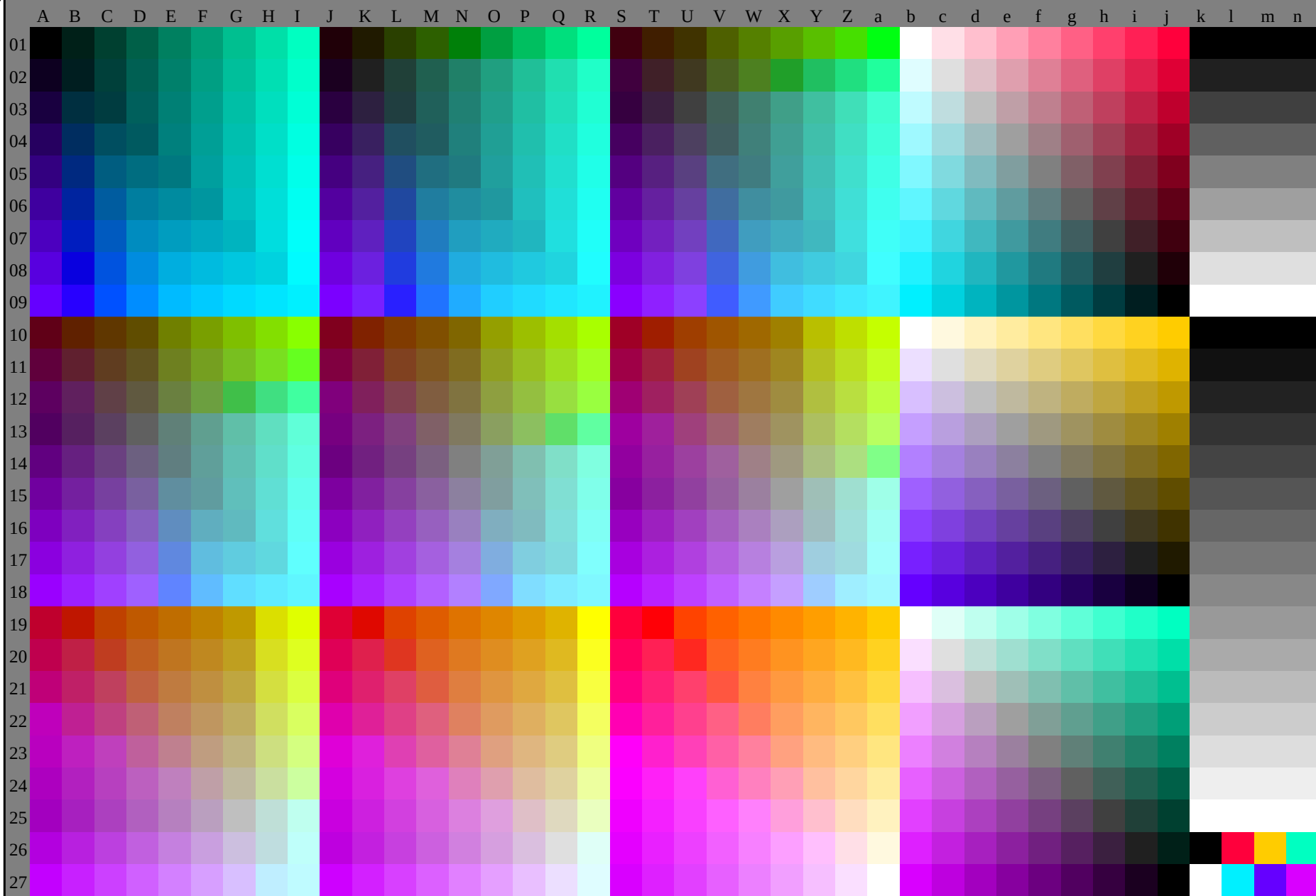
Test chart G with 40x27=1080 colours; equidistant 9 or 16 step colour scales; Colour data in column (A-n);, 3D = 0

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

1-013234-F0

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /PS
application for measurement of laser printer output, no separation rgb (RGB)
TUB material: code=rha4ta



RE710-71 1-013334-L0

TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872

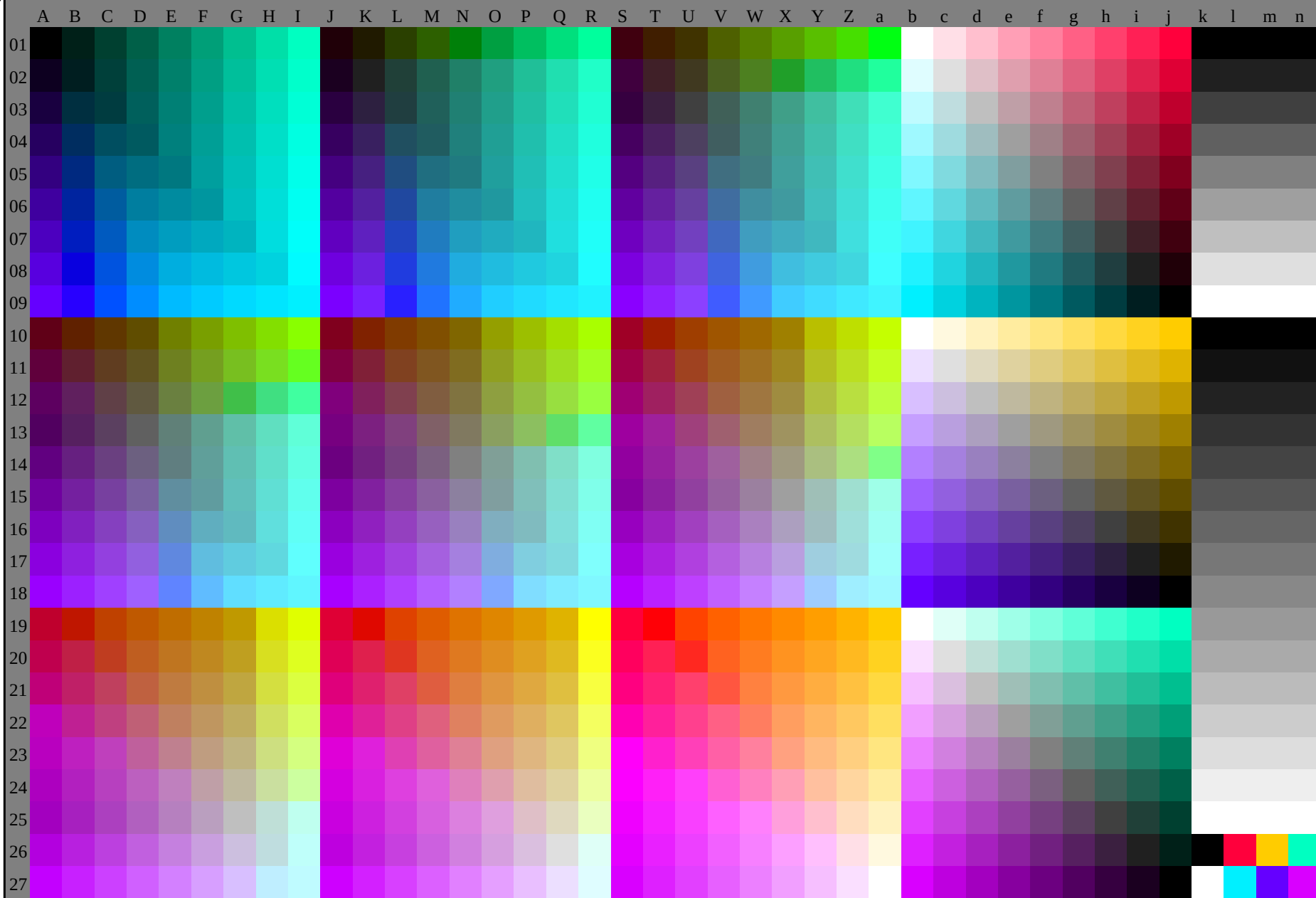
Test chart G with 40x27=1080 colours; equidistant 9 or 16 step colour scales; Colour data in column (A-n); 3D = 0

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

1-013334-F0

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /.PS TUB material: code=rha4ta
application for measurement of laser printer output, no separation rgb (RGB)



RE710-71 1-013434-L0

TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872

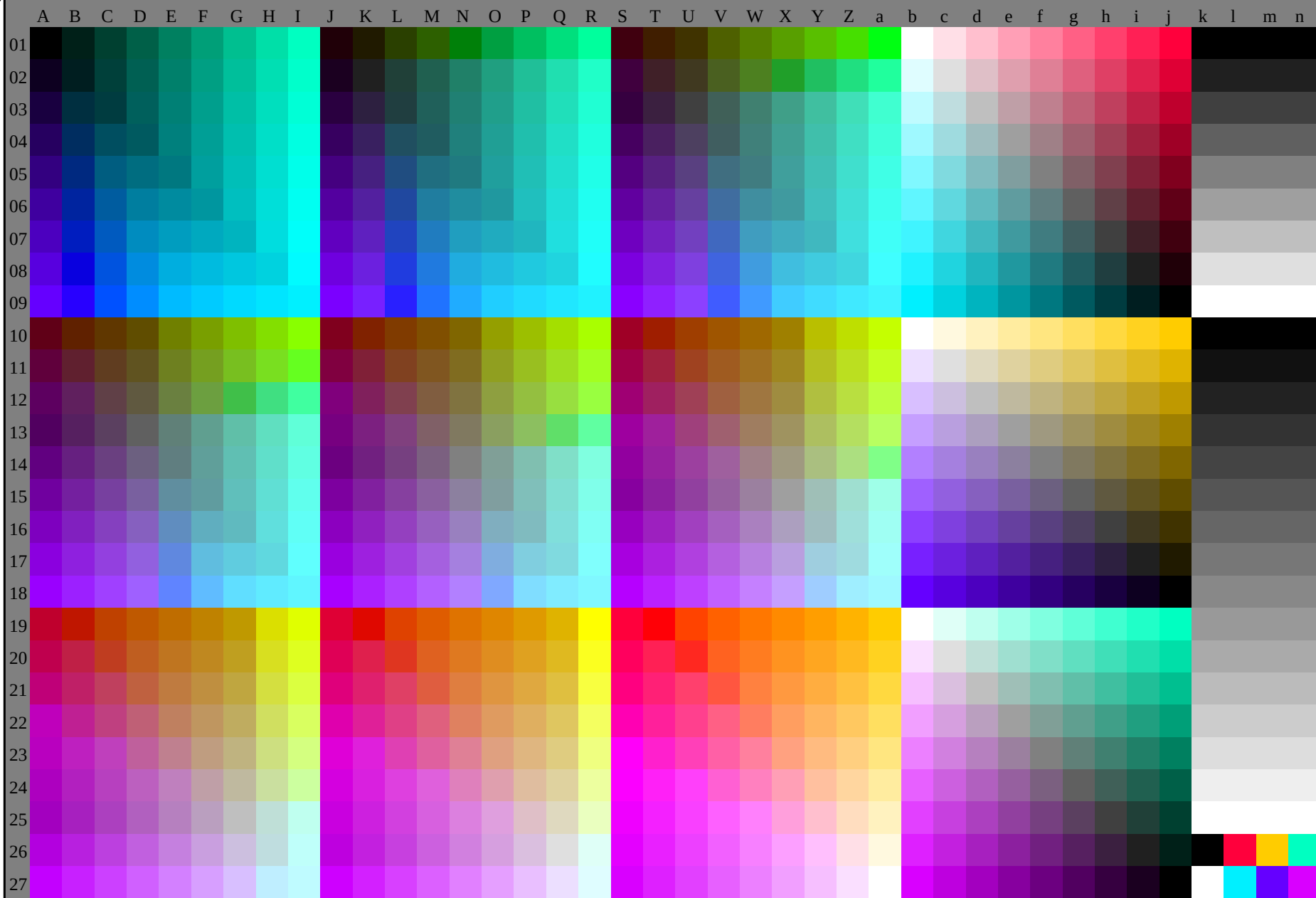
Test chart G with 40x27=1080 colours; equidistant 9 or 16 step colour scales; Colour data in column (A-n);, 3D = 0

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

1-013434-F0

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20150701-RE71/RE71L0NA.TXT /PS
application for measurement of laser printer output, no separation rgb (RGB)
TUB material: code=rha4ta



RE710-71 1-013534-L0

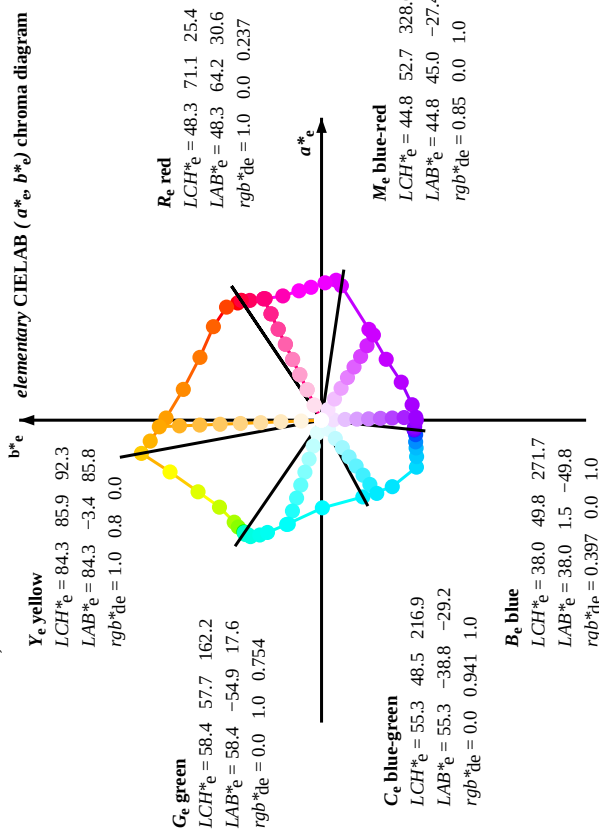
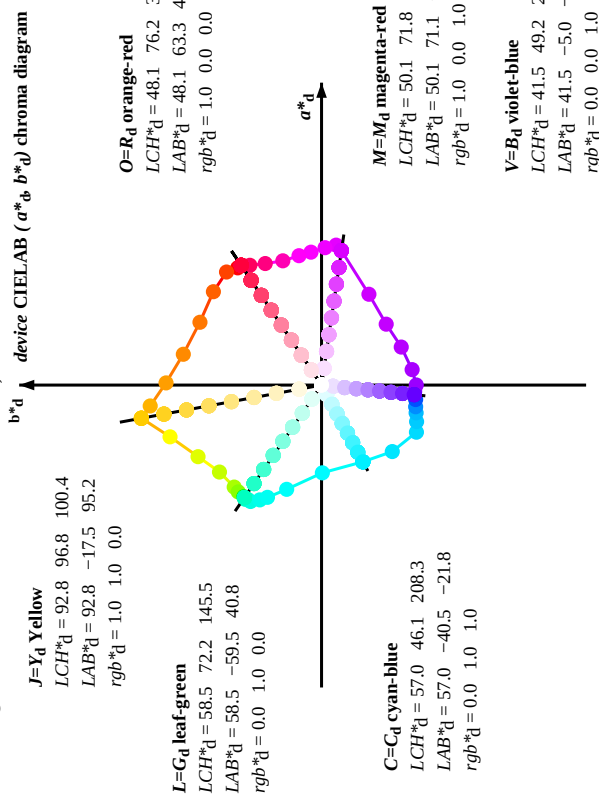
TUB-test chart RE71; 1080 standard colours, $cf=0,9$
Test chart according to DIN 33872

Test chart G with 40x27=1080 colours; equidistant 9 or 16 step colour scales; Colour data in column (A-n);, 3D = 0

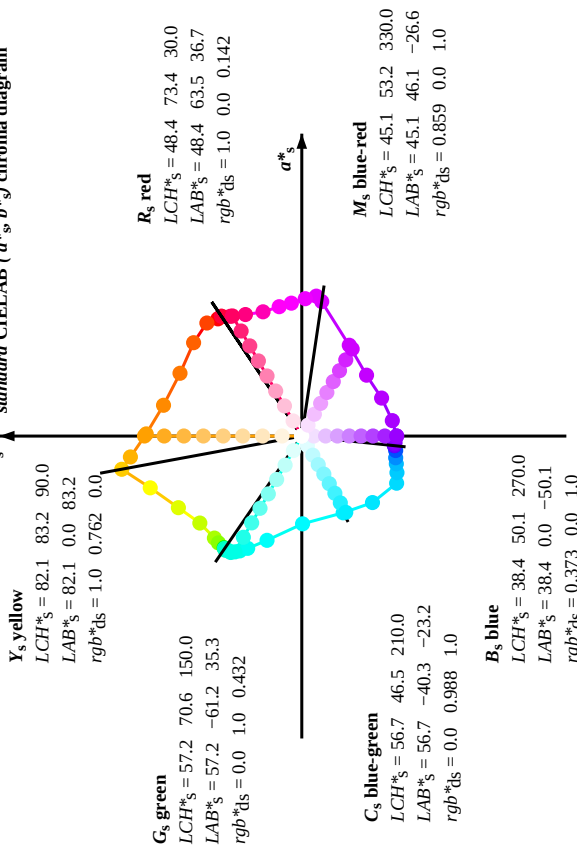
input: $rgb/cmyk \rightarrow rgb_e$
output: transfer to rgb_e

1-013534-F0

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 RYGCMB₆; $h_{ab,d} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours RYGCMB₆; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; Six hue angles of the elementary colours RYGCMB₆; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$



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:

$$h_{ab,s} = \arctan \left[\frac{r^*_d \cos(30) + g^*_d \sin(150)}{1 - r^*_d \sin(30) + g^*_d \sin(270)} \right] \quad (1)$$
- 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: $h_{ab,s} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0, 390.0$ ($i=0,6$) and the equations for a 48 and 360 step hue circle:

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

$$h_{360ab,sij} = h_{ab,si} + j \frac{h_{ab,si+1} - h_{ab,si}}{1/60} \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 59) \quad (3)$$
- 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: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6, 385.5$ ($i=0,6$) and the equations for a 48 and 360 step elementary hue circle:

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

$$h_{360ab,eij} = h_{ab,ei} + j \frac{h_{ab,ei+1} - h_{ab,ei}}{1/60} \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 59) \quad (5)$$
- 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

RE710-71 1-013634-L0

LAB*_{ab}, YN=0%, XY Zmw=2.0, 2.1, 2.1, 85.9, 90.9, 95.1, LAB*_{mw}=15.8, 0.0, 0.0, 96.4, 0.0, 0.0, not adapted=adapted

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

TUB-test chart RE71; 1080 standard colours, $cf=0.9$
48 step hue circles; $rgb-LabCh$ *tables

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

http://130.149.60.45/~farbmetrik/RE71/RE71L0NA.TXT /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 8/33

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*; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$;										Data of Maximum color M in colorimetric system Offset standard print; separation cmyk6*; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$									
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}	rgb^*_{d64M}
33.8	30.0	25.4	1.0	0.0	0.0	48.1	63.3	42.5	76.2	33.8	1.0	0.0	0.0	0.0	0.143	48.5	63.6	36.7	73.4
35.6	37.5	33.8	1.0	0.125	0.0	48.8	62.0	44.3	76.2	35.6	1.0	0.164	0.0	0.0	0.164	49.2	61.4	46.2	76.8
40.0	40.0	42.1	1.0	0.25	0.0	49.9	59.8	50.2	78.1	40.0	1.0	0.318	0.0	0.0	0.318	50.2	54.3	54.3	76.8
49.1	52.5	50.5	1.0	0.375	0.0	55.1	49.4	57.2	75.6	49.1	1.0	0.401	0.0	0.0	0.401	56.9	46.2	59.1	75.0
62.6	60.0	58.8	1.0	0.5	0.0	63.4	33.2	64.2	72.4	62.6	1.0	0.475	0.0	0.0	0.475	61.8	36.6	63.3	73.1
77.4	67.5	67.2	1.0	0.625	0.0	72.5	16.3	73.1	74.9	77.4	1.0	0.617	0.0	0.0	0.617	71.9	17.6	72.7	74.8
89.2	75.0	75.6	1.0	0.75	0.0	81.3	1.1	82.3	82.3	89.2	1.0	0.75	0.0	0.0	0.75	81.3	1.1	82.3	82.3
96.9	82.5	83.9	1.0	0.875	0.0	88.7	-11.0	90.6	91.3	96.9	1.0	0.867	0.0	0.0	0.867	88.3	-10.1	90.2	90.7
100.4	90.0	92.3	1.0	1.0	0.0	92.8	-17.5	95.2	96.8	100.4	1.0	1.0	0.0	0.0	0.929	-17.4	95.3	96.9	96.9
108.8	97.5	101.0	0.875	1.0	0.0	83.7	-27.3	80.1	84.7	108.8	0.883	1.0	0.0	0.0	0.844	-26.8	81.2	85.5	108.8
120.1	105.0	109.7	0.75	1.0	0.0	74.4	-37.9	65.2	75.5	120.1	0.75	1.0	0.0	0.0	0.744	-37.8	65.3	75.5	120.1
130.4	112.5	118.5	0.625	1.0	0.0	67.3	-45.9	53.9	70.9	130.4	0.633	1.0	0.0	0.0	0.678	-45.4	54.8	71.2	129.0
139.3	120.0	127.2	0.5	1.0	0.0	61.7	-53.9	46.2	71.0	139.3	0.5	1.0	0.0	0.0	0.618	-53.8	46.2	71.0	139.3
142.0	127.5	136.0	0.375	1.0	0.0	60.5	-56.5	44.0	71.6	142.0	0.383	1.0	0.0	0.0	0.606	-56.2	44.2	71.6	141.0
145.1	135.0	144.7	0.25	1.0	0.0	58.6	-59.0	41.1	71.9	145.1	0.25	1.0	0.0	0.0	0.587	-58.9	41.1	71.9	145.1
145.5	142.5	153.4	0.125	1.0	0.0	58.5	-59.5	40.8	72.2	145.5	0.133	1.0	0.0	0.0	0.585	-59.4	40.9	72.2	145.5
145.5	150.0	162.2	0.0	1.0	0.0	58.5	-59.5	40.8	72.2	145.5	0.0	1.0	0.0	0.0	0.585	-59.4	40.9	72.2	145.5
146.1	157.5	169.0	0.0	1.0	0.125	57.9	-60.4	40.4	72.7	146.1	0.0	1.0	0.125	0.0	0.579	-60.3	40.5	72.7	146.1
147.2	165.0	175.9	0.0	1.0	0.25	57.6	-60.6	38.9	72.0	147.2	0.0	1.0	0.25	0.0	0.576	-60.5	38.9	72.0	147.2
148.5	172.5	182.7	0.0	1.0	0.375	57.2	-61.5	37.6	72.1	148.5	0.0	1.0	0.375	0.0	0.572	-61.4	37.7	72.1	148.5
151.6	180.0	189.6	0.0	1.0	0.5	57.1	-60.7	37.2	68.9	151.6	0.0	1.0	0.5	0.0	0.571	-60.6	37.7	69.0	151.0
154.2	187.5	196.4	0.0	1.0	0.625	57.3	-59.4	28.6	65.9	154.2	0.0	1.0	0.625	0.0	0.573	-59.4	28.9	66.2	154.0
161.5	195.0	203.2	0.0	1.0	0.75	58.4	-55.1	18.4	58.1	161.5	0.0	1.0	0.75	0.0	0.584	-55.0	18.4	58.1	161.5
180.5	202.5	210.1	0.0	1.0	0.875	59.9	-46.4	-0.4	46.4	180.5	0.0	1.0	0.875	0.0	0.599	-46.4	-0.4	46.4	180.5
208.3	210.0	216.9	0.0	1.0	1.0	57.0	-40.5	-21.8	46.1	208.3	0.0	1.0	1.0	0.0	0.571	-40.5	-21.8	46.1	208.3
226.7	217.5	223.8	0.0	0.875	1.0	53.3	-35.2	-37.3	51.3	226.7	0.0	0.883	1.0	0.0	0.536	-35.7	-36.3	51.0	225.0
243.5	225.0	230.6	0.0	0.75	1.0	52.6	-24.9	-50.1	56.0	243.5	0.0	0.75	1.0	0.0	0.527	-24.8	-50.1	56.0	243.5
248.9	232.5	237.5	0.0	0.625	1.0	49.4	-19.3	-50.3	53.8	248.9	0.0	0.633	1.0	0.0	0.496	-19.6	-50.2	54.0	248.0
253.6	240.0	244.3	0.0	0.5	1.0	47.1	-14.6	-50.0	52.1	253.6	0.0	0.5	1.0	0.0	0.471	-14.6	-50.0	52.2	253.0
256.9	247.5	251.2	0.0	0.375	1.0	45.3	-11.4	-49.7	51.0	256.9	0.0	0.383	1.0	0.0	0.454	-11.6	-49.7	51.1	256.0
261.2	255.0	258.0	0.0	0.25	1.0	42.9	-7.6	-49.7	50.3	261.2	0.0	0.25	1.0	0.0	0.430	-7.6	-49.6	50.3	261.0
264.0	262.5	264.8	0.0	0.125	1.0	41.5	-5.0	-49.0	49.2	264.0	0.0	0.133	1.0	0.0	0.417	-5.1	-49.0	49.4	263.0
264.0	270.0	271.7	0.0	0.0	1.0	41.5	-5.0	-49.0	49.2	264.0	0.0	0.0	1.0	0.0	0.416	-5.0	-48.9	49.3	264.0
265.1	277.5	278.8	0.0	0.125	0.0	40.9	-4.1	-49.0	49.2	265.1	0.0	0.125	0.0	0.0	0.410	-4.2	-49.0	49.3	265.0
266.0	285.0	285.9	0.0	0.25	0.0	40.3	-3.3	-49.3	49.4	266.0	0.0	0.25	0.0	0.0	0.404	-3.3	-49.2	49.4	266.0
270.0	292.5	293.0	0.0	0.375	0.0	38.3	0.0	-50.1	50.1	270.0	0.0	0.375	0.0	0.0	0.385	-0.1	-50.0	50.0	269.0
279.6	300.0	300.1	0.0	0.5	0.0	36.4	8.1	-47.9	48.5	279.6	0.0	0.5	0.0	0.0	0.365	8.1	-48.6	47.9	279.6
295.4	307.5	307.2	0.0	0.625	0.0	37.3	20.1	-42.2	46.7	295.4	0.0	0.625	0.0	0.0	0.373	20.1	-42.2	46.7	295.4
313.1	315.0	314.3	0.0	0.75	0.0	41.4	32.1	-34.2	46.9	313.1	0.0	0.75	0.0	0.0	0.414	32.1	-34.2	47.0	313.1
332.4	322.5	321.4	0.0	0.875	0.0	45.7	48.0	-25.0	54.1	332.4	0.0	0.875	0.0	0.0	0.457	48.0	-25.0	54.1	332.4
351.5	330.0	328.6	1.0	0.0	1.0	50.1	71.1	-10.5	71.8	351.5	1.0	0.0	1.0	0.0	0.502	71.1	-10.4	71.9	351.5
354.0	337.5	335.7	1.0	0.0	0.875	48.7	74.0	-7.7	74.4	354.0	1.0	0.0	0.875	0.0	0.488	73.9	-7.8	74.4	354.0
358.5	345.0	342.8	1.0	0.0	0.75	48.3	72.7	-1.8	72.7	358.5	1.0	0.0	0.75	0.0	0.483	72.7	-1.8	72.7	358.5
364.5	352.5	349.9	1.0	0.0	0.625	48.3	70.3	5.5	70.5	364.5	1.0	0.0	0.625	0.0	0.483	70.3	5.5	70.5	364.5
369.8	360.0	357.0	1.0	0.0	0.5	48.3	68.4	11.9	69.5	369.8	1.0	0.0	0.5	0.0	0.483	68.4	11.9	69.5	369.8
377.3	367.5	364.1	1.0	0.0	0.375	48.4	65.6	20.4	68.8	377.3	1.0	0.0	0.375	0.0	0.484	65.6	20.4	68.8	377.3
384.8	375.0	371.2	1.0	0.0	0.25	48.3	64.2	29.8	70.8	384.8	1.0	0.0	0.25	0.0	0.483	64.2	29.8	70.8	384.8
390.8	382.5	378.3	1.0	0.0	0.125	48.4	63.4	37.8	73.8	390.8	1.0	0.0	0.125	0.0	0.484	63.4	37.8	73.8	390.8
393.8	390.0	385.4	1.0	0.0	0.0	48.1	63.3	42.5	76.2	393.8	1.0	0.0	0.0	0.0	0.481	63.3	42.5	76.3	393.8
RE710-71	1-013734-10																		

no contours
appropriate correction factors
 $0.15, 0.0, 0.0, 0.0, 0.0, 0.0$
input: $rgb/cmyk$ -> rgb_e
output: transfer to rgb_e

TUB-test chart RE71; 1080 standard colours, $cf=0.9$
48 step hue circles; $rgb-LabCh$ *tables

Six hue angles of the device colours RYGBCM _d ; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; Six hue angles of the elementary colours RYGBCM _s ; $h_{ab,s} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$										Six hue angles of the elementary colours RYGBCM _s ; $h_{ab,s} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$												
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,s}$	rgb^* _d 361M	LAB^* _d 361M	rgb^* _d 361MI	LAB^* _d 361MI (x=LabCh)	rgb^* _d 361MI	LAB^* _d 361MI (x=LabCh)	rgb^* _d 361MI	LAB^* _d 361MI (x=LabCh)	rgb^* _s 361M	LAB^* _s 361M	rgb^* _s 361MI	LAB^* _s 361MI (x=LabCh)	rgb^* _s 361MI	LAB^* _s 361MI (x=LabCh)	rgb^* _s 361MI	LAB^* _s 361MI (x=LabCh)	rgb^* _s 361MI	LAB^* _s 361MI (x=LabCh)		
89	75	75	1.0	0.75	0.0	81.3	1.1	82.3	82.3	89	1.0	0.605	0.0	71.1	19.3	72.0	74.6	75	1.0	0.75	0.0	
90	76	76	1.0	0.766	0.0	82.3	-0.3	83.5	83.5	90	1.0	0.613	0.0	71.7	18.1	72.5	74.7	76	1.0	0.767	0.0	
91	77	77	1.0	0.783	0.0	83.3	-1.8	84.7	84.7	91	1.0	0.622	0.0	72.3	16.9	73.0	74.9	77	1.0	0.783	0.0	
92	78	78	1.0	0.8	0.0	84.3	-3.4	85.8	85.9	92	1.0	0.631	0.0	73.0	15.7	73.7	75.3	78	1.0	0.8	0.0	
93	79	80	1.0	0.816	0.0	85.3	-5.0	86.9	87.1	93	1.0	0.642	0.0	73.7	14.5	74.6	76.0	79	1.0	0.817	0.0	
94	80	81	1.0	0.833	0.0	86.2	-6.7	88.0	88.3	94	1.0	0.652	0.0	74.5	13.3	75.4	76.6	80	1.0	0.833	0.0	
95	81	82	1.0	0.85	0.0	87.2	-8.4	89.1	89.5	95	1.0	0.663	0.0	75.2	12.1	76.3	77.2	81	1.0	0.85	0.0	
96	82	83	1.0	0.866	0.0	88.2	-10.1	90.1	90.7	96	1.0	0.674	0.0	76.0	10.8	77.1	77.8	82	1.0	0.867	0.0	
97	83	84	1.0	0.883	0.0	89.0	-11.4	90.9	91.7	97	1.0	0.684	0.0	76.7	9.6	77.9	78.5	83	1.0	0.883	0.0	
97	84	85	1.0	0.9	0.0	89.5	-12.2	91.6	92.4	97	1.0	0.695	0.0	77.5	8.3	78.7	79.1	84	1.0	0.9	0.0	
98	85	86	1.0	0.916	0.0	90.1	-13.1	92.2	93.1	98	1.0	0.705	0.0	78.2	6.9	79.4	79.7	85	1.0	0.917	0.0	
98	86	87	1.0	0.933	0.0	90.6	-14.0	92.8	93.9	98	1.0	0.716	0.0	79.0	5.6	80.1	80.3	86	1.0	0.933	0.0	
99	87	88	1.0	0.95	0.0	91.2	-14.8	93.4	94.6	99	1.0	0.727	0.0	79.7	4.2	80.8	81.0	87	1.0	0.95	0.0	
99	88	90	1.0	0.966	0.0	91.7	-15.7	94.0	95.4	99	1.0	0.737	0.0	80.4	2.8	81.5	81.6	88	1.0	0.967	0.0	
99	89	91	1.0	0.983	0.0	92.3	-16.6	94.6	96.1	99	1.0	0.748	0.0	81.2	1.4	82.2	82.2	89	1.0	0.983	0.0	
100	90	92	1.0	1.0	0.0	92.8	-17.5	95.2	96.8	100	1.0	0.763	0.0	82.1	0.0	83.3	83.3	90	1.0	1.0	0.0	
101	91	93	0.983	1.0	0.0	91.6	-19.0	93.3	95.2	101	1.0	0.779	0.0	83.1	-1.4	84.4	84.4	91	0.983	1.0	0.0	
102	92	94	0.966	1.0	0.0	90.4	-20.5	91.3	93.6	102	1.0	0.795	0.0	84.0	-2.9	85.5	85.6	92	0.967	1.0	0.0	
103	93	95	0.95	1.0	0.0	89.2	-21.9	89.3	92.0	103	1.0	0.811	0.0	85.0	-4.4	86.6	86.7	93	0.95	1.0	0.0	
104	94	96	0.933	1.0	0.0	88.0	-23.2	87.3	90.4	104	1.0	0.827	0.0	85.9	-6.0	87.7	87.9	94	0.933	1.0	0.0	
106	95	98	0.916	1.0	0.0	86.8	-24.5	85.3	88.7	106	1.0	0.844	0.0	86.9	-7.7	88.7	89.1	95	0.917	1.0	0.0	
107	96	99	0.9	1.0	0.0	85.5	-25.7	83.2	87.1	107	1.0	0.86	0.0	87.9	-9.3	89.7	90.2	96	0.9	1.0	0.0	
108	97	100	0.883	1.0	0.0	84.3	-26.8	81.2	85.5	108	1.0	0.877	0.0	88.8	-11.0	90.7	91.4	97	0.883	1.0	0.0	
109	98	101	0.866	1.0	0.0	83.1	-28.2	79.2	84.1	109	1.0	0.913	0.0	90.0	-12.8	92.1	93.0	98	0.867	1.0	0.0	
111	99	102	0.85	1.0	0.0	81.9	-29.8	77.3	82.8	111	1.0	0.949	0.0	91.2	-14.7	93.4	94.6	99	0.85	1.0	0.0	
112	100	103	0.833	1.0	0.0	80.6	-31.4	75.3	81.6	112	1.0	0.985	0.0	92.3	-16.6	94.7	96.2	100	0.833	1.0	0.0	
114	101	105	0.816	1.0	0.0	79.4	-32.8	73.4	80.4	114	1.0	0.992	1.0	0.0	92.2	-18.2	94.3	96.1	101	0.817	1.0	0.0
115	102	106	0.8	1.0	0.0	78.1	-34.2	71.4	79.1	115	1.0	0.977	1.0	0.0	91.2	-19.6	92.6	94.6	102	0.8	1.0	0.0
117	103	107	0.783	1.0	0.0	76.9	-35.5	69.3	77.9	117	1.0	0.962	1.0	0.0	90.1	-20.9	90.8	93.2	103	0.783	1.0	0.0
118	104	108	0.766	1.0	0.0	75.6	-36.7	67.3	76.7	118	1.0	0.947	1.0	0.0	89.0	-22.1	89.0	91.7	104	0.767	1.0	0.0
120	105	109	0.75	1.0	0.0	74.4	-37.9	65.2	75.5	120	1.0	0.932	1.0	0.0	87.9	-23.3	87.2	90.3	105	0.75	1.0	0.0
121	106	110	0.733	1.0	0.0	73.4	-39.1	63.8	74.8	121	1.0	0.917	1.0	0.0	86.9	-24.4	85.4	88.9	106	0.733	1.0	0.0
122	107	112	0.716	1.0	0.0	72.5	-40.3	62.3	74.2	122	1.0	0.903	1.0	0.0	85.8	-25.5	83.6	87.4	107	0.717	1.0	0.0
124	108	113	0.7	1.0	0.0	71.5	-41.4	60.8	73.6	124	1.0	0.888	1.0	0.0	84.7	-26.5	81.8	86.0	108	0.7	1.0	0.0
125	109	114	0.683	1.0	0.0	70.6	-42.5	59.3	73.0	125	1.0	0.873	1.0	0.0	83.7	-27.4	80.0	84.6	109	0.683	1.0	0.0
126	110	115	0.666	1.0	0.0	69.6	-43.5	57.8	72.4	126	1.0	0.862	1.0	0.0	82.8	-28.6	78.7	83.8	110	0.667	1.0	0.0
128	111	116	0.65	1.0	0.0	68.7	-44.5	56.3	71.8	128	1.0	0.851	1.0	0.0	82.0	-29.6	77.5	83.0	111	0.65	1.0	0.0
129	112	117	0.633	1.0	0.0	67.7	-45.5	54.7	71.2	129	1.0	0.84	1.0	0.0	81.2	-30.7	76.2	82.2	112	0.633	1.0	0.0
131	113	119	0.616	1.0	0.0	66.9	-46.5	53.5	70.9	131	1.0	0.829	1.0	0.0	80.3	-31.7	74.9	81.3	113	0.617	1.0	0.0
132	114	120	0.6	1.0	0.0	66.2	-47.5	52.5	70.9	132	1.0	0.818	1.0	0.0	79.5	-32.7	73.6	80.5	114	0.6	1.0	0.0
133	115	121	0.583	1.0	0.0	65.4	-48.7	51.5	70.9	133	1.0	0.807	1.0	0.0	78.7	-33.6	72.2	79.7	115	0.583	1.0	0.0
134	116	122	0.566	1.0	0.0	64.7	-49.8	50.5	70.9	134	1.0	0.796	1.0	0.0	77.9	-34.5	70.9	78.9	116	0.567	1.0	0.0
135	117	123	0.55	1.0	0.0	63.9	-50.8	49.4	70.9	135	1.0	0.785	1.0	0.0	77.0	-35.3	69.6	78.1	117	0.55	1.0	0.0
136	118	124	0.533	1.0	0.0	63.2	-51.9	48.4	71.0	136	1.0	0.774	1.0	0.0	76.2	-36.2	68.2	77.3	118	0.533	1.0	0.0
138	119	126	0.516	1.0	0.0	62.5	-52.9	47.3	71.0	138	1.0	0.763	1.0	0.0	75.4	-37.0	66.8	76.4	119	0.517	1.0	0.0
139	120	127	0.5	1.0	0.0	61.7	-53.9	46.2	71.0	139	1.0	0.752	1.0	0.0	74.5	-37.7	65.5	75.6	120	0.5	1.0	0.0
LAB* _a 0, YN=0%, XYZ _{mnw} =2.0, 2.1, 2.1, 85.9, 90.9, 95.1, LAB* _{mnw} =15.8, 0.0, 0.0, 96.4, 0.0, 0.0, not adapted-adapted																				Output: Offset standard print; separation cmy ₆ *; D65		
TUB-test chart RE71; 1080 standard colours, $cf=0.9$																				input: $rgb/cmyk \rightarrow rgb_e$		
48 step hue circles: $rgb-LabCh$ *tables																				output: transfer to rgb_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 RYGCBM; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; 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^*_s361M	$rgb^*_ds361Mi$	$rgb^*_ds361Mi$	$rgb^*_ds361Mi$	$rgb^*_ds361Mi$
139	120	127	0.5	1.0	0.0	61.7	-53.9	46.2	71.0
139	121	128	0.0	61.5	-54.2	45.9	71.1	139	139
140	122	129	0.466	1.0	0.0	61.4	-54.6	45.6	71.2
140	123	130	0.45	1.0	0.0	61.2	-54.9	45.4	71.2
140	124	131	0.433	1.0	0.0	61.0	-55.3	45.1	71.3
141	125	133	0.416	1.0	0.0	60.9	-55.6	44.8	71.4
141	126	134	0.4	1.0	0.0	60.7	-56.0	44.5	71.5
141	127	135	0.383	1.0	0.0	60.5	-56.3	44.2	71.6
142	128	136	0.366	1.0	0.0	60.3	-56.6	43.9	71.6
142	129	137	0.35	1.0	0.0	60.1	-57.0	43.5	71.7
143	130	138	0.333	1.0	0.0	59.8	-57.3	43.1	71.7
143	131	140	0.316	1.0	0.0	59.6	-57.7	42.7	71.8
143	132	141	0.3	1.0	0.0	59.3	-58.0	42.3	71.8
144	133	142	0.283	1.0	0.0	59.1	-58.3	41.9	71.8
144	134	143	0.266	1.0	0.0	58.9	-58.6	41.5	71.9
145	135	144	0.25	1.0	0.0	58.6	-59.0	41.1	71.9
145	136	145	0.233	1.0	0.0	58.6	-59.0	41.0	71.9
145	137	147	0.216	1.0	0.0	58.6	-59.1	41.0	72.0
145	138	148	0.2	1.0	0.0	58.5	-59.2	41.0	72.0
145	139	149	0.183	1.0	0.0	58.5	-59.3	40.9	72.0
145	140	150	0.166	1.0	0.0	58.5	-59.3	40.9	72.1
145	141	151	0.15	1.0	0.0	58.5	-59.4	40.9	72.1
145	142	152	0.133	1.0	0.0	58.5	-59.5	40.8	72.2
145	143	154	0.116	1.0	0.0	58.5	-59.5	40.8	72.2
145	144	155	0.1	1.0	0.0	58.5	-59.5	40.8	72.2
145	145	156	0.083	1.0	0.0	58.5	-59.5	40.8	72.2
145	146	157	0.066	1.0	0.0	58.5	-59.5	40.8	72.2
145	147	158	0.049	1.0	0.0	58.5	-59.5	40.8	72.2
145	148	159	0.033	1.0	0.0	58.5	-59.5	40.8	72.2
145	149	161	0.016	1.0	0.0	58.5	-59.5	40.8	72.2
145	150	162	0.0	1.0	0.0	58.5	-59.5	40.8	72.2
145	151	163	0.0	1.0	0.016	58.4	-59.6	40.8	72.2
145	152	164	0.0	1.0	0.033	58.3	-59.7	40.7	72.3
145	153	164	0.0	1.0	0.05	58.2	-59.9	40.7	72.4
145	154	165	0.0	1.0	0.066	58.2	-60.0	40.6	72.4
145	155	166	0.0	1.0	0.083	58.1	-60.1	40.5	72.5
146	156	167	0.0	1.0	0.1	58.0	-60.2	40.5	72.6
146	157	168	0.0	1.0	0.116	58.0	-60.3	40.4	72.6
146	158	169	0.0	1.0	0.133	57.9	-60.4	40.3	72.6
146	159	170	0.0	1.0	0.15	57.9	-60.4	40.1	72.5
146	160	171	0.0	1.0	0.166	57.8	-60.4	39.9	72.4
146	161	172	0.0	1.0	0.183	57.8	-60.5	39.7	72.4
146	162	173	0.0	1.0	0.2	57.7	-60.5	39.5	72.3
146	163	174	0.0	1.0	0.216	57.7	-60.5	39.3	72.2
146	164	175	0.0	1.0	0.233	57.6	-60.5	39.1	72.1
147	165	175	0.0	1.0	0.25	57.6	-60.6	38.9	72.0
147	166	175	0.0	1.0	0.27	57.6	-60.6	38.9	72.0
147	167	175	0.0	1.0	0.29	57.6	-60.6	38.9	72.0
147	168	175	0.0	1.0	0.31	57.6	-60.6	38.9	72.0
147	169	175	0.0	1.0	0.33	57.6	-60.6	38.9	72.0
147	170	175	0.0	1.0	0.35	57.6	-60.6	38.9	72.0
147	171	175	0.0	1.0	0.37	57.6	-60.6	38.9	72.0
147	172	175	0.0	1.0	0.39	57.6	-60.6	38.9	72.0
147	173	175	0.0	1.0	0.41	57.6	-60.6	38.9	72.0
147	174	175	0.0	1.0	0.43	57.6	-60.6	38.9	72.0
147	175	175	0.0	1.0	0.45	57.6	-60.6	38.9	72.0
147	176	175	0.0	1.0	0.47	57.6	-60.6	38.9	72.0
147	177	175	0.0	1.0	0.49	57.6	-60.6	38.9	72.0
147	178	175	0.0	1.0	0.51	57.6	-60.6	38.9	72.0
147	179	175	0.0	1.0	0.53	57.6	-60.6	38.9	72.0
147	180	175	0.0	1.0	0.55	57.6	-60.6	38.9	72.0
147	181	175	0.0	1.0	0.57	57.6	-60.6	38.9	72.0
147	182	175	0.0	1.0	0.59	57.6	-60.6	38.9	72.0
147	183	175	0.0	1.0	0.61	57.6	-60.6	38.9	72.0
147	184	175	0.0	1.0	0.63	57.6	-60.6	38.9	72.0
147	185	175	0.0	1.0	0.65	57.6	-60.6	38.9	72.0
147	186	175	0.0	1.0	0.67	57.6	-60.6	38.9	72.0
147	187	175	0.0	1.0	0.69	57.6	-60.6	38.9	72.0
147	188	175	0.0	1.0	0.71	57.6	-60.6	38.9	72.0
147	189	175	0.0	1.0	0.73	57.6	-60.6	38.9	72.0
147	190	175	0.0	1.0	0.75	57.6	-60.6	38.9	72.0
147	191	175	0.0	1.0	0.77	57.6	-60.6	38.9	72.0
147	192	175	0.0	1.0	0.79	57.6	-60.6	38.9	72.0
147	193	175	0.0	1.0	0.81	57.6	-60.6	38.9	72.0
147	194	175	0.0	1.0	0.83	57.6	-60.6	38.9	72.0
147	195	175	0.0	1.0	0.85	57.6	-60.6	38.9	72.0
147	196	175	0.0	1.0	0.87	57.6	-60.6	38.9	72.0
147	197	175	0.0	1.0	0.89	57.6	-60.6	38.9	72.0
147	198	175	0.0	1.0	0.91	57.6	-60.6	38.9	72.0
147	199	175	0.0	1.0	0.93	57.6	-60.6	38.9	72.0
147	200	175	0.0	1.0	0.95	57.6	-60.6	38.9	72.0
147	201	175	0.0	1.0	0.97	57.6	-60.6	38.9	72.0
147	202	175	0.0	1.0	0.99	57.6	-60.6	38.9	72.0
147	203	175	0.0	1.0	1.01	57.6	-60.6	38.9	72.0
147	204	175	0.0	1.0	1.03	57.6	-60.6	38.9	72.0
147	205	175	0.0	1.0	1.05	57.6	-60.6	38.9	72.0
147	206	175	0.0	1.0	1.07	57.6	-60.6	38.9	72.0
147	207	175	0.0	1.0	1.09	57.6	-60.6	38.9	72.0
147	208	175	0.0	1.0	1.11	57.6	-60.6	38.9	72.0
147	209	175	0.0	1.0	1.13	57.6	-60.6	38.9	72.0
147	210	175	0.0	1.0	1.15	57.6	-60.6	38.9	72.0
147	211	175	0.0	1.0	1.17	57.6	-60.6	38.9	72.0
147	212	175	0.0	1.0	1.19	57.6	-60.6	38.9	72.0
147	213	175	0.0	1.0	1.21	57.6	-60.6	38.9	72.0
147	214	175	0.0	1.0	1.23	57.6	-60.6	38.9	72.0
147	215	175	0.0	1.0	1.25	57.6	-60.6	38.9	72.0
147	216	175	0.0	1.0	1.27	57.6	-60.6	38.9	72.0
147	217	175	0.0	1.0	1.29	57.6	-60.6	38.9	72.0
147	218	175	0.0	1.0	1.31	57.6	-60.6	38.9	72.0
147	219	175	0.0	1.0	1.33	57.6	-60.6	38.9	72.0
147	220	175	0.0	1.0	1.35	57.6	-60.6	38.9	72.0
147	221	175	0.0	1.0	1.37	57.6	-60.6	38.9	72.0
147	222	175	0.0	1.0	1.39	57.6	-60.6	38.9	72.0
147	223	175	0.0	1.0	1.41	57.6	-60.6	38.9	72.0
147	224	175	0.0	1.0	1.43	57.6	-60.6	38.9	72.0
147	225	175	0.0	1.0	1.45	57.6	-60.6	38.9	72.0
147	226	175	0.0	1.0	1.47	57.6	-60.6	38.9	72.0
147	227	175	0.0	1.0	1.49	57.6	-60.6	38.9	72.0
147	228	175	0.0	1.0	1.51	57.6	-60.6	38.9	72.0
147	229	175	0.0	1.0	1.53	57.6	-60.6	38.9	72.0
147	230	175	0.0	1.0	1.55	57.6	-60.6	38.9	72.0
147	231	175	0.0	1.0	1.57	57.6	-60.6	38.9	72.0
147	232	175	0.0						

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

<http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT> /PS; transfer output N; no 3D-linearization (OL) in file (F) or PS-startup (S), page 13/33

TUTUB-test chart RE71; 1080 standard colours, $c_f=0,9$
 48 step hue circles; *rab-LabCh** tables
 input: *rgb/cmyk* $\rightarrow$ *rgb*
 output: transfer to *rab*₀

Data of Maximum color M in colorimetric system Offset standard print; separation cmyk0*, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBCM; h _{ab,s} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0; LAB* _{ab,s} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6										
h _{ab,d} h _{ab,s} h _{ab,e}		LAB* _{d361MI}		LAB* _{s361MI}		LAB* _{e361MI}		LAB* _{361MI}		
h _{ab,d}	h _{ab,s}	h _{ab,e}	L*	a*	b*	L*	a*	b*	L*	
147	165	175	0.0	1.0	0.773 58.7	-54.0 14.5	56.0 165	0.0	1.0	
147	147	165	0.0	1.0	0.773 58.7	-54.0 14.5	56.0 165	0.0	1.0	
147	166	176	0.0	1.0	0.779 58.8	-53.6 13.4	55.4 166	0.0	1.0	
147	167	177	0.0	1.0	0.786 58.9	-53.3 12.3	54.8 167	0.0	1.0	
147	168	178	0.0	1.0	0.793 58.9	-52.9 11.3	54.2 168	0.0	1.0	
147	169	179	0.0	1.0	0.799 59.0	-52.5 10.2	53.5 169	0.0	1.0	
148	170	180	0.0	1.0	0.806 59.1	-52.0 9.2	52.9 170	0.0	1.0	
148	171	181	0.0	1.0	0.812 59.2	-51.6 8.2	52.3 171	0.0	1.0	
148	172	182	0.0	1.0	0.819 59.3	-51.1 7.2	51.7 172	0.0	1.0	
148	173	183	0.0	1.0	0.825 59.3	-50.6 6.2	51.1 173	0.0	1.0	
149	174	184	0.0	1.0	0.832 59.4	-50.1 5.3	50.5 174	0.0	1.0	
149	175	185	0.0	1.0	0.839 59.5	-49.6 4.3	49.8 175	0.0	1.0	
150	176	185	0.0	1.0	0.845 59.6	-49.0 3.4	49.2 176	0.0	1.0	
150	177	186	0.0	1.0	0.852 59.6	-48.4 2.5	48.6 177	0.0	1.0	
150	178	187	0.0	1.0	0.858 59.7	-47.9 1.7	48.0 178	0.0	1.0	
151	179	188	0.0	1.0	0.865 59.8	-47.3 0.8	47.4 179	0.0	1.0	
151	180	189	0.0	1.0	0.871 59.9	-46.7 0.0	46.8 180	0.0	1.0	
152	181	190	0.0	1.0	0.877 59.9	-46.3 -0.7	46.4 181	0.0	1.0	
152	182	191	0.0	1.0	0.882 59.8	-46.3 -1.5	46.4 182	0.0	1.0	
152	183	192	0.0	1.0	0.886 59.7	-46.2 -2.3	46.4 183	0.0	1.0	
153	184	193	0.0	1.0	0.891 59.6	-46.2 -3.1	46.4 184	0.0	1.0	
153	185	194	0.0	1.0	0.895 59.5	-46.1 -3.9	46.4 185	0.0	1.0	
153	186	195	0.0	1.0	0.9	59.4	-46.0 -4.7	46.4 186	0.0	1.0
154	187	195	0.0	1.0	0.904 59.3	-45.9 -5.5	46.3 187	0.0	1.0	
154	188	196	0.0	1.0	0.909 59.2	-45.8 -6.3	46.3 188	0.0	1.0	
155	189	197	0.0	1.0	0.913 59.1	-45.7 -7.1	46.3 189	0.0	1.0	
156	190	198	0.0	1.0	0.918 59.0	-45.5 -7.9	46.3 190	0.0	1.0	
157	191	199	0.0	1.0	0.922 58.9	-45.4 -8.7	46.3 191	0.0	1.0	
158	192	200	0.0	1.0	0.926 58.8	-45.2 -9.5	46.3 192	0.0	1.0	
158	193	201	0.0	1.0	0.931 58.7	-45.0 -10.3	46.3 193	0.0	1.0	
160	194	202	0.0	1.0	0.935 58.6	-44.8 -11.1	46.3 194	0.0	1.0	
161	195	203	0.0	1.0	0.94	58.5	-44.6 -11.9	46.3 195	0.0	1.0
164	196	204	0.0	1.0	0.944 58.4	-44.4 -12.6	46.2 196	0.0	1.0	
166	197	205	0.0	1.0	0.949 58.3	-44.1 -13.4	46.2 197	0.0	1.0	
169	198	206	0.0	1.0	0.953 58.2	-43.9 -14.2	46.2 198	0.0	1.0	
171	199	206	0.0	1.0	0.958 58.0	-43.6 -14.9	46.2 199	0.0	1.0	
174	200	207	0.0	1.0	0.962 57.9	-43.3 -15.7	46.2 200	0.0	1.0	
176	201	208	0.0	1.0	0.967 57.8	-43.0 -16.5	46.2 201	0.0	1.0	
179	202	209	0.0	1.0	0.971 57.7	-42.7 -17.2	46.2 202	0.0	1.0	
182	203	210	0.0	1.0	0.976 57.6	-42.4 -17.9	46.2 203	0.0	1.0	
186	204	211	0.0	1.0	0.98	57.5	-42.1 -18.7	46.2 204	0.0	1.0
189	205	212	0.0	1.0	0.985 57.4	-41.7 -19.4	46.1 205	0.0	1.0	
193	206	213	0.0	1.0	0.989 57.3	-41.4 -20.1	46.1 206	0.0	1.0	
197	207	214	0.0	1.0	0.994 57.2	-41.0 -20.8	46.1 207	0.0	1.0	
200	208	215	0.0	1.0	0.998 57.1	-40.6 -21.5	46.1 208	0.0	1.0	
204	209	216	0.0	1.0	0.996 1.0	-40.4 -22.3	46.3 209	0.0	1.0	
208	210	216	0.0	1.0	0.989 1.0	-38.7 -29.1	48.6 210	0.0	1.0	
210	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0 28.3	48.3 216	0.0	1.0
216	216	216	0.0	1.0	0.941 1.0	55.3	-39.0			

<http://130.149.60.45/~farbmetrik/RE71/RE71L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 14/33

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 $RYGCBM_d$; $h_{ab,d} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours $RYGCBM_d$; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; Six hue angles of the elementary colours $RYGCBM_d$; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

$h_{ab,d}$		$h_{ab,s}$	$h_{ab,e}$	rgb^*_{d361M}	LAB^*_{d361M}	$RGB^*_{d361M}(x=LabCh)$	$LAB^*_{d361M}(x=LabCh)$	rgb^*_{d361M}	rgb^*_{d361M}	LAB^*_{d361M}	LAB^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	rgb^*_{d361M}	$rgb^$
------------	--	------------	------------	-----------------	-----------------	--------------------------	--------------------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	-----------------	--------

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; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; Six hue angles of the elementary colours RYGBCM; $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$									
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=L, aCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}(x=L, aCh)$	$rgb^*_{ds361MI}$	$LAB^*_{ds361MI}$	$rgb^*_{ds361MI}$
261	255	258	0.0	0.25	1.0	42.9	-7.6	-49.7	50.3
261	256	258	0.0	0.233	1.0	42.7	-7.3	-49.6	50.1
261	257	259	0.0	0.216	1.0	42.5	-6.9	-49.5	50.0
262	258	260	0.0	0.2	1.0	42.4	-6.6	-49.4	49.9
262	259	261	0.0	0.183	1.0	42.2	-6.2	-49.3	49.7
263	260	262	0.0	0.166	1.0	42.0	-5.9	-49.2	49.6
263	261	263	0.0	0.15	1.0	41.8	-5.5	-49.1	49.5
263	262	264	0.0	0.133	1.0	41.6	-5.2	-49.0	49.3
264	263	265	0.0	0.116	1.0	41.5	-5.0	-49.0	49.2
264	264	266	0.0	0.1	1.0	41.5	-5.0	-49.0	49.2
264	265	267	0.0	0.083	1.0	41.5	-5.0	-49.0	49.2
264	266	268	0.0	0.066	1.0	41.5	-5.0	-49.0	49.2
264	267	269	0.0	0.049	1.0	41.5	-5.0	-49.0	49.2
264	268	269	0.0	0.033	1.0	41.5	-5.0	-49.0	49.2
264	269	270	0.0	0.016	1.0	41.5	-5.0	-49.0	49.2
264	270	271	0.0	0.0	1.0	41.5	-5.0	-49.0	49.2
264	271	272	0.016	0.0	1.0	41.4	-4.9	-49.0	49.2
264	272	273	0.033	0.0	1.0	41.4	-4.8	-49.0	49.2
264	273	274	0.05	0.0	1.0	41.3	-4.7	-49.0	49.2
264	274	275	0.066	0.0	1.0	41.2	-4.6	-49.0	49.2
264	275	276	0.083	0.0	1.0	41.1	-4.4	-49.0	49.2
264	276	277	0.1	0.0	1.0	41.0	-4.3	-49.0	49.2
265	277	278	0.116	0.0	1.0	40.9	-4.2	-49.0	49.2
265	278	279	0.133	0.0	1.0	40.9	-4.1	-49.1	49.2
265	279	280	0.15	0.0	1.0	40.8	-4.0	-49.1	49.3
265	280	281	0.166	0.0	1.0	40.7	-3.9	-49.1	49.3
265	281	282	0.183	0.0	1.0	40.6	-3.8	-49.2	49.3
265	282	283	0.2	0.0	1.0	40.5	-3.7	-49.2	49.3
265	283	284	0.216	0.0	1.0	40.5	-3.5	-49.2	49.4
265	284	285	0.233	0.0	1.0	40.4	-3.4	-49.3	49.4
266	285	285	0.25	0.0	1.0	40.3	-3.3	-49.3	49.4
266	286	286	0.266	0.0	1.0	40.0	-2.9	-49.4	49.5
267	287	287	0.283	0.0	1.0	39.8	-2.4	-49.5	49.6
267	288	288	0.3	0.0	1.0	39.5	-2.0	-49.6	49.7
268	289	289	0.316	0.0	1.0	39.3	-1.5	-49.8	49.8
268	290	290	0.333	0.0	1.0	39.0	-1.1	-49.9	49.9
269	291	291	0.35	0.0	1.0	38.7	-0.6	-50.0	50.0
269	292	292	0.366	0.0	1.0	38.5	-0.1	-50.1	50.1
270	293	293	0.383	0.0	1.0	38.2	0.6	-50.0	50.0
271	294	294	0.4	0.0	1.0	38.0	1.7	-49.8	49.8
273	295	295	0.416	0.0	1.0	37.7	2.8	-49.5	49.6
274	296	296	0.433	0.0	1.0	37.4	3.8	-49.2	49.4
275	297	297	0.45	0.0	1.0	37.2	4.9	-49.0	49.2
277	298	298	0.466	0.0	1.0	36.9	6.0	-48.6	49.0
278	299	299	0.483	0.0	1.0	36.7	7.0	-48.2	48.8
279	300	300	0.5	0.0	1.0	36.4	8.1	-47.9	48.5
279	300	300	0.5	0.0	1.0	36.4	8.1	-47.9	48.5

TUB-test chart RE71; 1080 standard colours, $cf=0.9$
48 step hue circles; $rgb-LabCh$ *tables

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

Output: Offset standard print; separation cmyk6*, D65, page 15/38

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; $h_{ab,d65} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;									
Six hue angles of the device colours RYGBCM; $h_{ab,d} = 33.9, 100.4, 145.5, 208.3, 264.1, 351.6$; 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^*_{d361M}	rgb^*_{d361Mi}	$LAB^*_{d361Mi} (x=LabCh)$	rgb^*_{d361Mi}	$LAB^*_{d361Mi} (x=LabCh)$	rgb^*_{d361Mi}	$LAB^*_{d361Mi} (x=LabCh)$
279	300	300	0.5	0.0	1.0	36.4	8.1	-47.9	48.5
281	301	301	0.516	0.0	1.0	36.5	9.8	-47.3	48.3
283	302	302	0.533	0.0	1.0	36.6	11.5	-46.7	48.1
285	303	303	0.55	0.0	1.0	36.8	13.1	-46.0	47.8
288	304	304	0.566	0.0	1.0	36.9	14.7	-45.2	47.6
290	305	304	0.583	0.0	1.0	37.0	16.3	-44.4	47.3
292	306	305	0.6	0.0	1.0	37.1	17.8	-43.6	47.1
294	307	306	0.616	0.0	1.0	37.2	19.3	-42.6	46.8
296	308	307	0.633	0.0	1.0	37.5	20.9	-41.8	46.7
299	309	308	0.65	0.0	1.0	38.1	22.6	-40.9	46.8
301	310	309	0.666	0.0	1.0	38.6	24.3	-39.9	46.8
303	311	310	0.683	0.0	1.0	39.2	26.0	-38.9	46.8
306	312	311	0.7	0.0	1.0	39.7	27.6	-37.8	46.8
308	313	312	0.716	0.0	1.0	40.3	29.1	-36.7	46.9
310	314	313	0.733	0.0	1.0	40.8	30.6	-35.5	46.9
313	315	314	0.75	0.0	1.0	41.4	32.1	-34.2	46.9
315	316	315	0.766	0.0	1.0	42.0	34.3	-33.4	47.9
318	317	316	0.783	0.0	1.0	42.5	36.5	-32.5	48.9
320	318	317	0.8	0.0	1.0	43.1	38.6	-31.4	49.8
323	319	318	0.816	0.0	1.0	43.7	40.8	-30.2	50.8
326	320	319	0.833	0.0	1.0	44.3	42.9	-28.9	51.7
328	321	320	0.85	0.0	1.0	44.8	45.0	-27.4	52.7
331	322	321	0.866	0.0	1.0	45.4	47.0	-25.9	53.7
333	323	321	0.883	0.0	1.0	46.0	49.6	-24.5	55.3
336	324	322	0.9	0.0	1.0	46.6	52.8	-23.2	57.7
338	325	323	0.916	0.0	1.0	47.2	56.0	-21.7	60.0
341	326	324	0.933	0.0	1.0	47.8	59.1	-19.9	62.4
343	327	325	0.95	0.0	1.0	48.4	62.2	-17.9	64.8
346	328	326	0.966	0.0	1.0	48.9	65.3	-15.7	67.1
349	329	327	0.983	0.0	1.0	49.5	68.2	-13.2	69.5
351	330	328	1.0	0.0	1.0	50.1	71.1	-10.5	71.8
351	331	329	1.0	0.0	0.983	49.9	71.5	-10.1	72.2
352	332	330	1.0	0.0	0.966	49.7	71.9	-9.8	72.5
352	333	331	1.0	0.0	0.95	49.6	72.3	-9.4	72.9
352	334	332	1.0	0.0	0.933	49.4	72.7	-9.0	73.2
353	335	333	1.0	0.0	0.916	49.2	73.1	-8.6	73.6
353	336	334	1.0	0.0	0.9	49.0	73.4	-8.2	73.9
353	337	335	1.0	0.0	0.883	48.8	73.8	-7.9	74.3
354	338	336	1.0	0.0	0.866	48.6	74.0	-7.3	74.3
354	339	337	1.0	0.0	0.85	48.6	73.8	-6.5	74.1
355	340	338	1.0	0.0	0.833	48.5	73.6	-5.7	73.9
356	341	339	1.0	0.0	0.816	48.5	73.5	-4.9	73.6
356	342	339	1.0	0.0	0.8	48.4	73.3	-4.1	73.4
357	343	340	1.0	0.0	0.783	48.4	73.1	-3.3	73.2
357	344	341	1.0	0.0	0.766	48.3	72.9	-2.6	72.9
358	345	342	1.0	0.0	0.75	48.3	72.7	-1.8	72.7

<http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 20/33

n-j	HIC*Fe	rgb*Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe	Fe
-----	--------	--------	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/>

TUB-test chart RE71; 1080 s
colors and differences. ΔE^* ,

1: 1080 standard colours, $cf=0.9$

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

[illegible]

Mean color difference of this range:

 $\Delta E^* = 22.7$

see similar files: <http://130.149.60.45/~farbmatrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

TUB-test chart RE71; 1080 standard colours, $cf=0,9$

input: *rgb/cmyk* -> *rgbe*

n	HIC-Fe	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr^*Fe	h_{α}^*Fe	h_{α}^*Fe	DF^*Fe	rgb^*Fe	$LabCh^*Fe$	$LabCh^*Fe$	rgb^*Fe	lcr
-----	--------	-----------	-----------	------------------	-----------	-------------	-------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-----------	------------------	------------------	----------	-----------	-------------	-------------	-----------	-------

see similar files: <http://130.149.60.45/~farbmatrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmatrik>

http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 24/33

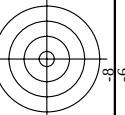
TUB-test chart RE71; 1080 standard colours, $cf=0.9$

input: *rgb/cmyk* -> *rgbe*

[illegible]

REF10-7N Page 34/33-E

1-0122224-F0



TUB material: code=rha4ta

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

1080 standard colours, $c^*/=0,9$ colors and differences, ΔE^* ,

L: 1080 stan

[illegible]

Mean color difference of this page:

DATE RECEIVED

or—good

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

<http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 29/33

n	H1C*Fe	rgb_Fe	ic1_Fe	hsa_Fa	rgb_Fe	LabCH*Fe	rgb_Fe	LabCH*Fe	DF*Fe	hsa_Fe	rgb_Fe	LabCH*Fe
729	NW_100%	0.875	1.0	1.0	0.875	0.992	1.0	96.3	0.5	360	1.0	96.3
730	G50B_100.012%	0.875	1.0	1.0	0.875	0.982	1.0	91.2	0.4	360	1.0	91.2
731	G50B_100.025%	0.875	1.0	1.0	0.875	0.965	1.0	86.1	0.3	360	1.0	86.1
732	G50B_100.037%	0.875	1.0	1.0	0.875	0.948	1.0	81.0	0.2	360	1.0	81.0
733	G50B_100.050%	0.875	1.0	1.0	0.875	0.931	1.0	75.8	0.1	360	1.0	75.8
734	G50B_100.062%	0.875	1.0	1.0	0.875	0.914	1.0	70.7	0.0	360	1.0	70.7
735	G50B_100.075%	0.875	1.0	1.0	0.875	0.897	1.0	65.5	0.0	360	1.0	65.5
736	G50B_100.087%	0.875	1.0	1.0	0.875	0.880	1.0	60.3	0.0	360	1.0	60.3
737	G50B_100.100%	0.875	1.0	1.0	0.875	0.863	1.0	55.3	0.0	360	1.0	55.3
738	ROY_100.012%	0.875	1.0	1.0	0.875	0.904	1.0	80.0	0.0	360	1.0	80.0
739	ROY_100.025%	0.875	1.0	1.0	0.875	0.887	1.0	74.8	0.0	360	1.0	74.8
740	ROY_100.037%	0.875	1.0	1.0	0.875	0.870	1.0	69.6	0.0	360	1.0	69.6
741	ROY_100.050%	0.875	1.0	1.0	0.875	0.853	1.0	64.4	0.0	360	1.0	64.4
742	ROY_100.062%	0.875	1.0	1.0	0.875	0.836	1.0	59.2	0.0	360	1.0	59.2
743	ROY_100.075%	0.875	1.0	1.0	0.875	0.819	1.0	54.0	0.0	360	1.0	54.0
744	ROY_100.087%	0.875	1.0	1.0	0.875	0.802	1.0	48.8	0.0	360	1.0	48.8
745	ROY_100.100%	0.875	1.0	1.0	0.875	0.785	1.0	43.6	0.0	360	1.0	43.6
746	ROY_100.012%	0.875	1.0	1.0	0.875	0.921	1.0	85.6	0.0	360	1.0	85.6
747	ROY_100.025%	0.875	1.0	1.0	0.875	0.904	1.0	80.4	0.0	360	1.0	80.4
748	ROY_100.037%	0.875	1.0	1.0	0.875	0.887	1.0	75.2	0.0	360	1.0	75.2
749	ROY_100.050%	0.875	1.0	1.0	0.875	0.870	1.0	69.9	0.0	360	1.0	69.9
750	ROY_100.062%	0.875	1.0	1.0	0.875	0.853	1.0	64.7	0.0	360	1.0	64.7
751	ROY_100.075%	0.875	1.0	1.0	0.875	0.836	1.0	59.4	0.0	360	1.0	59.4
752	ROY_100.087%	0.875	1.0	1.0	0.875	0.819	1.0	54.2	0.0	360	1.0	54.2
753	ROY_100.100%	0.875	1.0	1.0	0.875	0.802	1.0	48.9	0.0	360	1.0	48.9
754	ROY_100.012%	0.875	1.0	1.0	0.875	0.938	1.0	88.0	0.0	360	1.0	88.0
755	ROY_100.025%	0.875	1.0	1.0	0.875	0.921	1.0	82.8	0.0	360	1.0	82.8
756	ROY_100.037%	0.875	1.0	1.0	0.875	0.904	1.0	77.6	0.0	360	1.0	77.6
757	ROY_100.050%	0.875	1.0	1.0	0.875	0.887	1.0	72.4	0.0	360	1.0	72.4
758	ROY_100.062%	0.875	1.0	1.0	0.875	0.870	1.0	67.2	0.0	360	1.0	67.2
759	ROY_100.075%	0.875	1.0	1.0	0.875	0.853	1.0	62.0	0.0	360	1.0	62.0
760	ROY_100.087%	0.875	1.0	1.0	0.875	0.836	1.0	56.8	0.0	360	1.0	56.8
761	ROY_100.100%	0.875	1.0	1.0	0.875	0.819	1.0	51.6	0.0	360	1.0	51.6
762	ROY_100.012%	0.875	1.0	1.0	0.875	0.959	1.0	93.1	0.0	360	1.0	93.1
763	ROY_100.025%	0.875	1.0	1.0	0.875	0.942	1.0	87.9	0.0	360	1.0	87.9
764	ROY_100.037%	0.875	1.0	1.0	0.875	0.925	1.0	82.7	0.0	360	1.0	82.7
765	ROY_100.050%	0.875	1.0	1.0	0.875	0.908	1.0	77.5	0.0	360	1.0	77.5
766	ROY_100.062%	0.875	1.0	1.0	0.875	0.891	1.0	72.3	0.0	360	1.0	72.3
767	ROY_100.075%	0.875	1.0	1.0	0.875	0.874	1.0	67.1	0.0	360	1.0	67.1
768	ROY_100.087%	0.875	1.0	1.0	0.875	0.857	1.0	61.9	0.0	360	1.0	61.9
769	ROY_100.100%	0.875	1.0	1.0	0.875	0.840	1.0	56.7	0.0	360	1.0	56.7
770	ROY_100.012%	0.875	1.0	1.0	0.875	0.978	1.0	98.0	0.0	360	1.0	98.0
771	ROY_100.025%	0.875	1.0	1.0	0.875	0.961	1.0	92.8	0.0	360	1.0	92.8
772	ROY_100.037%	0.875	1.0	1.0	0.875	0.944	1.0	87.6	0.0	360	1.0	87.6
773	ROY_100.050%	0.875	1.0	1.0	0.875	0.927	1.0	82.4	0.0	360	1.0	82.4
774	ROY_100.062%	0.875	1.0	1.0	0.875	0.910	1.0	77.2	0.0	360	1.0	77.2
775	ROY_100.075%	0.875	1.0	1.0	0.875	0.893	1.0	72.0	0.0	360	1.0	72.0
776	ROY_100.087%	0.875	1.0	1.0	0.875	0.876	1.0	66.8	0.0	360	1.0	66.8
777	ROY_100.100%	0.875	1.0	1.0	0.875	0.859	1.0	61.6	0.0	360	1.0	61.6
778	ROY_100.012%	0.875	1.0	1.0	0.875	0.997	1.0	100.0	0.0	360	1.0	100.0
779	ROY_100.025%	0.875	1.0	1.0	0.875	0.980	1.0	94.8	0.0	360	1.0	94.8
780	ROY_100.037%	0.875	1.0	1.0	0.875	0.963	1.0	89.6	0.0	360	1.0	89.6
781	ROY_100.050%	0.875	1.0	1.0	0.875	0.946	1.0	84.4	0.0	360	1.0	84.4
782	ROY_100.062%	0.875	1.0	1.0	0.875	0.929	1.0	79.2	0.0	360	1.0	79.2
783	ROY_100.075%	0.875	1.0	1.0	0.875	0.912	1.0	74.0	0.0	360	1.0	74.0
784	ROY_100.087%	0.875	1.0	1.0	0.875	0.895	1.0	68.8	0.0	360	1.0	68.8
785	ROY_100.100%	0.875	1.0	1.0	0.875	0.878	1.0	63.6	0.0	360	1.0	63.6
786	ROY_100.012%	0.875	1.0	1.0	0.875	1.000	1.0	100.0	0.0	360	1.0	100.0
787	ROY_100.025%	0.875	1.0	1.0	0.875	0.983	1.0	94.8	0.0	360	1.0	94.8
788	ROY_100.037%	0.875	1.0	1.0	0.875	0.966	1.0	89.6	0.0	360	1.0	89.6
789	ROY_100.050%	0.875	1.0	1.0	0.875	0.949	1.0	84.4	0.0	360	1.0	84.4
790	ROY_100.062%	0.875	1.0	1.0	0.875	0.932	1.0	79.2	0.0	360	1.0	79.2
791	ROY_100.075%	0.875	1.0	1.0	0.875	0.915	1.0	74.0	0.0	360	1.0	74.0
792	ROY_100.087%	0.875	1.0	1.0	0.875	0.898	1.0	68.8	0.0	360	1.0	68.8
793	ROY_100.100%	0.875	1.0	1.0	0.875	0.881	1.0	63.6	0.0	360	1.0	63.6
794	ROY_100.012%	0.875	1.0	1.0	0.875	1.013	1.0	100.0	0.0	360	1.0	100.0
795	ROY_100.025%	0.875	1.0	1.0	0.875	0.996	1.0	94.8	0.0	360	1.0	94.8
796	ROY_100.037%	0.875	1.0	1.0	0.875	0.979	1.0	89.6	0.0	360	1.0	89.6
797	ROY_100.050%	0.875	1.0	1.0	0.875	0.962	1.0	84.4	0.0	360	1.0	84.4
798	ROY_100.062%	0.875	1.0	1.0	0.875	0.945	1.0	79.2	0.0	360	1.0	79.2
799	ROY_100.075%	0.875	1.0	1.0	0.875	0.928	1.0	74.0	0.0	360	1.0	74.0
800	ROY_100.087%	0.875	1.0	1.0	0.875	0.911	1.0	68.8	0.0	360	1.0	68.8
801	ROY_100.100%	0.875	1.0	1.0	0.875	0.894	1.0	63.6	0.0	360	1.0	63.6
802	ROY_100.012%	0.875	1.0	1.0	0.875	1.026	1.0	100.0	0.0	360	1.0	100.0
803	ROY_100.025%	0.875	1.0	1.0	0.875	1.009	1.0	94.8	0.0	360	1.0	94.8
804	ROY_100.037%	0.875	1.0	1.0	0.875	0.992	1.0	89.6	0.0	360	1.0	89.6
805	ROY_100.050%	0.875	1.0	1.0	0.875	0.975	1.0	84.4	0.0	360	1.0	84.4
806	ROY_100.062%	0.875	1.0	1.0	0.875	0.958	1.0	79.2	0.0	360	1.0	79.2
807	ROY_100.075%	0.875	1.0	1.0	0.875	0.941	1.0	74.0	0.0	360	1.0	74.0
808	ROY_100.087%	0.875	1.0	1.0	0.875	0.924	1.0	68.8	0.0	360	1.0	68.8
809	ROY_100.100%	0.875	1.0	1.0	0.875	0.907	1.0	63.6	0.0	360	1.0	63.6

RE710-7N, Page 29/33-F

TUB-test chart RE71; 1080 standard colours, cf=0.9
colors and differences, ΔE^*

input: $rgb/cm\text{y}k \rightarrow rgbe$
output: transfer to $rgbe$

Mean color difference of this page: $\Delta E^* = 12.5$

n	H1C*Fe	rgb_Fe	ic1_Fe	hs_Fe	LabCH*Fe	rgb*Fe	LabCH*Fe	rgb*Fe	DF*Fe	hs_Fe	rgb*Fe	LabCH*Fe
891	NW_100_0124	1.0	1.0	1.0	96.3	1.0	96.4	1.0	0.3	360	1.0	96.3
892	B50R_100_0124	1.0	0.875	1.0	89.9	5.6	83.5	1.0	272.1	10.8	322	44.8
893	B50R_100_0254	1.0	0.875	1.0	89.9	5.6	83.5	1.0	373.1	10.8	322	44.8
894	B50R_100_0374	1.0	0.875	1.0	89.9	5.6	83.5	1.0	474.1	10.8	322	44.8
895	B50R_100_0504	1.0	0.875	1.0	89.9	5.6	83.5	1.0	575.1	10.8	322	44.8
896	B50R_100_0624	1.0	0.875	1.0	89.9	5.6	83.5	1.0	676.1	10.8	322	44.8
897	B50R_100_0754	1.0	0.875	1.0	89.9	5.6	83.5	1.0	777.1	10.8	322	44.8
898	B50R_100_0874	1.0	0.875	1.0	89.9	5.6	83.5	1.0	878.1	10.8	322	44.8
899	B50R_100_1004	1.0	0.875	1.0	89.9	5.6	83.5	1.0	979.1	10.8	322	44.8
900	COB1_100_0124	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	1080.1	10.8	322	44.8
901	NW_0874	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	1181.1	10.8	322	44.8
902	B50R_087_0124	0.875	0.875	0.875	86.3	0.0	80.8	0.875	1282.1	10.8	322	44.8
903	B50R_087_0254	0.875	0.875	0.875	86.3	0.0	80.8	0.875	1383.1	10.8	322	44.8
904	B50R_087_0374	0.875	0.875	0.875	86.3	0.0	80.8	0.875	1484.1	10.8	322	44.8
905	B50R_087_0504	0.875	0.875	0.875	86.3	0.0	80.8	0.875	1585.1	10.8	322	44.8
906	B50R_087_0624	0.875	0.875	0.875	86.3	0.0	80.8	0.875	1686.1	10.8	322	44.8
907	B50R_087_0754	0.875	0.875	0.875	86.3	0.0	80.8	0.875	1787.1	10.8	322	44.8
908	B50R_087_0874	0.875	0.875	0.875	86.3	0.0	80.8	0.875	1888.1	10.8	322	44.8
909	COB1_100_0254	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	1989.1	10.8	322	44.8
910	COB1_100_0374	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	2090.1	10.8	322	44.8
911	NW_0754	0.875	0.875	0.875	86.3	0.0	80.8	0.875	2191.1	10.8	322	44.8
912	B50R_075_0124	0.875	0.875	0.875	86.3	0.0	80.8	0.875	2292.1	10.8	322	44.8
913	B50R_075_0254	0.875	0.875	0.875	86.3	0.0	80.8	0.875	2393.1	10.8	322	44.8
914	B50R_075_0374	0.875	0.875	0.875	86.3	0.0	80.8	0.875	2494.1	10.8	322	44.8
915	B50R_075_0504	0.875	0.875	0.875	86.3	0.0	80.8	0.875	2595.1	10.8	322	44.8
916	B50R_075_0624	0.875	0.875	0.875	86.3	0.0	80.8	0.875	2696.1	10.8	322	44.8
917	B50R_075_0754	0.875	0.875	0.875	86.3	0.0	80.8	0.875	2797.1	10.8	322	44.8
918	COB1_100_0374	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	2898.1	10.8	322	44.8
919	COB1_100_0504	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	2999.1	10.8	322	44.8
920	COB1_100_0624	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	3100.1	10.8	322	44.8
921	NW_0624	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	3201.1	10.8	322	44.8
922	B50R_062_0124	0.875	0.875	0.875	86.3	0.0	80.8	0.875	3302.1	10.8	322	44.8
923	B50R_062_0254	0.875	0.875	0.875	86.3	0.0	80.8	0.875	3403.1	10.8	322	44.8
924	B50R_062_0374	0.875	0.875	0.875	86.3	0.0	80.8	0.875	3504.1	10.8	322	44.8
925	B50R_062_0504	0.875	0.875	0.875	86.3	0.0	80.8	0.875	3605.1	10.8	322	44.8
926	B50R_062_0624	0.875	0.875	0.875	86.3	0.0	80.8	0.875	3706.1	10.8	322	44.8
927	COB1_100_0504	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	3807.1	10.8	322	44.8
928	COB1_100_0754	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	3908.1	10.8	322	44.8
929	COB1_100_0874	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	4009.1	10.8	322	44.8
930	COB1_100_1004	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	4110.1	10.8	322	44.8
931	NW_0504	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	4211.1	10.8	322	44.8
932	B50R_050_0124	0.875	0.875	0.875	86.3	0.0	80.8	0.875	4312.1	10.8	322	44.8
933	B50R_050_0254	0.875	0.875	0.875	86.3	0.0	80.8	0.875	4413.1	10.8	322	44.8
934	B50R_050_0374	0.875	0.875	0.875	86.3	0.0	80.8	0.875	4514.1	10.8	322	44.8
935	B50R_050_0504	0.875	0.875	0.875	86.3	0.0	80.8	0.875	4615.1	10.8	322	44.8
936	B50R_050_0624	0.875	0.875	0.875	86.3	0.0	80.8	0.875	4716.1	10.8	322	44.8
937	COB1_100_0624	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	4817.1	10.8	322	44.8
938	COB1_100_0874	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	4918.1	10.8	322	44.8
939	COB1_100_1004	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	5019.1	10.8	322	44.8
940	NW_0374	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	5120.1	10.8	322	44.8
941	B50R_037_0124	0.875	0.875	0.875	86.3	0.0	80.8	0.875	5221.1	10.8	322	44.8
942	B50R_037_0254	0.875	0.875	0.875	86.3	0.0	80.8	0.875	5322.1	10.8	322	44.8
943	B50R_037_0374	0.875	0.875	0.875	86.3	0.0	80.8	0.875	5423.1	10.8	322	44.8
944	B50R_037_0504	0.875	0.875	0.875	86.3	0.0	80.8	0.875	5524.1	10.8	322	44.8
945	COB1_100_0754	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	5625.1	10.8	322	44.8
946	COB1_100_0874	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	5726.1	10.8	322	44.8
947	COB1_100_1004	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	5827.1	10.8	322	44.8
948	COB1_050_0124	0.875	0.875	0.875	86.3	0.0	80.8	0.875	5928.1	10.8	322	44.8
949	COB1_050_0254	0.875	0.875	0.875	86.3	0.0	80.8	0.875	6029.1	10.8	322	44.8
950	COB1_050_0374	0.875	0.875	0.875	86.3	0.0	80.8	0.875	6130.1	10.8	322	44.8
951	NW_0254	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	6231.1	10.8	322	44.8
952	B50R_025_0124	0.875	0.875	0.875	86.3	0.0	80.8	0.875	6332.1	10.8	322	44.8
953	B50R_025_0254	0.875	0.875	0.875	86.3	0.0	80.8	0.875	6433.1	10.8	322	44.8
954	COB1_100_0874	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	6534.1	10.8	322	44.8
955	COB1_100_1004	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	6635.1	10.8	322	44.8
956	COB1_087_0754	0.875	0.875	0.875	86.3	0.0	80.8	0.875	6736.1	10.8	322	44.8
957	COB1_087_0874	0.875	0.875	0.875	86.3	0.0	80.8	0.875	6837.1	10.8	322	44.8
958	COB1_087_1004	0.875	0.875	0.875	86.3	0.0	80.8	0.875	6938.1	10.8	322	44.8
959	COB1_050_0504	0.875	0.875	0.875	86.3	0.0	80.8	0.875	7039.1	10.8	322	44.8
960	COB1_050_0624	0.875	0.875	0.875	86.3	0.0	80.8	0.875	7140.1	10.8	322	44.8
961	NW_0124	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	7241.1	10.8	322	44.8
962	B50R_012_0124	0.875	0.875	0.875	86.3	0.0	80.8	0.875	7342.1	10.8	322	44.8
963	COB1_100_1004	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	7443.1	10.8	322	44.8
964	COB1_087_0504	0.875	0.875	0.875	86.3	0.0	80.8	0.875	7544.1	10.8	322	44.8
965	COB1_087_0624	0.875	0.875	0.875	86.3	0.0	80.8	0.875	7645.1	10.8	322	44.8
966	COB1_087_0754	0.875	0.875	0.875	86.3	0.0	80.8	0.875	7746.1	10.8	322	44.8
967	COB1_087_0874	0.875	0.875	0.875	86.3	0.0	80.8	0.875	7847.1	10.8	322	44.8
968	COB1_087_1004	0.875	0.875	0.875	86.3	0.0	80.8	0.875	7948.1	10.8	322	44.8
969	COB1_050_0874	0.875	0.875	0.875	86.3	0.0	80.8	0.875	8049.1	10.8	322	44.8
970	COB1_050_1004	0.875	0.875	0.875	86.3	0.0	80.8	0.875	8150.1	10.8	322	44.8
971	NW_0004	0.875	1.0	0.875	91.6	-6.8	86.2	1.0	8251.1	10.8	322	44.8

<http://130.149.60.45/~farbmatrik/RE71/RE71L0NA.TXT> /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 32/33

n	HC*Fe	rgb_Fe	icL_Fe	hsa_Fa	rgb_Fe	LabCH*Fe	rgb_Fe	LabCH*Fe	DE**Fe	hsa_Fe	rgb_Fe	LabCH*Fe
972	NW 000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
973	NW 012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	3.6	360	1.0	1.0
974	NW 025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	10.7	360	1.0	1.0
975	NW 037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	24.4	360	1.0	1.0
976	NW 050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	48.6	360	1.0	1.0
977	NW 062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	74	360	1.0	1.0
978	NW 075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	111	360	1.0	1.0
979	NW 087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	163	360	1.0	1.0
980	NW 100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	215	360	1.0	1.0
981	NW 000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.6	360	1.0	1.0
982	NW 012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	10.7	360	1.0	1.0
983	NW 025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	24.4	360	1.0	1.0
984	NW 037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	48.6	360	1.0	1.0
985	NW 050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	74	360	1.0	1.0
986	NW 062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	111	360	1.0	1.0
987	NW 075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	163	360	1.0	1.0
988	NW 087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	215	360	1.0	1.0
989	NW 100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	267	360	1.0	1.0
990	NW 000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.6	360	1.0	1.0
991	NW 012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	10.7	360	1.0	1.0
992	NW 025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	24.4	360	1.0	1.0
993	NW 037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	48.6	360	1.0	1.0
994	NW 050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	74	360	1.0	1.0
995	NW 062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	111	360	1.0	1.0
996	NW 075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	163	360	1.0	1.0
997	NW 087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	215	360	1.0	1.0
998	NW 100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	267	360	1.0	1.0
999	NW 000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.6	360	1.0	1.0
1000	NW 012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	10.7	360	1.0	1.0
1001	NW 025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	24.4	360	1.0	1.0
1002	NW 037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	48.6	360	1.0	1.0
1003	NW 050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	74	360	1.0	1.0
1004	NW 062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	111	360	1.0	1.0
1005	NW 075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	163	360	1.0	1.0
1006	NW 087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	215	360	1.0	1.0
1007	NW 100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	267	360	1.0	1.0
1008	NW 000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.6	360	1.0	1.0
1009	NW 012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	10.7	360	1.0	1.0
1010	NW 025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	24.4	360	1.0	1.0
1011	NW 037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	48.6	360	1.0	1.0
1012	NW 050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	74	360	1.0	1.0
1013	NW 062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	111	360	1.0	1.0
1014	NW 075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	163	360	1.0	1.0
1015	NW 087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	215	360	1.0	1.0
1016	NW 100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	267	360	1.0	1.0
1017	NW 000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.6	360	1.0	1.0
1018	NW 012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	10.7	360	1.0	1.0
1019	NW 025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	24.4	360	1.0	1.0
1020	NW 037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	48.6	360	1.0	1.0
1021	NW 050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	74	360	1.0	1.0
1022	NW 062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	111	360	1.0	1.0
1023	NW 075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	163	360	1.0	1.0
1024	NW 087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	215	360	1.0	1.0
1025	NW 100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	267	360	1.0	1.0
1026	NW 000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.6	360	1.0	1.0
1027	NW 012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	10.7	360	1.0	1.0
1028	NW 025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	24.4	360	1.0	1.0
1029	NW 037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	48.6	360	1.0	1.0
1030	NW 050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	74	360	1.0	1.0
1031	NW 062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	111	360	1.0	1.0
1032	NW 075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	163	360	1.0	1.0
1033	NW 087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	215	360	1.0	1.0
1034	NW 100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	267	360	1.0	1.0
1035	NW 000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.6	360	1.0	1.0
1036	NW 012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	10.7	360	1.0	1.0
1037	NW 025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	24.4	360	1.0	1.0
1038	NW 037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	48.6	360	1.0	1.0
1039	NW 050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	74	360	1.0	1.0
1040	NW 062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	111	360	1.0	1.0
1041	NW 075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	163	360	1.0	1.0
1042	NW 087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	215	360	1.0	1.0
1043	NW 100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	267	360	1.0	1.0
1044	NW 000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.6	360	1.0	1.0
1045	NW 012a	0.125	0.125	0.125	0.125	0.125	0.125	0.125	10.7	360	1.0	1.0
1046	NW 025a	0.25	0.25	0.25	0.25	0.25	0.25	0.25	24.4	360	1.0	1.0
1047	NW 037a	0.375	0.375	0.375	0.375	0.375	0.375	0.375	48.6	360	1.0	1.0
1048	NW 050a	0.5	0.5	0.5	0.5	0.5	0.5	0.5	74	360	1.0	1.0
1049	NW 062a	0.625	0.625	0.625	0.625	0.625	0.625	0.625	111	360	1.0	1.0
1050	NW 075a	0.75	0.75	0.75	0.75	0.75	0.75	0.75	163	360	1.0	1.0
1051	NW 087a	0.875	0.875	0.875	0.875	0.875	0.875	0.875	215	360	1.0	1.0
1052	NW 100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	267	360	1.0	1.0

Mean color difference of this page: $\Delta E^* = 6.3$

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

TUB-test chart RE71; 1080 standard colours, $cf=0.9$
colors and differences, ΔE^*

RE710-7N, Page 32/33-F

TUB registration: 20150701-RE71/RE71L0NA.TXT /.PS
application for measurement of laser printer output, no separation rgb (RGB)

TUB material: code=rha4ta

see similar files: <http://130.149.60.45/~farbmetrik/RE71/RE71.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

<http://130.149.60.45/~farbmetrik/RE71/RE71L0NA.TXT> /.PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 33/33

n	HC*Fe	rgb_Fe	ic*_Fe	hsa_Fe	rgb*Fe	LabCh*Fe	LabCh*Fe	DE*Fe	rgb*Me	LabCh*Me	DE*Fe	rgb*Me	LabCh*Me	DE*Fe	rgb*Me	LabCh*Me	DE*Fe	rgb*Me	LabCh*Me
1053	NW_086e	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866
1054	NW_093e	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933
1055	NW_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	1.0	1.0	1.0	1.0	1.0	1.0
1056	NW_006e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1057	NW_006e	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066	0.066
1058	NW_013e	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133	0.133
1059	NW_020e	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2	0.2
1060	NW_026e	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266	0.266
1061	NW_033e	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333	0.333
1062	NW_040e	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4
1063	NW_046e	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466	0.466
1064	NW_053e	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533	0.533
1065	NW_060e	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6	0.6
1066	NW_066e	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666	0.666
1067	NW_073e	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734	0.734
1068	NW_080e	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8	0.8
1069	NW_086e	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866	0.866
1070	NW_093e	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933	0.933
1071	NW_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	1.0	1.0	1.0	1.0	1.0	1.0
1072	NW_100e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1073	RO07_100_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	1.0	1.0	1.0	1.0	1.0	1.0
1074	RO07_100_100e	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1075	Y06C_100_100e	0.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1076	Y06C_100_100e	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	0.0	0.0
1077	B00C_100_100e	0.0	0.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0
1078	B00C_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	0.0
1079	B50R_100_100e	1.0	0.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0

Mean color difference of this page:

delta E* = 8.0

RE710-7N, Page 33/33-F

TUB-test chart RE71; 1080 standard colours, cf=0.9
colors and differences, ΔE^*

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