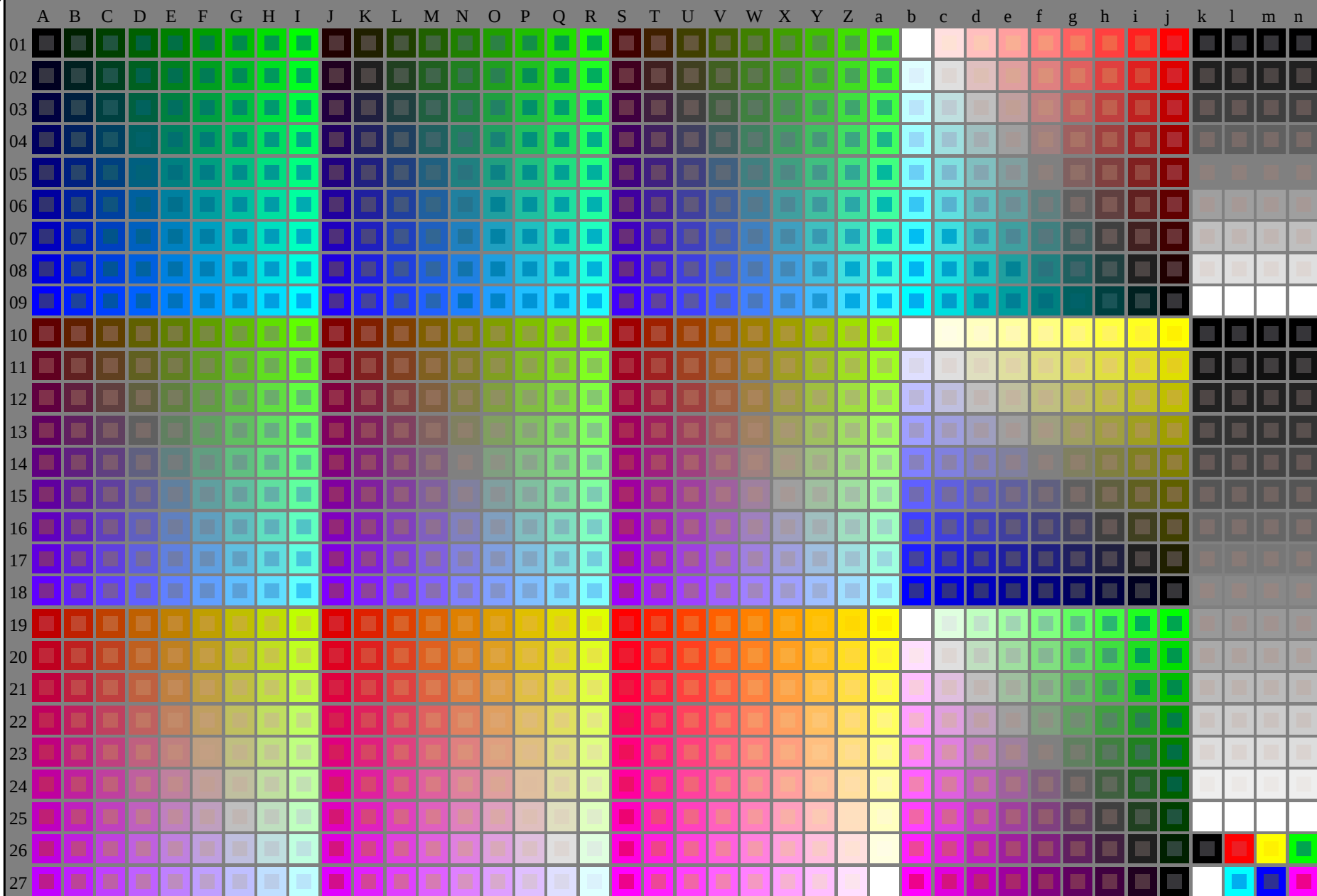


see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output

TUB material: code=rha4ta



1-003030-L0

SE140-7N

Test chart G with 40x27=1080 colours; digital equidistant 9 or 16 step colour scales; Colour data in column (A-n): **rgb + cmy0** (A-j + k26_n27), 000n (k), w (l), nnn0 (m), www (n), 3D = 0

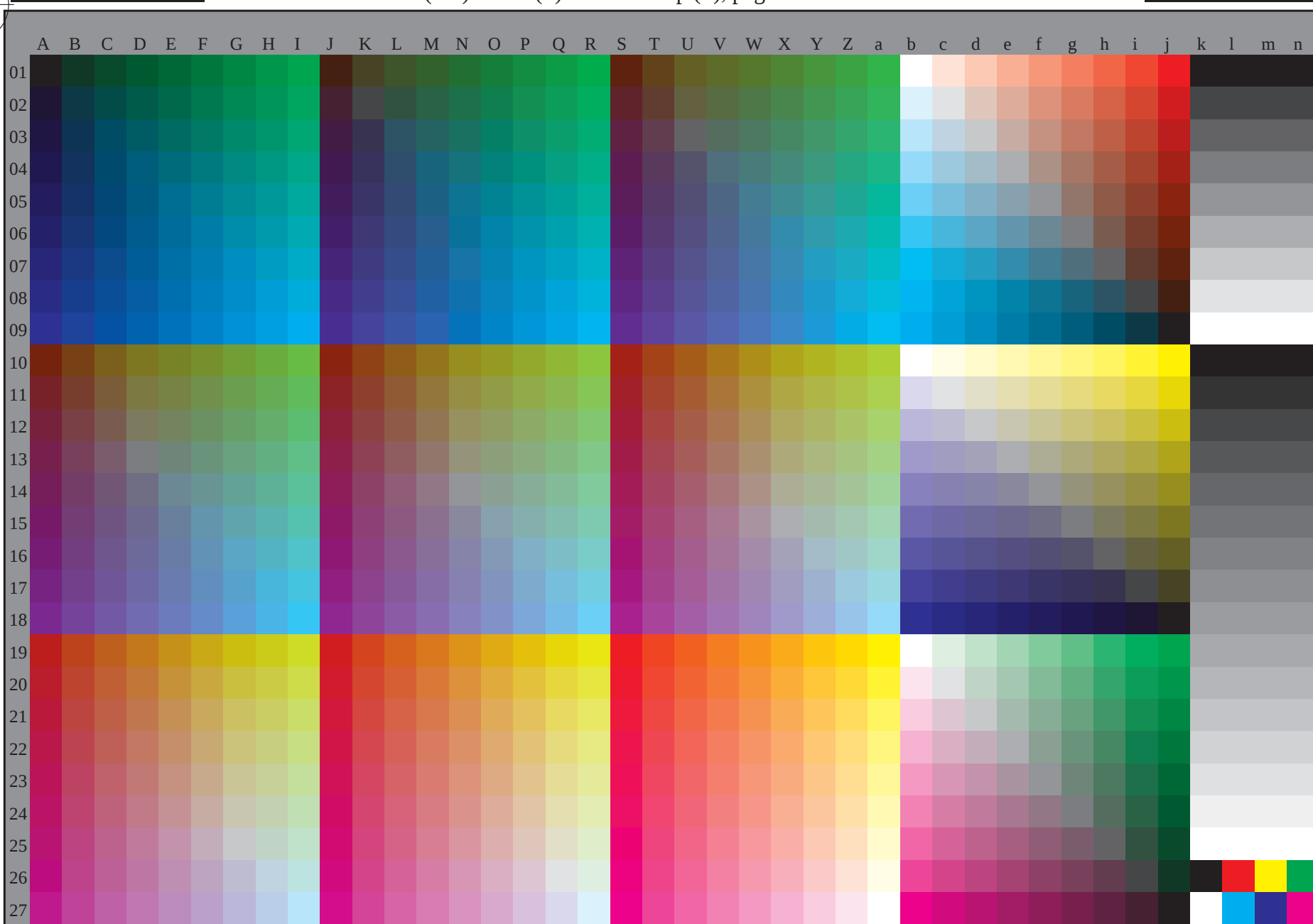
TUB-test chart SE14; 1080 colours, offset standard paper
Test chart according to DIN 33872

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

http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF /.PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 2/33

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output, separationcmykn6 (CMYK)
TUB material: code=rha4ta



1-003130-L0

SE140-70

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

TUB-test chart SE14; 1080 colours, offset standard paper
Test chart according to DIN 33872, 3D=0, de=0, cmyk

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

1-003130-F0

C

M

Y

O

L

V

C

http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF /.PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 3/33

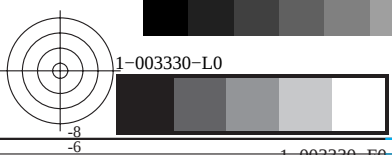
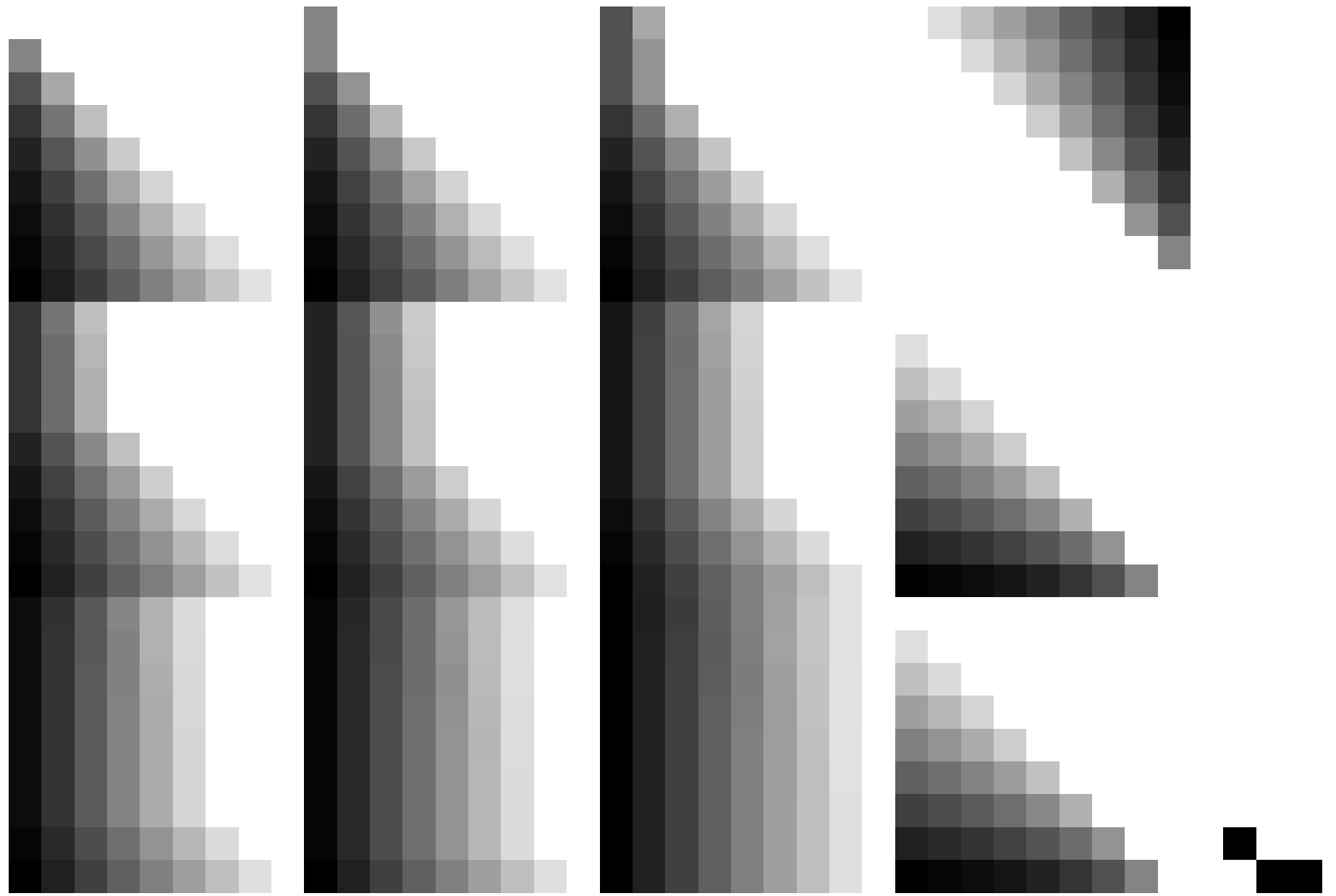
1-003230-L0 SE140-70
TUB-test chart SE14; 1080 colours, offset standard paper
Test chart according to DIN 33872, 3D=0, de=0, cmyk

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

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



<http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF> /.PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 5/33

TUB-test chart SE14; 1080 colours, offset standard paper
Test chart according to DIN 33872, 3D=0, de=0, cmyk

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

SE140-70

1-003430-L0

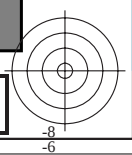
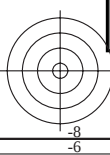
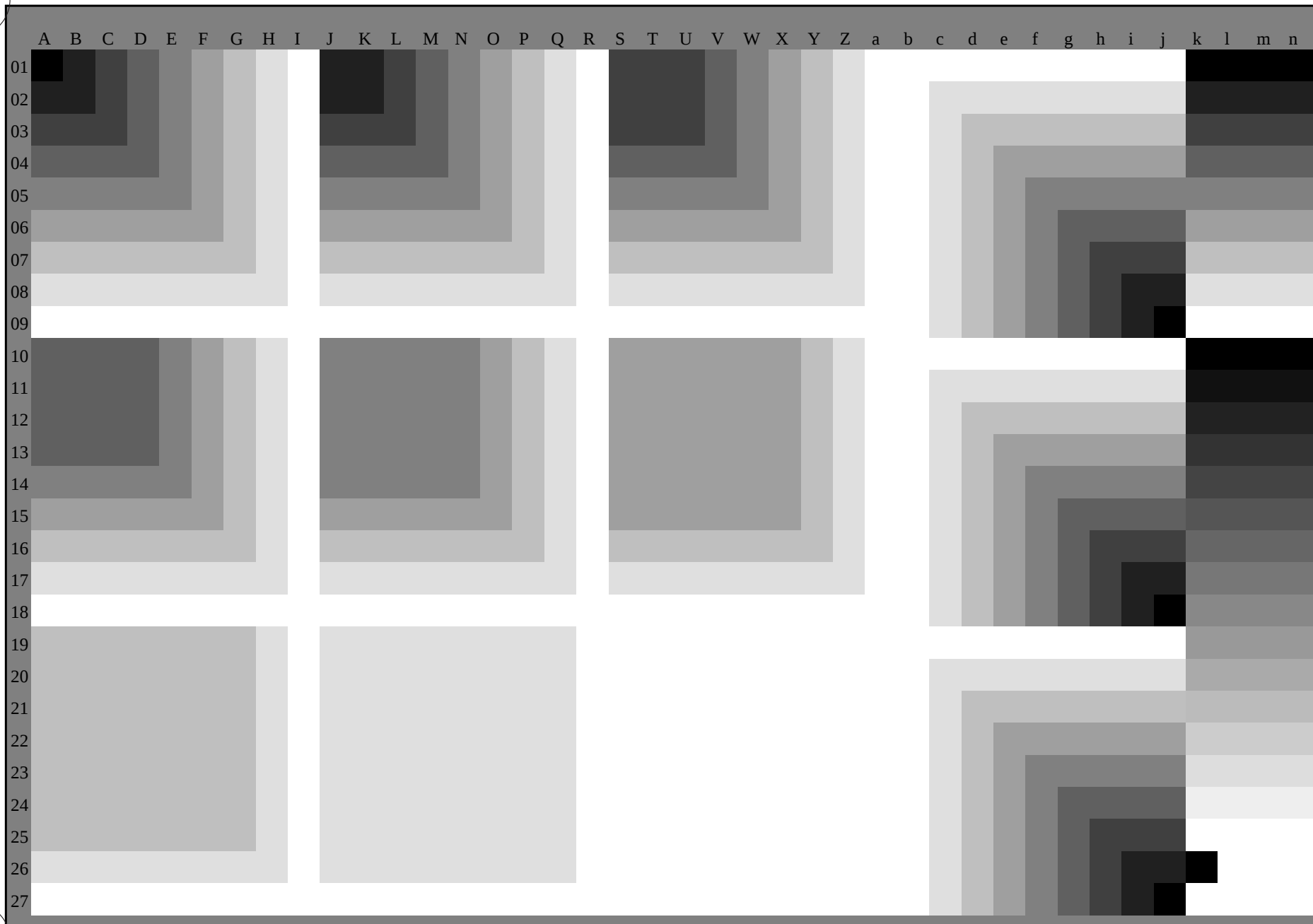
1-003430-E0

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>



see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output, separationcmykn6 (CMYK)
TUB material: code=rha4ta

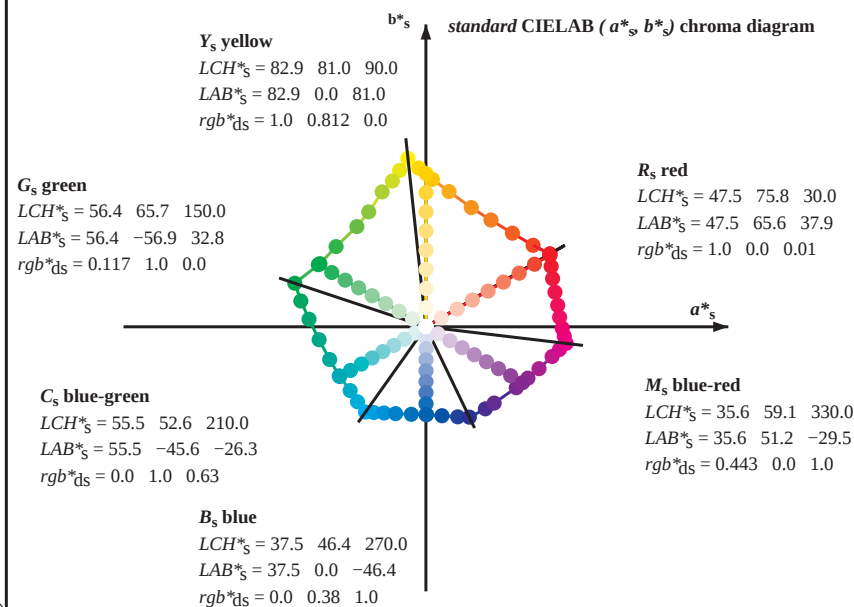
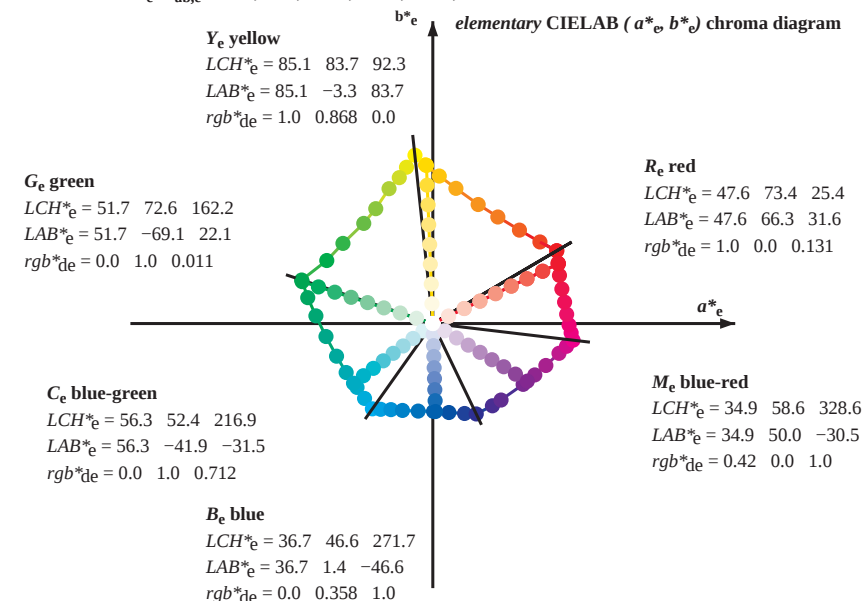
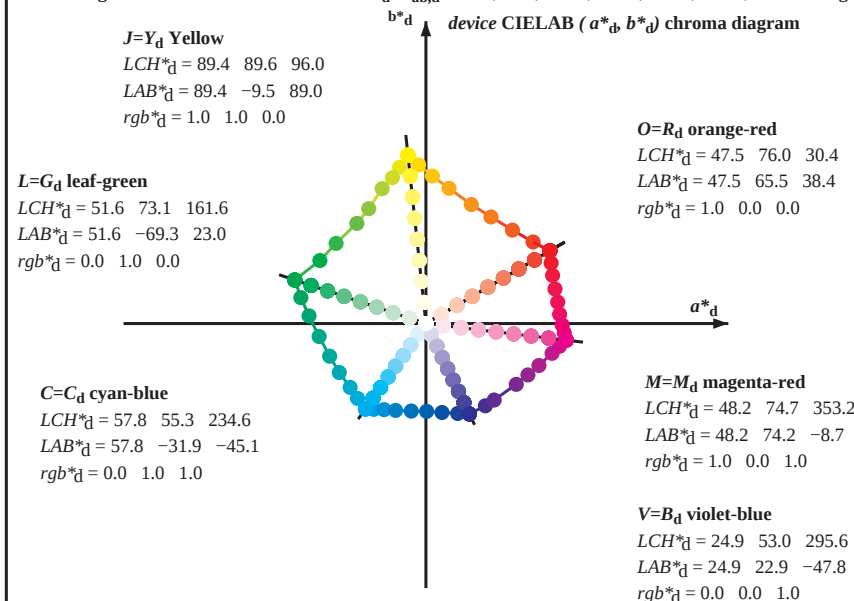


SE140-70
TUB-test chart SE14; 1080 colours, offset standard paper
Test chart according to DIN 33872, 3D=0, de=0, cmyk

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



Data of Maximum color M in colorimetric system Offset standard print; separation cmyn6*, D65 for input or output; Six hue angles of the 60 degree standard colours $RYGCBM_s$: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$; Six hue angles of the device colours $RYGCBM_d$: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours $RYGCBM_e$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$



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

- For the rgb^*_e -input values the CIELAB data LCH^*_e and LAB^*_e have been calculated.
- For the calculation of the standard hue angle $h_{ab,s}$ use for any device values rgb^*_e the equation:
$$h_{ab,s} = \text{atan} [r^*_d \cos(30) + g^*_d \cos(150)] / [r^*_d \sin(30) + g^*_d \sin(150) + b^*_d \sin(270)] \quad (1)$$
- For the 48 or 360 equally spaced standard hue angles $h_{ab,s}$ of the colours of maximum chroma use the seven hue angles of the 60 degree colours s : $h_{ab,si} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0, 390.0$ ($i=0,6$) and the equations for a 48 and 360 step hue circle:
$$h_{48ab,sij} = h_{ab,si} + j [h_{ab,si+1} - h_{ab,si}] / 8 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 7) \quad (2)$$
$$h_{360ab,sij} = h_{ab,si} + j [h_{ab,si+1} - h_{ab,si}] / 60 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 59) \quad (3)$$
- For the 48 or 360 elementary hue angles $h_{ab,e}$ of the colours of maximum chroma use the seven hue angles of the elementary colours e : $h_{ab,ei} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6, 385.5$ ($i=0,6$) and the equations for a 48 and 360 step elementary hue circle:
$$h_{48ab,eij} = h_{ab,ei} + j [h_{ab,ei+1} - h_{ab,ei}] / 8 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 7) \quad (4)$$
$$h_{360ab,eij} = h_{ab,ei} + j [h_{ab,ei+1} - h_{ab,ei}] / 60 \quad (i = 0, 1, \dots, 5; j = 0, 1, \dots, 59) \quad (5)$$
- For any elementary hue angle $h_{ab,e}$ there is a well defined device hue angle $h_{ab,d}$ see the following tables, columns 1 to 5 or 1 to 4.
- The values rgb^*_e produce the output of the device-independent elementary hues

1-003630-L0 SE140-70 LAB*la0, YN=0%, XYZnw=2.5, 2.6, 2.7, 86.0, 90.9, 95.9, LAB*nw=18.5, 0.0, 0.0, 96.4, 0.0, 0.0

Output: Offset standard print; separation cmyn6*, D65, page 7/33

TUB-test chart SE14; 1080 colours, offset standard paper
48 step hue circles; rgb - $LabCh^*$ tables, 3D=0, de=0, $cm\dot{y}k$

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

Data of Maximum color M in colorimetric system Offset standard print; separation cmykn6*, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM₆: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours RYGBM_d: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours RYGBM_e: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_{dd64M}			LAB^*_{ddx64M} (x=LabCh)					$rgb^*_{ddx361M}$			$LAB^*_{ddx361M}$ (x=LabCh)					$rgb^*_{dsx361M}$			$LAB^*_{dsx361M}$ (x=LabCh)					$rgb^*_{dex361M}$			$LAB^*_{dex361M}$				
30.4	30.0	25.4	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4	1.0	0.0	0.0	47.5	65.6	38.5	76.1	30	1.0	0.0	0.011	47.5	65.7	37.9	75.8	30	1.0	0.0	0.131	47.7	66.3	31.6	73.5	25
37.2	37.5	33.8	1.0	0.125	0.0	51.5	56.6	43.1	71.2	37.2	1.0	0.117	0.0	51.3	57.3	42.9	71.6	36	1.0	0.12	0.0	51.4	57.0	43.0	71.4	37	1.0	0.052	0.0	49.2	61.9	40.6	74.0	33
47.2	45.0	42.1	1.0	0.25	0.0	56.6	45.8	49.4	67.4	47.2	1.0	0.25	0.0	56.6	45.8	49.5	67.4	47	1.0	0.222	0.0	55.5	48.3	48.3	68.3	45	1.0	0.187	0.0	54.1	51.4	46.6	69.4	42
58.6	52.5	50.5	1.0	0.375	0.0	62.3	34.4	56.4	66.1	58.6	1.0	0.367	0.0	62.0	35.2	56.1	66.2	57	1.0	0.302	0.0	59.0	41.2	52.7	66.9	52	1.0	0.28	0.0	58.0	43.2	51.4	67.1	49
69.1	60.0	58.8	1.0	0.5	0.0	68.1	24.0	63.0	67.4	69.1	1.0	0.5	0.0	68.1	24.0	63.0	67.5	69	1.0	0.391	0.0	63.1	33.1	57.4	66.3	60	1.0	0.378	0.0	62.5	34.2	56.6	66.1	58
80.3	67.5	67.2	1.0	0.625	0.0	74.9	12.1	71.5	72.5	80.3	1.0	0.617	0.0	74.5	13.1	71.1	72.2	79	1.0	0.475	0.0	66.9	26.3	61.8	67.2	67	1.0	0.471	0.0	66.8	26.6	61.7	67.1	66
87.4	75.0	75.6	1.0	0.75	0.0	80.5	3.4	78.0	78.1	87.4	1.0	0.75	0.0	80.6	3.5	78.1	78.1	87	1.0	0.565	0.0	71.7	18.2	67.8	70.1	75	1.0	0.572	0.0	72.1	17.5	68.2	70.4	75
92.5	82.5	83.9	1.0	0.875	0.0	85.4	-3.7	84.0	84.0	92.5	1.0	0.867	0.0	85.1	-3.2	83.6	83.7	92	1.0	0.654	0.0	76.3	10.3	73.2	73.9	82	1.0	0.679	0.0	77.4	8.6	74.5	75.0	83
96.0	90.0	92.3	1.0	1.0	0.0	89.4	-9.5	89.0	89.6	96.0	1.0	1.0	0.0	89.4	-9.4	89.1	89.6	96	1.0	0.812	0.0	83.0	0.0	81.1	81.1	90	1.0	0.868	0.0	85.2	-3.3	83.7	83.8	92
99.5	97.5	101.0	0.875	1.0	0.0	86.7	-13.9	82.7	83.8	99.5	0.883	1.0	0.0	86.9	-13.6	83.1	84.3	99	0.967	1.0	0.0	88.7	-10.6	87.4	88.1	97	0.842	1.0	0.0	85.9	-14.9	81.3	82.6	100
102.9	105.0	109.7	0.75	1.0	0.0	83.7	-17.7	77.1	79.2	102.9	0.75	1.0	0.0	83.7	-17.7	77.2	79.2	102	0.7	1.0	0.0	81.4	-20.0	74.9	77.5	105	0.598	1.0	0.0	77.0	-24.8	69.2	73.5	109
107.9	112.5	118.5	0.625	1.0	0.0	77.9	-23.1	71.3	75.0	107.9	0.633	1.0	0.0	78.4	-22.7	71.7	75.3	107	0.566	1.0	0.0	75.7	-26.7	66.4	71.6	112	0.477	1.0	0.0	72.4	-31.4	59.4	67.3	111
116.4	120.0	127.2	0.5	1.0	0.0	73.1	-30.2	60.8	67.9	116.4	0.5	1.0	0.0	73.2	-30.1	60.8	67.9	116	0.445	1.0	0.0	71.3	-33.1	57.5	66.4	120	0.35	1.0	0.0	67.3	-38.8	51.1	64.3	121
124.5	127.5	136.0	0.375	1.0	0.0	68.8	-36.5	53.0	64.4	124.5	0.383	1.0	0.0	69.1	-36.1	53.6	64.6	124	0.353	1.0	0.0	67.4	-38.6	51.3	64.3	127	0.276	1.0	0.0	62.5	-45.4	44.8	63.9	133
138.2	135.0	144.7	0.25	1.0	0.0	60.8	-47.5	42.4	63.7	138.2	0.25	1.0	0.0	60.9	-47.4	42.4	63.7	138	0.28	1.0	0.0	62.8	-45.1	45.2	63.9	135	0.176	1.0	0.0	58.4	-52.7	37.3	64.6	144
149.2	142.5	153.4	0.125	1.0	0.0	56.7	-56.1	33.3	65.2	149.2	0.133	1.0	0.0	57.0	-55.5	34.0	65.2	148	0.207	1.0	0.0	59.5	-50.5	39.6	64.2	142	0.088	1.0	0.0	55.2	-60.1	30.8	67.6	153
161.6	150.0	162.2	0.0	1.0	0.0	51.6	-69.3	23.0	73.1	161.6	0.0	1.0	0.0	51.7	-69.3	23.1	73.1	161	0.117	1.0	0.0	56.4	-56.8	32.9	65.8	150	0.0	1.0	0.011	51.7	-69.0	22.2	72.6	163
168.3	157.5	169.0	0.0	1.0	0.125	52.3	-66.1	13.6	67.5	168.3	0.0	1.0	0.117	52.3	-66.3	14.3	67.9	167	0.047	1.0	0.0	53.5	-64.5	27.4	70.2	157	0.0	1.0	0.129	52.4	-65.9	13.3	67.3	168
176.2	165.0	175.9	0.0	1.0	0.25	53.0	-61.8	4.0	61.9	176.2	0.0	1.0	0.25	53.0	-61.8	4.0	62.0	176	0.0	1.0	0.063	52.0	-67.8	18.2	70.3	165	0.0	1.0	0.244	53.0	-62.0	4.4	62.2	173
186.9	172.5	182.7	0.0	1.0	0.375	53.8	-56.5	-6.8	56.9	186.9	0.0	1.0	0.367	53.8	-56.9	-6.1	57.3	186	0.0	1.0	0.183	52.7	-64.2	9.0	65.0	172	0.0	1.0	0.321	53.5	-59.0	-2.3	59.1	181
198.8	180.0	189.6	0.0	1.0	0.5	54.6	-50.8	-17.3	53.7	198.8	0.0	1.0	0.5	54.7	-50.8	-17.2	53.7	198	0.0	1.0	0.294	53.3	-60.1	0.0	60.2	180	0.0	1.0	0.403	54.0	-55.4	-9.3	56.2	189
209.5	187.5	196.4	0.0	1.0	0.625	55.4	-45.8	-25.9	52.6	209.5	0.0	1.0	0.617	55.4	-46.1	-25.3	52.7	208	0.0	1.0	0.376	53.9	-56.4	-6.8	56.9	187	0.0	1.0	0.47	54.5	-52.3	-14.9	54.5	193
220.1	195.0	203.2	0.0	1.0	0.75	56.6	-40.0	-33.7	52.4	220.1	0.0	1.0	0.75	56.7	-40.0	-33.7	52.4	220	0.0	1.0	0.46	54.4	-52.8	-14.1	54.8	195	0.0	1.0	0.552	55.0	-48.9	-21.0	53.3	205
227.6	202.5	210.1	0.0	1.0	0.875	57.2	-36.1	-39.6	53.6	227.6	0.0	1.0	0.867	57.3	-36.3	-39.2	53.6	227	0.0	1.0	0.537	54.9	-49.4	-19.9	53.4	202	0.0	1.0	0.627	55.5	-45.7	-26.0	52.7	209
234.6	210.0	216.9	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6	0.0	1.0	1.0	57.9	-31.9	-45.0	55.3	234	0.0	1.0	0.631	55.5	-45.5	-26.2	52.7	210	0.0	1.0	0.713	56.3	-41.8	-31.5	52.5	210
238.7	217.5	223.8	0.0	0.875	1.0	54.9	-27.5	-45.3	53.0	238.7	0.0	0.883	1.0	55.1	-27.8	-45.3	53.2	238	0.0	1.0	0.713	56.3	-41.8	-31.5	52.5	217	0.0	1.0	0.804	56.9	-38.4	-36.3	52.9	223
244.0	225.0	230.6	0.0	0.75	1.0	51.3	-22.1	-45.6	50.7	244.0	0.0	0.75	1.0	51.3	-22.1	-45.5	50.7	244	0.0	1.0	0.831	57.1	-37.5	-37.5	53.2	225	0.0	1.0	0.929	57.5	-34.4	-41.9	54.4	230
250.7	232.5	237.5	0.0	0.625	1.0	47.2	-16.0	-45.9	48.7	250.7	0.0	0.633	1.0	47.5	-16.3	-45.9	48.8	250	0.0	1.0	0.952	57.7	-33.6	-43.0	54.7	232	0.0	0.927	1.0	56.1	-29.3	-45.2	54.0	237
260.4	240.0	244.3	0.0	0.5	1.0	42.3	-7.7	-46.3	46.9	260.4	0.0	0.5	1.0	42.4	-7.7	-46.2	47.0	260	0.0	0.845	1.0	54.1	-26.2	-45.4	52.5	240	0.0	0.745	1.0	51.2	-21.8	-45.6	50.6	244
270.4	247.5	251.2	0.0	0.375	1.0	37.3	0.3	-46.4	46.4	270.4	0.0	0.383	1.0	37.7	-0.1	-46.4	46.5	269	0.0	0.695	1.0	49.6	-19.4	-45.8	49.8	247	0.0	0.625	1.0	47.3	-16.0	-45.9	48.7	250
280.2	255.0	258.0	0.0	0.25	1.0	32.7	8.5	-47.0	47.8	280.2	0.0	0.25	1.0	32.8	8.5	-47.0	47.8	280	0.0	0.57	1.0	45.1	-12.3	-46.2	47.9	255	0.0	0.531	1.0	43.6	-9.7	-46.3	47.4	258
289.3	262.5	264.8	0.0	0.125	1.0	28.1	16.7	-47.6	50.4	289.3	0.0	0.133	1.0	28.4	16.2	-47.5	50.3	288	0.0	0.481	1.0	41.6	-6.4	-46.3	46.9	262	0.0	0.45	1.0	40.3	-4.4	-46.5	46.8	264
295.6	270.0	271.7	0.0	0.0	1.0	24.9	22.9	-47.8	53.0	295.6	0.0	0.0	1.0	25.0	23.0	-47.7	53.1	295	0.0	0.38	1.0	37.5	0.0	-46.4	46.5	270	0.0	0.358	1.0	36.7	1.4	-4		

Data of Maximum color M in colorimetric system Offset standard print; separation cmyn6*, D65 for input or output; Six hue angles of the 60 degree standard colours $RYGBM_s$: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours $RYGBM_d$: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours $RYGBM_e$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_{dd64M}			$LAB^*_{ddx64M} (x=LabCh)$						$rgb^*_{dex361M}$					$LAB^*_{dex361M}$						rgb^*_{dd}			rgb^*_{ds}	rgb^*_{de}
30.4	30.0	25.4	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4	30.4	1.0	0.0	0.131	47.7	66.3	31.6	73.5	25								
37.2	37.5	33.8	1.0	0.125	0.0	51.5	56.6	43.1	71.2	37.2	37.2	1.0	0.052	0.0	49.2	61.9	40.6	74.0	33								
47.2	45.0	42.1	1.0	0.25	0.0	56.6	45.8	49.4	67.4	47.2	47.2	1.0	0.187	0.0	54.1	51.4	46.6	69.4	42								
58.6	52.5	50.5	1.0	0.375	0.0	62.3	34.4	56.4	66.1	58.6	58.6	1.0	0.28	0.0	58.0	43.2	51.4	67.1	49								
69.1	60.0	58.8	1.0	0.5	0.0	68.1	24.0	63.0	67.4	69.1	69.1	1.0	0.378	0.0	62.5	34.2	56.6	66.1	58								
80.3	67.5	67.2	1.0	0.625	0.0	74.9	12.1	71.5	72.5	80.3	80.3	1.0	0.471	0.0	66.8	26.6	61.7	67.1	66								
87.4	75.0	75.6	1.0	0.75	0.0	80.5	3.4	78.0	78.1	87.4	87.4	1.0	0.572	0.0	72.1	17.5	68.2	70.4	75								
92.5	82.5	83.9	1.0	0.875	0.0	85.4	-3.7	84.0	84.0	92.5	92.5	1.0	0.679	0.0	77.4	8.6	74.5	75.0	83								
96.0	90.0	92.3	1.0	1.0	0.0	89.4	-9.5	89.0	89.6	96.0	96.0	1.0	0.868	0.0	85.2	-3.3	83.7	83.8	92								
99.5	97.5	101.0	0.875	1.0	0.0	86.7	-13.9	82.7	83.8	99.5	99.5	0.842	1.0	0.0	85.9	-14.9	81.3	82.6	100								
102.9	105.0	109.7	0.75	1.0	0.0	83.7	-17.7	77.1	79.2	102.9	102.9	0.598	1.0	0.0	77.0	-24.8	69.2	73.5	109								
107.9	112.5	118.5	0.625	1.0	0.0	77.9	-23.1	71.3	75.0	107.9	107.9	0.477	1.0	0.0	72.4	-31.4	59.4	67.3	117								
116.4	120.0	127.2	0.5	1.0	0.0	73.1	-30.2	60.8	67.9	116.4	116.4	0.35	1.0	0.0	67.3	-38.8	51.1	64.3	127								
124.5	127.5	136.0	0.375	1.0	0.0	68.8	-36.5	53.0	64.4	124.5	124.5	0.276	1.0	0.0	62.5	-45.4	44.8	63.9	135								
138.2	135.0	144.7	0.25	1.0	0.0	60.8	-47.5	42.4	63.7	138.2	138.2	0.176	1.0	0.0	58.4	-52.7	37.3	64.6	144								
149.2	142.5	153.4	0.125	1.0	0.0	56.7	-56.1	33.3	65.2	149.2	149.2	0.088	1.0	0.0	55.2	-60.1	30.8	67.6	152								
161.6	150.0	162.2	0.0	1.0	0.0	51.6	-69.3	23.0	73.1	161.6	161.6	0.0	1.0	0.011	51.7	-69.0	22.2	72.6	162								
168.3	157.5	169.0	0.0	1.0	0.125	52.3	-66.1	13.6	67.5	168.3	168.3	0.0	1.0	0.129	52.4	-65.9	13.3	67.3	168								
176.2	165.0	175.9	0.0	1.0	0.25	53.0	-61.8	4.0	61.9	176.2	176.2	0.0	1.0	0.244	53.0	-62.0	4.4	62.2	175								
186.9	172.5	182.7	0.0	1.0	0.375	53.8	-56.5	-6.8	56.9	186.9	186.9	0.0	1.0	0.321	53.5	-59.0	-2.3	59.1	182								
198.8	180.0	189.6	0.0	1.0	0.5	54.6	-50.8	-17.3	53.7	198.8	198.8	0.0	1.0	0.403	54.0	-55.4	-9.3	56.2	189								
209.5	187.5	196.4	0.0	1.0	0.625	55.4	-45.8	-25.9	52.6	209.5	209.5	0.0	1.0	0.47	54.5	-52.3	-14.9	54.5	195								
220.1	195.0	203.2	0.0	1.0	0.75	56.6	-40.0	-33.7	52.4	220.1	220.1	0.0	1.0	0.552	55.0	-48.9	-21.0	53.3	203								
227.6	202.5	210.1	0.0	1.0	0.875	57.2	-36.1	-39.6	53.6	227.6	227.6	0.0	1.0	0.627	55.5	-45.7	-26.0	52.7	209								
234.6	210.0	216.9	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6	234.6	0.0	1.0	0.713	56.3	-41.8	-31.5	52.5	216								
238.7	217.5	223.8	0.0	0.875	1.0	54.9	-27.5	-45.3	53.0	238.7	238.7	0.0	1.0	0.804	56.9	-38.4	-36.3	52.9	223								
244.0	225.0	230.6	0.0	0.75	1.0	51.3	-22.1	-45.6	50.7	244.0	244.0	0.0	1.0	0.929	57.5	-34.4	-41.9	54.4	230								
250.7	232.5	237.5	0.0	0.625	1.0	47.2	-16.0	-45.9	48.7	250.7	250.7	0.0	0.927	1.0	56.1	-29.3	-45.2	54.0	237								
260.4	240.0	244.3	0.0	0.5	1.0	42.3	-7.7	-46.3	46.9	260.4	260.4	0.0	0.745	1.0	51.2	-21.8	-45.6	50.6	244								
270.4	247.5	251.2	0.0	0.375	1.0	37.3	0.3	-46.4	46.4	270.4	270.4	0.0	0.625	1.0	47.3	-16.0	-45.9	48.7	250								
280.2	255.0	258.0	0.0	0.25	1.0	32.7	8.5	-47.0	47.8	280.2	280.2	0.0	0.531	1.0	43.6	-9.7	-46.3	47.4	258								
289.3	262.5	264.8	0.0	0.125	1.0	28.1	16.7	-47.6	50.4	289.3	289.3	0.0	0.45	1.0	40.3	-4.4	-46.5	46.8	264								
295.6	270.0	271.7	0.0	0.0	1.0	24.9	22.9	-47.8	53.0	295.6	295.6	0.0	0.358	1.0	36.7	1.4	-46.5	46.7	271								
305.9	277.5	278.8	0.125	0.0	1.0	27.8	31.4	-43.4	53.6	305.9	305.9	0.0	0.274	1.0	33.7	6.9	-47.0	47.6	278								
311.7	285.0	285.9	0.25	0.0	1.0	29.9	36.0	-40.4	54.1	311.7	311.7	0.0	0.172	1.0	29.9	13.6	-47.5	49.5	285								
325.9	292.5	293.0	0.375	0.0	1.0	33.7	47.7	-32.2	57.5	325.9	325.9	0.0	0.061	1.0	26.5	19.9	-47.7	51.8	292								
333.2	300.0	300.1	0.5	0.0	1.0	37.0	53.9	-27.1	60.4	333.2	333.2	0.055	0.0	1.0	26.3	26.8	-46.0	53.3	300								
339.6	307.5	307.2	0.625	0.0	1.0	40.2	59.7	-22.1	63.7	339.6	339.6	0.144	0.0	1.0	28.2	32.2	-42.9	53.7	306								
346.7	315.0	314.3	0.75	0.0	1.0	43.3	66.7	-15.7	68.5	346.7	346.7	0.273	0.0	1.0	30.7	38.3	-39.1	54.8	314								
350.3	322.5	321.4	0.875	0.0	1.0	45.9	70.7	-12.0	71.7	350.3	350.3	0.332	0.0	1.0	32.5	43.9	-35.4	56.4	321								
353.2	330.0	328.6	1.0	0.0	1.0	48.2	74.2	-8.7	74.7	353.2	353.2	0.42	0.0	1.0	35.0	50.0	-30.4	58.6	328								
356.1	337.5	335.7	1.0	0.0	0.875	48.2	73.1	-4.9	73.3	356.1	356.1	0.538	0.0	1.0	38.1	55.8	-25.6	61.4	335								
359.3	345.0	342.8	1.0	0.0	0.75	48.1	72.1	-0.7	72.1	359.3	359.3	0.681	0.0	1.0	41.6	63.0	-19.4	65.9	342								
364.0	352.5	349.9	1.0	0.0	0.625	48.0	70.7	4.9	70.9	364.0	364.0	0.844	0.0	1.0	45.3	69.7	-12.9	70.9	349								
369.2	360.0	357.0	1.0	0.0	0.5	47.8	69.7	11.3	70.6	369.2	369.2	0.949	0.0	1.0	47.3	72.8	-10.1	73.5	352								
375.0	367.5	364.1	1.0	0.0	0.375	47.8	68.2	18.3	70.6	375.0	375.0	1.0	0.0	0.737	48.1	72.0	-0.1	72.0	359								
380.8	375.0	371.2	1.0	0.0	0.25	47.8	67.0	25.4	71.7	380.8	380.8	1.0	0.0	0.512	47.9	69.8	10.8	70.7	368								
385.7	382.5	378.3	1.0	0.0	0.125	47.6	66.2	31.9	73.5	385.7	385.7	1.0	0.0	0.342	47.9	68.0	20.2	70.9	376								
390.4	390.0	385.4	1.0	0.0	0.0	47.5	65.5	38.4	76.0	390.4	390.4	1.0	0.0	0.131	47.7	66.3	31.6	73.5	385								

1-003830-L0 SE140-70 LAB*la0, YN=0%, XYZnw=2.5, 2.6, 2.7, 86.0, 90.9, 95.9, LAB*nmw=18.5, 0.0, 0.0, 96.4, 0.0, 0.0

TUB-test chart SE14; 1080 colours, offset standard paper
48 step hue circles; $rgb-LabCh^*$ tables, 3D=0, de=0, $cmYk$

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

Output: Offset standard print; separation cmyn6*, D65, page 9/33

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output, separation cmyn6 (CMYK)
TUB material: code=rha4ta

Data of Maximum color M in colorimetric system Offset standard print; separation cmy6*, D65 for input or output; Six hue angles of the 60 degree standard colours $RYGCBM_s$: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;

Six hue angles of the device colours $RYGCBM_d$: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours $RYGCBM_e$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_{dd361M}	$LAB^*_{ddx361Mi}(x=LabCh)$	R_d	$rgb^*_{ds361Mi}$	$LAB^*_{dsx361Mi}(x=LabCh)$	R_s	$rgb^*_{dd361Mi}$	$LAB^*_{dex361Mi}(x=LabCh)$	R_e	$rgb^*_{dd361Mi}$	rgb^*_{dd}	rgb^*_{ds}	rgb^*_{de}
30	30	25	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30	1.0	0.0	0.0	0.0	0.0
31	31	26	1.0	0.016	0.0	48.0	64.4	39.2	75.4	31	1.0	0.017	0.0		
32	32	27	1.0	0.033	0.0	48.5	63.2	39.8	74.7	32	1.0	0.033	0.0		
33	33	28	1.0	0.05	0.0	49.1	62.0	40.5	74.1	33	1.0	0.05	0.0		
34	34	29	1.0	0.066	0.0	49.6	60.8	41.1	73.4	34	1.0	0.067	0.0		
34	35	31	1.0	0.083	0.0	50.2	59.6	41.7	72.8	34	1.0	0.083	0.0		
35	36	32	1.0	0.1	0.0	50.7	58.4	42.3	72.1	35	1.0	0.1	0.0		
36	37	33	1.0	0.116	0.0	51.2	57.2	42.8	71.5	36	1.0	0.117	0.0		
37	38	34	1.0	0.133	0.0	51.8	55.9	43.6	70.9	37	1.0	0.133	0.0		
39	39	35	1.0	0.15	0.0	52.5	54.5	44.5	70.4	39	1.0	0.15	0.0		
40	40	36	1.0	0.166	0.0	53.2	53.1	45.5	69.9	40	1.0	0.167	0.0		
41	41	37	1.0	0.183	0.0	53.9	51.7	46.3	69.4	41	1.0	0.183	0.0		
43	42	38	1.0	0.2	0.0	54.5	50.2	47.2	68.9	43	1.0	0.2	0.0		
44	43	39	1.0	0.216	0.0	55.2	48.7	48.0	68.4	44	1.0	0.217	0.0		
45	44	41	1.0	0.233	0.0	55.9	47.3	48.7	67.9	45	1.0	0.233	0.0		
47	45	42	1.0	0.25	0.0	56.6	45.8	49.4	67.4	47	1.0	0.25	0.0		
48	46	43	1.0	0.266	0.0	57.3	44.3	50.5	67.2	48	1.0	0.267	0.0		
50	47	44	1.0	0.283	0.0	58.1	42.8	51.5	67.0	50	1.0	0.283	0.0		
51	48	45	1.0	0.3	0.0	58.9	41.4	52.5	66.9	51	1.0	0.3	0.0		
53	49	46	1.0	0.316	0.0	59.6	39.8	53.5	66.7	53	1.0	0.317	0.0		
54	50	47	1.0	0.333	0.0	60.4	38.3	54.3	66.5	54	1.0	0.333	0.0		
56	51	48	1.0	0.35	0.0	61.2	36.7	55.2	66.3	56	1.0	0.35	0.0		
57	52	49	1.0	0.366	0.0	62.0	35.2	56.0	66.2	57	1.0	0.367	0.0		
59	53	51	1.0	0.383	0.0	62.7	33.7	56.9	66.2	59	1.0	0.383	0.0		
60	54	52	1.0	0.4	0.0	63.5	32.4	57.9	66.3	60	1.0	0.4	0.0		
62	55	53	1.0	0.416	0.0	64.2	31.1	58.8	66.5	62	1.0	0.417	0.0		
63	56	54	1.0	0.433	0.0	65.0	29.7	59.7	66.7	63	1.0	0.433	0.0		
64	57	55	1.0	0.45	0.0	65.8	28.3	60.6	66.9	64	1.0	0.45	0.0		
66	58	56	1.0	0.466	0.0	66.5	26.9	61.4	67.0	66	1.0	0.467	0.0		
67	59	57	1.0	0.483	0.0	67.3	25.4	62.2	67.2	67	1.0	0.483	0.0		
69	60	58	1.0	0.5	0.0	68.1	24.0	63.0	67.4	69	1.0	0.5	0.0		
70	61	60	1.0	0.516	0.0	69.0	22.5	64.2	68.1	70	1.0	0.517	0.0		
72	62	61	1.0	0.533	0.0	69.9	21.1	65.5	68.8	72	1.0	0.533	0.0		
73	63	62	1.0	0.55	0.0	70.8	19.6	66.6	69.5	73	1.0	0.55	0.0		
75	64	63	1.0	0.566	0.0	71.7	18.0	67.8	70.1	75	1.0	0.567	0.0		
76	65	64	1.0	0.583	0.0	72.6	16.4	68.9	70.8	76	1.0	0.583	0.0		
78	66	65	1.0	0.6	0.0	73.6	14.7	70.0	71.5	78	1.0	0.6	0.0		
79	67	66	1.0	0.616	0.0	74.5	13.0	71.0	72.2	79	1.0	0.617	0.0		
80	68	67	1.0	0.633	0.0	75.3	11.6	72.0	72.9	80	1.0	0.633	0.0		
81	69	68	1.0	0.65	0.0	76.0	10.5	72.9	73.6	81	1.0	0.65	0.0		
82	70	70	1.0	0.666	0.0	76.8	9.4	73.8	74.4	82	1.0	0.667	0.0		
83	71	71	1.0	0.683	0.0	77.5	8.3	74.7	75.1	83	1.0	0.683	0.0		
84	72	72	1.0	0.7	0.0	78.3	7.1	75.5	75.9	84	1.0	0.7	0.0		
85	73	73	1.0	0.716	0.0	79.0	5.9	76.4	76.6	85	1.0	0.717	0.0		
86	74	74	1.0	0.733	0.0	79.8	4.7	77.2	77.3	86	1.0	0.733	0.0		
87	75	75	1.0	0.75	0.0	80.5	3.4	78.0	78.1	87	1.0	0.75	0.0		

1-003930-L0 SE140-70 LAB*la0, YN=0%, XYZnw=2.5, 2.6, 2.7, 86.0, 90.9, 95.9, LAB*nmw=18.5, 0.0, 0.0, 96.4, 0.0, 0.0

Output: Offset standard print; separation cmy6*, D65, page 10/33

TUB-test chart SE14; 1080 colours, offset standard paper
48 step hue circles; $rgb-LabCh$ *tables, 3D=0, de=0, $cmyk$

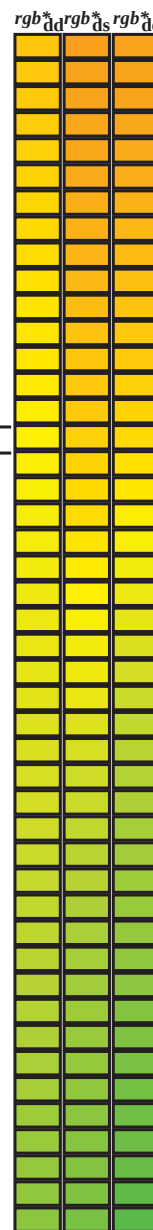
input: $rgb/cmyk \rightarrow rgb_d$
output: transfer to $cmyk_d$

TUB registration: 20130201-SE14/SE14L0NP.PDF /PS
application for measurement of offset print output, separationcmy6 (CMYK)
TUB material: code=rha4ta

Data of Maximum color M in colorimetric system Offset standard print; separation cmykn6*, D65 for input or output; Six hue angles of the 60 degree standard colours $RYGCBM_c$: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;

Six hue angles of the device colours $RYGCBM_d$: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours $RYGCBM_c$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_{dd361M}	$LAB^*_{dd361MI} (x=LabCh)$	$rgb^*_{ds361Mi}$	$LAB^*_{dsx361MI} (x=LabCh)$	$rgb^*_{dd361Mi}$	$rgb^*_{de361Mi}$	$LAB^*_{dex361MI} (x=LabCh)$	$rgb^*_{dd361Mi}$	Y_d	Y_s	Y_e												
87	75	75	1.0	0.75 0.0	80.5	3.4	78.0	78.1	87	1.0	0.565 0.0	71.7	18.2	67.8	70.1	75	1.0	0.75 0.0	72.1	17.5	68.2	70.4	75	1.0	0.75 0.0
88	76	76	1.0	0.766 0.0	81.2	2.5	78.8	78.9	88	1.0	0.577 0.0	72.3	17.1	68.5	70.6	76	1.0	0.767 0.0	72.8	16.3	69.0	70.9	76	1.0	0.767 0.0
88	77	77	1.0	0.783 0.0	81.8	1.6	79.7	79.7	88	1.0	0.588 0.0	72.9	16.0	69.2	71.1	77	1.0	0.783 0.0	73.5	15.1	69.8	71.4	77	1.0	0.783 0.0
89	78	78	1.0	0.8 0.0	82.4	0.6	80.5	80.5	89	1.0	0.599 0.0	73.6	14.9	70.0	71.5	78	1.0	0.8 0.0	74.1	13.8	70.6	72.0	78	1.0	0.8 0.0
90	79	80	1.0	0.816 0.0	83.1	-0.2	81.3	81.3	90	1.0	0.61 0.0	74.2	13.7	70.7	72.0	79	1.0	0.817 0.0	74.8	12.5	71.4	72.5	80	1.0	0.817 0.0
90	80	81	1.0	0.833 0.0	83.7	-1.2	82.0	82.1	90	1.0	0.621 0.0	74.8	12.6	71.3	72.4	80	1.0	0.833 0.0	75.6	11.2	72.4	73.2	81	1.0	0.833 0.0
91	81	82	1.0	0.85 0.0	84.4	-2.2	82.8	82.8	91	1.0	0.637 0.0	75.5	11.4	72.2	73.1	81	1.0	0.85 0.0	76.5	9.9	73.4	74.1	82	1.0	0.85 0.0
92	82	83	1.0	0.866 0.0	85.0	-3.2	83.6	83.6	92	1.0	0.654 0.0	76.3	10.3	73.2	73.9	82	1.0	0.867 0.0	77.4	8.6	74.5	75.0	83	1.0	0.867 0.0
92	83	84	1.0	0.883 0.0	85.6	-4.1	84.3	84.4	92	1.0	0.672 0.0	77.1	9.1	74.1	74.7	83	1.0	0.883 0.0	78.3	7.2	75.5	75.8	84	1.0	0.883 0.0
93	84	85	1.0	0.9 0.0	86.2	-4.8	85.0	85.1	93	1.0	0.689 0.0	77.9	7.9	75.0	75.4	84	1.0	0.9 0.0	79.1	5.8	76.5	76.7	85	1.0	0.9 0.0
93	85	86	1.0	0.916 0.0	86.7	-5.6	85.7	85.9	93	1.0	0.707 0.0	78.6	6.6	75.9	76.2	85	1.0	0.917 0.0	80.0	4.4	77.5	77.6	86	1.0	0.917 0.0
94	86	87	1.0	0.933 0.0	87.2	-6.3	86.4	86.6	94	1.0	0.725 0.0	79.4	5.4	76.8	77.0	86	1.0	0.933 0.0	80.9	2.9	78.5	78.6	87	1.0	0.933 0.0
94	87	88	1.0	0.95 0.0	87.8	-7.1	87.1	87.3	94	1.0	0.742 0.0	80.2	4.1	77.7	77.8	87	1.0	0.95 0.0	82.0	1.4	79.9	79.9	88	1.0	0.95 0.0
95	88	90	1.0	0.966 0.0	88.3	-7.9	87.7	88.1	95	1.0	0.763 0.0	81.1	2.7	78.7	78.8	88	1.0	0.967 0.0	83.0	0.0	81.2	81.2	90	1.0	0.967 0.0
95	89	91	1.0	0.983 0.0	88.8	-8.7	88.4	88.8	95	1.0	0.788 0.0	82.0	1.4	79.9	79.9	89	1.0	0.983 0.0	84.1	-1.6	82.5	82.5	91	1.0	0.983 0.0
96	90	92	1.0	1.0 0.0	89.4	-9.5	89.0	89.6	96	1.0	0.812 0.0	83.0	0.0	81.1	81.1	90	1.0	1.0 0.0	85.2	-3.3	83.7	83.8	92	1.0	1.0 0.0
96	91	93	0.983	1.0 0.0	89.0	-10.1	88.2	88.8	96	1.0	0.836 0.0	83.9	-1.3	82.2	82.2	91	0.983	1.0 0.0	86.4	-5.1	85.3	85.5	93	0.983	1.0 0.0
97	92	94	0.966	1.0 0.0	88.6	-10.7	87.4	88.0	97	1.0	0.861 0.0	84.9	-2.8	83.4	83.4	92	0.967	1.0 0.0	87.8	-7.0	87.0	87.3	94	0.967	1.0 0.0
97	93	95	0.95	1.0 0.0	88.3	-11.3	86.5	87.3	97	1.0	0.89 0.0	85.9	-4.3	84.6	84.7	93	0.95	1.0 0.0	89.1	-8.9	88.7	89.2	95	0.95	1.0 0.0
97	94	96	0.933	1.0 0.0	87.9	-11.9	85.7	86.5	97	1.0	0.925 0.0	87.0	-5.9	86.1	86.3	94	0.933	1.0 0.0	88.7	-10.6	87.5	88.1	96	0.933	1.0 0.0
98	95	98	0.916	1.0 0.0	87.6	-12.5	84.8	85.7	98	1.0	0.961 0.0	88.2	-7.6	87.6	87.9	95	0.917	1.0 0.0	89.6	-12.1	85.3	86.2	98	0.917	1.0 0.0
98	96	99	0.9	1.0 0.0	87.2	-13.0	84.0	85.0	98	1.0	0.997 0.0	89.3	-9.3	89.0	89.5	96	0.9	1.0 0.0	88.4	-13.5	83.2	84.3	99	0.9	1.0 0.0
99	97	100	0.883	1.0 0.0	86.9	-13.6	83.1	84.2	99	0.967	1.0 0.0	88.7	-10.6	87.4	88.1	97	0.883	1.0 0.0	85.9	-14.9	81.3	82.6	100	0.883	1.0 0.0
99	98	101	0.866	1.0 0.0	86.5	-14.2	82.3	83.5	99	0.931	1.0 0.0	87.9	-11.9	85.6	86.4	98	0.867	1.0 0.0	84.9	-16.2	79.4	81.0	101	0.867	1.0 0.0
100	99	102	0.85	1.0 0.0	86.1	-14.7	81.6	82.9	100	0.895	1.0 0.0	87.2	-13.2	83.7	84.8	99	0.85	1.0 0.0	83.9	-17.5	77.5	79.5	102	0.85	1.0 0.0
100	100	103	0.833	1.0 0.0	85.7	-15.2	80.8	82.3	100	0.859	1.0 0.0	86.3	-14.4	82.0	83.3	100	0.833	1.0 0.0	82.6	-18.8	76.1	78.4	103	0.833	1.0 0.0
101	101	105	0.816	1.0 0.0	85.3	-15.8	80.1	81.6	101	0.822	1.0 0.0	85.5	-15.5	80.4	81.9	101	0.817	1.0 0.0	81.3	-20.1	74.7	77.4	105	0.817	1.0 0.0
101	102	106	0.8	1.0 0.0	84.9	-16.3	79.4	81.0	101	0.786	1.0 0.0	84.6	-16.6	78.8	80.5	102	0.8	1.0 0.0	79.9	-21.3	73.4	76.4	106	0.8	1.0 0.0
102	103	107	0.783	1.0 0.0	84.5	-16.8	78.6	80.4	102	0.75	1.0 0.0	83.7	-17.7	77.2	79.2	103	0.783	1.0 0.0	78.6	-22.5	72.0	75.5	107	0.783	1.0 0.0
102	104	108	0.766	1.0 0.0	84.1	-17.3	77.9	79.8	102	0.725	1.0 0.0	82.5	-18.9	76.0	78.4	104	0.767	1.0 0.0	77.6	-23.7	70.6	74.5	108	0.767	1.0 0.0
102	105	109	0.75	1.0 0.0	83.7	-17.7	77.1	79.2	102	0.7	1.0 0.0	81.4	-20.0	74.9	77.5	105	0.75	1.0 0.0	77.0	-24.8	69.2	73.5	109	0.75	1.0 0.0
103	106	110	0.733	1.0 0.0	82.9	-18.5	76.4	78.6	103	0.675	1.0 0.0	80.3	-21.0	73.7	76.7	106	0.733	1.0 0.0	76.3	-25.8	67.7	72.5	110	0.733	1.0 0.0
104	107	112	0.716	1.0 0.0	82.1	-19.3	75.6	78.0	104	0.65	1.0 0.0	79.1	-22.1	72.5	75.9	107	0.717	1.0 0.0	75.6	-26.8	66.3	71.5	112	0.717	1.0 0.0
104	108	113	0.7	1.0 0.0	81.4	-20.0	74.8	77.5	104	0.625	1.0 0.0	78.0	-23.1	71.3	75.0	108	0.7	1.0 0.0	75.0	-27.8	64.8	70.6	113	0.7	1.0 0.0
105	109	114	0.683	1.0 0.0	80.6	-20.7	74.1	76.9	105	0.61	1.0 0.0	77.4	-24.0	70.1	74.2	109	0.683	1.0 0.0	74.3	-28.7	63.3	69.6	114	0.683	1.0 0.0
106	110	115	0.666	1.0 0.0	79.8	-21.4	73.3	76.4	106	0.595	1.0 0.0	76.8	-25.0	68.9	73.3	110	0.667	1.0 0.0	73.6	-29.6	61.8	68.6	115	0.667	1.0 0.0
106	111	116	0.65	1.0 0.0	79.1	-22.1	72.5	75.8	106	0.58	1.0 0.0	76.3	-25.9	67.7	72.5	111	0.65	1.0 0.0	73.0	-30.4	60.5	67.8	116	0.65	1.0 0.0
107	112	117	0.633	1.0 0.0	78.3	-22.8	71.7	75.2	107	0.566	1.0 0.0	75.7	-26.7	66.4	71.6	112	0.633	1.0 0.0	72.4	-31.4	59.4	67.3	117	0.633	1.0 0.0
108	113	119	0.616	1.0 0.0	77.6	-23.7	70.6	74.5	108	0.551	1.0 0.0	75.1	-27.6	65.2	70.8	113	0.617	1.0 0.0	71.8	-32.4	58.3	66.8	119	0.617	1.0 0.0
109	114	120	0.6	1.0 0.0	77.0	-24.7	69.2	73.5	109	0.536	1.0 0.0	74.6	-28.4	63.9	70.0	114	0.6	1.0 0.0	71.1	-33.3	57.2	66.3	120	0.6	1.0 0.0
110	115	121	0.583	1.0 0.0	76.3	-25.8	67.9	72.6	110	0.521	1.0 0.0	74.0	-29.1	62.6	69.1	115	0.583	1.0 0.0	70.5	-34.2	56.1	65.8	121	0.583	1.0 0.0
111	116	122	0.566	1.0 0.0	75.7	-26.7	66.5	71.7	111	0.506	1.0 0.0	73.4	-29.8	61.4	68.3	116	0.567	1.0 0.0	69.9	-35.1	55.0	65.3	122	0.567	1.0 0.0
113	117	123	0.55	1.0 0.0	75.1	-27.6	65.1	70.7	113	0.491	1.0 0.0	72.9	-30.6	60.3	67.7	117	0.55	1.0 0.0	69.3	-35.9	53.8	64.8	123	0.55	1.0 0.0
114	118	124	0.533	1.0 0.0	74.4	-28.5	63.6	69.8	114	0.476	1.0 0.0	72.3	-31.5	59.4	67.2	118	0.533	1.0 0.0	68.6	-36.8	52.8	64.4	124	0.533	1.0 0.0
115	119	126	0.516	1.0																					



1-0031030-L0 SE140-70 LAB*la0, YN=0%, XYZnw=2.5, 2.6, 2.7, 86.0, 90.9, 95.9, LAB*nmw=18.5, 0.0, 0.0, 96.4, 0.0, 0.0

Output: Offset standard print; separation cmykn6*, D65, page 11/33

TUB-test chart SE14; 1080 colours, offset standard paper
48 step hue circles; $rgb-LabCh^*$ tables, 3D=0, de=0, $cmyk$

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

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or

Data of Maximum color M in colorimetric system Offset standard print; separation cmykn6*, D65 for input or output; Six hue angles of the 60 degree standard colours $RYGCBM_c$: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours $RYGCBM_d$: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours $RYGCBM_c$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

h _{ab,d} h _{ab,s} h _{ab,e}			rgb* _{dd361M}				LAB* _{dd361Mi} (x=LabCh)				rgb* _{ds361Mi}				LAB* _{dsx361Mi} (x=LabCh)				rgb* _{dd361Mi}				rgb* _{de361Mi}				LAB* _{dex361Mi} (x=LabCh)				rgb* _{dd361Mi}				rgb* _{ds361Mi}				rgb* _{de}			
116	120	127	0.5	1.0	0.0	73.1	-30.2	60.8	67.9	116	0.445	1.0	0.0	71.3	-33.1	57.5	66.4	120	0.5	1.0	0.0	0.35	1.0	0.0	67.3	-38.8	51.1	64.3	127	0.5	1.0	0.0										
117	121	128	0.483	1.0	0.0	72.6	-31.1	59.8	67.4	117	0.43	1.0	0.0	70.8	-33.9	56.5	65.9	121	0.483	1.0	0.0	0.34	1.0	0.0	66.6	-39.8	50.3	64.2	128	0.483	1.0	0.0										
118	122	129	0.466	1.0	0.0	72.0	-32.0	58.8	66.9	118	0.415	1.0	0.0	70.2	-34.6	55.6	65.5	122	0.467	1.0	0.0	0.329	1.0	0.0	65.9	-40.8	49.4	64.2	129	0.467	1.0	0.0										
119	123	130	0.45	1.0	0.0	71.4	-32.9	57.7	66.5	119	0.399	1.0	0.0	69.7	-35.3	54.6	65.1	123	0.45	1.0	0.0	0.319	1.0	0.0	65.2	-41.7	48.5	64.1	130	0.45	1.0	0.0										
120	124	131	0.433	1.0	0.0	70.8	-33.7	56.7	66.0	120	0.384	1.0	0.0	69.2	-36.1	53.6	64.7	124	0.433	1.0	0.0	0.308	1.0	0.0	64.6	-42.7	47.6	64.0	131	0.433	1.0	0.0										
121	125	133	0.416	1.0	0.0	70.2	-34.6	55.6	65.5	121	0.371	1.0	0.0	68.6	-36.8	52.7	64.4	125	0.417	1.0	0.0	0.297	1.0	0.0	63.9	-43.6	46.7	64.0	133	0.417	1.0	0.0										
122	126	134	0.4	1.0	0.0	69.7	-35.4	54.6	65.1	122	0.362	1.0	0.0	68.0	-37.7	52.0	64.3	126	0.4	1.0	0.0	0.287	1.0	0.0	63.2	-44.5	45.8	63.9	134	0.4	1.0	0.0										
124	127	135	0.383	1.0	0.0	69.1	-36.1	53.5	64.6	124	0.353	1.0	0.0	67.4	-38.6	51.3	64.3	127	0.383	1.0	0.0	0.276	1.0	0.0	62.5	-45.4	44.8	63.9	135	0.383	1.0	0.0										
125	128	136	0.366	1.0	0.0	68.3	-37.3	52.3	64.3	125	0.344	1.0	0.0	66.9	-39.4	50.6	64.2	128	0.367	1.0	0.0	0.265	1.0	0.0	61.8	-46.2	43.8	63.8	136	0.367	1.0	0.0										
127	129	137	0.35	1.0	0.0	67.2	-38.9	51.1	64.2	127	0.335	1.0	0.0	66.3	-40.3	49.9	64.2	129	0.35	1.0	0.0	0.255	1.0	0.0	61.2	-47.1	42.9	63.7	137	0.35	1.0	0.0										
129	130	138	0.333	1.0	0.0	66.1	-40.5	49.7	64.1	129	0.326	1.0	0.0	65.7	-41.1	49.1	64.1	130	0.333	1.0	0.0	0.243	1.0	0.0	60.6	-48.0	41.9	63.8	138	0.333	1.0	0.0										
130	131	140	0.316	1.0	0.0	65.1	-42.0	48.3	64.0	130	0.316	1.0	0.0	65.1	-41.9	48.4	64.1	131	0.317	1.0	0.0	0.229	1.0	0.0	60.2	-49.0	41.0	64.0	140	0.317	1.0	0.0										
132	132	141	0.3	1.0	0.0	64.0	-43.4	46.9	63.9	132	0.307	1.0	0.0	64.5	-42.7	47.6	64.0	132	0.3	1.0	0.0	0.216	1.0	0.0	59.7	-49.9	40.1	64.1	141	0.3	1.0	0.0										
134	133	142	0.283	1.0	0.0	63.0	-44.8	45.4	63.8	134	0.298	1.0	0.0	63.9	-43.5	46.8	64.0	133	0.283	1.0	0.0	0.203	1.0	0.0	59.3	-50.9	39.2	64.3	142	0.283	1.0	0.0										
136	134	143	0.266	1.0	0.0	61.9	-46.2	43.9	63.8	136	0.289	1.0	0.0	63.4	-44.3	46.0	63.9	134	0.267	1.0	0.0	0.19	1.0	0.0	58.9	-51.8	38.3	64.5	143	0.267	1.0	0.0										
138	135	144	0.25	1.0	0.0	60.8	-47.5	42.4	63.7	138	0.28	1.0	0.0	62.8	-45.1	45.2	63.9	135	0.25	1.0	0.0	0.176	1.0	0.0	58.4	-52.7	37.3	64.6	144	0.25	1.0	0.0										
139	136	145	0.233	1.0	0.0	60.3	-48.7	41.3	63.9	139	0.271	1.0	0.0	62.2	-45.8	44.3	63.8	136	0.233	1.0	0.0	0.163	1.0	0.0	58.0	-53.6	36.3	64.8	145	0.233	1.0	0.0										
141	137	147	0.216	1.0	0.0	59.7	-49.9	40.1	64.1	141	0.262	1.0	0.0	61.6	-46.5	43.5	63.8	137	0.217	1.0	0.0	0.15	1.0	0.0	57.6	-54.4	35.3	65.0	147	0.217	1.0	0.0										
142	138	148	0.2	1.0	0.0	59.2	-51.1	39.0	64.3	142	0.252	1.0	0.0	61.0	-47.3	42.6	63.7	138	0.2	1.0	0.0	0.137	1.0	0.0	57.1	-55.3	34.3	65.1	148	0.2	1.0	0.0										
144	139	149	0.183	1.0	0.0	58.6	-52.3	37.8	64.5	144	0.242	1.0	0.0	60.6	-48.1	41.9	63.8	139	0.183	1.0	0.0	0.123	1.0	0.0	56.7	-56.2	33.3	65.4	149	0.183	1.0	0.0										
145	140	150	0.166	1.0	0.0	58.1	-53.4	36.5	64.7	145	0.23	1.0	0.0	60.2	-48.9	41.1	64.0	140	0.167	1.0	0.0	0.112	1.0	0.0	56.2	-57.5	32.5	66.1	150	0.167	1.0	0.0										
147	141	151	0.15	1.0	0.0	57.5	-54.5	35.3	64.9	147	0.219	1.0	0.0	59.8	-49.7	40.3	64.1	141	0.15	1.0	0.0	0.1	1.0	0.0	55.7	-58.8	31.7	66.9	151	0.15	1.0	0.0										
148	142	152	0.133	1.0	0.0	57.0	-55.5	34.0	65.1	148	0.207	1.0	0.0	59.5	-50.5	39.6	64.2	142	0.133	1.0	0.0	0.088	1.0	0.0	55.2	-60.1	30.8	67.6	152	0.133	1.0	0.0										
150	143	154	0.116	1.0	0.0	56.3	-57.0	32.8	65.8	150	0.196	1.0	0.0	59.1	-51.3	38.8	64.4	143	0.117	1.0	0.0	0.076	1.0	0.0	54.8	-61.3	29.9	68.3	154	0.117	1.0	0.0										
151	144	155	0.1	1.0	0.0	55.7	-58.8	31.6	66.8	151	0.185	1.0	0.0	58.7	-52.1	37.9	64.5	144	0.1	1.0	0.0	0.065	1.0	0.0	54.3	-62.6	28.9	69.1	155	0.1	1.0	0.0										
153	145	156	0.083	1.0	0.0	55.0	-60.6	30.4	67.8	153	0.173	1.0	0.0	58.3	-52.9	37.1	64.7	145	0.083	1.0	0.0	0.053	1.0	0.0	53.8	-63.9	27.9	69.8	156	0.083	1.0	0.0										
155	146	157	0.066	1.0	0.0	54.3	-62.4	29.1	68.9	155	0.162	1.0	0.0	58.0	-53.6	36.2	64.8	146	0.067	1.0	0.0	0.041	1.0	0.0	53.3	-65.1	26.9	70.5	157	0.067	1.0	0.0										
156	147	158	0.049	1.0	0.0	53.6	-64.2	27.7	69.9	156	0.151	1.0	0.0	57.6	-54.4	35.4	65.0	147	0.05	1.0	0.0	0.029	1.0	0.0	52.8	-66.3	25.9	71.3	158	0.05	1.0	0.0										
158	148	159	0.033	1.0	0.0	53.0	-65.9	26.2	71.0	158	0.139	1.0	0.0	57.2	-55.1	34.5	65.1	148	0.033	1.0	0.0	0.017	1.0	0.0	52.4	-67.5	24.8	72.0	159	0.033	1.0	0.0										
159	149	161	0.016	1.0	0.0	52.3	-67.7	24.6	72.0	159	0.128	1.0	0.0	56.8	-55.8	33.6	65.2	149	0.017	1.0	0.0	0.006	1.0	0.0	51.9	-68.7	23.6	72.8	161	0.017	1.0	0.0										
161	150	162	0.0	1.0	0.0	51.6	-69.3	23.0	73.1	161	G_d 0.117	1.0	0.0	56.4	-56.8	32.9	65.8	150	G_s 0.0	1.0	0.0	0.0	1.0	0.011	51.7	-69.0	22.2	72.6	162	G_c 0.0	1.0	0.0										
162	151	163	0.0	1.0	0.016	51.7	-69.0	21.7	72.3	162	0.107	1.0	0.0	56.0	-58.0	32.2	66.4	151	0.0	1.0	0.017	0.0	1.0	0.028	51.8	-68.7	20.8	71.8	163	0.0	1.0	0.017										

Data of Maximum color M in colorimetric system Offset standard print; separation cmyrn6*, D65 for input or output; Six hue angles of the 60 degree standard colours $RYGCBM_s$: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours $RYGCBM_d$: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours $RYGCBM_e$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

Lengths of the three colours (x=LabCh)										Lengths of the elementary colours (x=LabCh)										Lengths of the elementary colours (x=LabCh)										Lengths of the elementary colours (x=LabCh)									
h _{ab,d}	h _{ab,s}	h _{ab,e}	rgb* _{dd361Mi}			LAB* _{dsx361Mi} (x=LabCh)			rgb* _{ds361Mi}			LAB* _{dsx361Mi} (x=LabCh)			rgb* _{dd361Mi}			LAB* _{dex361Mi} (x=LabCh)			rgb* _{dd361Mi}			rgb* _{dd}			rgb* _{ds}			rgb* _{de}									
176	165	175	0.0	1.0	0.25	53.0	-61.8	4.0	61.9	176	0.0	1.0	0.063	52.0	-67.8	18.2	70.3	165	0.0	1.0	0.25	0.0	1.0	0.244	53.0	-62.0	4.4	62.2	175	0.0	1.0	0.25							
177	166	176	0.0	1.0	0.266	53.1	-61.2	2.4	61.3	177	0.0	1.0	0.082	52.1	-67.3	16.8	69.5	166	0.0	1.0	0.267	0.0	1.0	0.257	53.1	-61.5	3.4	61.7	176	0.0	1.0	0.267							
179	167	177	0.0	1.0	0.283	53.2	-60.6	0.9	60.6	179	0.0	1.0	0.1	52.2	-66.8	15.4	68.6	167	0.0	1.0	0.283	0.0	1.0	0.267	53.2	-61.2	2.4	61.3	177	0.0	1.0	0.283							
180	168	178	0.0	1.0	0.3	53.3	-59.9	-0.5	59.9	180	0.0	1.0	0.119	52.3	-66.2	14.1	67.8	168	0.0	1.0	0.3	0.0	1.0	0.278	53.2	-60.8	1.4	60.9	178	0.0	1.0	0.3							
181	169	179	0.0	1.0	0.316	53.4	-59.2	-2.0	59.3	181	0.0	1.0	0.136	52.4	-65.7	12.8	67.1	169	0.0	1.0	0.317	0.0	1.0	0.289	53.3	-60.3	0.5	60.4	179	0.0	1.0	0.317							
183	170	180	0.0	1.0	0.333	53.5	-58.5	-3.4	58.6	183	0.0	1.0	0.151	52.5	-65.3	11.5	66.4	170	0.0	1.0	0.333	0.0	1.0	0.299	53.4	-59.9	-0.4	60.0	180	0.0	1.0	0.333							
184	171	181	0.0	1.0	0.35	53.7	-57.7	-4.8	57.9	184	0.0	1.0	0.167	52.6	-64.8	10.3	65.7	171	0.0	1.0	0.35	0.0	1.0	0.31	53.4	-59.5	-1.3	59.6	181	0.0	1.0	0.35							
186	172	182	0.0	1.0	0.366	53.8	-56.9	-6.1	57.3	186	0.0	1.0	0.183	52.7	-64.2	9.0	65.0	172	0.0	1.0	0.367	0.0	1.0	0.321	53.5	-59.0	-2.3	59.1	182	0.0	1.0	0.367							
187	173	183	0.0	1.0	0.383	53.9	-56.2	-7.6	56.7	187	0.0	1.0	0.199	52.8	-63.7	7.8	64.3	173	0.0	1.0	0.383	0.0	1.0	0.332	53.6	-58.5	-3.2	58.7	183	0.0	1.0	0.383							
189	174	184	0.0	1.0	0.4	54.0	-55.5	-9.0	56.3	189	0.0	1.0	0.214	52.9	-63.1	6.6	63.6	174	0.0	1.0	0.4	0.0	1.0	0.342	53.7	-58.0	-4.1	58.3	184	0.0	1.0	0.4							
190	175	185	0.0	1.0	0.416	54.1	-54.8	-10.5	55.8	190	0.0	1.0	0.23	52.9	-62.5	5.5	62.9	175	0.0	1.0	0.417	0.0	1.0	0.353	53.7	-57.5	-5.0	57.9	185	0.0	1.0	0.417							
192	176	185	0.0	1.0	0.433	54.2	-54.1	-11.9	55.4	192	0.0	1.0	0.246	53.0	-61.9	4.3	62.2	176	0.0	1.0	0.433	0.0	1.0	0.364	53.8	-57.0	-5.9	57.4	185	0.0	1.0	0.433							
194	177	186	0.0	1.0	0.45	54.3	-53.3	-13.3	55.0	194	0.0	1.0	0.259	53.1	-61.5	3.2	61.6	177	0.0	1.0	0.45	0.0	1.0	0.374	53.9	-56.5	-6.7	57.0	186	0.0	1.0	0.45							
195	178	187	0.0	1.0	0.466	54.4	-52.5	-14.7	54.6	195	0.0	1.0	0.27	53.2	-61.0	2.1	61.2	178	0.0	1.0	0.467	0.0	1.0	0.384	53.9	-56.1	-7.6	56.7	187	0.0	1.0	0.467							
197	179	188	0.0	1.0	0.483	54.5	-51.7	-16.0	54.1	197	0.0	1.0	0.282	53.3	-60.6	1.1	60.7	179	0.0	1.0	0.483	0.0	1.0	0.394	54.0	-55.7	-8.4	56.5	188	0.0	1.0	0.483							
198	180	189	0.0	1.0	0.5	54.6	-50.8	-17.3	53.7	198	0.0	1.0	0.294	53.3	-60.1	0.0	60.2	180	0.0	1.0	0.5	0.0	1.0	0.403	54.0	-55.4	-9.3	56.2	189	0.0	1.0	0.5							
200	181	190	0.0	1.0	0.516	54.7	-50.2	-18.5	53.6	200	0.0	1.0	0.306	53.4	-59.7	-0.9	59.8	181	0.0	1.0	0.517	0.0	1.0	0.413	54.1	-55.0	-10.1	56.0	190	0.0	1.0	0.517							
201	182	191	0.0	1.0	0.533	54.8	-49.6	-19.7	53.4	201	0.0	1.0	0.317	53.5	-59.2	-2.0	59.3	182	0.0	1.0	0.533	0.0	1.0	0.422	54.2	-54.5	-10.9	55.7	191	0.0	1.0	0.533							
203	183	192	0.0	1.0	0.55	54.9	-49.0	-20.9	53.3	203	0.0	1.0	0.329	53.6	-58.6	-3.0	58.8	183	0.0	1.0	0.55	0.0	1.0	0.432	54.2	-54.1	-11.8	55.5	192	0.0	1.0	0.55							
204	184	193	0.0	1.0	0.566	55.0	-48.3	-22.0	53.1	204	0.0	1.0	0.341	53.6	-58.1	-4.0	58.3	184	0.0	1.0	0.567	0.0	1.0	0.442	54.3	-53.7	-12.6	55.3	193	0.0	1.0	0.567							
205	185	194	0.0	1.0	0.583	55.1	-47.6	-23.1	53.0	205	0.0	1.0	0.352	53.7	-57.6	-4.9	57.9	185	0.0	1.0	0.583	0.0	1.0	0.451	54.4	-53.2	-13.4	55.0	194	0.0	1.0	0.583							
207	186	195	0.0	1.0	0.6	55.2	-46.9	-24.3	52.8	207	0.0	1.0	0.364	53.8	-57.0	-5.9	57.4	186	0.0	1.0	0.6	0.0	1.0	0.461	54.4	-52.8	-14.1	54.8	195	0.0	1.0	0.6							
208	187	195	0.0	1.0	0.616	55.3	-46.2	-25.4	52.7	208	0.0	1.0	0.376	53.9	-56.4	-6.8	56.9	187	0.0	1.0	0.617	0.0	1.0	0.47	54.5	-52.3	-14.9	54.5	195	0.0	1.0	0.617							
210	188	196	0.0	1.0	0.633	55.5	-45.4	-26.5	52.6	210	0.0	1.0	0.386	53.9	-56.0	-7.8	56.7	188	0.0	1.0	0.633	0.0	1.0	0.48	54.5	-51.8	-15.7	54.3	196	0.0	1.0	0.633							
211	189	197	0.0	1.0	0.65	55.6	-44.7	-27.5	52.6	211	0.0	1.0	0.397	54.0	-55.6	-8.7	56.4	189	0.0	1.0	0.65	0.0	1.0	0.49	54.6	-51.3	-16.4	54.0	197	0.0	1.0	0.65							
213	190	198	0.0	1.0	0.666	55.8	-44.0	-28.6	52.5	213	0.0	1.0	0.407	54.1	-55.2	-9.6	56.1	190	0.0	1.0	0.667	0.0	1.0	0.499	54.7	-50.8	-17.2	53.8	198	0.0	1.0	0.667							
214	191	199	0.0	1.0	0.683	56.0	-43.3	-29.7	52.5	214	0.0	1.0	0.418	54.1	-54.7	-10.6	55.9	191	0.0	1.0	0.683	0.0	1.0	0.51	54.7	-50.4	-17.9	53.7	199	0.0	1.0	0.683							
215	192	200	0.0	1.0	0.7	56.1	-42.5	-30.7	52.5	215	0.0	1.0	0.428	54.2	-54.3	-11.5	55.6	192	0.0	1.0	0.7	0.0	1.0	0.52	54.8	-50.1	-18.7	53.6	200	0.0	1.0	0.7							
217	193	201	0.0	1.0	0.716	56.3	-41.7	-31.8	52.4	217	0.0	1.0	0.439	54.3	-53.8	-12.3	55.3	193	0.0	1.0	0.717	0.0	1.0	0.531	54.9	-49.7	-19.5	53.5	201	0.0	1.0	0.717							
218	194	202	0.0	1.0	0.733	56.5	-40.9	-32.8	52.4	218	0.0	1.0	0.449	54.3	-53.3	-13.2	55.1	194	0.0	1.0	0.733	0.0	1.0	0.542	54.9	-49.3	-20.2	53.4	202	0.0	1.0	0.733							
220	195	203	0.0	1.0	0.75	56.6	-40.0	-33.7	52.4	220	0.0	1.0	0.46	54.4	-52.8	-14.1	54.8	195	0.0	1.0	0.75	0.0	1.0	0.552	55.0	-48.9	-21.0	53.3	203	0.0	1.0	0.75							
221	196	204	0.0	1.0	0.766	56.7	-39.6	-34.5	52.5	221	0.0	1.0	0.471	54.5	-52.3	-14.9	54.5	196	0.0	1.0	0.767	0.0	1.0	0.563	55.1	-48.4	-21.7	53.2	204	0.0	1.0	0.767							
222	197	205	0.0	1.0	0.783	56.8	-39.1	-35.3	52.7	222	0.0	1.0	0.481	54.5	-51.8	-15.8	54.2	197	0.0	1.0	0.783	0.0	1.0	0.574	55.1	-48.0	-22.4	53.1	205	0.0	1.0	0.783							
223	198	206	0.0	1.0	0.8	56.9	-38.6	-36.1	52.9	223	0.0	1.0	0.492	54.6	-51.2	-16.6	54.0	198	0.0	1.0	0.8	0.0	1.0	0.584	55.2	-47.5	-23.2	53.0	206	0.0	1.0	0.8							
224	199	206	0.0	1.0	0.816	56.9	-38.0	-36.9	53.0	224	0.0	1.0	0.502	54.7	-50.7	-17.4	53.7	199	0.0	1.0	0.817	0.0	1.0	0.595	55.3	-47.1	-23.9	52.9	206	0.0	1.0	0.817							
225	200	207	0.0	1.0	0.833	57.0	-37.5	-37.7	53.2	225	0.0	1.0	0.514	54.8	-50.3	-18.2	53.6	200	0.0	1.0	0.833	0.0	1.0	0.606	55.3	-46.6	-24.6	52.8	207	0.0	1.0	0.833							
226	201	208	0.0	1.0	0.85	57.1	-36.9	-38.5	53.3	226	0.0	1.0	0.526	54.8	-49.9	-19.1	53.5	201	0.0	1.0	0.85	0.0	1.0	0.616	55.4	-46.1	-25.3	52.7	208	0.0	1.0	0.85							
227	202	209	0.0	1.0	0.866	57.2	-36.4	-39.2	53.5	227	0.0	1.0	0.537	54.9	-49.4	-19.9	53.4	202	0.0	1.0	0.867	0.0	1.0	0.627	55.5	-45.7	-26.0	52.7	209	0.0	1.0	0.867							
228	203	210	0.0	1.0	0.883	57.3	-35.8	-40.0	53.7	228	0.0	1.0	0.549	55.0	-49.0	-20.7	53.3	203	0.0	1.0	0.883	0.0	1.0	0.638	55.6	-45.2	-26.7	52.6	210	0.0	1.0	0.883							
229	204	211	0.0	1.0	0.9	57.4	-35.3	-40.7	53.9	229	0.0	1.0	0.561	55.0	-48.5	-21.5	53.2	204	0.0	1.0	0.9	0.0	1.0	0.648	55.7	-44.8	-27.4	52.6	211	0.0	1.0	0.9							
230	205	212	0.0	1.0	0.916	57.4	-34.8	-41.5	54.1</																														

Output: Offset standard print; separation cmyrn6*, D65, page 13/33

TUB-test chart SE14; 1080 colours, offset standard paper
48 step hue circles; $rgb-LabCh$ *tables, 3D=0, de=0, $cmyk$
input: $rgb/cmyk \rightarrow rgb_d$
output: transfer to $cmyk_d$

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output, separation cmyrn6 (CMYK)

TUB material: code=rha4ta

Data of Maximum color M in colorimetric system Offset standard print; separation cmykn6*, D65 for input or output; Six hue angles of the 60 degree standard colours $RYGCBM_s$: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours $RYGCBM_d$: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours $RYGCBM_e$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

Six hue angles of the device colours $RYGCBM_d$: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours $RYGCBM_e$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$																					
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_{dd361M}	$LAB^*_{dsx361Mi}$ (x=LabCh)	C_d	$rgb^*_{ds361Mi}$	$LAB^*_{dsx361Mi}$ (x=LabCh)	$210C_s$	$rgb^*_{dd361Mi}$	$LAB^*_{de361Mi}$	$rgb^*_{de361Mi}$	$LAB^*_{dex361Mi}$ (x=LabCh)	$216C_e$	$rgb^*_{dd361Mi}$	rgb^*_{dd}	rgb^*_{ds}	rgb^*_{de}				
234	210	216	0.0	1.0	1.0	57.8	-31.9 -45.1 55.3	234	0.0	1.0	0.631 55.5	-45.5 -26.2 52.7	210	0.0	1.0	1.0	0.0				
235	211	217	0.0	0.983	1.0	57.4	-31.3 -45.1 55.0	235	0.0	1.0	0.643 55.6	-45.0 -27.0 52.6	211	0.0	0.983	1.0	0.0				
235	212	218	0.0	0.966	1.0	57.0	-30.7 -45.2 54.7	235	0.0	1.0	0.654 55.7	-44.5 -27.8 52.6	212	0.0	0.967	1.0	0.0				
236	213	219	0.0	0.95	1.0	56.6	-30.1 -45.2 54.4	236	0.0	1.0	0.666 55.8	-44.0 -28.5 52.6	213	0.0	0.95	1.0	0.0				
236	214	220	0.0	0.933	1.0	56.2	-29.6 -45.3 54.1	236	0.0	1.0	0.678 56.0	-43.5 -29.3 52.6	214	0.0	0.933	1.0	0.0				
237	215	221	0.0	0.916	1.0	55.9	-29.0 -45.3 53.8	237	0.0	1.0	0.69	56.1	-42.9 -30.0 52.5	215	0.0	0.917	1.0	0.0			
237	216	222	0.0	0.9	1.0	55.5	-28.4 -45.3 53.5	237	0.0	1.0	0.701 56.2	-42.4 -30.8 52.5	216	0.0	0.9	1.0	0.0				
238	217	223	0.0	0.883	1.0	55.1	-27.8 -45.3 53.2	238	0.0	1.0	0.713 56.3	-41.8 -31.5 52.5	217	0.0	0.883	1.0	0.0				
239	218	224	0.0	0.866	1.0	54.6	-27.2 -45.4 52.9	239	0.0	1.0	0.725 56.4	-41.2 -32.2 52.5	218	0.0	0.867	1.0	0.0				
239	219	225	0.0	0.85	1.0	54.1	-26.4 -45.4 52.6	239	0.0	1.0	0.737 56.5	-40.7 -32.9 52.4	219	0.0	0.85	1.0	0.0				
240	220	226	0.0	0.833	1.0	53.7	-25.7 -45.5 52.3	240	0.0	1.0	0.749 56.6	-40.1 -33.6 52.4	220	0.0	0.833	1.0	0.0				
241	221	227	0.0	0.816	1.0	53.2	-25.0 -45.5 51.9	241	0.0	1.0	0.765 56.7	-39.6 -34.4 52.6	221	0.0	0.817	1.0	0.0				
241	222	227	0.0	0.8	1.0	52.7	-24.3 -45.5 51.6	241	0.0	1.0	0.781 56.8	-39.1 -35.2 52.7	222	0.0	0.8	1.0	0.0				
242	223	228	0.0	0.783	1.0	52.2	-23.5 -45.6 51.3	242	0.0	1.0	0.798 56.9	-38.6 -36.0 52.9	223	0.0	0.783	1.0	0.0				
243	224	229	0.0	0.766	1.0	51.8	-22.8 -45.6 51.0	243	0.0	1.0	0.814 57.0	-38.1 -36.7 53.0	224	0.0	0.767	1.0	0.0				
244	225	230	0.0	0.75	1.0	51.3	-22.1 -45.6 50.7	244	0.0	1.0	0.831 57.1	-37.5 -37.5 53.2	225	0.0	0.75	1.0	0.0				
244	226	231	0.0	0.733	1.0	50.7	-21.3 -45.7 50.4	244	0.0	1.0	0.847 57.2	-37.0 -38.3 53.4	226	0.0	0.733	1.0	0.0				
245	227	232	0.0	0.716	1.0	50.2	-20.5 -45.7 50.1	245	0.0	1.0	0.864 57.2	-36.4 -39.1 53.5	227	0.0	0.717	1.0	0.0				
246	228	233	0.0	0.7	1.0	49.7	-19.6 -45.8 49.9	246	0.0	1.0	0.881 57.3	-35.8 -39.8 53.7	228	0.0	0.7	1.0	0.0				
247	229	234	0.0	0.683	1.0	49.1	-18.8 -45.9 49.6	247	0.0	1.0	0.899 57.4	-35.3 -40.6 54.0	229	0.0	0.683	1.0	0.0				
248	230	235	0.0	0.666	1.0	48.6	-18.0 -45.9 49.3	248	0.0	1.0	0.917 57.5	-34.7 -41.4 54.2	230	0.0	0.667	1.0	0.0				
249	231	236	0.0	0.65	1.0	48.0	-17.2 -45.9 49.1	249	0.0	1.0	0.934 57.6	-34.2 -42.2 54.4	231	0.0	0.65	1.0	0.0				
250	232	237	0.0	0.633	1.0	47.5	-16.4 -45.9 48.8	250	0.0	1.0	0.952 57.7	-33.6 -43.0 54.7	232	0.0	0.633	1.0	0.0				
251	233	237	0.0	0.616	1.0	46.9	-15.4 -46.0 48.5	251	0.0	1.0	0.97	57.7	-32.9 -43.8 54.9	233	0.0	0.617	1.0	0.0			
252	234	238	0.0	0.6	1.0	46.2	-14.3 -46.1 48.3	252	0.0	1.0	0.988 57.8	-32.3 -44.5 55.2	234	0.0	0.6	1.0	0.0				
253	235	239	0.0	0.583	1.0	45.6	-13.2 -46.2 48.1	253	0.0	0.99	1.0	57.6	-31.5 -45.1 55.1	235	0.0	0.583	1.0	0.0			
255	236	240	0.0	0.566	1.0	44.9	-12.1 -46.3 47.8	255	0.0	0.959	1.0	56.9	-30.4 -45.2 54.6	236	0.0	0.567	1.0	0.0			
256	237	241	0.0	0.55	1.0	44.3	-11.0 -46.3 47.6	256	0.0	0.928	1.0	56.2	-29.3 -45.2 54.0	237	0.0	0.55	1.0	0.0			
257	238	242	0.0	0.533	1.0	43.6	-9.9 -46.3 47.4	257	0.0	0.897	1.0	55.4	-28.2 -45.3 53.5	238	0.0	0.533	1.0	0.0			
259	239	243	0.0	0.516	1.0	43.0	-8.8 -46.3 47.2	259	0.0	0.868	1.0	54.7	-27.2 -45.3 53.0	239	0.0	0.517	1.0	0.0			
260	240	244	0.0	0.5	1.0	42.3	-7.7 -46.3 46.9	260	0.0	0.845	1.0	54.1	-26.2 -45.4 52.5	240	0.0	0.5	1.0	0.0			
261	241	245	0.0	0.483	1.0	41.6	-6.7 -46.4 46.9	261	0.0	0.822	1.0	53.4	-25.2 -45.5 52.1	241	0.0	0.483	1.0	0.0			
263	242	246	0.0	0.466	1.0	41.0	-5.6 -46.4 46.8	263	0.0	0.798	1.0	52.7	-24.1 -45.5 51.6	242	0.0	0.467	1.0	0.0			
264	243	247	0.0	0.45	1.0	40.3	-4.5 -46.5 46.7	264	0.0	0.775	1.0	52.1	-23.1 -45.5 51.2	243	0.0	0.45	1.0	0.0			
265	244	248	0.0	0.433	1.0	39.6	-3.4 -46.5 46.7	265	0.0	0.752	1.0	51.4	-22.2 -45.5 50.8	244	0.0	0.433	1.0	0.0			
267	245	248	0.0	0.416	1.0	38.9	-2.3 -46.5 46.6	267	0.0	0.733	1.0	50.8	-21.2 -45.6 50.4	245	0.0	0.417	1.0	0.0			
268	246	249	0.0	0.4	1.0	38.3	-1.2 -46.5 46.5	268	0.0	0.714	1.0	50.2	-20.3 -45.7 50.1	246	0.0	0.4	1.0	0.0			
269	247	250	0.0	0.383	1.0	37.6	-0.2 -46.5 46.5	269	0.0	0.695	1.0	49.6	-19.4 -45.8 49.8	247	0.0	0.383	1.0	0.0			
271	248	251	0.0	0.366	1.0	37.0	0.8 -46.5 46.5	271	0.0	0.677	1.0	49.0	-18.5 -45.8 49.5	248	0.0	0.367	1.0	0.0			
272	249	252	0.0	0.35	1.0	36.4	1.9 -46.7 46.7	272	0.0	0.658	1.0	48.3	-17.5 -45.9 49.2	249	0.0	0.35	1.0	0.0			
273	250	253	0.0	0.333	1.0	35.8	3.0 -46.8 46.9	273	0.0	0.639	1.0	47.7	-16.6 -45.9 48.9	250	0.0	0.333	1.0	0.0			
275	251	254	0.0	0.316	1.0	35.1	4.1 -46.9 47.1	275	0.0	0.622	1.0	47.2	-15.7 -45.9 48.7	251	0.0	0.317	1.0	0.0			
276	252	255	0.0	0.3	1.0	34.5	5.2 -46.9 47.2	276	0.0	0.609	1.0	46.6	-14.9 -46.0 48.5	252	0.0	0.3	1.0	0.0			
277	253	256	0.0	0.283	1.0	33.9	6.3 -47.0 47.4	277	0.0	0.596	1.0	46.1	-14.0 -46.1 48.3	253	0.0	0.283	1.0	0.0			
278	254	257	0.0	0.266	1.0	33.3	7.4 -47.0 47.6	278	0.0	0.583	1.0	45.6	-13.2 -46.2 48.1	254	0.0	0.267	1.0	0.0			
280	255	258	0.0	0.25	1.0	32.7	8.5 -47.0 47.8	280	0.0	0.57	1.0	45.1	-12.3 -46.2 47.9	255	0.0	0.25	1.0	0.0			

see similar files: <http://130.149.60.45/~farbmeterik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmeterik>

Data of Maximum color M in colorimetric system Offset standard print; separation cmyn6*, D65 for input or output; Six hue angles of the 60 degree standard colours $RYGCBM_s$: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$; Six hue angles of the device colours $RYGCBM_d$: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours $RYGCBM_e$: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

Table 1. The ranges of the three colours (RGB) and the ranges of the elementary colour (LAB) for the 361Mi and 361Mi (x=LabCh) galaxies.										Table 2. The ranges of the three colours (RGB) and the ranges of the elementary colour (LAB) for the 361Mi and 361Mi (x=LabCh) galaxies.																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																										
Lab _d	hab _s	hab _e	rgb*	dd361Mi	LAB*	dd361Mi (x=LabCh)	rgb*	ds361Mi	LAB*	ds361Mi (x=LabCh)	rgb*	dd361Mi	rgb*	de361Mi	LAB*	de361Mi (x=LabCh)	rgb*	dd361Mi	rgb*	de361Mi																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																
280	255	258	0.0	0.25	1.0	32.7	8.5	-47.0	47.8	280	0.0	0.57	1.0	45.1	-12.3	-46.2	47.9	255	0.0	0.25	1.0																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															</

1-0031430-L0

SE140-70

LAB*la0, YN=0%, XYZnw=2.5, 2.6, 2.7, 86.0, 90.9, 95.9, LAB*nw=18.5, 0.0, 0.0, 96.4, 0.0, 0.0

Output: Offset standard print; separation cmyn6*, D65, page 15/33

TUB-test chart SE14; 1080 colours, offset standard paper
48 step hue circles; *rgb-LabCh** tables, 3D=0, de=0, *cmv*

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

~~1-0031430-F0~~

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
+ application for measurement of offset print output, separa-

TUB material: code=rha4ta

Data of Maximum color M in colorimetric system Offset standard print; separation cmycn6*, D65 for input or output; Six hue angles of the 60 degree standard colours RYGBM_s: $h_{ab,ds} = 30.0, 90.0, 150.0, 210.0, 270.0, 330.0$;
Six hue angles of the device colours RYGBM_d: $h_{ab,d} = 30.4, 96.1, 161.6, 234.7, 295.7, 353.2$; Six hue angles of the elementary colours RYGBM_e: $h_{ab,e} = 25.5, 92.3, 162.2, 217.0, 271.7, 328.6$

Six hue angles of the device colours $\lambda_{FCB}m_e$, $h_{ab,d}$, $h_{ab,s}$, $h_{ab,e}$, $h_{ab,e} - 23.3$, 32.3 , 101.0 , 234.7 , 239.7 , 335.2 , Six hue angles of the elementary colours $\lambda_{FCB}m_e$, $h_{ab,e} - 23.3$, 32.3 , 101.0 , 234.7 , 239.7 , 335.2										Six hue angles of the device colours $\lambda_{FCB}m_e$, $h_{ab,d}$, $h_{ab,s}$, $h_{ab,e}$, $h_{ab,e} - 23.3$, 32.3 , 101.0 , 234.7 , 239.7 , 335.2																																																																																																																																																																																																																																																																																																												
$h_{ab,d}$	$h_{ab,s}$	$h_{ab,e}$	rgb^*_{dd361M}	$LAB^*_{d361M}(x=LabCh)$	$rgb^*_{ds361Mi}$	$LAB^*_{dsx361Mi}(x=LabCh)$	$rgb^*_{dd361Mi}$	$LAB^*_{de361Mi}$	$LAB^*_{dex361Mi}(x=LabCh)$	$rgb^*_{dd361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$	$rgb^*_{ds361Mi}$

output: transfer to $cmyk_d$

http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF /.PS; transfer output

N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 18/33

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

n/j	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsi_Md	rgb*Md	LabCh*Md				
0/648	R00Y_100_100d	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4
1/657	R13Y_100_100d	1.0	0.125	0.0	1.0	1.0	0.5	37	1.0	0.116	0.0	51.2	57.2	42.8	71.5	36.8
2/666	R25Y_100_100d	1.0	0.25	0.0	1.0	1.0	0.5	44	1.0	0.233	0.0	55.9	47.3	48.7	67.9	45.8
3/675	R38Y_100_100d	1.0	0.375	0.0	1.0	1.0	0.5	52	1.0	0.366	0.0	62.0	35.2	56.0	66.2	57.8
4/684	R50Y_100_100d	1.0	0.5	0.0	1.0	1.0	0.5	60	1.0	0.5	0.0	68.1	24.0	63.0	67.4	69.1
5/693	R63Y_100_100d	1.0	0.625	0.0	1.0	1.0	0.5	68	1.0	0.633	0.0	75.3	11.6	72.0	72.9	80.8
6/702	R75Y_100_100d	1.0	0.75	0.0	1.0	1.0	0.5	76	1.0	0.766	0.0	81.2	2.5	78.8	78.9	88.1
7/711	R88Y_100_100d	1.0	0.875	0.0	1.0	1.0	0.5	83	1.0	0.883	0.0	85.6	-4.1	84.3	84.4	92.8
8/720	Y00G_100_100d	1.0	1.0	0.0	1.0	1.0	0.5	90	1.0	1.0	0.0	89.4	-9.5	89.0	89.6	96.0
9/639	Y13G_100_100d	0.875	1.0	0.0	1.0	1.0	0.5	97	0.883	1.0	0.0	86.9	-13.6	83.1	84.2	99.3
10/558	Y25G_100_100d	0.75	1.0	0.0	1.0	1.0	0.5	104	0.766	1.0	0.0	84.1	-17.3	77.9	79.8	102.5
11/477	Y38G_100_100d	0.625	1.0	0.0	1.0	1.0	0.5	112	0.633	1.0	0.0	78.3	-22.8	71.7	75.2	107.6
12/396	Y50G_100_100d	0.5	1.0	0.0	1.0	1.0	0.5	120	0.5	1.0	0.0	73.1	-30.2	60.8	67.9	116.4
13/315	Y63G_100_100d	0.375	1.0	0.0	1.0	1.0	0.5	128	0.366	1.0	0.0	68.3	-37.3	52.3	64.3	125.4
14/234	Y75G_100_100d	0.25	1.0	0.0	1.0	1.0	0.5	136	0.233	1.0	0.0	60.3	-48.7	41.3	63.9	139.7
15/153	Y88G_100_100d	0.125	1.0	0.0	1.0	1.0	0.5	143	0.116	1.0	0.0	56.3	-57.0	32.8	65.8	150.0
16/72	G00C_100_100d	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	51.6	-69.3	23.0	73.1	161.6
17/73	G13C_100_100d	0.0	1.0	0.125	1.0	1.0	0.5	157	0.0	1.0	0.116	52.3	-66.3	14.2	67.9	167.8
18/74	G25C_100_100d	0.0	1.0	0.25	1.0	1.0	0.5	164	0.0	1.0	0.233	52.9	-62.5	5.2	62.7	175.2
19/75	G38C_100_100d	0.0	1.0	0.375	1.0	1.0	0.5	172	0.0	1.0	0.366	53.8	-56.9	-6.1	57.3	186.2
20/76	G50C_100_100d	0.0	1.0	0.5	1.0	1.0	0.5	180	0.0	1.0	0.5	54.6	-50.8	-17.3	53.7	198.8
21/77	G63C_100_100d	0.0	1.0	0.625	1.0	1.0	0.5	188	0.0	1.0	0.633	55.5	-45.4	-26.5	52.6	210.2
22/78	G75C_100_100d	0.0	1.0	0.75	1.0	1.0	0.5	196	0.0	1.0	0.766	56.7	-39.6	-34.5	52.5	221.1
23/79	G88C_100_100d	0.0	1.0	0.875	1.0	1.0	0.5	203	0.0	1.0	0.883	57.3	-35.8	-40.0	53.7	228.1
24/80	C00B_100_100d	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6
25/71	C13B_100_100d	0.0	0.875	1.0	1.0	1.0	0.5	217	0.0	0.883	1.0	55.1	-27.8	-45.3	53.2	238.4
26/62	C25B_100_100d	0.0	0.75	1.0	1.0	1.0	0.5	224	0.0	0.766	1.0	51.8	-22.8	-45.6	51.0	243.3
27/53	C38B_100_100d	0.0	0.625	1.0	1.0	1.0	0.5	232	0.0	0.633	1.0	47.5	-16.4	-45.9	48.8	250.3
28/44	C50B_100_100d	0.0	0.5	1.0	1.0	1.0	0.5	240	0.0	0.5	1.0	42.3	-7.7	-46.3	46.9	260.4
29/35	C63B_100_100d	0.0	0.375	1.0	1.0	1.0	0.5	248	0.0	0.366	1.0	37.0	0.8	-46.4	46.4	271.0
30/26	C75B_100_100d	0.0	0.25	1.0	1.0	1.0	0.5	256	0.0	0.233	1.0	32.1	9.5	-47.2	48.1	281.4
31/17	C88B_100_100d	0.0	0.125	1.0	1.0	1.0	0.5	263	0.0	0.116	1.0	27.8	17.1	-47.6	50.6	289.8
32/8	B00M_100_100d	0.0	0.0	1.0	1.0	1.0	0.5	270	0.0	0.0	1.0	24.9	22.9	-47.8	53.0	295.6
33/89	B13M_100_100d	0.125	0.0	1.0	1.0	1.0	0.5	277	0.116	0.0	1.0	27.6	30.9	-43.8	53.6	305.2
34/170	B25M_100_100d	0.25	0.0	1.0	1.0	1.0	0.5	284	0.233	0.0	1.0	29.6	35.4	-40.8	54.1	310.9
35/251	B38M_100_100d	0.375	0.0	1.0	1.0	1.0	0.5	292	0.366	0.0	1.0	33.5	47.0	-32.8	57.3	325.0
36/332	B50M_100_100d	0.5	0.0	1.0	1.0	1.0	0.5	300	0.5	0.0	1.0	37.0	53.9	-27.1	60.4	333.2
37/413	B63M_100_100d	0.625	0.0	1.0	1.0	1.0	0.5	308	0.633	0.0	1.0	40.4	60.2	-21.7	64.0	340.1
38/494	B75M_100_100d	0.75	0.0	1.0	1.0	1.0	0.5	316	0.766	0.0	1.0	43.6	67.3	-15.2	69.0	347.2
39/575	B88M_100_100d	0.875	0.0	1.0	1.0	1.0	0.5	323	0.883	0.0	1.0	46.0	70.9	-11.8	71.9	350.5
40/656	M00R_100_100d	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	48.2	74.2	-8.7	74.7	353.2
41/655	M13R_100_100d	1.0	0.0	0.875	1.0	1.0	0.5	337	1.0	0.0	0.883	48.2	73.2	-5.1	73.4	355.9
42/654	M25R_100_100d	1.0	0.0	0.75	1.0	1.0	0.5	344	1.0	0.0	0.766	48.1	72.2	-1.3	72.2	358.9
43/653	M38R_100_100d	1.0	0.0	0.625	1.0	1.0	0.5	352	1.0	0.0	0.633	48.0	70.8	4.5	71.0	3.7
44/652	M50R_100_100d	1.0	0.0	0.5	1.0	1.0	0.5	360	1.0	0.0	0.5	47.8	69.7	11.3	70.6	9.2
45/651	M63R_100_100d	1.0	0.0	0.375	1.0	1.0	0.5	368	1.0	0.0	0.366	47.8	68.1	18.7	70.7	15.4
46/650	M75R_100_100d	1.0	0.0	0.25	1.0	1.0	0.5	376	1.0	0.0	0.233	47.8	66.9	26.3	71.9	21.4
47/649	M88R_100_100d	1.0	0.0	0.125	1.0	1.0	0.5	383	1.0	0.0	0.116	47.6	66.2	32.3	73.7	26.0
48/648	R00Y_100_100d	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4
49/0	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	51.5	56.6	43.1	71.2	397.2
50/91	NW_013d	0.125	0.125	0.125	0.125	0.0	0.0	360	0.125	0.125	0.125	18.5	0.0	0.0	0.0	9.7
51/182	NW_025d	0.25	0.25	0.25	0.25	0.0	0.0	360	0.25	0.25	0.25	30.8	-0.2	-1.2	1.3	25.7
52/273	NW_038d	0.375	0.375	0.375	0.375	0.0	0.0	360	0.375	0.375	0.375	45.2	-0.4	-2.1	2.2	25.9
53/364	NW_050d	0.5	0.5	0.5	0.5	0.0	0.0	360	0.5	0.5	0.5	57.6	-0.3	-2.0	2.0	26.1
54/455	NW_063d	0.625	0.625	0.625	0.625	0.0	0.0	360	0.625	0.625	0.625	65.9	-0.2	-2.0	2.0	26.1
55/546	NW_075d	0.75	0.75	0.75	0.75	0.0	0.0	360	0.75	0.75	0.75	75.0	-0.2	-1.6	1.6	26.1
56/637	NW_088d	0.875	0.875	0.875	0.875	0.0	0.0	360	0.875	0.875	0.875	82.6	-0.1	-1.1	1.1	26.3
57/728	NW_100d	1.0	1.0	1.0	1.0	0.0	0.0	360	1.0	1.0	1.0	90.9	0.0	-0.4	0.4	25.9

Mean color difference of this page:

delta E* = 2.4

TUB-test chart SE14; 1080 colours, offset standard paper
colors and differences, ΔE^* , 3D=0, de=0, *cmYk*input: *rgb/cmyk* -> *rgb*_d
output: transfer to *cmYk*_dTUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output, separationcmyk6 (CMYK)
TUB material: code=rha4ta

http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF /.PS; transfer output

N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 19/33

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

n/f	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md						
0/648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4		
1/666	R25Y_100_100a	1.0	0.25	0.0	1.0	1.0	0.5	44	1.0	0.233	0.0	47.5	65.5	38.7	67.9	45.8		
2/684	R50Y_100_100a	1.0	0.5	0.0	1.0	1.0	0.5	60	1.0	0.5	0.0	68.1	24.0	63.0	67.4	69.1		
3/702	R75Y_100_100a	1.0	0.75	0.0	1.0	1.0	0.5	76	1.0	0.766	0.0	81.2	2.5	78.8	78.9	88.1		
4/720	Y00G_100_100a	1.0	1.0	0.0	1.0	1.0	0.5	90	1.0	1.0	0.0	89.4	-9.5	89.0	89.6	96.0		
5/558	Y25G_100_100a	0.75	1.0	0.0	1.0	1.0	0.5	104	0.766	1.0	0.0	84.1	-17.3	77.9	79.8	102.5		
6/396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	120	0.5	1.0	0.0	73.1	-30.2	60.8	67.9	116.4		
7/234	Y75G_100_100a	0.25	1.0	0.0	1.0	1.0	0.5	136	0.233	1.0	0.0	60.3	-48.7	41.3	63.9	139.7		
8/72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	51.6	-69.3	23.0	73.1	161.6		
9/72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	51.6	-69.3	23.0	73.1	161.6		
10/76	G25B_100_100a	0.0	1.0	0.5	1.0	1.0	0.5	180	0.0	1.0	0.5	54.6	-50.8	-17.3	53.7	198.8		
11/80	G50B_100_100a	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
12/44	G75B_100_100a	0.0	0.5	1.0	1.0	1.0	0.5	240	0.0	0.5	1.0	42.3	-7.7	-46.3	46.9	260.4		
13/8	B00M_100_100a	0.0	0.0	1.0	1.0	1.0	0.5	270	0.0	0.0	1.0	24.9	22.9	-47.8	53.0	295.6		
14/332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	300	0.5	0.0	1.0	37.0	53.9	-27.1	60.4	333.2		
15/656	B50R_100_100a	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	48.2	74.2	-8.7	74.7	353.2		
16/652	B75R_100_100a	1.0	0.0	0.5	1.0	1.0	0.5	360	1.0	0.0	0.5	47.8	69.7	11.3	70.6	9.2		
17/648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4		
18/688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	390	1.0	0.5	0.5	70.7	26.4	21.0	33.8	38.5	6.6	389
19/706	R50Y_100_050a	1.0	0.75	0.5	1.0	0.5	0.75	60	1.0	0.75	0.5	82.2	7.9	28.6	29.6	74.5	5.0	59
20/724	Y00G_100_050a	1.0	1.0	0.5	1.0	0.5	0.75	90	1.0	1.0	0.5	92.8	-6.1	35.6	36.2	99.7	8.9	89
21/562	Y50G_100_050a	0.75	1.0	0.5	1.0	0.5	0.75	120	0.75	1.0	0.5	86.5	-13.2	24.9	28.2	117.8	6.0	119
22/400	G00B_100_050a	0.5	1.0	0.5	1.0	0.5	0.75	150	0.5	1.0	0.5	76.1	-23.7	13.0	27.0	151.2	11.2	149
23/404	G50B_100_050a	0.5	1.0	1.0	1.0	0.5	0.75	210	0.5	1.0	1.0	80.1	-13.2	-19.2	23.3	235.4	5.2	210
24/368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	270	0.5	0.5	1.0	59.3	14.9	-24.3	28.5	301.5	3.6	270
25/692	B50R_100_050a	1.0	0.5	1.0	1.0	0.5	0.75	330	1.0	0.5	1.0	73.1	31.3	-7.2	32.1	346.9	6.5	330
26/688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.75	390	1.0	0.5	0.5	70.7	26.4	21.0	33.8	38.5	6.6	389
27/506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	390	0.75	0.25	0.25	54.5	30.7	22.3	37.9	36.0	4.2	389
28/524	R50Y_075_050a	0.75	0.5	0.25	0.75	0.5	0.5	60	0.75	0.5	0.25	68.4	8.1	31.3	32.3	75.4	6.8	59
29/542	Y00G_075_050a	0.75	0.75	0.25	0.75	0.5	0.5	90	0.75	0.75	0.25	78.8	-7.3	39.0	39.7	100.6	8.1	89
30/380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	120	0.5	0.75	0.25	70.5	-15.6	29.0	33.0	118.2	5.0	119
31/218	G00B_075_050a	0.25	0.75	0.25	0.75	0.5	0.5	150	0.25	0.75	0.25	50.3	-18.7	14.4	32.2	153.3	6.0	149
32/222	G50B_075_050a	0.25	0.75	0.75	0.75	0.5	0.5	210	0.25	0.75	0.75	53.3	-15.8	-23.1	28.0	235.6	5.6	210
33/186	B00R_075_050a	0.25	0.25	0.75	0.75	0.5	0.5	270	0.25	0.25	0.75	42.6	13.7	-27.5	30.7	296.4	4.4	270
34/510	B50R_075_050a	0.75	0.25	0.75	0.75	0.5	0.5	330	0.75	0.25	0.75	56.6	36.1	-8.5	37.1	346.7	5.6	330
35/506	R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	390	0.75	0.25	0.25	54.5	30.7	22.3	37.9	36.0	4.2	389
36/324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.25	390	0.5	0.0	0.0	35.3	37.1	22.2	43.3	30.9	5.7	389
37/342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	60	0.5	0.25	0.0	50.3	8.4	35.9	36.9	76.7	8.9	59
38/360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	90	0.5	0.5	0.0	61.1	-8.1	47.0	47.7	98.8	8.3	89
39/198	Y50G_050_050a	0.25	0.5	0.0	0.5	0.5	0.25	120	0.25	0.5	0.0	51.9	-19.0	34.0	39.0	119.2	8.1	119
40/36	G00B_050_050a	0.0	0.5	0.0	0.5	0.5	0.25	150	0.0	0.5	0.0	41.1	-36.9	16.6	40.5	155.7	8.2	149
41/40	G50B_050_050a	0.0	0.5	0.5	0.5	0.5	0.25	210	0.0	0.5	0.5	44.9	-18.7	-26.5	32.4	234.7	8.2	210
42/4	B00R_050_050a	0.0	0.0	0.5	0.5	0.5	0.25	270	0.0	0.0	0.5	22.5	17.7	-29.3	34.3	301.0	8.3	270
43/328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.25	330	0.5	0.0	0.5	36.1	44.2	-8.4	45.0	349.2	8.6	330
44/324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.25	390	0.5	0.0	0.0	35.3	37.1	22.2	43.3	30.9	5.7	389
45/0	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	18.5	0.0	0.0	0.0	0.0	0.0	0.0
46/91	NW_013a	0.125	0.125	0.125	0.125	0.0	0.125	360	0.125	0.125	0.125	30.8	-0.2	-1.2	1.3	257.7	2.9	360
47/182	NW_025a	0.25	0.25	0.25	0.25	0.0	0.25	360	0.25	0.25	0.25	45.2	-0.4	-2.1	2.2	259.4	7.5	360
48/273	NW_038a	0.375	0.375	0.375	0.375	0.0	0.375	360	0.375	0.375	0.375	57.6	-0.3	-2.0	2.0	261.2	10.1	360
49/364	NW_050a	0.5	0.5	0.5	0.5	0.0	0.5	360	0.5	0.5	0.5	65.9	-0.2	-2.0	2.0	261.9	8.7	360
50/455	NW_063a	0.625	0.625	0.625	0.625	0.0	0.625	360	0.625	0.625	0.625	75.0	-0.2	-1.6	1.6	261.9	8.0	360
51/546	NW_075a	0.75	0.75	0.75	0.75	0.0	0.75	360	0.75	0.75	0.75	82.6	-0.1	-1.1	1.1	263.4	5.8	360
52/637	NW_088a	0.875	0.875	0.875	0.875	0.0	0.875	360	0.875	0.875	0.875	90.9	0.0	-0.4	0.4	259.7	4.2	360
53/728	NW_100a	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0	0.0	360
													Mean color difference of this page: delta E* = 4.3					

1-0031830-F0

SE140-7N, Page 19/33-F

TUB-test chart SE14; 1080 colours, offset standard paper
colors and differences, ΔE^* , 3D=0, de=0, *cm*ykinput: *rgb/cmyk* -> *rgb*_d
output: transfer to *cm*yk_d

1-0031830-F0

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output, separationcmykn6 (CMYK)
TUB material: code=rha4ta

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF> / .PS
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output, separationcnymn6 (CMYK)
TUB material: code=rha4ta

n=j	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsi_d	rgb*Md	LabCh*Md
0	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	1.0
1	B00R_012_012a	0.0	0.0	0.125	0.125	0.125	0.062	0.062	2.8	2.7	0.0	0.0
2	B00R_025_025a	0.0	0.0	0.25	0.25	0.25	0.125	0.125	6.5	6.5	0.0	0.0
3	B00R_037_037a	0.0	0.0	0.375	0.375	0.375	0.187	0.187	7.9	7.9	0.0	0.0
4	B00R_050_050a	0.0	0.0	0.5	0.5	0.5	0.25	0.25	8.3	8.3	0.0	0.0
5	B00R_062_062a	0.0	0.0	0.625	0.625	0.625	0.312	0.312	7.3	7.3	0.0	0.0
6	B00R_075_075a	0.0	0.0	0.75	0.75	0.75	0.375	0.375	5.7	5.7	0.0	0.0
7	B00R_087_087a	0.0	0.0	0.875	0.875	0.875	0.437	0.437	3.9	3.9	0.0	0.0
8	B00R_100_100a	0.0	0.0	1.0	1.0	1.0	0.5	0.5	2.7	2.7	0.0	0.0
9	G00B_012_012a	0.0	0.125	0.0	0.125	0.125	0.062	0.062	2.6	2.6	0.0	0.0
10	G50B_012_012a	0.0	0.125	0.125	0.125	0.125	0.062	0.062	3.1	3.1	0.0	0.0
11	G75B_025_025a	0.0	0.125	0.25	0.25	0.25	0.125	0.125	6.4	6.4	0.0	0.0
12	G84B_037_037a	0.0	0.125	0.375	0.375	0.375	0.187	0.187	6.8	6.8	0.0	0.0
13	G88B_050_050a	0.0	0.125	0.5	0.5	0.5	0.25	0.25	25.1	25.1	0.0	0.0
14	G90B_062_062a	0.0	0.125	0.625	0.625	0.625	0.312	0.312	5.2	5.2	0.0	0.0
15	G92B_075_075a	0.0	0.125	0.75	0.75	0.75	0.375	0.375	4.2	4.2	0.0	0.0
16	G93B_087_087a	0.0	0.125	0.875	0.875	0.875	0.437	0.437	2.9	2.9	0.0	0.0
17	G94B_100_100a	0.0	0.125	1.0	1.0	1.0	0.5	0.5	2.6	2.6	0.0	0.0
18	G00B_025_025a	0.0	0.25	0.0	0.25	0.25	0.125	0.125	7.8	7.8	0.0	0.0
19	G25B_025_025a	0.0	0.25	0.125	0.25	0.25	0.125	0.125	7.6	7.6	0.0	0.0
20	G50B_025_025a	0.0	0.25	0.25	0.25	0.25	0.125	0.125	8.4	8.4	0.0	0.0
21	G65B_037_037a	0.0	0.25	0.375	0.375	0.375	0.187	0.187	8.4	8.4	0.0	0.0
22	G75B_050_050a	0.0	0.25	0.5	0.5	0.5	0.25	0.25	7.3	7.3	0.0	0.0
23	G80B_062_062a	0.0	0.25	0.625	0.625	0.625	0.312	0.312	6.4	6.4	0.0	0.0
24	G84B_075_075a	0.0	0.25	0.75	0.75	0.75	0.375	0.375	4.6	4.6	0.0	0.0
25	G86B_087_087a	0.0	0.25	0.875	0.875	0.875	0.437	0.437	3.1	3.1	0.0	0.0
26	G88B_100_100a	0.0	0.25	1.0	1.0	1.0	0.5	0.5	2.5	2.5	0.0	0.0
27	G00B_037_037a	0.0	0.375	0.0	0.375	0.375	0.187	0.187	15.7	15.7	0.0	0.0
28	G15B_037_037a	0.0	0.375	0.125	0.375	0.375	0.187	0.187	7.6	7.6	0.0	0.0
29	G34B_037_037a	0.0	0.375	0.25	0.375	0.375	0.187	0.187	8.0	8.0	0.0	0.0
30	G50B_037_037a	0.0	0.375	0.375	0.375	0.375	0.187	0.187	9.3	9.3	0.0	0.0
31	G61B_050_050a	0.0	0.375	0.5	0.5	0.5	0.25	0.25	22.2	22.2	0.0	0.0
32	G69B_062_062a	0.0	0.375	0.625	0.625	0.625	0.312	0.312	6.3	6.3	0.0	0.0
33	G75B_075_075a	0.0	0.375	0.75	0.75	0.75	0.375	0.375	4.7	4.7	0.0	0.0
34	G79B_087_087a	0.0	0.375	0.875	0.875	0.875	0.437	0.437	3.2	3.2	0.0	0.0
35	G81B_100_100a	0.0	0.375	1.0	1.0	1.0	0.5	0.5	2.4	2.4	0.0	0.0
36	G00B_050_050a	0.0	0.5	0.0	0.5	0.5	0.25	0.25	8.2	8.2	0.0	0.0
37	G11B_050_050a	0.0	0.5	0.125	0.5	0.5	0.25	0.25	6.8	6.8	0.0	0.0
38	G25B_050_050a	0.0	0.5	0.25	0.5	0.5	0.25	0.25	7.1	7.1	0.0	0.0
39	G38B_050_050a	0.0	0.5	0.375	0.5	0.5	0.25	0.25	7.7	7.7	0.0	0.0
40	G50B_050_050a	0.0	0.5	0.5	0.5	0.5	0.25	0.25	8.2	8.2	0.0	0.0
41	G59B_062_062a	0.0	0.5	0.625	0.625	0.625	0.312	0.312	6.8	6.8	0.0	0.0
42	G65B_075_075a	0.0	0.5	0.75	0.75	0.75	0.375	0.375	4.5	4.5	0.0	0.0
43	G70B_087_087a	0.0	0.5	0.875	0.875	0.875	0.437	0.437	3.4	3.4	0.0	0.0
44	G75B_100_100a	0.0	0.5	1.0	1.0	1.0	0.5	0.5	2.4	2.4	0.0	0.0
45	G00B_062_062a	0.0	0.625	0.0	0.625	0.625	0.312	0.312	15.9	15.9	0.0	0.0
46	G09B_062_062a	0.0	0.625	0.125	0.625	0.625	0.312	0.312	6.4	6.4	0.0	0.0
47	G19B_062_062a	0.0	0.625	0.25	0.625	0.625	0.312	0.312	6.8	6.8	0.0	0.0
48	G30B_062_062a	0.0	0.625	0.375	0.625	0.625	0.312	0.312	4.7	4.7	0.0	0.0
49	G40B_062_062a	0.0	0.625	0.5	0.625	0.625	0.312	0.312	3.9	3.9	0.0	0.0
50	G50B_062_062a	0.0	0.625	0.625	0.625	0.625	0.312	0.312	6.3	6.3	0.0	0.0
51	G57B_075_075a	0.0	0.625	0.75	0.75	0.75	0.375	0.375	2.9	2.9	0.0	0.0
52	G63B_087_087a	0.0	0.625	0.875	0.875	0.875	0.437	0.437	2.6	2.6	0.0	0.0
53	G68B_100_100a	0.0	0.625	1.0	1.0	1.0	0.5	0.5	2.3	2.3	0.0	0.0
54	G00B_075_075a	0.0	0.75	0.0	0.75	0.75	0.375	0.375	15.0	15.0	0.0	0.0
55	G07B_075_075a	0.0	0.75	0.125	0.75	0.75	0.375	0.375	7.5	7.5	0.0	0.0
56	G15B_075_075a	0.0	0.75	0.25	0.75	0.75	0.375	0.375	8.2	8.2	0.0	0.0
57	G25B_075_075a	0.0	0.75	0.375	0.75	0.75	0.375	0.375	8.5	8.5	0.0	0.0
58	G34B_075_075a	0.0	0.75	0.5	0.75	0.75	0.375	0.375	7.7	7.7	0.0	0.0
59	G42B_075_075a	0.0	0.75	0.625	0.75	0.75	0.375	0.375	6.8	6.8	0.0	0.0
60	G50B_075_075a	0.0	0.75	0.75	0.75	0.75	0.375	0.375	8.2	8.2	0.0	0.0
61	G56B_087_087a	0.0	0.75	0.875	0.875	0.875	0.437	0.437	4.9	4.9	0.0	0.0
62	G61B_100_100a	0.0	0.75	1.0	1.0	1.0	0.5	0.5	2.4	2.4	0.0	0.0
63	G00B_087_087a	0.0	0.875	0.0	0.875	0.875	0.437	0.437	15.0	15.0	0.0	0.0
64	G06B_087_087a	0.0	0.875	0.125	0.875	0.875	0.437	0.437	7.6	7.6	0.0	0.0
65	G13B_087_087a	0.0	0.875	0.25	0.875	0.875	0.437	0.437	8.2	8.2	0.0	0.0
66	G20B_087_087a	0.0	0.875	0.375	0.875	0.875	0.437	0.437	7.5	7.5	0.0	0.0
67	G29B_087_087a	0.0	0.875	0.5	0.875	0.875	0.437	0.437	6.8	6.8	0.0	0.0
68	G36B_087_087a	0.0	0.875	0.625	0.875	0.875	0.437	0.437	5.2	5.2	0.0	0.0
69	G43B_087_087a	0.0	0.875	0.75	0.875	0.875	0.437	0.437	4.2	4.2	0.0	0.0
70	G50B_087_087a	0.0	0.875	0.875	0.875	0.875	0.437	0.437	6.3	6.3	0.0	0.0
71	G55B_100_100a	0.0	0.875	1.0	1.0	1.0	0.5	0.5	2.1	2.1	0.0	0.0
72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	0.5	15.0	15.0	0.0	0.0
73	G05B_100_100a	0.0	1.0	0.125	1.0	1.0	0.5	0.5	15.7	15.7	0.0	0.0
74	G11B_100_100a	0.0	1.0	0.25	1.0	1.0	0.5	0.5	16.4	16.4	0.0	0.0
75	G18B_100_100a	0.0	1.0	0.375	1.0	1.0	0.5	0.5	17.2	17.2	0.0	0.0
76	G25B_100_100a	0.0	1.0	0.5	1.0	1.0	0.5	0.5	18.0	18.0	0.0	0.0
77	G31B_100_100a	0.0	1.0	0.625	1.0	1.0	0.5	0.5	18.8	18.8	0.0	0.0
78	G38B_100_100a	0.0	1.0	0.75	1.0	1.0	0.5	0.5	19.6	19.6	0.0	0.0
79	G44B_100_100a	0.0	1.0	0.875	1.0	1.0	0.5	0.5	20.3	20.3	0.0	0.0
80	G50B_100_100a	0.0	1.0	1.0	1.0	1.0	0.5	0.5	21.0	21.0	0.0	0.0

Mean color difference of this page: $\Delta E^*_{ab} = 4.4$

1-0031930-F0

SE140-7N, Page 20/33-F

TUB-test chart SE14; 1080 colours, offset standard paper
colors and differences, ΔE^* , 3D=0, de=0, cmYk

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

1-0031930-F0

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsi.d	rgb**Md	LabCh**Md		
81	R00Y_012_012a	0.125 0.0 0.0	0.125 0.125 0.062	390	0.125 0.0 0.0	22.1 8.1 4.8	9.5 30.4	0.125 0.0 0.0	24.2 6.7 5.6	8.7 39.9 2.7	389	1.0 0.0 0.0	47.5 65.5 38.4	76.0 30.4
82	B50R_012_012a	0.125 0.0 0.125	0.125 0.125 0.062	330	0.125 0.0 0.125	22.2 9.2 -1.0	9.3 353.2	0.125 0.0 0.125	24.3 9.5 -2.7	9.9 343.7 2.7	330	1.0 0.0 0.0	48.2 74.2 -8.7	74.7 353.2
83	B25R_025_025a	0.125 0.0 0.25	0.25 0.25 0.125	300	0.125 0.0 0.25	23.1 13.4 -6.7	15.1 333.2	0.125 0.0 0.25	27.3 16.8 -10.1	19.6 328.9 6.3	300	0.5 0.0 0.0	37.0 53.9 -27.1	60.4 333.2
84	B15R_037_037a	0.125 0.0 0.375	0.375 0.375 0.187	289	0.118 0.0 0.375	23.5 15.9 -13.6	20.9 319.3	0.125 0.0 0.375	26.9 21.9 -17.4	28.0 321.6 7.8	288	0.316 0.0 0.0	32.0 42.4 -36.4	55.9 319.3
85	B11R_050_050a	0.125 0.0 0.5	0.5 0.5 0.25	284	0.116 0.0 0.5	24.1 17.7 -20.4	27.0 310.9	0.125 0.0 0.5	26.4 25.3 -23.7	34.7 316.9 8.6	282	0.233 0.0 0.0	29.6 35.4 -40.8	54.1 310.9
86	B09R_062_062a	0.125 0.0 0.625	0.625 0.625 0.312	281	0.114 0.0 0.625	24.9 21.0 -26.3	33.6 308.6	0.125 0.0 0.625	26.5 27.3 -30.1	40.7 312.2 7.5	279	0.183 0.0 0.0	28.8 33.6 -42.1	53.9 308.6
87	B07R_075_075a	0.125 0.0 0.75	0.75 0.75 0.375	279	0.112 0.0 0.75	25.8 24.3 -32.1	40.3 307.0	0.125 0.0 0.75	26.9 29.1 -35.3	45.8 309.4 5.8	278	0.15 0.0 0.0	28.2 32.4 -42.8	53.7 307.0
88	B06R_087_087a	0.125 0.0 0.875	0.875 0.875 0.437	278	0.116 0.0 0.875	26.8 27.8 -37.8	46.9 306.2	0.125 0.0 0.875	27.6 31.4 -40.0	50.9 308.0 4.3	277	0.133 0.0 0.0	28.0 31.7 -43.2	53.7 306.2
89	B05R_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	27.6 30.9 -43.8	53.6 305.2	0.125 0.0 1.0	27.8 31.4 -43.4	53.6 305.9 0.6	276	0.116 0.0 0.0	27.6 30.9 -43.8	53.6 305.2
90	Y00G_012_012a	0.125 0.125 0.0	0.125 0.125 0.062	90	0.125 0.125 0.0	27.3 -1.1 11.1	11.2 96.0	0.125 0.125 0.0	30.3 -3.0 8.7	9.2 108.8 4.1	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
91	NW_012a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	28.2 0.0 0.0	0.0 0.0	0.125 0.125 0.125	30.8 -0.2 -1.2	1.3 257.7 2.9	360	1.0 1.0 1.0	96.3 0.0 0.0	0.0 0.0
92	B00R_025_012a	0.125 0.125 0.25	0.25 0.125 0.187	270	0.124 0.124 0.25	29.0 2.8 -5.9	6.6 295.6	0.125 0.125 0.25	32.9 4.5 -10.7	11.6 292.8 6.4	270	0.0 0.0 0.0	24.9 22.9 -47.8	53.0 295.6
93	B00R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.124 0.375	29.8 5.7 -11.9	13.2 295.6	0.125 0.125 0.375	33.5 8.8 -17.1	19.3 297.4 7.0	270	0.0 0.0 0.0	24.9 22.9 -47.8	53.0 295.6
94	B00R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.124 0.5	30.6 8.6 -17.9	19.9 295.6	0.125 0.125 0.5	32.6 13.0 -23.2	26.7 299.3 7.2	270	0.0 0.0 0.0	24.9 22.9 -47.8	53.0 295.6
95	B00R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	31.4 11.4 -23.9	26.5 295.6	0.125 0.125 0.625	32.5 16.1 -28.9	33.1 299.2 6.9	270	0.0 0.0 0.0	24.9 22.9 -47.8	53.0 295.6
96	B00R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	32.2 14.3 -29.9	33.1 295.6	0.125 0.125 0.75	32.1 18.9 -34.3	39.2 298.9 6.3	270	0.0 0.0 0.0	24.9 22.9 -47.8	53.0 295.6
97	B00R_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	33.1 17.2 -35.8	39.8 295.6	0.125 0.125 0.875	32.6 22.2 -39.1	45.0 299.5 5.9	270	0.0 0.0 0.0	24.9 22.9 -47.8	53.0 295.6
98	B00R_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	33.9 20.1 -41.8	46.4 295.6	0.125 0.125 1.0	32.2 23.3 -42.8	48.8 298.5 3.8	270	0.0 0.0 0.0	24.9 22.9 -47.8	53.0 295.6
99	Y50G_025_025a	0.125 0.25 0.0	0.25 0.25 0.125	120	0.125 0.25 0.0	32.2 -7.5 15.2	16.9 116.4	0.125 0.25 0.0	38.4 -10.6 16.0	19.2 123.5 7.0	119	0.5 1.0 0.0	73.1 -60.2 60.8	67.9 116.4
100	G00B_025_012a	0.125 0.25 0.125	0.25 0.125 0.187	150	0.124 0.25 0.124	32.4 -8.6 2.8	9.1 161.6	0.125 0.25 0.125	37.8 -9.7 3.0	10.1 162.7 5.5	149	0.0 1.0 0.0	51.6 -69.3 23.0	73.1 161.6
101	G50B_025_012a	0.125 0.25 0.25	0.25 0.125 0.187	210	0.124 0.25 0.25	33.1 -3.9 -5.6	6.9 234.6	0.125 0.25 0.25	38.8 -5.7 -9.2	10.8 237.9 6.9	210	0.0 1.0 0.0	57.8 -31.9 -45.1	55.3 234.6
102	G75B_037_025a	0.125 0.25 0.375	0.375 0.25 0.25	240	0.124 0.25 0.375	34.2 -1.9 -11.5	11.7 260.4	0.125 0.25 0.375	40.1 -1.9 -15.6	15.7 263.0 7.1	240	0.0 0.5 1.0	42.3 -7.7 -46.3	49.6 260.4
103	G84B_050_037a	0.125 0.25 0.5	0.5 0.375 0.312	251	0.124 0.243 0.5	34.5 1.5 -17.5	17.6 275.0	0.125 0.25 0.5	39.3 1.8 -21.9	21.9 274.8 6.4	251	0.0 0.316 1.0	35.1 4.1 -46.9	47.1 275.0
104	G88B_062_050a	0.125 0.25 0.625	0.625 0.5 0.375	256	0.125 0.241 0.625	35.0 4.7 -23.6	24.0 281.4	0.125 0.25 0.625	38.8 5.5 -27.5	28.1 281.3 5.5	257	0.0 0.233 1.0	32.1 9.5 -47.2	48.1 281.4
105	G90B_075_062a	0.125 0.25 0.75	0.75 0.625 0.437	259	0.125 0.239 0.75	35.6 8.0 -29.7	30.7 285.1	0.125 0.25 0.75	37.4 9.6 -33.3	34.6 286.1 4.3	260	0.0 0.183 1.0	30.2 12.8 -47.5	49.2 285.1
106	G92B_087_075a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.237 0.875	36.1 11.3 -35.7	37.4 287.5	0.125 0.25 0.875	37.5 12.7 -37.9	40.0 288.5 3.0	262	0.0 0.15 1.0	29.0 15.0 -47.6	49.9 287.5
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.241 1.0	36.9 14.1 -41.6	40.0 288.7	0.125 0.25 1.0	36.8 14.9 -41.8	44.4 289.5 0.7	262	0.0 0.133 1.0	28.4 16.1 -47.6	50.3 288.7
108	Y68G_037_037a	0.125 0.375 0.0	0.375 0.375 0.187	131	0.118 0.375 0.0	36.0 -15.7 18.1	24.0 130.9	0.125 0.375 0.0	42.5 -19.0 20.7	28.1 132.6 7.7	131	0.316 1.0 0.0	65.1 -42.0 48.3	64.0 130.9
109	G00B_037_025a	0.125 0.375 0.125	0.375 0.25 0.25	150	0.124 0.375 0.124	36.5 -17.3 5.7	18.2 161.6	0.125 0.375 0.125	43.0 -17.2 7.4	18.7 156.4 6.7	149	0.0 1.0 0.0	51.6 -69.3 23.0	73.1 161.6
110	G25B_037_025a	0.125 0.375 0.25	0.375 0.25 0.25	180	0.124 0.375 0.25	37.3 -12.7 -4.3	13.4 198.8	0.125 0.375 0.25	44.0 -13.3 -5.3	14.3 202.0 6.8	180	0.0 1.0 0.5	54.6 -50.8 -17.3	53.7 198.8
111	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	38.1 -7.9 -11.2	13.8 234.6	0.125 0.375 0.375	45.2 -9.7 -14.5	17.5 236.2 8.0	210	0.0 1.0 1.0	57.8 -31.9 -45.1	55.3 234.6
112	G65B_050_037a	0.125 0.375 0.5	0.5 0.375 0.312	229	0.124 0.381 0.5	39.7 -7.0 -17.2	18.6 247.6	0.125 0.375 0.5	45.0 -7.4 -21.0	22.2 250.5 6.5	228	0.0 0.683 1.0	49.1 -18.8 -45.9	49.6 247.6
113	G75B_062_050a	0.125 0.375 0.625	0.625 0.5 0.375	240	0.125 0.375 0.625	40.1 -3.8 -23.1	23.4 260.4	0.125 0.375 0.625	44.8 -4.3 -26.7	27.1 260.6 5.9	240	0.0 0.5 1.0	42.3 -7.7 -46.3	49.6 260.4
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.364 0.75	40.2 -0.1 -29.0	29.0 269.7	0.125 0.375 0.75	43.0 0.4 -32.4	32.4 270.7 4.4	247	0.0 0.383 1.0	37.6 -0.2 -46.5	46.5 269.7
115	G84B_087_075a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.362 0.875	40.7 3.0 -35.1	33.5 275.0	0.125 0.375 0.875	43.1 3.2 -37.4	37.5 274.9 3.2	251	0.0 0.316 1.0	35.1 4.1 -46.9	47.1 275.0
116	G86B_100_087a	0.125 0.375 1.0	1.0 0.875 0.562	254	0.125 0.358 1.0	41.2 6.4 -41.1	41.6 278.9	0.125 0.375 1.0	40.9 7.3 -41.3	41.9 280.1 0.9	255	0.0 0.266 1.0	33.3 7.4 -47.0	47.6 278.9
117	Y76G_050_050a	0.125 0.5 0.0	0.5 0.5 0.25	136	0.116 0.5 0.0	39.4 -24.3 20.6	31.9 139.7	0.125 0.5 0.0	45.8 -27.0 24.1	36.2 138.2 7.7	137	0.233 1.0 0.0	60.3 -48.7 41.3	63.9 139.7
118	G00B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.124	40.6 -26.0 8.6	27.4 161.6	0.125 0.5 0.125	46.3 -24.7 11.1	27.1 155.7 6.3	149	0.0 1.0 0.0	51.6 -69.3 23.0	73.1 161.6
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.243	41.3 -22.2 -0.7	22.2 181.9	0.125 0.5 0.25	47.4 -21.0 -1.1	21.0 183.0 6.2	168	0.0 1.0 0.316	53.4 -59.2 -2.0	59.3 181.9
120	G34B_050_037a	0.125 0.5 0.375	0.5 0.375 0.312	191	0.124 0.5 0.381	42.3 -16.2 -11.1	19.7 214.4	0.125 0.5 0.375	48.9 -16.6 -12.5	21.0 216.5 6.7	191	0.0 1.0 0.683	56.0 -43.3 -29.7	52.5 214.4
121	G50B_050_037a	0.125 0.5 0.5	0.5 0.375 0.312	210	0.124 0.5 0.5	43.0 -11.9 -16.9	20.7 234.6	0.125 0.5 0.5	49.3 -13.9 -20.0	24.2 235.7 7.1	210	0.0 1.0 1.0	57.8 -31.9 -45.1	55.3 234.6
122	G61B_062_050a	0.125 0.5 0.625	0.625 0.5 0.375	224	0.125 0.508 0.625	44.9 -11.4 -22.8	25.5 243.3	0.125 0.5 0.625	49.9 -11.9 -25.8	28.5 245.2 5.9	222	0.0 0.766 1.0	51.8 -22.8 -45.6	51.0 243.3
123	G69B_075_062a	0.125 0.5 0.75	0.75 0.625 0.437	233	0.125 0.51 0.75	46.0 -9.6 -28.7	30.3 251.4	0.125 0.5 0.75	49.2 -9.0 -31.4	32.6 253.9 4.1	232	0.0 0.616 1.0	46.9 -15.4 -46.0	48.5 251.4
124	G75B_087_075a	0.125 0.5 0.875	0.875 0.75 0.5	240	0.125 0.5 0.875	46.1 -5.8 -34.7	35.2 260.4	0.125 0.5 0.875	48.8 -5.8 -36.5	36.9 260.8 3.2	240	0.0 0.5 1.0	42.3 -7.7 -46.3	46.9 260.4
125	G79B_100_087a	0.125 0.5 1.0	1.0 0.875 0.562	245	0.125 0.489 1.0	46.1 -2.0 -40.7	40.8 267.0	0.125 0.5 1.0	46.1 -1.2 -40.6	40.6 268.2 0.8	245	0.0 0.416 1.0	38.9 -2.3 -46.5	46.6 267.0
126	Y81G_062_062a	0.125 0.625 0.0	0.625 0.625 0.312	139	0.114 0.625 0.0	43.6 -32.6 23.6	40.3 144.1	0.125 0.625 0.0	49.1 -35.0 26.8	44.1 142.5 6.7	149	0.183 1.0 0.0	58.6 -52.3 37.8	64.5 144.1
127	G00B_062_050a	0.125 0.625 0.125	0.625 0.5 0.375	150	0.125 0.625 0.125	44.8 -34.6 11.5	36.5 161.6	0.125 0.625 0.125	49.8 -32.5 14.5	35.6 155.8 6.2	149	0.0 1.0 0.0	51.6 -69.3 23.0	73.1 161.6
128	G11B_062_050a	0.125 0.625 0.25	0.625 0.5 0.375	164	0.125 0.625 0.241	45.4 -31.2 2.6	31.3 175.2	0.125 0.625 0.25	51.0 -28.8 2.9	28.9 174.1 6.0	162	0.0 1.0 0.233	52.9 -62.5 5.2	62.7 175.2
129	G25B_062_050a	0.125 0.625 0.375	0.625 0.5 0.375	180	0.125 0.625 0.375	46.3 -25.4 -8.6	26.8 198.8	0.125 0.625 0.375	52.1 -2					

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n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md	
162	R00Y_025_025a	0.25 0.0 0.0	0.25 0.25 0.125	390	0.25 0.0 0.0	25.7 16.3 9.6	19.0 30.4	0.25 0.0 0.0	29.3 16.0 12.5	20.3 37.9 4.5	389	1.0 0.0 0.0	47.5 65.5 38.4 76.0 30.4
163	R00Y_025_025a	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.125	25.8 17.4 2.8	17.6 9.2	0.25 0.0 0.125	29.6 18.6 1.9	18.7 6.0 4.1	360	1.0 0.0 0.5	47.8 69.7 11.3 70.6 9.2
164	B50R_025_025a	0.25 0.0 0.25	0.25 0.25 0.125	330	0.25 0.0 0.25	25.9 18.5 -2.1	18.6 353.2	0.25 0.0 0.25	30.0 21.2 -5.5	21.9 345.2 5.9	330	1.0 0.0 1.0	48.2 74.2 -8.7 74.7 353.2
165	B34R_037_037a	0.25 0.0 0.375	0.375 0.375 0.187	311	0.256 0.0 0.375	27.2 23.6 -7.2	24.7 342.9	0.25 0.0 0.375	31.0 27.7 -11.0	29.8 338.2 6.7	311	0.683 0.0 1.0	41.6 63.1 -19.3 66.0 342.9
166	B25R_050_050a	0.25 0.0 0.5	0.5 0.5 0.25	300	0.25 0.0 0.5	27.8 26.9 -13.9	30.2 333.2	0.25 0.0 0.5	30.1 31.5 -17.8	36.2 330.5 6.6	300	0.5 0.0 1.0	37.0 53.9 -27.1 60.4 333.2
167	B19R_062_062a	0.25 0.0 0.625	0.625 0.625 0.312	293	0.239 0.0 0.625	28.2 30.1 -19.5	36.1 326.4	0.25 0.0 0.625	30.2 33.7 -24.1	41.4 324.4 5.9	292	0.383 0.0 1.0	34.0 48.1 -31.9 57.7 326.4
168	B15R_075_075a	0.25 0.0 0.75	0.75 0.75 0.375	289	0.237 0.0 0.75	28.6 31.8 -27.3	41.9 319.3	0.25 0.0 0.75	30.2 35.4 -30.0	46.4 319.7 4.7	288	0.316 0.0 1.0	32.0 42.4 -36.4 55.9 319.3
169	B13R_087_087a	0.25 0.0 0.875	0.875 0.875 0.437	286	0.233 0.0 0.875	28.9 32.9 -34.5	47.8 313.6	0.25 0.0 0.875	30.2 36.7 -35.9	51.4 315.6 4.2	284	0.266 0.0 1.0	30.4 37.7 -39.5 54.6 313.6
170	B11R_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	29.6 35.4 -40.8	54.1 310.9	0.25 0.0 1.0	29.9 36.0 -40.4	54.1 311.7 0.7	282	0.233 0.0 1.0	29.6 35.4 -40.8 54.1 310.9
171	R50Y_025_025a	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.125 0.0	30.9 6.0 15.7	16.8 69.1	0.25 0.125 0.0	37.3 2.7 18.2	18.4 81.3 7.6	59	1.0 0.5 0.0	68.1 24.0 63.0 67.4 69.1
172	R00Y_025_012a	0.25 0.125 0.125	0.25 0.125 0.187	390	0.25 0.124 0.124	31.8 8.1 4.8	9.5 30.4	0.25 0.125 0.125	36.7 7.1 6.8	9.8 43.9 5.3	389	1.0 0.0 0.0	47.5 65.5 38.4 76.0 30.4
173	B50R_025_012a	0.25 0.125 0.25	0.25 0.125 0.187	330	0.25 0.124 0.25	31.9 9.2 -1.0	9.3 353.2	0.25 0.125 0.25	37.4 10.0 -4.4	10.9 335.9 6.4	330	1.0 0.0 1.0	48.2 74.2 -8.7 74.7 353.2
174	B25R_037_025a	0.25 0.125 0.375	0.375 0.25 0.25	300	0.25 0.124 0.375	32.9 13.4 -6.7	15.1 333.2	0.25 0.125 0.375	38.5 14.3 -10.6	17.8 323.3 6.9	300	0.5 0.0 1.0	37.0 53.9 -27.1 60.4 333.2
175	B15R_050_037a	0.25 0.125 0.5	0.5 0.375 0.312	289	0.243 0.124 0.5	33.3 15.9 -13.6	20.9 319.3	0.25 0.125 0.5	36.9 18.1 -17.3	25.1 316.1 5.6	288	0.316 0.0 1.0	32.0 42.4 -36.4 55.9 319.3
176	B11R_062_050a	0.25 0.125 0.625	0.625 0.5 0.375	284	0.241 0.125 0.625	33.8 17.7 -20.4	27.0 310.9	0.25 0.125 0.625	36.9 20.9 -23.3	31.3 311.8 5.2	282	0.233 0.0 1.0	29.6 35.4 -40.8 54.1 310.9
177	B09R_075_062a	0.25 0.125 0.75	0.75 0.625 0.437	281	0.239 0.125 0.75	34.7 21.0 -26.3	33.6 308.6	0.25 0.125 0.75	36.0 23.8 -29.1	37.6 309.3 4.2	279	0.183 0.0 1.0	28.8 33.6 -42.1 53.9 308.6
178	B07R_087_075a	0.25 0.125 0.875	0.875 0.75 0.5	279	0.237 0.125 0.875	35.5 24.3 -32.1	40.3 307.0	0.25 0.125 0.875	35.5 26.6 -35.2	44.2 307.0 3.9	278	0.15 0.0 1.0	28.2 32.4 -42.8 53.7 307.0
179	B06R_100_087a	0.25 0.125 1.0	1.0 0.875 0.562	278	0.241 0.125 1.0	36.5 27.8 -37.8	46.9 306.2	0.25 0.125 1.0	34.9 27.3 -39.6	48.1 304.6 2.4	277	0.133 0.0 1.0	28.0 31.7 -43.2 53.7 306.2
180	Y00G_025_025a	0.25 0.25 0.0	0.25 0.25 0.125	90	0.25 0.25 0.0	36.2 -2.3 22.2	22.4 96.0	0.25 0.25 0.0	42.5 -5.4 22.3	23.0 103.7 7.0	89	1.0 1.0 0.0	89.4 -9.5 89.0 89.6 96.0
181	Y00G_025_012a	0.25 0.25 0.125	0.25 0.125 0.187	90	0.25 0.25 0.124	37.1 -1.1 11.1	11.2 96.0	0.25 0.25 0.125	43.9 -3.4 10.7	11.2 107.9 7.1	89	1.0 1.0 0.0	89.4 -9.5 89.0 89.6 96.0
182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	37.9 0.0 0.0	0.0 0.0	0.25 0.25 0.25	45.2 -0.4 -2.1	2.2 259.4 7.5	360	1.0 1.0 1.0	96.3 0.0 0.0 0.0 0.0
183	B00R_037_012a	0.25 0.25 0.375	0.375 0.125 0.312	270	0.249 0.249 0.375	38.8 2.8 -5.9	6.6 295.6	0.25 0.25 0.375	45.0 3.5 -9.6	10.3 290.2 7.2	270	1.0 0.0 1.0	24.9 22.9 -47.8 53.0 295.6
184	B00R_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.249 0.5	39.6 5.7 -11.9	13.2 295.6	0.25 0.25 0.5	44.0 6.6 -16.3	17.6 292.2 6.2	270	1.0 0.0 1.0	24.9 22.9 -47.8 53.0 295.6
185	B00R_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	40.4 8.6 -17.9	19.9 295.6	0.25 0.25 0.625	43.6 10.2 -21.5	23.8 295.3 5.1	270	1.0 0.0 1.0	24.9 22.9 -47.8 53.0 295.6
186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	41.2 11.4 -23.9	26.5 295.6	0.25 0.25 0.75	42.6 13.7 -27.5	30.7 296.4 4.4	270	1.0 0.0 1.0	24.9 22.9 -47.8 53.0 295.6
187	B00R_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.0 14.3 -29.9	33.1 295.6	0.25 0.25 0.875	41.6 16.9 -33.7	37.7 296.6 4.6	270	1.0 0.0 1.0	24.9 22.9 -47.8 53.0 295.6
188	B00R_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	42.8 17.2 -35.8	39.8 295.6	0.25 0.25 1.0	40.3 17.7 -38.3	42.2 294.8 3.4	270	1.0 0.0 1.0	24.9 22.9 -47.8 53.0 295.6
189	Y31G_037_037a	0.25 0.375 0.0	0.375 0.375 0.187	109	0.256 0.375 0.0	41.8 -7.7 27.7	28.8 105.6	0.25 0.375 0.0	48.6 -11.7 29.5	31.7 111.6 8.0	108	0.683 1.0 0.0	80.6 -20.7 74.1 76.9 105.6
190	Y50G_037_025a	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	41.9 -7.5 15.2	16.9 116.4	0.25 0.375 0.125	49.3 -10.3 15.6	18.7 123.4 7.9	119	0.5 1.0 0.0	73.1 -30.2 60.8 67.9 116.4
191	G00B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	42.1 -8.6 2.8	9.1 161.6	0.25 0.375 0.25	49.7 -8.0 2.5	8.4 162.1 7.6	149	0.0 1.0 0.0	51.6 -69.3 23.0 73.1 161.6
192	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	42.9 -3.9 -5.6	6.9 234.6	0.25 0.375 0.375	51.3 -4.8 -8.1	9.5 239.2 8.8	210	0.0 1.0 1.0	57.8 -31.9 -45.1 55.3 234.6
193	G75B_050_025a	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.375 0.5	43.9 -1.9 -11.5	11.7 260.4	0.25 0.375 0.5	50.5 -1.8 -14.6	14.7 262.7 7.2	240	0.0 0.5 1.0	42.3 -7.7 -46.3 46.9 260.4
194	G84B_062_037a	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.368 0.625	44.2 1.5 -17.5	17.6 275.0	0.25 0.375 0.625	50.0 1.0 -20.5	20.5 273.0 6.5	251	0.0 0.316 1.0	35.1 4.1 -46.9 47.1 275.0
195	G88B_075_050a	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.366 0.75	44.8 4.7 -23.6	24.0 281.4	0.25 0.375 0.75	48.3 4.9 -26.5	27.0 280.5 4.6	257	0.0 0.233 1.0	32.1 9.5 -47.2 48.1 281.4
196	G90B_087_062a	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.364 0.875	45.3 8.0 -29.7	30.7 285.1	0.25 0.375 0.875	47.3 7.6 -32.4	33.3 283.2 3.4	260	0.0 0.183 1.0	30.2 12.8 -47.5 49.2 285.1
197	G92B_100_075a	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.362 1.0	45.8 11.3 -35.7	37.4 287.5	0.25 0.375 1.0	44.6 10.8 -37.4	39.0 286.1 2.2	262	0.0 0.15 1.0	29.0 15.0 -47.6 49.9 287.5
198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	45.8 -11.5 30.4	33.9 116.4	0.25 0.5 0.0	51.9 -19.0 34.0	39.0 119.2 8.1	119	0.5 1.0 0.0	73.1 -30.2 60.8 67.9 116.4
199	Y68G_050_037a	0.25 0.5 0.125	0.5 0.375 0.312	131	0.243 0.5 0.124	45.7 -15.7 18.1	24.0 130.9	0.25 0.5 0.125	52.0 -17.9 19.5	26.5 132.5 6.8	131	0.316 1.0 0.0	65.1 -42.0 48.3 64.0 130.9
200	G00B_050_025a	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.249	46.2 -17.3 5.7	18.2 161.6	0.25 0.5 0.25	52.9 -15.1 6.6	16.5 156.4 7.0	149	0.0 1.0 0.0	51.6 -69.3 23.0 73.1 161.6
201	G25B_050_025a	0.25 0.5 0.375	0.5 0.25 0.375	180	0.249 0.5 0.375	47.0 -12.7 -4.3	13.4 198.8	0.25 0.5 0.375	54.4 -11.9 -4.6	12.7 201.1 7.4	180	0.0 1.0 0.5	54.6 -50.8 -17.3 53.7 198.8
202	G50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	47.8 -7.9 -11.2	13.8 234.6	0.25 0.5 0.5	55.7 -8.8 -13.5	16.1 236.8 8.2	210	0.0 1.0 1.0	57.8 -31.9 -45.1 55.3 234.6
203	G65B_062_037a	0.25 0.5 0.625	0.625 0.375 0.437	229	0.25 0.506 0.625	49.4 -7.0 -17.2	18.6 247.6	0.25 0.5 0.625	56.0 -6.7 -19.3.				

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

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md				
243	R00Y_037_037a	0.375 0.0 0.0	0.375 0.375 0.187	390	0.375 0.0 0.0	29.4 24.5 14.4	28.5 30.4	0.375 0.0 0.0	32.3 26.8 18.3	32.5 34.3 5.4	389	1.0 0.0 0.0	47.5 65.5 38.4	76.0 30.4
244	R18Y_037_037a	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.118	29.5 25.4 8.1	26.6 17.7	0.375 0.0 0.125	32.5 26.8 8.6	29.8 16.7 4.4	371	1.0 0.0 0.316	47.8 67.7 21.6	71.1 17.7
245	B65R_037_037a	0.375 0.0 0.25	0.375 0.375 0.187	349	0.375 0.0 0.256	29.6 26.7 0.8	26.8 1.8	0.375 0.0 0.25	33.1 31.2 -0.8	31.2 35.8 5.9	348	1.0 0.0 0.683	48.1 71.4 2.3	71.4 1.8
246	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	29.6 27.8 -3.2	28.0 353.2	0.375 0.0 0.375	33.1 -7.1 33.9	347.9 7.4 330	330	1.0 0.0 1.0	48.2 74.2 -8.7	74.7 353.2
247	B38R_050_050a	0.375 0.0 0.5	0.5 0.5 0.25	316	0.383 0.0 0.5	31.1 33.6 -7.6	34.5 347.2	0.375 0.0 0.5	33.1 39.7 -11.9	41.5 343.3 7.7	317	0.766 0.0 1.0	43.6 67.3 -15.2	69.0 347.2
248	B30R_062_062a	0.375 0.0 0.625	0.625 0.625 0.312	307	0.385 0.0 0.625	31.9 37.1 -14.0	39.7 339.2	0.375 0.0 0.625	33.8 44.0 -17.2	47.3 338.6 7.8	307	0.616 0.0 1.0	40.0 59.4 -22.5	63.5 339.2
249	B25R_075_075a	0.375 0.0 0.75	0.75 0.75 0.375	300	0.375 0.0 0.75	32.4 40.4 -20.3	45.3 333.2	0.375 0.0 0.75	33.7 45.5 -23.0	51.0 333.2 5.8	300	0.5 0.0 1.0	37.0 53.9 -27.1	60.4 333.2
250	B20R_087_087a	0.375 0.0 0.875	0.875 0.875 0.437	295	0.364 0.0 0.875	32.8 43.6 -26.8	51.2 328.4	0.375 0.0 0.875	33.8 47.5 -28.5	55.4 329.0 4.3	294	0.416 0.0 1.0	34.8 49.8 -30.6	58.5 328.4
251	B18R_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	33.5 47.0 -32.8	57.3 325.0	0.375 0.0 1.0	33.7 47.7 -32.2	57.5 325.9 1.0	291	0.366 0.0 1.0	33.5 47.0 -32.8	57.3 325.0
252	R31Y_037_037a	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.118 0.0	33.9 14.9 20.0	25.0 53.2	0.375 0.125 0.0	39.7 12.8 23.8	27.1 61.6 7.2	48	1.0 0.316 0.0	59.6 39.8 53.5	66.7 53.2
253	R00Y_037_025a	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.124	35.5 16.3 9.6	19.0 30.4	0.375 0.125 0.125	39.8 15.7 12.9	20.4 39.4 5.5	389	1.0 0.0 0.0	47.5 65.5 38.4	76.0 30.4
254	R00Y_037_025a	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.25	35.5 17.4 2.8	17.6 9.2	0.375 0.125 0.25	40.4 17.9 1.9	18.0 6.0 4.9	360	1.0 0.0 0.5	47.8 69.7 11.3	70.6 9.2
255	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.124 0.375	35.6 18.5 -2.1	18.6 353.2	0.375 0.125 0.375	40.8 20.2 -6.3	21.1 342.6 6.7	330	1.0 0.0 1.0	48.2 74.2 -8.7	74.7 353.2
256	B34R_050_037a	0.375 0.125 0.5	0.5 0.375 0.312	311	0.381 0.124 0.5	36.9 23.6 -7.2	24.7 342.9	0.375 0.125 0.5	40.0 26.8 -11.7	29.3 336.3 6.3	311	0.683 0.0 1.0	41.6 63.1 -19.3	66.0 342.9
257	B25R_062_050a	0.375 0.125 0.625	0.625 0.5 0.375	300	0.375 0.125 0.625	37.5 26.9 -13.5	30.2 333.2	0.375 0.125 0.625	40.0 30.6 -17.1	35.1 330.8 5.7	300	0.5 0.0 1.0	37.0 53.9 -27.1	60.4 333.2
258	B19R_075_062a	0.375 0.125 0.75	0.75 0.625 0.437	293	0.364 0.125 0.75	37.9 30.1 -19.9	36.1 326.4	0.375 0.125 0.75	39.0 33.8 -22.8	40.8 326.0 4.8	292	0.383 0.0 1.0	34.0 48.1 -31.9	57.7 326.4
259	B15R_087_075a	0.375 0.125 0.875	0.875 0.75 0.5	289	0.362 0.125 0.875	38.3 31.8 -27.3	41.9 319.3	0.375 0.125 0.875	39.1 37.2 -27.8	46.4 323.2 5.4	288	0.316 0.0 1.0	32.0 42.4 -36.4	55.9 319.3
260	B13R_100_087a	0.375 0.125 1.0	1.0 0.875 0.562	286	0.358 0.125 1.0	38.7 32.9 -34.5	47.8 313.6	0.375 0.125 1.0	38.3 37.9 -31.9	49.5 319.8 5.5	284	0.266 0.0 1.0	30.4 37.7 -39.5	54.6 313.6
261	R68Y_037_037a	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.256 0.0	40.6 3.1 28.0	28.1 83.6	0.375 0.25 0.0	49.0 0.5 31.4	31.4 88.9 9.3	71	1.0 0.683 0.0	77.5 8.3 74.7	75.1 83.6
262	R50Y_037_025a	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.25 0.124	40.6 6.0 15.7	16.8 69.1	0.375 0.25 0.125	48.9 3.5 17.7	18.0 78.7 8.8	59	1.0 0.5 0.0	68.1 24.0 63.0	67.4 69.1
263	R00Y_037_012a	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.249	41.6 8.1 4.8	9.5 30.4	0.375 0.25 0.25	49.4 6.5 5.7	8.7 41.3 8.0	389	1.0 0.0 0.0	47.5 65.5 38.4	76.0 30.4
264	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.249 0.375	41.7 9.2 -1.0	9.3 353.2	0.375 0.25 0.375	50.3 9.1 -4.4	10.1 334.0 9.3	330	1.0 0.0 1.0	48.2 74.2 -8.7	74.7 353.2
265	B25R_050_025a	0.375 0.25 0.5	0.5 0.25 0.375	300	0.375 0.249 0.5	42.6 13.4 -6.7	15.1 333.2	0.375 0.25 0.5	48.8 14.4 -10.8	18.0 323.1 7.4	300	0.5 0.0 1.0	37.0 53.9 -27.1	60.4 333.2
266	B15R_062_037a	0.375 0.25 0.625	0.625 0.375 0.437	289	0.368 0.25 0.625	43.0 15.9 -13.6	20.9 319.3	0.375 0.25 0.625	47.4 18.8 -16.8	25.2 318.2 6.1	288	0.316 0.0 1.0	32.0 42.4 -36.4	55.9 319.3
267	B11R_075_050a	0.375 0.25 0.75	0.75 0.5 0.5	284	0.366 0.25 0.75	43.5 17.7 -20.4	27.0 310.9	0.375 0.25 0.75	46.2 22.8 -22.1	31.8 315.8 6.0	282	0.233 0.0 1.0	29.6 35.4 -40.8	54.1 310.9
268	B09R_087_062a	0.375 0.25 0.875	0.875 0.625 0.562	281	0.364 0.25 0.875	44.4 21.0 -26.3	33.6 308.6	0.375 0.25 0.875	46.0 26.4 -26.8	37.6 314.6 5.6	279	0.183 0.0 1.0	28.8 33.6 -42.1	53.9 308.6
269	B07R_100_075a	0.375 0.25 1.0	1.0 0.75 0.625	279	0.362 0.25 1.0	45.3 24.3 -32.1	40.3 307.0	0.375 0.25 1.0	45.0 27.6 -30.5	41.2 312.1 3.7	278	0.15 0.0 1.0	28.2 32.4 -42.8	53.7 307.0
270	Y00G_037_037a	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.375 0.0	45.1 -3.5 33.4	33.6 96.0	0.375 0.375 0.0	54.3 -7.1 35.8	36.5 101.2 10.1	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
271	Y00G_037_025a	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.375 0.124	45.9 -2.3 22.2	22.4 96.0	0.375 0.375 0.125	55.5 -5.3 21.8	22.4 103.8 10.0	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
272	Y00G_037_012a	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.375 0.249	46.8 -1.1 11.1	11.2 96.0	0.375 0.375 0.25	56.4 -2.9 8.8	9.3 108.5 10.0	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
273	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	47.7 0.0 0.0	0.0 0.0	0.375 0.375 0.375	57.6 -0.3 -2.0	2.0 261.2 10.1	360	1.0 1.0 1.0	96.3 0.0 0.0	0.0 0.0
274	B00R_050_012a	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.375 0.5	48.5 2.8 -5.9	6.6 295.6	0.375 0.375 0.5	56.0 4.3 -8.8	9.9 296.2 8.2	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
275	B00R_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	49.3 5.7 -11.9	13.2 295.6	0.375 0.375 0.625	54.9 8.3 -14.7	16.9 299.3 6.7	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
276	B00R_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	50.1 8.6 -17.9	19.9 295.6	0.375 0.375 0.75	52.9 12.5 -20.5	24.1 301.4 5.5	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
277	B00R_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	50.9 11.4 -23.9	26.5 295.6	0.375 0.375 0.875	52.8 16.1 -24.9	29.6 302.9 5.1	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
278	B00R_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	51.7 14.3 -29.9	33.1 295.6	0.375 0.375 1.0	50.7 19.0 -29.1	34.8 303.2 4.9	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
279	Y23G_050_050a													

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

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb**Md	LabCh**Md	
324	R00Y_050_050a	0.5	0.0	0.0	0.5	0.5	0.0	0.0	33.0	32.7	19.2	38.0	30.4
325	R26Y_050_050a	0.5	0.0	0.125	0.5	0.5	0.116	33.1	33.4	13.1	35.9	21.4	30.4
326	R00Y_050_050a	0.5	0.0	0.25	0.5	0.5	0.25	33.1	34.8	5.6	35.3	9.2	30.4
327	B61R_050_050a	0.5	0.0	0.375	0.5	0.5	0.383	33.3	36.1	-0.6	36.1	358.9	30.4
328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.5	33.3	37.1	-4.3	37.3	353.2	30.4
329	B40R_062_062a	0.5	0.0	0.625	0.625	0.625	0.625	34.9	43.0	-8.6	43.9	348.6	30.4
330	B34R_075_075a	0.5	0.0	0.75	0.75	0.75	0.75	35.9	47.3	-14.4	49.5	342.9	30.4
331	B29R_087_087a	0.5	0.0	0.875	0.875	0.875	0.875	36.6	50.6	-20.9	54.8	337.5	30.4
332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	1.0	37.0	53.9	-27.1	60.4	333.2	30.4
333	R23Y_050_050a	0.5	0.125	0.0	0.5	0.5	0.116	0.0	37.2	23.6	24.3	33.9	45.8
334	R00Y_050_037a	0.5	0.125	0.125	0.5	0.375	0.312	39.0	0.5	0.124	0.124	39.1	24.5
335	R18Y_050_037a	0.5	0.125	0.25	0.5	0.375	0.312	37.1	0.5	0.124	0.243	39.2	25.4
336	B65R_050_037a	0.5	0.125	0.375	0.5	0.375	0.312	34.9	0.5	0.124	0.381	39.3	26.7
337	B50R_050_037a	0.5	0.125	0.5	0.5	0.375	0.312	33.0	0.5	0.124	0.5	39.4	27.8
338	B38R_062_050a	0.5	0.125	0.625	0.625	0.5	0.375	31.6	0.508	0.125	0.625	40.8	33.6
339	B30R_075_062a	0.5	0.125	0.75	0.75	0.625	0.437	30.7	0.51	0.125	0.75	41.7	37.1
340	B25R_087_075a	0.5	0.125	0.875	0.875	0.75	0.5	30.0	0.5	0.125	0.875	42.1	40.4
341	B20R_100_087a	0.5	0.125	1.0	1.0	0.875	0.562	29.5	0.489	0.125	1.0	42.5	43.6
342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	6.0	0.5	0.25	0.0	43.0	31.5
343	R31Y_050_037a	0.5	0.25	0.125	0.5	0.375	0.312	4.9	0.5	0.243	0.124	43.7	14.9
344	R00Y_050_025a	0.5	0.25	0.25	0.5	0.25	0.375	39.0	0.5	0.249	0.249	45.2	16.3
345	R00Y_050_025a	0.5	0.25	0.375	0.5	0.25	0.375	36.0	0.5	0.249	0.375	45.3	17.4
346	B50R_050_025a	0.5	0.25	0.5	0.5	0.25	0.375	33.0	0.5	0.249	0.5	45.4	18.6
347	B34R_062_037a	0.5	0.25	0.625	0.625	0.375	0.437	31.1	0.506	0.25	0.625	46.6	23.6
348	B25R_075_050a	0.5	0.25	0.75	0.75	0.5	0.5	30.0	0.5	0.25	0.75	47.2	26.9
349	B19R_087_062a	0.5	0.25	0.875	0.875	0.625	0.562	29.3	0.489	0.25	0.875	47.6	30.1
350	B15R_100_075a	0.5	0.25	1.0	1.0	0.75	0.625	28.9	0.487	0.25	1.0	48.1	31.8
351	R76Y_050_050a	0.5	0.375	0.0	0.5	0.5	0.25	7.6	0.5	0.383	0.0	49.8	1.2
352	R68Y_050_037a	0.5	0.375	0.125	0.5	0.375	0.312	7.1	0.5	0.381	0.124	50.4	3.1
353	R50Y_050_025a	0.5	0.375	0.25	0.5	0.25	0.375	6.0	0.5	0.375	0.249	50.3	6.0
354	R00Y_050_012a	0.5	0.375	0.375	0.5	0.125	0.437	39.0	0.5	0.375	0.375	51.3	8.1
355	B50R_050_012a	0.5	0.375	0.5	0.5	0.125	0.437	33.0	0.5	0.375	0.5	51.4	9.2
356	B25R_062_025a	0.5	0.375	0.625	0.625	0.25	0.5	30.0	0.5	0.375	0.625	52.3	13.4
357	B15R_075_037a	0.5	0.375	0.75	0.75	0.375	0.562	28.9	0.493	0.375	0.75	52.7	15.9
358	B11R_087_050a	0.5	0.375	0.875	0.875	0.5	0.625	28.4	0.491	0.375	0.875	53.3	17.7
359	B09R_100_062a	0.5	0.375	1.0	1.0	0.625	0.687	28.1	0.489	0.375	1.0	54.1	21.0
360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	9.0	0.5	0.5	0.0	53.9	-4.7
361	Y00G_050_037a	0.5	0.5	0.125	0.5	0.375	0.312	9.0	0.5	0.5	0.124	54.8	-3.5
362	Y00G_050_025a	0.5	0.5	0.25	0.5	0.25	0.375	9.0	0.5	0.5	0.249	55.7	-2.3
363	Y00G_050_012a	0.5	0.5	0.375	0.5	0.125	0.437	9.0	0.5	0.5	0.375	56.5	-1.1
364	NW_050a	0.5	0.5	0.5	0.5	0.0	0.5	36.0	0.5	0.5	0.5	57.4	0.0
365	B00R_062_012a	0.5	0.5	0.625	0.625	0.125	0.562	27.0	0.5	0.625	0.125	60.2	2.8
366	B00R_075_025a	0.5	0.5	0.75	0.75	0.25	0.625	27.0	0.5	0.5	0.75	59.0	5.7
367	B00R_087_037a	0.5	0.5	0.875	0.875	0.375	0.687	27.0	0.5	0.5	0.875	59.8	8.6
368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	27.0	0.5	0.5	1.0	60.6	11.4
369	Y18G_062_062a	0.5	0.625	0.0	0.625	0.625	0.312	10.1	0.51	0.625	0.0	60.2	-9.8
370	Y23G_062_050a	0.5	0.625	0.125	0.625	0.5	0.375	10.4	0.508	0.625	0.125	61.0	-8.6
371	Y31G_062_037a	0.5	0.625	0.25	0.625	0.375	0.437	10.9	0.506	0.625	0.25	61.2	-7.7
372	Y50G_062_025a	0.5	0.625	0.375	0.625	0.25	0.5	12.0	0.5	0.625	0.375	61.3	-7.5
373	G00B_062_012a	0.5	0.625	0.5	0.625	0.125	0.562	15.0	0.5	0.625	0.5	61.5	-8.6
374	G50B_062_012a	0.5	0.625	0.625	0.625	0.125	0.562	21.0	0.5	0.625	0.625	62.3	-3.9
375	G75B_075_025a	0.5	0.625	0.75	0.75	0.25	0.625	24.0	0.5	0.625	0.75	63.4	-1.9
376	G84B_087_037a	0.5	0.625	0.875	0.875	0.375	0.687	25.1	0.5	0.618	0.875	63.7	1.5
377	G88B_100_050a	0.5	0.625	1.0	1.0	0.5	0.75	25.6	0.5	0.616	1.0	64.2	4.7
378	Y31G_075_075a	0.5	0.75	0.0	0.75	0.75	0.375	10.9	0.512	0.75	0.0	65.1	-15.5
379	Y38G_075_062a	0.5	0.75	0.125	0.75	0.625	0.437	11.3	0.51	0.75	0.125	65.2	-14.8
380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	12.0	0.5	0.75	0.25	65.3	-15.1
381	Y68G_075_037a	0.5	0.75	0.375	0.75	0.375	0.562	13.1	0.493	0.75	0.375	65.2	-15.7
382	G00B_075_025a	0.5	0.75	0.5	0.75	0.25	0.625	15.0	0.5	0.75	0.5	65.7	-17.3
383	G25B_075_025a	0.5	0.75	0.625	0.75	0.25	0.625	18.0	0.5	0.75	0.625	66.4	-12.7
384	G50B_075_025a	0.5	0.75	0.75	0.75	0.25	0.625	21.0	0.5	0.75	0.75	67.2	-7.9
385	G65B_087_037a	0.5	0.75	0.875	0.875	0.375	0.687	22.9	0.5	0.756	0.875	68.9	-7.0
386	G75B_100_050a	0.5	0.75	1.0	1.0	0.5	0.75	24.0	0.5	0.75	1.0	69.3	-3.8
387	Y41G_087_087a	0.5	0.875	0.0	0.875	0.875	0.437	11.5	0.51	0.875	0.0	69.1	-22.5
388	Y50G_087_075a	0.5	0.875	0.125	0.875	0.75	0.5	12.0	0.5	0.875	0.125	69.2	-22.6
389	Y61G_087_062a	0.5	0.875	0.25	0.875	0.625	0.562	12.7	0.489	0.875	0.25	69.6	-22.6
390	Y76G_087_050a	0.5	0.875	0.375	0.875	0.5	0.625	13.6	0.491	0.875	0.375	68.6	-24.3
391	G00B_087_037a	0.5	0.875	0.5	0.875	0.375	0.687	15.0	0.5	0.875	0.5	69.8	-26.0
392	G15B_087_037a	0.5	0.875	0.625	0.875	0.375	0.687	16.9	0.5	0.875	0.625	70.5	-22.2
393	G34B_087_037a	0.5	0.875	0.75	0.875	0.375	0.687	19.1	0.5	0.875	0.75	71.5	-16.2
394	G50B_087_037a	0.5	0.875	0.875	0.875	0.375	0.687	21.0	0.5	0.875	0.875	72.2	-11.9
395	G61B_100_050a	0.5	0.875	1.0	1.0	0.5	0.75	22.4	0.5	0.883	1.0	74.0	-11.4
396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	12.0	0.5	1.0	0.0	73.1	-30.2
397	Y58G_100_087a	0.5	1.0	0.125	1.0	0.875	0.562	12.5	0.489	1.0	0.125	73.5	-30.2
398	Y68G_100_075a	0.5	1.0	0.25	1.0	0.75	0.625	13.1	0.487	1.0	0.25	72.9	-31.5
399	Y81G_100_062a	0.5	1.0	0.375	1.0	0.625	0.687	13.9	0.489	1.0	0.375	72.8	-32.6
400	G00B_100_050a	0.5	1.0	0.5	1.0	0.5	0.75	15.0	0.5	1.0	0.5	74.0	-34.6
401	G11B_100_050a	0.5	1.0	0.625	1.0	0.5	0.75	16.4	0.5	1.0	0.625	74.6	-31.2
402	G25B_100_050a	0.5	1.0	0.75	1.0	0.5	0.75	18.0	0.5	1.0	0.75	75.5	-25.4
403	G38B_100_050a	0.5	1.0	0.875	1.								

Mean color difference of this page: $\Delta E^* = 6.0$

TUB-test chart SE14; 1080 colours, offset standard paper
colors and differences, ΔE^* , 3D=0, de=0, *cmYk*

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

TUB registration: 20130201-SE14/SE14L0NP.PDF / .PS
application for measurement of offset print output, separationcnmy6 (CMYK)
TUB material: code=rha4ta

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF> / .PS
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output, separationcnymn6 (CMYK)

TUB material: code=rh4ta

		HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb**Md	LabCh**Md			
405	R00Y_062_062a	0.625	0.0	0.0	0.625	0.625	0.312	390	0.625	0.0	0.0	47.5	65.5	38.4	76.0	30.4
406	R31Y_062_062a	0.625	0.0	0.125	0.625	0.625	0.312	379	0.625	0.0	0.183	47.7	66.7	28.9	72.7	23.4
407	R11Y_062_062a	0.625	0.0	0.25	0.625	0.625	0.312	367	0.625	0.0	0.383	47.8	68.3	17.8	70.6	14.6
408	B69R_062_062a	0.625	0.0	0.375	0.625	0.625	0.312	353	0.625	0.0	0.616	48.0	70.7	5.3	70.9	4.3
409	B59R_062_062a	0.625	0.0	0.5	0.625	0.625	0.312	341	0.625	0.0	0.816	48.1	72.7	-2.9	72.7	357.6
410	B50R_062_062a	0.625	0.0	0.625	0.625	0.625	0.312	330	0.625	0.0	1.0	48.2	74.2	-8.7	74.7	353.2
411	B42R_075_075a	0.625	0.0	0.75	0.75	0.75	0.375	321	0.637	0.0	1.0	45.4	69.9	-12.7	71.0	349.6
412	B36R_087_087a	0.625	0.0	0.875	0.875	0.875	0.437	314	0.641	0.0	0.875	39.8	57.6	-14.5	59.4	345.8
413	B31R_100_100a	0.625	0.0	1.0	1.0	1.0	0.5	308	0.633	0.0	1.0	40.4	60.2	-21.7	64.0	340.1
414	R18Y_062_062a	0.625	0.125	0.0	0.625	0.625	0.312	41	0.625	0.114	0.0	40.6	32.3	28.9	43.4	41.9
415	R00Y_062_050a	0.625	0.125	0.125	0.625	0.5	0.375	390	0.625	0.125	0.125	42.7	32.7	19.2	38.0	30.4
416	R26Y_062_050a	0.625	0.125	0.25	0.625	0.5	0.375	376	0.625	0.125	0.241	42.9	33.4	13.1	35.9	21.4
417	R00Y_062_050a	0.625	0.125	0.375	0.625	0.5	0.375	360	0.625	0.125	0.375	42.9	34.8	5.6	35.3	9.2
418	B61R_062_050a	0.625	0.125	0.5	0.625	0.5	0.375	344	0.625	0.125	0.508	43.0	36.1	-0.6	36.1	358.9
419	B50R_062_050a	0.625	0.125	0.625	0.625	0.5	0.375	330	0.625	0.125	0.625	43.1	37.1	-4.3	37.3	353.2
420	B40R_075_062a	0.625	0.125	0.75	0.75	0.625	0.437	319	0.635	0.125	0.75	44.6	43.0	-8.6	43.9	348.6
421	B34R_087_075a	0.625	0.125	0.875	0.875	0.75	0.5	311	0.637	0.125	0.875	45.6	47.3	-14.4	49.5	342.9
422	B29R_100_087a	0.625	0.125	1.0	1.0	0.875	0.562	305	0.635	0.125	1.0	46.3	50.6	-20.9	54.8	337.5
423	R38Y_062_062a	0.625	0.25	0.0	0.625	0.625	0.312	53	0.625	0.239	0.0	46.1	21.1	35.5	41.3	59.3
424	R23Y_062_050a	0.625	0.25	0.125	0.625	0.5	0.375	44	0.625	0.241	0.125	46.9	23.6	24.3	33.9	45.8
425	R00Y_062_037a	0.625	0.25	0.25	0.625	0.375	0.437	390	0.625	0.25	0.25	48.8	24.5	14.4	28.5	30.4
426	R18Y_062_037a	0.625	0.25	0.375	0.625	0.375	0.437	371	0.625	0.25	0.368	48.9	25.4	8.1	26.6	17.7
427	B65R_062_037a	0.625	0.25	0.5	0.625	0.375	0.437	349	0.625	0.25	0.506	49.0	26.7	0.8	26.8	1.8
428	B50R_062_037a	0.625	0.25	0.625	0.625	0.375	0.437	330	0.625	0.25	0.625	49.1	27.8	-3.2	28.0	353.2
429	B38R_075_050a	0.625	0.25	0.75	0.75	0.5	0.5	316	0.633	0.25	0.75	50.5	33.6	-7.6	34.5	347.2
430	B30R_087_062a	0.625	0.25	0.875	0.875	0.625	0.562	307	0.635	0.25	0.875	51.4	37.1	-14.0	39.7	339.2
431	B25R_100_075a	0.625	0.25	1.0	1.0	0.75	0.625	300	0.625	0.25	1.0	51.9	40.4	-20.3	45.3	333.2
432	R61Y_062_062a	0.625	0.375	0.0	0.625	0.625	0.312	67	0.625	0.385	0.0	53.5	8.1	44.4	45.1	79.5
433	R50Y_062_050a	0.625	0.375	0.125	0.625	0.5	0.375	60	0.625	0.375	0.125	53.0	12.0	31.5	33.7	69.1
434	R31Y_062_037a	0.625	0.375	0.25	0.625	0.375	0.437	49	0.625	0.368	0.25	53.4	14.9	20.0	25.0	53.2
435	R00Y_062_025a	0.625	0.375	0.375	0.625	0.25	0.5	390	0.625	0.375	0.375	54.9	16.3	9.6	19.0	30.4
436	R00Y_062_025a	0.625	0.375	0.5	0.625	0.25	0.5	360	0.625	0.375	0.5	55.0	17.4	2.8	17.6	9.2
437	B50R_062_025a	0.625	0.375	0.625	0.625	0.25	0.5	330	0.625	0.375	0.625	55.1	18.5	-2.1	18.6	353.2
438	B34R_075_037a	0.625	0.375	0.75	0.75	0.375	0.562	311	0.631	0.375	0.75	56.4	23.6	-7.2	24.7	342.9
439	B25R_087_050a	0.625	0.375	0.875	0.875	0.5	0.625	300	0.625	0.375	0.875	57.0	26.9	-13.5	30.2	333.2
440	B19R_100_062a	0.625	0.375	1.0	1.0	0.625	0.687	293	0.614	0.375	1.0	57.4	30.1	-19.9	36.1	326.4
441	R81Y_062_062a	0.625	0.5	0.0	0.625	0.625	0.312	79	0.625	0.51 0.0	0.0	58.9	-0.1	50.8	50.8	90.1
442	R76Y_062_050a	0.625	0.5	0.125	0.625	0.5	0.375	76	0.625	0.508 0.125	0.0	59.6	1.2	39.4	39.4	88.1
443	R68Y_062_037a	0.625	0.5	0.25	0.625	0.375	0.437	71	0.625	0.506 0.25	0.0	60.1	3.1	28.0	28.1	83.6
444	R50Y_062_025a	0.625	0.5	0.375	0.625	0.25	0.5	60	0.625	0.5 0.375	0.0	60.1	6.0	15.7	16.8	69.1
445	R00Y_062_012a	0.625	0.5	0.625	0.625	0.125	0.562	330	0.625	0.5 0.5	0.0	61.0	8.1	4.8	9.5	30.4
446	B50R_062_012a	0.625	0.5	0.625	0.625	0.125	0.562	330	0.625	0.5 0.625	0.1	61.1	9.2	-1.0	9.3	353.2
447	B25R_075_025a	0.625	0.5	0.75	0.75	0.25	0.625	300	0.625	0.5 0.75	0.0	62.0	13.4	-6.7	15.1	333.2
448	B15R_087_037a	0.625	0.5	0.875	0.875	0.375	0.687	289	0.618	0.5 0.875	0.625	62.5	15.9	-13.6	20.9	319.3
449	B11R_100_050a	0.625	0.5	1.0	1.0	0.5	0.75	284	0.616	0.5 1.0	0.0	63.0	17.7	-20.4	27.0	310.9
450	Y00G_062_062a	0.625	0.625	0.0	0.625	0.625	0.312	90	0.625	0.625 0.0	0.0	62.8	-5.9	55.6	56.0	96.0
451	Y00G_062_050a	0.625	0.625	0.125	0.625	0.5	0.375	90	0.625	0.625 0.125	0.63	63.7	-4.7	44.5	44.8	96.0
452	Y00G_062_037a	0.625	0.625	0.25	0.625	0.375	0.437	90	0.625	0.625 0.25	0.645	-3.5	33.4	33.6	96.0	
453	Y00G_062_025a	0.625	0.625	0.375	0.625	0.25	0.5	90	0.625	0.625 0.375	0.654	-2.3	22.2	22.4	96.0	
454	Y00G_062_012a	0.625	0.625	0.5	0.625	0.125	0.562	90	0.625	0.625 0.5	0.663	-1.1	11.1	11.2	96.0	
455	NW_062a	0.625	0.625	0.625	0.625	0.0	0.625	360	0.625	0.625 0.625	0.671	0.0	0.0	0.0	0.0	
456	B00R_075_012a	0.625	0.625	0.75	0.75	0.125	0.687	270	0.625	0.625 0.75	0.679	2.8	-5.9	6.6	295.6	
457	B00R_087_025a	0.625	0.625	0.875	0.875	0.25	0.75	270	0.625	0.625 0.875	0.688	5.7	-11.9	13.2	295.6	
458	B00R_100_037a	0.625	0.625	1.0	1.0	0.37										

http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF /PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 26/33

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF> /PS
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md		
486	R00Y_075_075a	0.75 0.0 0.0	0.75 0.75 0.375	390	0.75 0.0 0.0	40.2 49.1 28.8	57.0 30.4	0.75 0.0 0.0	41.5 52.5 31.0	61.0 30.6 4.2	389	1.0 0.0 0.0	47.5 65.5 38.4	76.0 30.4
487	R35Y_075_075a	0.75 0.0 0.125	0.75 0.75 0.375	381	0.75 0.0 0.112	40.4 49.8 22.9	54.9 24.7	0.75 0.0 0.125	41.9 53.1 24.2	58.4 24.4 3.8	382	1.0 0.0 0.15	47.6 66.4 30.6	73.2 24.7
488	R18Y_075_075a	0.75 0.0 0.25	0.75 0.75 0.375	371	0.75 0.0 0.237	40.5 50.8 16.2	53.3 17.7	0.75 0.0 0.25	42.0 54.5 16.4	56.9 16.7 4.0	371	1.0 0.0 0.316	47.8 67.7 21.6	71.1 17.7
489	R00Y_075_075a	0.75 0.0 0.375	0.75 0.75 0.375	360	0.75 0.0 0.375	40.5 52.2 8.5	52.9 9.2	0.75 0.0 0.375	42.1 56.3 7.4	56.8 7.4 4.5	360	1.0 0.0 0.5	47.8 69.7 11.3	70.6 9.2
490	B65R_075_075a	0.75 0.0 0.5	0.75 0.75 0.375	349	0.75 0.0 0.517	40.7 53.5 1.7	53.6 1.8	0.75 0.0 0.5	41.9 58.1 0.1	58.1 0.1 4.9	348	1.0 0.0 0.683	48.1 71.4 2.3	71.4 1.8
491	B57R_075_075a	0.75 0.0 0.625	0.75 0.75 0.375	339	0.75 0.0 0.632	40.7 54.7 -3.0	54.8 356.7	0.75 0.0 0.625	42.1 59.5 -5.2	59.8 354.9 5.4	337	1.0 0.0 0.85	48.1 72.9 -4.0	73.0 356.7
492	B50R_075_075a	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	40.8 55.6 -6.5	56.0 353.2	0.75 0.0 0.75	42.2 60.7 -9.4	61.4 351.1 5.9	330	1.0 0.0 1.0	48.2 74.2 -8.7	74.7 353.2
493	B43R_087_087a	0.75 0.0 0.875	0.875 0.875 0.437	322	0.758 0.0 0.875	42.3 61.6 -10.7	62.5 350.1	0.75 0.0 0.875	43.3 65.5 -12.5	66.7 349.1 4.4	322	0.866 0.0 1.0	45.7 70.4 -12.2	71.5 350.1
494	B38R_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	43.6 67.3 -15.2	69.0 347.2	0.75 0.0 1.0	43.3 66.7 -15.7	68.5 346.7 0.7	317	0.766 0.0 1.0	43.6 67.3 -15.2	69.0 347.2
495	R15Y_075_075a	0.75 0.125 0.0	0.75 0.75 0.375	399	0.75 0.112 0.0	44.0 49.9 33.4	52.8 39.2	0.75 0.125 0.0	46.1 41.9 35.9	55.2 40.5 3.3	37	1.0 0.15 0.0	52.5 54.5 44.5	70.4 39.2
496	R00Y_075_062a	0.75 0.125 0.125	0.75 0.625 0.437	390	0.75 0.125 0.125	46.3 40.9 24.0	47.5 30.4	0.75 0.125 0.125	47.0 41.6 27.3	49.8 33.3 3.4	389	1.0 0.0 0.0	47.5 65.5 38.4	76.0 30.4
497	R31Y_075_062a	0.75 0.125 0.25	0.75 0.625 0.437	379	0.75 0.125 0.239	46.5 41.6 18.0	45.4 23.4	0.75 0.125 0.25	47.3 42.7 18.9	46.7 23.9 1.5	380	1.0 0.0 0.183	47.7 66.7 28.9	72.7 23.4
498	R11Y_075_062a	0.75 0.125 0.375	0.75 0.625 0.437	367	0.75 0.125 0.364	46.6 42.7 11.1	44.1 14.6	0.75 0.125 0.375	47.8 43.8 10.0	45.0 12.8 2.0	367	1.0 0.0 0.383	47.8 68.3 17.8	70.6 14.6
499	B69R_075_062a	0.75 0.125 0.5	0.75 0.625 0.437	353	0.75 0.125 0.51	46.7 44.1 3.3	44.3 4.3	0.75 0.125 0.5	48.0 45.7 1.4	45.7 1.7 2.8	352	1.0 0.0 0.616	48.0 70.7 5.3	70.9 4.3
500	B59R_075_062a	0.75 0.125 0.625	0.75 0.625 0.437	341	0.75 0.125 0.635	46.7 45.4 -1.8	45.4 357.6	0.75 0.125 0.625	48.3 47.0 -4.5	47.2 354.4 3.4	339	1.0 0.0 0.816	48.1 72.7 -2.9	72.7 357.6
501	B50R_075_062a	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	46.8 46.3 -5.4	46.7 353.2	0.75 0.125 0.75	48.5 48.3 -9.3	49.2 349.0 4.6	330	1.0 0.0 1.0	48.2 74.2 -8.7	74.7 353.2
502	B42R_087_075a	0.75 0.125 0.875	0.875 0.75 0.5	321	0.762 0.125 0.875	48.4 52.4 -9.5	53.3 349.6	0.75 0.125 0.875	49.1 54.0 -12.6	55.4 346.8 3.5	322	0.85 0.0 1.0	45.4 69.9 -12.7	71.0 349.6
503	B36R_100_087a	0.75 0.125 1.0	1.0 0.875 0.562	314	0.766 0.125 1.0	49.6 57.6 -14.5	59.4 345.8	0.75 0.125 1.0	48.5 56.4 -15.9	58.7 344.1 2.1	315	0.733 0.0 1.0	42.9 65.8 -16.6	67.9 345.8
504	R31Y_075_075a	0.75 0.25 0.0	0.75 0.75 0.375	49	0.75 0.237 0.0	49.4 29.9 40.1	50.0 53.2	0.75 0.25 0.0	53.1 29.5 42.8	52.0 55.4 4.6	48	1.0 0.316 0.0	59.6 39.8 53.5	66.7 53.2
505	R18Y_075_062a	0.75 0.25 0.125	0.75 0.625 0.437	49	0.75 0.239 0.125	50.3 32.3 28.9	43.4 41.9	0.75 0.25 0.125	53.4 30.6 32.0	44.3 46.2 4.6	39	1.0 0.183 0.0	53.9 51.7 46.3	69.4 41.9
506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	52.4 32.7 19.2	38.0 30.4	0.75 0.25 0.25	54.5 30.7 22.3	37.9 36.0 4.2	389	1.0 0.0 0.0	47.5 65.5 38.4	76.0 30.4
507	R26Y_075_050a	0.75 0.25 0.375	0.75 0.5 0.5	376	0.75 0.25 0.366	52.6 33.4 13.1	35.9 21.4	0.75 0.25 0.375	55.2 31.6 13.3	34.3 22.8 3.2	377	1.0 0.0 0.233	47.8 66.9 26.3	71.9 21.4
508	R00Y_075_050a	0.75 0.25 0.5	0.75 0.5 0.5	360	0.75 0.25 0.5	52.6 34.8 5.6	35.3 9.2	0.75 0.25 0.5	55.5 33.6 4.2	33.8 7.1 3.4	360	1.0 0.0 0.5	47.8 69.7 11.3	70.6 9.2
509	B61R_075_050a	0.75 0.25 0.625	0.75 0.5 0.5	344	0.75 0.25 0.633	52.8 36.1 -0.6	36.1 358.9	0.75 0.25 0.625	56.2 34.9 -2.9	35.0 355.1 4.2	342	1.0 0.0 0.766	48.1 72.2 -1.3	72.2 358.9
510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	52.8 37.1 -4.3	37.3 353.2	0.75 0.25 0.75	56.6 36.1 -8.5	37.1 346.7 5.6	330	1.0 0.0 1.0	48.2 74.2 -8.7	74.7 353.2
511	B40R_087_062a	0.75 0.25 0.875	0.875 0.625 0.562	319	0.76 0.25 0.875	54.3 43.0 -8.6	43.9 348.6	0.75 0.25 0.875	56.6 42.0 -12.2	43.7 343.7 4.3	320	0.816 0.0 1.0	44.7 68.8 -13.7	70.2 348.6
512	B34R_100_075a	0.75 0.25 1.0	1.0 0.75 0.625	311	0.762 0.25 1.0	55.3 47.3 -14.4	49.5 342.9	0.75 0.25 1.0	55.2 44.5 -15.7	47.2 340.5 3.0	311	0.683 0.0 1.0	41.6 63.1 -19.3	66.0 342.9
513	R50Y_075_075a	0.75 0.375 0.0	0.75 0.75 0.375	60	0.75 0.375 0.0	55.7 18.0 47.2	50.5 69.1	0.75 0.375 0.0	60.6 15.9 50.6	53.0 72.5 6.2	59	1.0 0.5 0.0	68.1 24.0 63.0	67.4 69.1
514	R38Y_075_062a	0.75 0.375 0.125	0.75 0.625 0.437	53	0.75 0.364 0.125	55.9 21.1 35.5	41.3 59.3	0.75 0.375 0.125	60.0 18.4 37.4	41.7 63.7 5.2	52	1.0 0.383 0.0	62.7 33.7 56.9	66.2 59.3
515	R23Y_075_050a	0.75 0.375 0.25	0.75 0.5 0.5	44	0.75 0.366 0.25	56.7 23.6 24.3	33.9 45.8	0.75 0.375 0.25	61.1 19.3 26.1	32.5 53.4 6.4	42	1.0 0.233 0.0	55.9 47.3 48.7	67.9 45.8
516	R00Y_075_037a	0.75 0.375 0.375	0.75 0.375 0.562	390	0.75 0.375 0.375	58.5 24.5 14.4	28.5 30.4	0.75 0.375 0.375	62.2 20.1 16.0	25.7 38.6 5.9	389	1.0 0.0 0.0	47.5 65.5 38.4	76.0 30.4
517	R18Y_075_037a	0.75 0.375 0.5	0.75 0.375 0.562	371	0.75 0.375 0.493	58.7 25.4 8.1	26.6 17.7	0.75 0.375 0.5	62.8 21.5 7.4	22.8 19.0 5.6	371	1.0 0.0 0.316	47.8 67.7 21.6	71.1 17.7
518	B65R_075_037a	0.75 0.375 0.625	0.75 0.375 0.562	349	0.75 0.375 0.631	58.8 26.7 0.8	26.8 1.8	0.75 0.375 0.625	63.3 23.1 -0.9	23.2 357.5 6.1	348	1.0 0.0 0.683	48.1 71.4 2.3	71.4 1.8
519	B50R_075_037a	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	58.8 27.8 -3.2	28.0 353.2	0.75 0.375 0.75	64.1 24.5 -6.9	25.5 344.1 7.1	330	1.0 0.0 1.0	48.2 74.2 -8.7	74.7 353.2
520	B38R_087_050a	0.75 0.375 0.875	0.875 0.5 0.625	316	0.758 0.375 0.875	60.3 33.6 -7.6	34.5 347.2	0.75 0.375 0.875	63.5 30.1 -11.0	32.1 339.8 5.8	317	0.766 0.0 1.0	43.6 67.3 -15.2	69.0 347.2
521	B30R_100_062a	0.75 0.375 1.0	1.0 0.625 0.687	307	0.76 0.375 1.0	61.1 37.1 -14.0	39.7 339.2	0.75 0.375 1.0	61.4 34.3 -14.7	37.3 336.8 2.8	307	0.616 0.0 1.0	40.0 59.4 -22.5	63.5 339.2
522	R68Y_075_075a	0.75												

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n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsi.d	rgb**Md	LabCh**Md
567	R00Y_087_087a	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.0	43.9 57.3 33.6	0.875 0.0 0.0	45.0 60.9 34.2	69.9 29.3 3.7	389	1.0 0.0 0.0	47.5 65.5 38.4 76.0 30.4
568	R36Y_087_087a	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.116	44.0 58.0 27.5	0.875 0.0 0.125	45.2 61.6 27.5	67.5 24.0 3.7	382	1.0 0.0 0.133	47.6 66.3 31.5 73.4 25.4
569	R23Y_087_087a	0.875 0.0 0.25	0.875 0.875 0.437	374	0.875 0.0 0.233	44.1 58.8 21.4	0.875 0.0 0.25	45.4 62.5 20.9	65.9 18.4 3.9	375	1.0 0.0 0.266	47.8 67.2 24.5 71.5 20.0
570	R08Y_087_087a	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.364	44.2 60.2 14.0	0.875 0.0 0.375	45.4 63.8 13.1	65.2 11.6 3.9	365	1.0 0.0 0.416	47.8 68.8 16.0 70.6 13.1
571	B70R_087_087a	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.51	44.3 61.6 6.2	0.875 0.0 0.5	45.4 65.6 5.6	65.8 4.8 4.2	354	1.0 0.0 0.583	48.0 70.4 7.1 70.8 5.7
572	B63R_087_087a	0.875 0.0 0.625	0.875 0.875 0.437	346	0.875 0.0 0.641	44.4 62.9 0.0	0.875 0.0 0.625	45.5 66.8 -0.5	66.8 359.4 4.1	344	1.0 0.0 0.733	48.1 71.9 0.0 71.9 359.9
573	B56R_087_087a	0.875 0.0 0.75	0.875 0.875 0.437	338	0.875 0.0 0.758	44.4 63.9 -4.0	0.875 0.0 0.75	45.8 68.1 -5.3	68.3 355.5 4.5	337	1.0 0.0 0.866	48.2 73.1 -4.6 73.2 356.3
574	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	44.5 64.9 -7.6	0.875 0.0 0.875	45.7 69.3 -9.3	69.9 352.2 4.8	330	1.0 0.0 1.0	48.2 74.2 -8.7 74.7 353.2
575	B44R_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	46.0 70.9 -11.8	0.875 0.0 1.0	45.9 70.7 -12.0	71.7 350.3 0.3	323	0.883 0.0 1.0	46.0 70.9 -11.8 71.9 350.5
576	R13Y_087_087a	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.116 0.0	47.7 48.9 38.1	0.875 0.125 0.0	49.5 50.6 39.6	64.3 38.0 2.9	37	1.0 0.133 0.0	51.8 55.9 43.6 70.9 37.9
577	R00Y_087_075a	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.125	50.0 49.1 28.8	0.875 0.125 0.125	50.2 50.6 31.4	59.6 31.8 2.9	389	1.0 0.0 0.0	47.5 65.5 38.4 76.0 30.4
578	R35Y_087_075a	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.237	50.1 49.8 22.9	0.875 0.125 0.25	50.6 51.2 23.7	56.5 24.8 1.6	382	1.0 0.0 0.15	47.6 66.4 30.6 73.2 24.7
579	R18Y_087_075a	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.362	50.2 50.8 16.2	0.875 0.125 0.375	50.9 52.4 15.4	54.6 16.4 1.9	371	1.0 0.0 0.316	47.8 67.7 21.6 71.1 17.7
580	R00Y_087_075a	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.5	50.2 52.2 8.5	0.875 0.125 0.5	51.0 54.0 7.1	54.5 7.5 2.4	360	1.0 0.0 0.5	47.8 69.7 11.3 70.6 9.2
581	B65R_087_075a	0.875 0.125 0.625	0.875 0.75 0.5	349	0.875 0.125 0.637	50.4 53.5 1.7	0.875 0.125 0.625	51.5 55.2 0.1	55.2 0.1 2.5	348	1.0 0.0 0.683	48.1 71.4 2.3 71.4 1.8
582	B57R_087_075a	0.875 0.125 0.75	0.875 0.75 0.5	339	0.875 0.125 0.762	50.5 54.7 -3.0	0.875 0.125 0.75	51.7 56.5 -5.4	56.8 354.5 3.1	337	1.0 0.0 0.85	48.1 72.9 -4.0 73.0 356.7
583	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	50.5 55.6 -6.5	0.875 0.125 0.875	51.8 57.6 -9.6	58.4 350.4 3.8	330	1.0 0.0 1.0	48.2 74.2 -8.7 74.7 353.2
584	B43R_100_087a	0.875 0.125 1.0	1.0 0.875 0.562	322	0.883 0.125 1.0	52.0 61.6 -10.7	0.875 0.125 1.0	51.3 60.0 -12.8	61.4 347.9 2.7	322	0.866 0.0 1.0	45.7 70.4 -12.2 71.5 350.1
585	R26Y_087_087a	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.233 0.0	52.5 38.8 44.2	0.875 0.25 0.0	55.2 38.6 46.0	60.1 50.0 3.3	44	1.0 0.266 0.0	57.3 44.3 50.5 67.2 48.2
586	R15Y_087_075a	0.875 0.25 0.125	0.875 0.75 0.5	39	0.875 0.237 0.125	53.7 40.9 33.4	0.875 0.25 0.125	55.8 39.0 36.0	53.1 42.7 3.8	37	1.0 0.15 0.0	52.5 54.5 44.5 70.4 39.2
587	R00Y_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	56.1 40.9 24.0	0.875 0.25 0.25	56.7 39.0 27.2	47.6 34.9 3.7	389	1.0 0.0 0.0	47.5 65.5 38.4 76.0 30.4
588	R31Y_087_062a	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.364	56.2 41.6 18.0	0.875 0.25 0.375	57.3 39.8 18.7	44.0 25.2 2.2	380	1.0 0.0 0.183	47.7 66.7 28.9 72.7 23.4
589	R11Y_087_062a	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.489	56.3 42.7 11.1	0.875 0.25 0.5	57.6 41.2 10.0	42.4 13.6 2.2	367	1.0 0.0 0.383	47.8 68.3 17.8 70.6 14.6
590	B69R_087_062a	0.875 0.25 0.625	0.875 0.625 0.562	353	0.875 0.25 0.635	56.4 44.1 3.3	0.875 0.25 0.625	58.1 42.8 1.9	42.8 2.5 2.6	352	1.0 0.0 0.616	48.0 70.7 5.3 70.9 4.3
591	B59R_087_062a	0.875 0.25 0.75	0.875 0.625 0.562	341	0.875 0.25 0.76	56.5 45.4 -1.8	0.875 0.25 0.75	58.4 44.2 -4.1	44.4 354.6 3.2	339	1.0 0.0 0.816	48.1 72.7 -2.9 72.7 357.6
592	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	56.5 46.3 -5.4	0.875 0.25 0.875	59.0 45.0 -9.0	45.9 348.6 4.4	330	1.0 0.0 1.0	48.2 74.2 -8.7 74.7 353.2
593	B42R_100_075a	0.875 0.25 1.0	1.0 0.75 0.625	321	0.887 0.25 1.0	58.1 52.4 -9.5	0.875 0.25 1.0	58.1 48.2 -11.9	49.7 346.0 4.8	322	0.85 0.0 1.0	45.4 69.9 -12.7 71.0 349.6
594	R41Y_087_087a	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.364 0.0	58.5 27.2 51.4	0.875 0.375 0.0	61.7 26.3 53.2	59.4 63.7 3.7	54	1.0 0.416 0.0	64.2 31.1 58.8 66.5 62.1
595	R31Y_087_075a	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.362 0.125	59.1 29.9 40.1	0.875 0.375 0.125	62.3 26.8 41.9	49.8 57.3 4.7	48	1.0 0.316 0.0	59.6 39.8 53.5 66.7 53.2
596	R18Y_087_062a	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.364 0.25	60.1 32.3 28.9	0.875 0.375 0.25	63.1 27.3 31.8	41.9 49.3 6.4	39	1.0 0.183 0.0	53.9 51.7 46.3 69.4 41.9
597	R00Y_087_050a	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	62.2 32.7 19.2	0.875 0.375 0.375	64.0 27.8 21.9	35.5 38.2 5.9	389	1.0 0.0 0.0	47.5 65.5 38.4 76.0 30.4
598	R26Y_087_050a	0.875 0.375 0.5	0.875 0.5 0.625	376	0.875 0.375 0.491	62.3 33.4 13.1	0.875 0.375 0.5	64.6 29.1 13.2	32.0 24.4 4.9	377	1.0 0.0 0.233	47.8 66.9 26.3 71.9 21.4
599	R00Y_087_050a	0.875 0.375 0.625	0.875 0.5 0.625	360	0.875 0.375 0.625	62.3 34.8 5.6	0.875 0.375 0.625	65.3 30.4 4.6	30.8 8.6 5.4	360	1.0 0.0 0.5	47.8 69.7 11.3 70.6 9.2
600	B61R_087_050a	0.875 0.375 0.75	0.875 0.5 0.625	344	0.875 0.375 0.758	62.5 36.1 -0.6	0.875 0.375 0.75	66.0 31.8 -2.3	31.9 355.8 5.7	342	1.0 0.0 0.766	48.1 72.2 -1.3 72.2 358.9
601	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	62.5 37.1 -4.3	0.875 0.375 0.875	66.2 33.2 -7.7	34.1 346.8 6.3	330	1.0 0.0 1.0	48.2 74.2 -8.7 74.7 353.2
602	B40R_100_062a	0.875 0.375 1.0	1.0 0.625 0.687	319	0.885 0.375 1.0	64.0 43.0 -8.6	0.875 0.375 1.0	64.0 37.8 -11.5	39.5 343.0 5.9	320	0.816 0.0 1.0	44.7 68.8 -13.7 70.2 348.6
603	R58Y_087_087a	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.51 0.0	65.9 14.3 60.3	0.875 0.5 0.0	69.3 12.9 61.9	63.3 78.1 4.0	65	1.0 0.583 0.0	72.6 16.4 68.9 70.8 76.5
604	R50Y_087_075a	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.5 0.125	65.4 18.0 47.2	0.875 0.5 0.125	69.4 14.5 48.4	50.6 73.2 5.4	59	1.0 0.5 0.0	68.1 24.0 63.0 67.4 69.1
605	R38Y_087_062a	0.875 0.5 0.25	0.875 0.625 0.562	53	0.875 0.489 0.25	65.6 21.1 35.5	0.875 0.5 0.25	69.7 16.2 37.0				

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE**Fd	hsi_Md	rgb*Md	LabCh*Md
648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	47.5 65.5 38.4	76.0 30.4	1.0 0.0 0.0	47.5 65.5 38.4	76.0 30.4	1.0 0.0 0.0	47.5 65.5 38.4
649	R38Y_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	47.6 66.2 32.3	73.7 26.0	1.0 0.0 0.125	47.6 66.2 31.9	73.5 25.7	1.0 0.0 0.116	47.6 66.2 32.3
650	R26Y_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	47.8 66.9 26.3	71.9 21.4	1.0 0.0 0.25	47.8 67.0 25.4	71.7 20.8	1.0 0.0 0.233	47.8 66.9 26.3
651	R13Y_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	47.8 68.1 18.7	70.7 15.4	1.0 0.0 0.375	47.8 68.2 18.3	70.6 15.0	1.0 0.0 0.366	47.8 68.1 18.7
652	R00Y_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	47.8 69.7 11.3	70.6 9.2	1.0 0.0 0.5	47.8 69.7 11.3	70.6 9.2	1.0 0.0 0.5	47.8 69.7 11.3
653	B68R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	48.0 70.8 4.5	71.0 3.7	1.0 0.0 0.625	48.0 70.7 4.9	70.9 4.0	1.0 0.0 0.633	48.0 70.8 4.5
654	B61R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	48.1 72.2 -1.3	72.2 358.9	1.0 0.0 0.75	48.1 72.1 -0.7	72.1 359.3	1.0 0.0 0.766	48.1 72.2 -1.3
655	B55R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	48.2 73.2 -5.1	73.4 355.9	1.0 0.0 0.875	48.2 73.1 -4.9	73.3 356.1	1.0 0.0 0.883	48.2 73.2 -5.1
656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	48.2 74.2 -8.7	74.7 353.2	1.0 0.0 1.0	48.2 74.2 -8.7	74.7 353.2	1.0 0.0 1.0	48.2 74.2 -8.7
657	R11Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	51.2 57.2 42.8	71.5 36.8	1.0 0.125 0.0	51.5 56.6 43.1	71.2 37.2	1.0 0.116 0.0	51.2 57.2 42.8
658	R00Y_100_087a	1.0 0.125 0.125	1.0 0.875 0.562	390	1.0 0.125 0.125	53.6 57.3 33.6	66.5 30.4	1.0 0.125 0.125	52.3 56.2 35.5	66.5 32.3	1.0 0.0 0.875	48.1 72.2 -1.3
659	R36Y_100_087a	1.0 0.125 0.25	1.0 0.875 0.562	382	1.0 0.125 0.241	53.7 58.0 27.5	64.2 25.4	1.0 0.125 0.25	52.5 56.7 27.8	63.2 26.1	1.0 0.0 0.875	48.1 72.2 -1.3
660	R23Y_100_087a	1.0 0.125 0.375	1.0 0.875 0.562	374	1.0 0.125 0.358	53.9 58.8 21.4	62.6 20.0	1.0 0.125 0.375	52.7 57.8 20.9	61.4 19.9	1.0 0.0 0.875	48.1 72.2 -1.3
661	R08Y_100_087a	1.0 0.125 0.5	1.0 0.875 0.562	365	1.0 0.125 0.489	53.9 60.2 14.0	61.8 13.1	1.0 0.125 0.5	52.8 59.2 13.1	60.6 12.5	1.0 0.0 0.875	48.1 72.2 -1.3
662	B70R_100_087a	1.0 0.125 0.625	1.0 0.875 0.562	355	1.0 0.125 0.635	54.0 61.6 6.2	61.9 5.7	1.0 0.125 0.625	53.4 59.9 5.3	60.2 5.1	1.0 0.0 0.875	48.1 72.2 -1.3
663	B63R_100_087a	1.0 0.125 0.75	1.0 0.875 0.562	346	1.0 0.125 0.766	54.1 62.9 0.0	62.9 359.9	1.0 0.125 0.75	53.5 61.3 -0.7	61.3 359.2	1.0 0.0 0.875	48.1 72.2 -1.3
664	B56R_100_087a	1.0 0.125 0.875	1.0 0.875 0.562	338	1.0 0.125 0.883	54.2 63.9 -4.0	64.0 356.3	1.0 0.125 0.875	53.7 62.3 -5.2	62.5 355.1	1.0 0.0 0.875	48.1 72.2 -1.3
665	B50R_100_087a	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	54.2 64.9 -7.6	65.3 353.2	1.0 0.125 1.0	53.9 63.2 -9.6	64.0 351.3	1.0 0.0 0.875	48.1 72.2 -1.3
666	R23Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	55.9 47.3 48.7	67.9 45.8	1.0 0.25 0.0	56.6 45.8 49.4	67.4 47.2	1.0 0.233 0.0	55.9 47.3 48.7
667	R13Y_100_100a	1.0 0.25 0.125	1.0 0.875 0.562	38	1.0 0.241 0.125	57.4 48.9 38.1	62.1 37.9	1.0 0.25 0.125	57.5 45.2 40.5	60.7 41.8	1.0 0.133 0.0	51.8 55.9 43.6
668	R00Y_100_075a	1.0 0.25 0.25	1.0 0.75 0.625	390	1.0 0.25 0.25	59.7 49.1 28.8	57.0 30.4	1.0 0.25 0.25	58.4 44.9 31.9	55.1 35.4	1.0 0.0 0.75	48.1 72.2 -1.3
669	R35Y_100_075a	1.0 0.25 0.375	1.0 0.75 0.625	381	1.0 0.25 0.362	59.8 49.8 22.9	54.9 24.7	1.0 0.25 0.375	58.6 45.9 24.3	52.0 27.9	1.0 0.0 0.75	48.1 72.2 -1.3
670	R18Y_100_075a	1.0 0.25 0.5	1.0 0.75 0.625	371	1.0 0.25 0.487	59.9 50.8 16.2	53.3 17.7	1.0 0.25 0.5	59.0 46.6 16.2	49.4 19.2	1.0 0.0 0.75	48.1 72.2 -1.3
671	R00Y_100_075a	1.0 0.25 0.625	1.0 0.75 0.625	360	1.0 0.25 0.625	59.9 52.2 8.5	52.9 9.2	1.0 0.25 0.625	59.8 47.3 8.0	48.0 9.6	1.0 0.0 0.75	48.1 72.2 -1.3
672	B65R_100_075a	1.0 0.25 0.75	1.0 0.75 0.625	349	1.0 0.25 0.762	60.1 53.5 1.7	53.6 1.8	1.0 0.25 0.75	60.0 49.0 0.9	49.0 1.1	1.0 0.0 0.75	48.1 72.2 -1.3
673	B57R_100_075a	1.0 0.25 0.875	1.0 0.75 0.625	339	1.0 0.25 0.887	60.2 54.7 -3.0	54.8 356.7	1.0 0.25 0.875	60.4 50.1 -4.4	50.3 354.8	1.0 0.0 0.75	48.1 72.2 -1.3
674	B50R_100_075a	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	60.2 55.6 -6.5	56.0 353.2	1.0 0.25 1.0	60.4 51.5 -9.1	52.4 349.8	1.0 0.0 0.75	48.1 72.2 -1.3
675	R36Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	62.0 35.2 56.0	66.2 57.8	1.0 0.375 0.0	62.3 34.4 56.4	66.1 58.6	1.0 0.366 0.0	62.0 35.2 56.0
676	R26Y_100_100a	1.0 0.375 0.125	1.0 0.875 0.562	46	1.0 0.358 0.125	62.2 38.8 44.2	58.8 48.7	1.0 0.375 0.125	62.3 36.0 45.1	57.7 51.4	1.0 0.266 0.0	57.3 44.3 50.5
677	R15Y_100_100a	1.0 0.375 0.25	1.0 0.75 0.625	39	1.0 0.362 0.25	63.5 40.9 33.4	52.8 39.2	1.0 0.375 0.25	63.1 35.8 35.8	50.7 44.9	1.0 0.15 0.0	52.5 44.5 44.5
678	R00Y_100_062a	1.0 0.375 0.375	1.0 0.625 0.687	390	1.0 0.375 0.375	65.8 40.9 24.0	47.5 30.4	1.0 0.375 0.375	64.4 35.1 27.2	44.5 37.7	1.0 0.0 0.625	48.0 70.7 5.3
679	R31Y_100_062a	1.0 0.375 0.5	1.0 0.625 0.687	379	1.0 0.375 0.489	65.9 41.6 18.0	45.4 23.4	1.0 0.375 0.5	64.8 36.2 18.7	40.8 27.3	1.0 0.0 0.625	48.0 70.7 5.3
680	R11Y_100_062a	1.0 0.375 0.625	1.0 0.625 0.687	367	1.0 0.375 0.614	66.0 42.7 11.1	44.1 14.6	1.0 0.375 0.625	65.6 36.9 10.7	38.4 16.2	1.0 0.0 0.625	48.0 70.7 5.3
681	B69R_100_062a	1.0 0.375 0.75	1.0 0.625 0.687	353	1.0 0.375 0.76	66.1 44.1 3.3	44.3 4.3	1.0 0.375 0.75	66.1 38.5 2.6	38.6 3.9	1.0 0.0 0.625	48.0 70.7 5.3
682	B59R_100_062a	1.0 0.375 0.875	1.0 0.625 0.687	341	1.0 0.375 0.885	66.2 45.4 -1.8	45.4 357.6	1.0 0.375 0.875	66.8 39.4 -3.4	39.5 355.0	1.0 0.0 0.625	48.0 70.7 5.3
683	B50R_100_062a	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	66.2 46.3 -5.4	46.7 353.2	1.0 0.375 1.0	66.9 40.7 -8.3	41.5 348.4	1.0 0.0 0.625	48.0 70.7 5.3
684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	68.1 24.0 63.0	67.4 69.1	1.0 0.5 0.0	68.1 24.0 63.0	67.4 69.1	1.0 0.5 0.0	68.1 24.0 63.0
685	R41Y_100_087a	1.0 0.5 0.125	1.0 0.875 0.562	55	1.0 0.489 0.125	68.3 27.2 51.4	58.2 62.1	1.0 0.5 0.125	68.3 25.0 51.0	56.8 63.8	1.0 0.416 0.0	64.2 31.1 58.8
686	R31Y_100_075a	1.0 0.5 0.25	1.0 0.75 0.625	49	1.0 0.487 0.25	68.8 29.9 40.1	50.0 53.2	1.0 0.5 0.25	68.8 25.6 40.2	47.7 57.5	1.0 0.316 0.0	59.6 39.8 53.5
687	R18Y_100_062a	1.0 0.5 0.375	1.0 0.625 0.687	41	1.0 0.489 0.375	69.8 32.3 28.9	43.4 41.9	1.0 0.5 0.375	69.5 26.4 30.5	40.4 49.1	1.0 0.183 0.0	53.9 51.7 46.3
688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	71.9 32.7 19.2	38.0 30.4	1.0 0.5 0.5	70.7 26.4 21.0	33.8 38.5	1.0 0.0 0.5	47.5 65.5 38.4
689	R26Y_100_050a	1.0 0.5 0.625	1.0 0.5 0.75	376	1.0 0.5 0.616	72.0 33.4 13.1	35.9 21.4	1.0 0.5 0.625	71.4 27.6 12.8	30.4 24.8	1.0 0.233 0.0	47.8 66.9 26.3
690	R00Y_100_050a	1.0 0.5 0.75	1.0 0.5 0.75	360	1.0 0.5 0.75	72.1 34.8 5.6	35.3 9.2	1.0 0.5 0.75	72.1 28.5 5.2	29.0 10.3	1.0 0.0 0.5	47.8 69.7 11.3
691	B61R_100_050a	1.0 0.5 0.875	1.0 0.5 0.75	344	1.0 0.5 0.883	72.2 36.1 -0.6	36.1 358.9	1.0 0.5 0.875	72.7 30.0 -1.5	30.0 357.0	1.0 0.0 0.766	48.1 72.2 -1.3
692	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	72.3 37.1 -4.3	37.3 353.2	1.0 0.5 1.0	73.1 31.3 -7.2	32.1 346.9	1.0 0.0 1.0	48.2 74.2 -8.7
693	R63Y_100_100a	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	75.3 11.6 72.0	72.9 80.8	1.0 0.625 0.0	74.9 12.1 71.5	72.5 80.3	1.0 0.633 0.0	75.3 11.6 72.0
694	R58Y_100_087a	1.0 0.625 0.125	1.0 0.875 0.562	65	1.0 0.635 0.125	75.6 14.3 60.3	62.0 76.5	1.0 0.625 0.125	75.7 12.4 58.9	60.1 78.1	1.0 0.583 0.0	72.6 11.6 68.9
695	R50Y_100_075a	1.0 0.625 0.25	1.0 0.75 0.625	60	1.0 0.625 0.25	75.1 18.0 47.2	50.5 69.1	1.0 0.625 0.25	76.2 13.4 46.8	48.7 73.9	1.0 0.5 0.0	68.1 24.0 63.0
696	R38Y_100_062a	1.0 0.625 0.375	1.0 0.625 0.687	53	1.0 0.614 0.375	75.3 21.1 35.5	41.3 59.3	1.0 0.625 0.375	76.4 15.0 35.5	38.5 67.0	1.0 0.383 0.0	62.7 33.7 56.9
697	R23Y_100_050a	1.0 0.625 0.5	1.0 0.5 0.75	44	1.0 0.616 0.5	76.1 23.6 24.3	33.9 45.8	1.0 0.625 0.5	77.2 16.1 25.2	29.9 57.3	1.0 0.233 0.0	55.9 47.3 48.7
698	R00Y_100_037a	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	78.0 24.5 14.4	28.5 30.4	1.0 0.625 0.625	78.3 16.8 16.1	23.3 43.6	1.0 0.0 0.625	48.1 72.2 -1.3
699	R18Y_100_037a	1.0 0.625 0.75	1.0 0.375 0.812	371	1.0 0.625 0.743	78.1 25.4 8.1	26.6 17.7	1.0 0.625 0.75	78.8 18.3 8.3	20.1 24.5	1.0 0.0 0.625	48.1 72.2 -1.3
700	B65R_100_037a	1.0 0.625 0.875	1.0 0.375 0.812	349	1.0 0.625 0.881	78.2 26.7 0.8	26.8 1.8	1.0 0.625 0.875	79.6 19.7 0.6	19.7 1.8	1.0 0.0 0.625	48.1 72.2 -1.3
701	B50R_100_037a	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	78.3 27.8 -3.2	28.0 353.2	1.0 0.625 1.0	80.2 21.1 -5.4	21.7 345.5	1.0 0.0 1.0	48.2 74.2 -8.7
702	R76Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	81.2 2.5 78.8	78.9 88.1	1.0 0.75 0.0	80.5 3.4 78.0	78.1 87.4	1.0 0.766 0.0	81.2 2.5 78.8
703	R73Y_100_087a	1.0 0.75 0.125	1.0 0.875 0.562	74	1.0 0.766 0.125	81.8 4.1 67.5	87.7 86.5	1.0 0.75 0.125	81.3 3.7 64.6	64.7 86.6	1.0 0.733 0.0	79.8 4.7 77.2
704	R68Y_100_075a	1.0 0.75 0.25	1.0 0.75 0.625	71	1.0 0.762 0.25	82.2 6.2 56.0	56.3 83.6	1.0 0.75				

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

		HIC*Fd			rgb_Fd			ict_Fd			hsi_Fd			rgb*Fd			LabCh*Fd			rgb**Fd			LabCh**Fd			DE**Fd			hsi_Md			rgb*Md			LabCh*Md		
729	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0	0.0	1.0	1.0	1.0	96.4	0.0	0.0	0.0	227.6	0.0	0.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0	
730	G50B_100_012d	0.875	1.0	1.0	1.0	0.125	0.937	210	0.875	1.0	1.0	91.5	-3.9	-5.6	6.9	234.6	0.875	1.0	1.0	93.1	-3.0	-4.1	5.1	233.7	2.4	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
731	G50B_100_025d	0.75	1.0	1.0	1.0	0.25	0.875	210	0.75	1.0	1.0	86.7	-7.9	-11.2	13.8	234.6	0.75	1.0	1.0	89.4	-6.0	-8.6	10.5	234.7	4.2	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
732	G50B_100_037d	0.625	1.0	1.0	1.0	0.375	0.812	210	0.625	1.0	1.0	81.9	-11.9	-16.9	20.7	234.6	0.625	1.0	1.0	84.9	-9.4	-13.6	16.5	235.3	5.1	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
733	G50B_100_050d	0.5	1.0	1.0	1.0	0.5	0.75	210	0.5	1.0	1.0	77.1	-15.9	-22.5	27.6	234.6	0.5	1.0	1.0	79.6	-13.6	-19.7	24.0	235.3	4.3	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
734	G50B_100_062d	0.375	1.0	1.0	1.0	0.625	0.687	210	0.375	1.0	1.0	72.3	-19.9	-28.2	34.5	234.6	0.375	1.0	1.0	74.4	-17.7	-25.7	31.2	235.4	3.9	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
735	G50B_100_075d	0.25	1.0	1.0	1.0	0.75	0.625	210	0.25	1.0	1.0	67.5	-23.9	-33.8	41.4	234.6	0.25	1.0	1.0	68.6	-22.7	-32.5	39.7	235.0	2.1	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
736	G50B_100_087d	0.125	1.0	1.0	1.0	0.875	0.562	210	0.125	1.0	1.0	62.6	-27.9	-39.4	48.4	234.6	0.125	1.0	1.0	63.2	-26.8	-38.6	47.0	235.1	1.5	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
737	G50B_100_100d	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6	0.0	1.0	1.0	57.0	-31.9	-45.6	55.7	235.0	1.0	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
738	ROOY_100_012d	1.0	0.875	0.875	1.0	0.125	0.937	390	1.0	0.875	0.875	90.2	8.1	4.8	9.5	30.4	1.0	0.875	0.875	91.1	4.0	5.7	7.0	55.0	4.3	389	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4			
739	NW_087d	0.875	0.875	0.875	0.875	0.0	0.875	360	0.875	0.875	0.875	86.6	0.0	0.0	0.0	0.0	0.875	0.875	0.875	91.0	-0.1	-0.5	0.5	258.7	4.4	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0			
740	G50B_087_012d	0.75	0.875	0.875	0.875	0.125	0.812	210	0.75	0.875	0.875	81.8	-3.9	-5.6	6.9	234.6	0.75	0.875	0.875	87.6	-3.1	-4.6	5.6	235.8	5.9	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
741	G50B_087_025d	0.625	0.875	0.875	0.875	0.25	0.75	210	0.625	0.875	0.875	77.0	-7.9	-11.2	13.8	234.6	0.625	0.875	0.875	83.4	-6.3	-9.3	11.2	235.9	6.9	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
742	G50B_087_037d	0.5	0.875	0.875	0.875	0.375	0.687	210	0.5	0.875	0.875	72.2	-11.9	-16.9	20.7	234.6	0.5	0.875	0.875	78.0	-10.5	-15.4	18.6	235.7	6.2	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
743	G50B_087_050d	0.375	0.875	0.875	0.875	0.5	0.625	210	0.375	0.875	0.875	67.3	-15.9	-22.5	27.6	234.6	0.375	0.875	0.875	72.9	-14.5	-21.1	25.7	235.4	5.9	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
744	G50B_087_062d	0.25	0.875	0.875	0.875	0.625	0.562	210	0.25	0.875	0.875	62.5	-19.9	-28.2	34.5	234.6	0.25	0.875	0.875	66.8	-19.6	-28.2	34.4	235.1	4.2	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
745	G50B_087_075d	0.125	0.875	0.875	0.875	0.75	0.5	210	0.125	0.875	0.875	57.7	-23.9	-33.8	41.4	234.6	0.125	0.875	0.875	61.2	-24.2	-34.8	42.4	235.1	3.6	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
746	G50B_087_087d	0.0	0.875	0.875	0.875	0.875	0.437	210	0.0	0.875	0.875	52.9	-27.9	-39.4	48.4	234.6	0.0	0.875	0.875	55.2	-29.9	-41.9	51.5	234.5	3.8	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
747	ROOY_100_025d	1.0	0.75	0.75	1.0	0.25	0.875	390	1.0	0.75	0.75	84.1	16.3	9.6	19.0	30.4	1.0	0.75	0.75	84.1	10.5	11.3	15.4	47.2	6.1	389	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4			
748	ROOY_087_012d	0.875	0.75	0.75	0.875	0.125	0.812	390	0.875	0.75	0.75	80.5	8.1	4.8	9.5	30.4	0.875	0.75	0.75	85.5	4.0	5.7	7.0	54.5	6.5	389	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4			
749	NW_075d	0.75	0.75	0.75	0.75	0.0	0.75	360	0.75	0.75	0.75	76.9	0.0	0.0	0.0	0.0	0.75	0.75	0.75	83.2	-0.1	-1.1	1.1	262.6	6.4	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0			
750	G50B_075_012d	0.625	0.75	0.75	0.75	0.125	0.687	210	0.625	0.75	0.75	72.1	-3.9	-5.6	6.9	234.6	0.625	0.75	0.75	79.4	-3.3	-5.4	6.4	238.1	7.4	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
751	G50B_075_025d	0.5	0.75	0.75	0.75	0.25	0.625	210	0.5	0.75	0.75	67.2	-7.9	-11.2	13.8	234.6	0.5	0.75	0.75	74.9	-6.8	-10.4	12.4	236.8	7.8	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
752	G50B_075_037d	0.375	0.75	0.75	0.75	0.375	0.562	210	0.375	0.75	0.75	62.4	-11.9	-16.9	20.7	234.6	0.375	0.75	0.75	69.4	-11.2	-16.6	20.0	236.1	7.0	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
753	G50B_075_050d	0.25	0.75	0.75	0.75	0.5	0.5	210	0.25	0.75	0.75	57.6	-15.9	-22.5	27.6	234.6	0.25	0.75	0.75	63.8	-15.8	-23.1	28.0	235.5	6.2	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
754	G50B_075_062d	0.125	0.75	0.75	0.75	0.625	0.437	210	0.125	0.75	0.75	52.8	-19.9	-28.2	34.5	234.6	0.125	0.75	0.75	57.4	-21.1	-30.3	36.9	235.1	5.2	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
755	G50B_075_075d	0.0	0.75	0.75	0.75	0.75	0.375	210	0.0	0.75	0.75	48.0	-23.9	-33.8	41.4	234.6	0.0	0.75	0.75	51.7	-26.4	-37.1	45.6	234.5	5.5	210	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6		
756	ROOY_100_037d	1.0	0.625	0.625	1.0	0.375	0.812	390	1.0	0.625	0.625	78.0	24.5	14.4	28.5	30.4	1.0	0.625	0.625	78.0	16.7	17.1	23.9	45.7	8.3	389	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4			
757	ROOY_087_025d	0.875	0.625	0.625	0.875	0.25	0.75	390	0.875	0.625	0.625	74.4	16.3	9.6	19.0	30.4	0.875	0.625	0.625	78.0	11.2	11.6	16.1	46.0	6.6	389	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4			
758	ROOY_075_012d	0.75	0.625	0.625	0.75	0.125	0.687	390	0.75	0.625	0.625	70.8	8.1	4.8	9.5	30.4	0.75	0.625	0.625	77.4	4.5	5.4	7.0	50.1	7.6	389	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4			
759	NW_062d	0.625	0.625	0.625	0.625	0.0	0.625	360	0.625	0.625	0.625	67.1	0.0	0.0	0.0	0.0	0.6																				

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF> / .PS
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output, separationcnymn6 (CMYK)
TUB material: code=rha4ta

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb*Md	LabCh*Md		
810	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	96.3 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	96.5 0.0 0.0	216.0 0.1 360	1.0 1.0 1.0	96.3 0.0 0.0	0.0 0.0 0.0
811	B00R_100_012d	0.875 0.875 1.0	1.0 0.125 0.937	270	0.875 0.875 1.0	87.4 2.8 -5.9	6.6 295.6	0.875 0.875 1.0	88.8 2.6 -5.8	6.3 294.7 1.4	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
812	B00R_100_025d	0.75 0.75 1.0	1.0 0.25 0.875	270	0.75 0.75 1.0	78.5 5.7 -11.9	13.2 295.6	0.75 0.75 1.0	79.2 6.9 -11.8	13.7 300.4 1.4	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
813	B00R_100_037d	0.625 0.625 1.0	1.0 0.375 0.812	270	0.625 0.625 1.0	69.6 8.6 -17.9	19.9 295.6	0.625 0.625 1.0	70.2 9.8 -17.6	20.2 299.1 1.4	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
814	B00R_100_050d	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	60.6 11.4 -23.9	26.5 295.6	0.5 0.5 1.0	58.7 15.0 -24.7	28.9 301.3 4.1	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
815	B00R_100_062d	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	51.7 14.3 -29.9	33.1 295.6	0.375 0.375 1.0	48.9 18.6 -30.6	35.8 301.2 5.1	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
816	B00R_100_075d	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	42.8 17.2 -35.8	39.8 295.6	0.25 0.25 1.0	39.8 21.0 -36.7	42.3 299.7 4.8	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
817	B00R_100_087d	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	33.9 20.1 -41.8	46.4 295.6	0.125 0.125 1.0	31.0 24.9 -42.3	49.1 300.5 5.6	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
818	B00R_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6	0.0 0.0 1.0	24.2 24.1 -47.5	53.2 296.8 1.3	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
819	Y00G_100_012d	1.0 1.0 0.875	1.0 0.125 0.937	90	1.0 1.0 0.875	95.5 -1.1 11.1	11.2 96.0	1.0 1.0 0.875	95.6 -1.7 7.9	8.1 102.7 3.2	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
820	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.6 0.0 0.0	0.0 0.0	0.875 0.875 0.875	91.1 0.0 -0.4	0.4 260.2 4.4	360	1.0 1.0 1.0	96.3 0.0 0.0	0.0 0.0 0.0
821	B00R_087_012d	0.75 0.75 0.875	0.875 0.125 0.812	270	0.75 0.75 0.875	77.7 2.8 -5.9	6.6 295.6	0.75 0.75 0.875	83.1 2.5 -6.4	6.9 292.0 5.4	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
822	B00R_087_025d	0.625 0.625 0.875	0.875 0.25 0.75	270	0.625 0.625 0.875	68.8 5.7 -11.9	13.2 295.6	0.625 0.625 0.875	72.9 7.4 -12.4	14.4 300.9 4.5	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
823	B00R_087_037d	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.5 0.875	59.8 8.6 -17.9	19.9 295.6	0.5 0.5 0.875	62.4 10.1 -19.4	21.9 297.4 3.3	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
824	B00R_087_050d	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	50.9 11.4 -23.9	26.5 295.6	0.375 0.375 0.875	51.5 15.2 -25.7	29.9 300.5 4.2	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
825	B00R_087_062d	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.0 14.3 -29.9	33.1 295.6	0.25 0.25 0.875	40.8 19.2 -32.7	37.9 300.5 5.7	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
826	B00R_087_075d	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	33.1 17.2 -35.8	39.8 295.6	0.125 0.125 0.875	31.1 23.3 -38.6	45.1 301.0 6.9	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
827	B00R_087_087d	0.0 0.0 0.875	0.875 0.875 0.437	270	0.0 0.0 0.875	24.1 20.1 -41.8	46.4 295.6	0.0 0.0 0.875	23.9 23.3 -44.1	49.9 297.8 3.9	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
828	Y00G_100_025d	1.0 1.0 0.75	1.0 0.25 0.875	90	1.0 1.0 0.75	94.6 -2.3 22.2	22.4 96.0	1.0 1.0 0.75	94.6 -3.2 16.6	16.9 101.1 5.6	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
829	Y00G_087_012d	0.875 0.875 0.75	0.875 0.125 0.812	90	0.875 0.875 0.75	85.7 -1.1 11.1	11.2 96.0	0.875 0.875 0.75	90.2 -1.9 8.1	8.3 103.7 5.4	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
830	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	76.9 0.0 0.0	0.0 0.0	0.75 0.75 0.75	83.3 -0.1 -1.1	1.1 261.9 6.5	360	1.0 1.0 1.0	96.3 0.0 0.0	0.0 0.0 0.0
831	B00R_075_012d	0.625 0.625 0.75	0.75 0.125 0.687	270	0.625 0.625 0.75	67.9 2.8 -5.9	6.6 295.6	0.625 0.625 0.75	74.5 3.0 -7.2	7.9 292.7 6.6	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
832	B00R_075_025d	0.5 0.5 0.75	0.75 0.25 0.625	270	0.5 0.5 0.75	59.0 5.7 -11.9	13.2 295.6	0.5 0.5 0.75	63.9 7.4 -13.5	15.5 298.8 5.4	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
833	B00R_075_037d	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	50.1 8.6 -17.9	19.9 295.6	0.375 0.375 0.75	52.6 11.8 -20.5	23.6 299.8 4.7	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
834	B00R_075_050d	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	41.2 11.4 -23.9	26.5 295.6	0.25 0.25 0.75	42.0 16.3 -26.7	31.3 301.3 5.6	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
835	B00R_075_062d	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	32.2 14.3 -29.9	33.1 295.6	0.125 0.125 0.75	31.5 20.3 -33.5	39.2 301.2 7.0	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
836	B00R_075_075d	0.0 0.0 0.75	0.75 0.75 0.375	270	0.0 0.0 0.75	23.3 17.2 -35.8	39.8 295.6	0.0 0.0 0.75	23.3 22.0 -39.3	45.0 299.2 5.8	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
837	Y00G_100_037d	1.0 1.0 0.625	1.0 0.375 0.812	90	1.0 1.0 0.625	93.7 -3.5 33.4	33.6 96.0	1.0 1.0 0.625	93.6 -4.5 26.0	26.4 100.0 7.4	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
838	Y00G_087_025d	0.875 0.875 0.625	0.875 0.25 0.75	90	0.875 0.875 0.625	84.9 -2.3 22.2	22.4 96.0	0.875 0.875 0.625	89.2 -3.5 17.3	17.7 101.5 6.6	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
839	Y00G_075_012d	0.75 0.75 0.625	0.75 0.125 0.687	90	0.75 0.75 0.625	76.0 -1.1 11.1	11.2 96.0	0.75 0.75 0.625	82.5 -2.1 7.8	8.1 105.3 7.3	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
840	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	67.1 0.0 0.0	0.0 0.0	0.625 0.625 0.625	76.1 -0.2 -1.5	1.5 261.4 9.0	360	1.0 1.0 1.0	96.3 0.0 0.0	0.0 0.0 0.0
841	B00R_062_012d	0.5 0.5 0.625	0.625 0.125 0.562	270	0.5 0.5 0.625	58.2 2.8 -5.9	6.6 295.6	0.5 0.5 0.625	66.0 3.4 -8.2	8.9 292.4 8.0	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
842	B00R_062_025d	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	49.3 5.7 -11.9	13.2 295.6	0.375 0.375 0.625	55.2 7.5 -14.6	16.4 297.1 6.7	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
843	B00R_062_037d	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	40.4 8.6 -17.9	19.9 295.6	0.25 0.25 0.625	43.7 12.1 -21.3	24.5 299.7 5.9	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
844	B00R_062_050d	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	31.4 11.4 -23.9	26.5 295.6	0.125 0.125 0.625	32.1 17.0 -28.5	33.2 300.8 7.2	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
845	B00R_062_062d	0.0 0.0 0.625	0.625 0.625 0.312	270	0.0 0.0 0.625	22.5 14.3 -29.9	33.1 295.6	0.0 0.0 0.625	22.7 20.0 -34.8	40.2 299.9 7.5	270	0.0 0.0 1.0	24.9 22.9 -47.8	53.0 295.6
846	Y00G_100_050d	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	92.8 -4.7 44.5	44.8 96.0	1.0 1.0 0.5	92.4 -5.8 36.9	37.3 99.0 7.7	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
847	Y00G_087_037d	0.875 0.875 0.5	0.875 0.375 0.687	90	0.875 0.875 0.5	84.0 -3.5 33.4	33.6 96.0	0.875 0.875 0.5	88.2 -5.0 27.6	28.0 100.3 7.2	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
848	Y00G_075_025d	0.75 0.75 0.5	0.75 0.25 0.625	90	0.75 0.75 0.5	75.1 -2.3 22.2	22.4 96.0	0.75 0.75 0.5	81.4 -3.8 17.5	17.9 102.2 7.9	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
849	Y00G_062_012d	0.625 0.625 0.5	0.625 0.125 0.562	90	0.625 0.625 0.5	66.3 -1.1 11.1	11.2 96.0	0.625 0.625 0.5	75.1 -2.3 7.8	8.2 106.4 9.4	89	1.0 1.0 0.0	89.4 -9.5 89.0	89.6 96.0
850	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	57.4 0.0 0.0	0.0 0.0	0.5 0.5 0.5	67.9 -0.2 -1.8	1.9 261.9 10.6	360	1.0 1.0 1.0	96.3 0.0 0.0	0.0 0.

http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF /.PS; transfer output
N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 31/33

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF> / .PS
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsi_d	rgb**Md	LabCh**Md
891	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	96.3 0.0 0.0	1.0 1.0 1.0	96.4 0.0 0.0	207.2 0.0	360	1.0 1.0 1.0	96.3 0.0 0.0
892	B50R_100_012d	1.0 0.875 1.0	1.0 0.125 0.937	330	1.0 0.875 1.0	90.3 9.2 -1.0	1.0 0.875 1.0	91.9 5.7 -1.8	6.0 342.2 3.9	330	1.0 0.0 1.0	48.2 74.2 -8.7
893	B50R_100_025d	1.0 0.75 1.0	1.0 0.25 0.875	330	1.0 0.75 1.0	84.3 18.5 -2.1	1.0 0.75 1.0	86.0 13.3 -3.6	13.8 344.4 5.7	330	1.0 0.0 1.0	48.2 74.2 -8.7
894	B50R_100_037d	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	78.3 27.8 -3.2	1.0 0.625 1.0	80.5 20.7 -5.3	21.4 345.6 7.6	330	1.0 0.0 1.0	48.2 74.2 -8.7
895	B50R_100_050d	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	72.3 37.1 -4.3	1.0 0.5 1.0	72.8 31.7 -7.1	32.5 347.3 6.0	330	1.0 0.0 1.0	48.2 74.2 -8.7
896	B50R_100_062d	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	66.2 46.3 -5.4	1.0 0.375 1.0	66.0 41.9 -8.1	42.7 349.0 5.1	330	1.0 0.0 1.0	48.2 74.2 -8.7
897	B50R_100_075d	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	60.2 55.6 -6.5	1.0 0.25 1.0	59.4 52.7 -8.4	53.4 350.8 3.5	330	1.0 0.0 1.0	48.2 74.2 -8.7
898	B50R_100_087d	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	54.2 64.9 -7.6	1.0 0.125 1.0	52.4 65.2 -8.3	65.7 352.6 1.9	330	1.0 0.0 1.0	48.2 74.2 -8.7
899	B50R_100_100d	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	48.2 74.2 -8.7	1.0 0.0 1.0	47.2 75.2 -7.1	75.5 354.5 2.1	330	1.0 0.0 1.0	48.2 74.2 -8.7
900	GO0B_100_012d	0.875 1.0 0.875	1.0 0.125 0.937	150	0.875 1.0 0.875	90.7 -8.6 2.8	0.875 1.0 0.875	92.2 -4.9 3.7	6.2 142.9 4.0	149	0.0 1.0 0.0	51.6 -69.3
901	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.6 0.0 0.0	0.875 0.875 0.875	91.0 0.0 -0.4	0.4 260.7 4.4	360	1.0 1.0 1.0	96.3 0.0 0.0
902	B50R_087_012d	0.875 0.75 0.875	0.875 0.125 0.812	330	0.875 0.75 0.875	80.6 9.2 -1.0	0.875 0.75 0.875	86.5 5.8 -2.3	6.3 337.9 6.9	330	1.0 0.0 1.0	48.2 74.2 -8.7
903	B50R_087_025d	0.875 0.625 0.875	0.875 0.25 0.75	330	0.875 0.625 0.875	74.6 18.5 -2.1	0.875 0.625 0.875	79.8 14.3 -4.4	15.0 342.9 7.0	330	1.0 0.0 1.0	48.2 74.2 -8.7
904	B50R_087_037d	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	68.6 27.8 -3.2	0.875 0.5 0.875	73.9 22.3 -5.8	23.1 345.2 8.0	330	1.0 0.0 1.0	48.2 74.2 -8.7
905	B50R_087_050d	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	62.5 37.1 -4.3	0.875 0.375 0.875	66.1 33.4 -7.4	34.3 347.4 5.9	330	1.0 0.0 1.0	48.2 74.2 -8.7
906	B50R_087_062d	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	56.5 46.3 -5.4	0.875 0.25 0.875	58.1 46.0 -8.5	46.8 349.5 3.4	330	1.0 0.0 1.0	48.2 74.2 -8.7
907	B50R_087_075d	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	50.5 55.6 -6.5	0.875 0.125 0.875	51.0 58.4 -8.5	59.1 351.6 3.4	330	1.0 0.0 1.0	48.2 74.2 -8.7
908	B50R_087_087d	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	44.5 64.9 -7.6	0.875 0.0 0.875	44.8 70.2 -7.8	70.6 353.5 5.3	330	1.0 0.0 1.0	48.2 74.2 -8.7
909	GO0B_100_025d	0.75 1.0 0.75	1.0 0.25 0.875	150	0.75 1.0 0.75	85.1 -17.3 5.7	0.75 1.0 0.75	87.3 -9.9 7.4	12.3 143.1 7.9	149	0.0 1.0 0.0	51.6 -69.3
910	GO0B_087_012d	0.75 0.875 0.75	0.875 0.125 0.812	150	0.75 0.875 0.75	81.0 -8.6 2.8	0.75 0.875 0.75	86.7 -5.3 3.6	6.4 145.3 6.6	149	0.0 1.0 0.0	51.6 -69.3
911	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	76.9 0.0 0.0	0.75 0.75 0.75	83.5 -0.1 -1.0	1.0 261.1 6.7	360	1.0 1.0 1.0	96.3 0.0 0.0
912	B50R_075_012d	0.75 0.625 0.75	0.75 0.125 0.687	330	0.75 0.625 0.75	70.9 9.2 -1.0	0.75 0.625 0.75	78.4 6.3 -3.0	7.0 334.1 8.3	330	1.0 0.0 1.0	48.2 74.2 -8.7
913	B50R_075_025d	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	64.8 18.5 -2.1	0.75 0.5 0.75	71.7 14.7 -4.8	15.5 341.6 8.3	330	1.0 0.0 1.0	48.2 74.2 -8.7
914	B50R_075_037d	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	58.8 27.8 -3.2	0.75 0.375 0.75	64.2 24.9 -6.7	25.8 344.8 7.0	330	1.0 0.0 1.0	48.2 74.2 -8.7
915	B50R_075_050d	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	52.8 37.1 -4.3	0.75 0.25 0.75	56.2 37.0 -7.9	37.8 347.8 4.9	330	1.0 0.0 1.0	48.2 74.2 -8.7
916	B50R_075_062d	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	46.8 46.3 -5.4	0.75 0.125 0.75	48.6 49.4 -8.4	50.1 350.3 4.5	330	1.0 0.0 1.0	48.2 74.2 -8.7
917	B50R_075_075d	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	40.8 55.6 -6.5	0.75 0.0 0.75	42.1 61.9 -8.3	62.5 352.3 6.6	330	1.0 0.0 1.0	48.2 74.2 -8.7
918	GO0B_100_037d	0.625 1.0 0.625	1.0 0.375 0.812	150	0.625 1.0 0.625	79.6 -26.0 8.6	0.625 1.0 0.625	81.9 -15.7 10.8	19.1 145.4 10.7	149	0.0 1.0 0.0	51.6 -69.3
919	GO0B_087_025d	0.625 0.875 0.625	0.875 0.25 0.75	150	0.625 0.875 0.625	75.4 -17.3 5.7	0.625 0.875 0.625	81.6 -10.7 7.4	13.0 145.1 9.1	149	0.0 1.0 0.0	51.6 -69.3
920	GO0B_075_012d	0.625 0.75 0.625	0.75 0.125 0.687	150	0.625 0.75 0.625	71.3 -8.6 2.8	0.625 0.75 0.625	78.9 -5.6 3.2	6.4 149.8 8.2	149	0.0 1.0 0.0	51.6 -69.3
921	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	67.1 0.0 0.0	0.625 0.625 0.625	76.6 -0.2 -1.4	1.5 261.2 9.5	360	1.0 1.0 1.0	96.3 0.0 0.0
922	B50R_062_012d	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	61.1 9.2 -1.0	0.625 0.5 0.625	70.3 7.0 -3.6	7.9 332.9 9.7	330	1.0 0.0 1.0	48.2 74.2 -8.7
923	B50R_062_025d	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	55.1 18.5 -2.1	0.625 0.375 0.625	63.6 15.7 -5.4	16.6 341.0 9.5	330	1.0 0.0 1.0	48.2 74.2 -8.7
924	B50R_062_037d	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	49.1 27.8 -3.2	0.625 0.25 0.625	55.4 27.4 -7.0	28.3 345.5 7.3	330	1.0 0.0 1.0	48.2 74.2 -8.7
925	B50R_062_050d	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	43.1 37.1 -4.3	0.625 0.125 0.625	46.9 40.7 -8.0	41.5 348.8 6.4	330	1.0 0.0 1.0	48.2 74.2 -8.7
926	B50R_062_062d	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	37.1 46.3 -5.4	0.625 0.0 0.625	39.6 54.2 -8.0	54.8 351.5 8.6	330	1.0 0.0 1.0	48.2 74.2 -8.7
927	GO0B_100_050d	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	74.0 -34.6 11.5	0.5 1.0 0.5	75.7 -22.9 13.7	26.7 149.0 12.0	149	0.0 1.0 0.0	51.6 -69.3
928	GO0B_087_037d	0.5 0.875 0.5	0.875 0.375 0.687	150	0.5 0.875 0.5	69.8 -26.0 8.6	0.5 0.875 0.5	75.1 -17.9 10.1	20.5 150.5 9.7	149	0.0 1.0 0.0	51.6 -69.3
929	GO0B_075_025d	0.5 0.75 0.5	0.75 0.25 0.625	150	0.5 0.75 0.5	65.7 -17.3 5.7	0.5 0.75 0.5	73.3 -11.5 7.0	13.5 148.6 9.6	149	0.0 1.0 0.0	51.6 -69.3
930	GO0B_062_012d	0.5 0.625 0.5	0.625 0.125 0.562	150	0.5 0.625 0.5	61.5 -8.6 2.8	0.5 0.625 0.5	71.3 -6.1 2.7	6.7 156.2 10.0	149	0.0 1.0 0.0	51.6 -69.3
931	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	57.4 0.0 0.0	0.5 0.5 0.5	68.5 -0.2 -1.8	1.8 261.0 11.2	360	1.0 1.0 1.0	96.3 0.0 0.0
932	B50R_050_012d	0.5 0.375 0.5	0.5 0.125 0.437	330	0.5 0.375 0.5	51.4 9.2 -1.0	0.5 0.375 0.5	61.7 8.3 -4.1	9.3 333.6 10.7	330	1.0 0.0 1.0	48.2 74.2 -8.7
933	B50R_050_025d	0.5 0.25 0.5	0.5 0.25 0.375	330	0.5 0.249 0.5	45.4 18.5 -2.1	0.5 0.25 0.5	53.8 18.7 -6.0	19.6 342.2 9.2	330	1.0 0.0 1.0	48.2 74.2 -8.7
934	B50R_050_037d	0.5 0.125 0.5	0.5 0.375 0.312	330	0.5 0.124 0.5	39.4 27.8 -3.2	0.5 0.125 0.5	44.5 32.3 -7.3	33.1 347.1 7.9	330	1.0 0.0 1.0	48.2 74.2 -8.7
935	B50R_050_050d	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	33.3 37.1 -4.3	0.5 0.0 0.5	36.8 45.8 -7.6	46.4 350.5 9.9	330	1.0 0.0 1.0	48.2 74.2 -8.7
936	GO0B_100_062d	0.375 1.0 0.375	1.0 0.625 0.687	150	0.375 1.0 0.375	68.4 -43.3 14.4	0.375 1.0 0.375	69.1 -31.2 17.0	35.5 151.3 12.4	149	0.0 1.0 0.0	51.6 -69.3
937	GO0B_087_050d	0.375 0.875 0.375	0.875 0.5 0.625	150	0.375 0.875 0.375	64.2 -34.6 11.5	0.375 0.875 0.375	68.9 -25.6 14.1	29.3 151.2 10.4	149	0.0 1.0 0.0	51.6 -69.3
938	GO0B_075_037d	0.375 0.75 0.375										

http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF /.PS; transfer output

N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 32/33

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		HIC*Fd			rgb_Fd			ict_Fd			hsi_Fd			rgb*Fd			LabCh*Fd						rgb**Fd			LabCh**Fd			DE**Fd			hsiMd	rgb**Md			LabCh**Md						
972	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	18.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.3	0.3	76.3	1.9	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0				
973	NW_012d	0.125	0.125	0.125	0.125	0.125	0.125	0.125	360	0.125	0.125	0.125	28.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.125	0.125	0.125	33.4	-0.2	-1.2	1.2	259.3	5.3	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
974	NW_025d	0.25	0.25	0.25	0.25	0.25	0.25	0.25	360	0.25	0.25	0.25	37.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.25	0.25	0.25	49.1	-0.3	-1.9	2.0	259.1	11.3	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
975	NW_037d	0.375	0.375	0.375	0.375	0.375	0.375	0.375	360	0.375	0.375	0.375	47.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.375	0.375	0.375	59.5	-0.3	-2.0	2.0	260.3	12.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
976	NW_050d	0.5	0.5	0.5	0.5	0.5	0.5	0.5	360	0.5	0.5	0.5	57.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.5	0.5	0.5	68.4	-0.2	-1.8	1.8	261.7	11.1	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
977	NW_062d	0.625	0.625	0.625	0.625	0.625	0.625	0.625	360	0.625	0.625	0.625	67.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.625	0.625	0.625	76.7	-0.2	-1.5	1.5	262.5	9.7	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
978	NW_075d	0.75	0.75	0.75	0.75	0.75	0.75	0.75	360	0.75	0.75	0.75	76.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.75	0.75	0.75	84.2	-0.1	-1.1	1.1	261.9	7.4	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
979	NW_087d	0.875	0.875	0.875	0.875	0.875	0.875	0.875	360	0.875	0.875	0.875	86.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.875	0.875	0.875	91.5	0.0	-0.4	0.4	260.0	4.9	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
980	NW_100d	1.0	1.0	1.0	1.0	1.0	1.0	1.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	1.0	1.0	96.5	0.0	0.0	0.0	251.4	0.2	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
981	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	18.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	20.1	0.0	0.1	0.1	78.3	1.6	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
982	NW_012d	0.125	0.125	0.125	0.125	0.125	0.125	0.125	360	0.125	0.125	0.125	28.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.125	0.125	0.125	33.3	-0.2	-1.3	1.3	258.6	5.2	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
983	NW_025d	0.25	0.25	0.25	0.25	0.25	0.25	0.25	360	0.25	0.25	0.25	37.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.25	0.25	0.25	48.4	-0.3	-2.0	2.0	259.6	10.6	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
984	NW_037d	0.375	0.375	0.375	0.375	0.375	0.375	0.375	360	0.375	0.375	0.375	47.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.375	0.375	0.375	59.7	-0.3	-2.0	2.0	260.4	12.2	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
985	NW_050d	0.5	0.5	0.5	0.5	0.5	0.5	0.5	360	0.5	0.5	0.5	57.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.5	0.5	0.5	68.4	-0.2	-1.8	1.8	261.5	11.1	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
986	NW_062d	0.625	0.625	0.625	0.625	0.625	0.625	0.625	360	0.625	0.625	0.625	67.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.625	0.625	0.625	76.6	-0.2	-1.5	1.5	262.7	9.6	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
987	NW_075d	0.75	0.75	0.75	0.75	0.75	0.75	0.75	360	0.75	0.75	0.75	76.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.75	0.75	0.75	83.9	-0.1	-1.1	1.1	262.2	7.1	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
988	NW_087d	0.875	0.875	0.875	0.875	0.875	0.875	0.875	360	0.875	0.875	0.875	86.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.875	0.875	0.875	91.5	0.0	-0.4	0.4	262.2	4.8	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
989	NW_100d	1.0	1.0	1.0	1.0	1.0	1.0	1.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	1.0	1.0	96.5	0.0	0.0	0.0	217.9	0.1	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
990	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	18.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	19.4	0.0	0.0	0.0	59.7	0.9	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
991	NW_012d	0.125	0.125	0.125	0.125	0.125	0.125	0.125	360	0.125	0.125	0.125	28.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.125	0.125	0.125	32.8	-0.2	-1.4	1.4	258.4	4.7	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
992	NW_025d	0.25	0.25	0.25	0.25	0.25	0.25	0.25	360	0.25	0.25	0.25	37.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.25	0.25	0.25	48.5	-0.4	-2.1	2.1	259.2	10.7	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
993	NW_037d	0.375	0.375	0.375	0.375	0.375	0.375	0.375	360	0.375	0.375	0.375	47.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.375	0.375	0.375	59.4	-0.3	-2.1	2.2	261.3	11.8	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
994	NW_050d	0.5	0.5	0.5	0.5	0.5	0.5	0.5	360	0.5	0.5	0.5	57.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.5	0.5	0.5	67.9	-0.2	-2.0	2.0	262.2	10.7	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
995	NW_062d	0.625	0.625	0.625	0.625	0.625	0.625	0.625	360	0.625	0.625	0.625	67.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.625	0.625	0.625	76.1	-0.2	-1.7	1.7	262.8	9.1	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
996	NW_075d	0.75	0.75	0.75	0.75	0.75	0.75	0.75	360	0.75	0.75	0.75	76.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.75	0.75	0.75	83.5	-0.1	-1.3	1.3	263.1	6.7	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
997	NW_087d	0.875	0.875	0.875	0.875	0.875	0.875	0.875	360	0.875	0.875	0.875	86.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.875	0.875	0.875	91.1	0.0	-0.7	0.7	264.8	4.5	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
998	NW_100d	1.0	1.0	1.0	1.0	1.0	1.0	1.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	1.0	1.0	96.2	0.0	-0.2	0.2	267.4	0.2	360	1.0	1.0	1.0	96				

http://130.149.60.45/~farbmetrik/SE14/SE14L0NP.PDF /.PS; transfer output

N: no 3D-linearization (OL) in file (F) or PS-startup (S), page 33/33

see similar files: <http://130.149.60.45/~farbmetrik/SE14/SE14.HTM>
technical information: <http://www.ps.bam.de> or <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd		ict_Fd		hsi_Fd		rgb*Fd		LabCh*Fd		rgb**Fd		LabCh**Fd		DE**Fd		hsiMd	rgb*Md		LabCh*Md													
1053	NW_086d	0.866	0.866	0.866	0.866	0.0	0.866	360	0.866	0.866	0.866	85.9	0.0	0.0	0.0	0.0	0.866	0.866	0.866	91.0	0.0	-0.5	0.5	264.4	5.1	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1054	NW_093d	0.933	0.933	0.933	0.933	0.0	0.933	360	0.933	0.933	0.933	91.1	0.0	0.0	0.0	0.0	0.933	0.933	0.933	93.5	0.0	-0.1	0.3	271.8	2.4	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1055	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0	1.0	1.0	1.0	96.3	0.0	-0.1	0.1	284.8	0.1	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1056	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	18.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	19.2	0.0	0.0	0.0	6.0	0.7	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1057	NW_006d	0.066	0.066	0.066	0.066	0.0	0.066	360	0.066	0.066	0.066	23.6	0.0	0.0	0.0	0.0	0.066	0.066	0.066	24.0	0.0	-0.5	0.5	260.1	0.6	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1058	NW_013d	0.133	0.133	0.133	0.133	0.0	0.133	360	0.133	0.133	0.133	28.8	0.0	0.0	0.0	0.0	0.133	0.133	0.133	33.3	-0.3	-1.5	1.5	258.6	4.7	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1059	NW_020d	0.2	0.2	0.2	0.2	0.0	0.2	360	0.2	0.2	0.2	34.1	0.0	0.0	0.0	0.0	0.2	0.2	0.2	42.5	-0.3	-2.0	2.0	259.3	8.6	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1060	NW_026d	0.266	0.266	0.266	0.266	0.0	0.266	360	0.266	0.266	0.266	39.2	0.0	0.0	0.0	0.0	0.266	0.266	0.266	48.9	-0.3	-2.2	2.2	260.4	9.9	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1061	NW_033d	0.333	0.333	0.333	0.333	0.0	0.333	360	0.333	0.333	0.333	44.4	0.0	0.0	0.0	0.0	0.333	0.333	0.333	55.2	-0.3	-2.2	2.3	260.3	11.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1062	NW_040d	0.4	0.4	0.4	0.4	0.0	0.4	360	0.4	0.4	0.4	49.6	0.0	0.0	0.0	0.0	0.4	0.4	0.4	60.4	-0.3	-2.1	2.2	261.8	11.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1063	NW_046d	0.466	0.466	0.466	0.466	0.0	0.466	360	0.466	0.466	0.466	54.8	0.0	0.0	0.0	0.0	0.466	0.466	0.466	65.1	-0.2	-2.0	2.2	261.9	10.5	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1064	NW_053d	0.533	0.533	0.533	0.533	0.0	0.533	360	0.533	0.533	0.533	60.0	0.0	0.0	0.0	0.0	0.533	0.533	0.533	70.0	-0.2	-1.9	1.9	262.3	10.2	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1065	NW_060d	0.6	0.6	0.6	0.6	0.0	0.6	360	0.6	0.6	0.6	65.2	0.0	0.0	0.0	0.0	0.6	0.6	0.6	74.7	-0.2	-1.6	1.6	262.4	9.6	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1066	NW_066d	0.666	0.666	0.666	0.666	0.0	0.666	360	0.666	0.666	0.666	70.3	0.0	0.0	0.0	0.0	0.666	0.666	0.666	78.8	-0.1	-1.4	1.4	262.5	8.6	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1067	NW_073d	0.734	0.734	0.734	0.734	0.0	0.734	360	0.734	0.734	0.734	75.6	0.0	0.0	0.0	0.0	0.734	0.734	0.734	82.8	-0.1	-1.2	1.2	262.2	7.2	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1068	NW_080d	0.8	0.8	0.8	0.8	0.0	0.8	360	0.8	0.8	0.8	80.8	0.0	0.0	0.0	0.0	0.8	0.8	0.8	86.6	0.0	-0.9	0.9	264.2	5.8	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1069	NW_086d	0.866	0.866	0.866	0.866	0.0	0.866	360	0.866	0.866	0.866	85.9	0.0	0.0	0.0	0.0	0.866	0.866	0.866	90.9	0.0	-0.6	0.6	268.4	5.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1070	NW_093d	0.933	0.933	0.933	0.933	0.0	0.933	360	0.933	0.933	0.933	91.1	0.0	0.0	0.0	0.0	0.933	0.933	0.933	93.4	0.0	-0.3	0.3	272.6	2.3	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1071	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0	1.0	1.0	1.0	96.3	0.0	0.0	0.0	289.1	0.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1072	NW_000d	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	18.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	20.7	0.1	0.3	0.3	65.1	2.1	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1073	NW_100d	1.0	1.0	1.0	1.0	0.0	1.0	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0	1.0	1.0	1.0	96.6	0.0	0.0	0.0	196.8	0.2	360	1.0	1.0	1.0	96.3	0.0	0.0	0.0	0.0
1074	R00Y_100_100d	1.0	0.0	0.0	1.0	1.0	0.5	390	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4	1.0	0.0	0.0	45.6	67.7	39.1	78.2	30.0	2.8	389	1.0	0.0	0.0	47.5	65.5	38.4	76.0	30.4
1075	G50B_100_100d	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6	0.0	1.0	1.0	56.4	-31.6	-46.1	55.9	235.5	1.7	210	0.0	1.0	1.0	57.8	-31.9	-45.1	55.3	234.6
1076	Y00G_100_100d	1.0	1.0	0.0	1.0	1.0	0.5	90	1.0	1.0	0.0	89.4	-9.5	89.0	89.6	96.0	1.0	1.0	0.0	87.9	-7.9	89.2	89.6	95.0	2.1	89	1.0	1.0	0.0	89.4	-9.5	89.0	89.6	96.0
1077	B00R_100_100d	0.0	0.0	1.0	1.0	1.0	0.5	270	0.0	0.0	1.0	24.9	22.9	-47.8	53.0	295.6	0.0	0.0	1.0	23.1	24.3	-46.5	52.5	297.6	2.6	270	0.0	0.0	1.0	24.9	22.9	-47.8	53.0	295.6
1078	G00B_100_100d	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	51.6	-69.3	23.0	73.1	161.6	0.0	1.0	0.0	48.7	-69.8	21.6	73.1	162.7	3.2	149	0.0	1.0	0.0	51.6	-69.3	23.0	73.1	161.6
1079	B50R_100_100d	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	48.2	74.2	-8.7	74.7	353.2	1.0	0.0	1.0	45.8	75.8	-5.0	76.0	356.2	4.7	330	1.0	0.0	1.0	48.2	74.2	-8.7	74.7	353.2

Mean color difference of this page:

delta E* = 4.9

TUB registration: 20130201-SE14/SE14L0NP.PDF /.PS
application for measurement of offset print output, separationcnmy6 (CMYK)
TUB material: code=rha4taTUB-test chart SE14; 1080 colours, offset standard paper
colours and differences, ΔE^* , 3D=0, de=0, *cmYk*input: *rgb/cmyk* -> *rgb_d*
output: transfer to *cmYk_d*