

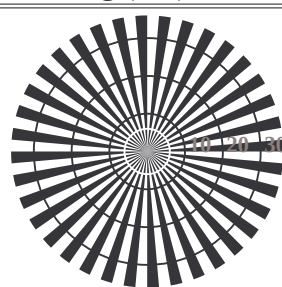
TG771-5, Bild C6W-: Element F: Linienraster unter 90° (oder 0°); PS-Operator: *rgb/cmyc*

Siehe ähnliche Dateien: <http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT /.PS>
Technische Information: <http://www.ps.bam.de> oder <http://130.149.60.45/~farbmetrik>

TUB-Registrierung: 20150901-TG77/TG77L0NA.TXT /.PS TUB-Material: Code=rh4ta
Anwendung für Messung von Offsetdruck-Ausgabe, Separation cmyk0 (CMY0)



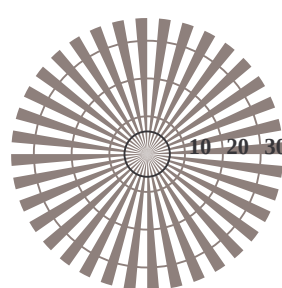
Radialgitter (Siemenssterne) N-W



Radialgitter (Siemenssterne) W-N



Radialgitter (Siemenssterne) N-Z



Radialgitter (Siemenssterne) W-Z

TG770-3, Bild C1Wd: Element A: Radialgitter N-W, W-N, N-Z und W-Z; PS-Operator: *rgb/cmy0*

L^*/Y_{Eingabe} (absolut)	18.0/2.5	37.3/9.7	56.7/24.6	76.1/49.9	95.4/88.6	N_0 (min.)	W_1 (max.)
$w^*=l^*_{\text{CIELAB}, r}$ (relativ)							
w^*_{Eingabe}	0,000	0,250	0,500	0,750	1,000	N_0 (min.)	W_1 (max.)

TG770-5, Bild C2Wd: Element B: 5 visuell gleichabständige L^* -Graustufen + N_0 + W_1 ; PS-Operator: *rgb/cmy0*

L^*/Y_{Eingabe} (absolut)	18.0/2.5	23.2/3.8	28.3/5.6	33.5/7.8	38.6/10.5	43.8/13.7	49.0/17.6	54.1/22.1	59.3/27.3	64.4/33.3	69.6/40.2	74.8/47.9	79.9/56.5	85.1/66.2	90.2/76.8	95.4/88.6
Nr. und Hex-Code	00;F	01;E	02;D	03;C	04;B	05;A	06;9	07;8	08;7	09;6	10;5	11;4	12;3	13;2	14;1	15;0
$w^*=l^*_{\text{CIELAB}, r}$ (relativ)																
w^*_{Eingabe}	0,000	0,067	0,133	0,200	0,267	0,333	0,400	0,467	0,533	0,600	0,667	0,733	0,800	0,867	0,933	1,000

TG770-7, Bild C3Wd: Element C: 16 visuell gleichabständige L^* -Graustufen; PS-Operator: *rgb/cmy0*



Prüfvorlage TG77; ME16(ISO 9241-306), 3(ISO/IEC 15775)
Achromatische Prüfvorlage N, 3D=0, de=0, cmyk

Eingabe: *rgb/cmyk* -> *rgb_d*
Ausgabe: Transfer nach *cmyk_d*



Umfeldstufe Hex-Code	0	7	E	2	8	F
Ringstufe Hex-Code	0-1	7-8	E-F	2-0	8-6	F-D
Landoltringe W-N						
Code: Umfeld-Ring						

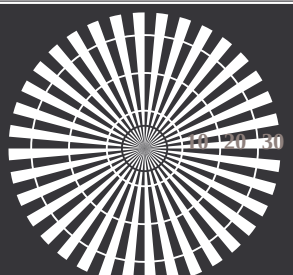
TG771-1, Bild C4Wd: Element D: Landoltringe W-N; PS-Operator: *rgb/cmy0*

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	
Rasterweite in lpi																	

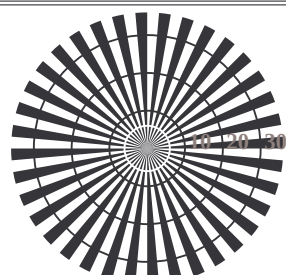
TG771-3, Bild C5Wd: Element E: Linienraster unter 45° (oder 135°); PS-Operator: *rgb/cmy0*

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	
Rasterweite in lpi																	

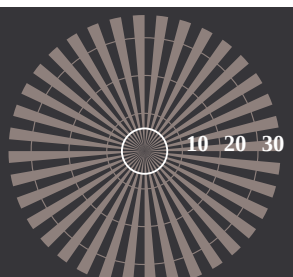
TG771-5, Bild C6Wd: Element F: Linienraster unter 90° (oder 0°); PS-Operator: *rgb/cmy0*



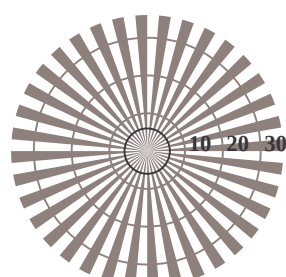
Radialgitter (Siemenssterne) N-W



Radialgitter (Siemenssterne) W-N



Radialgitter (Siemenssterne) N-Z



Radialgitter (Siemenssterne) W-Z

TG770-3, Bild C1Wd: Element A: Radialgitter N-W, W-N, N-Z und W-Z; PS-Operator: *rgb/cmy0*

L^*/Y_{Eingabe} (absolut)	18.0/2.5	37.3/9.7	56.7/24.6	76.1/49.9	95.4/88.6	N_0 (min.)	W_1 (max.)
$w^*=l^*_{\text{CIE LAB}, r}$ (relativ)						N_0 (min.)	W_1 (max.)
w^*_{Eingabe}	0,000	0,250	0,500	0,750	1,000	N_0 (min.)	W_1 (max.)

TG770-5, Bild C2Wd: Element B: 5 visuell gleichabständige L^* -Graustufen + N_0 + W_1 ; PS-Operator: *rgb/cmy0*

L^*/Y_{Eingabe} (absolut)	18.0/2.5	23.2/3.8	28.3/5.6	33.5/7.8	38.6/10.5	43.8/13.7	49.0/17.6	54.1/22.1	59.3/27.3	64.4/33.3	69.6/40.2	74.8/47.9	79.9/56.5	85.1/66.2	90.2/76.8	95.4/88.6
Nr. und Hex-Code	00;F	01;E	02;D	03;C	04;B	05;A	06;9	07;8	08;7	09;6	10;5	11;4	12;3	13;2	14;1	15;0
$w^*=l^*_{\text{CIE LAB}, r}$ (relativ)																
w^*_{Eingabe}	0,000	0,067	0,133	0,200	0,267	0,333	0,400	0,467	0,533	0,600	0,667	0,733	0,800	0,867	0,933	1,000

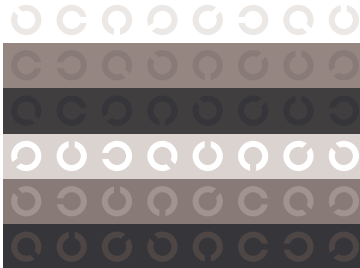
TG770-7, Bild C3Wd: Element C: 16 visuell gleichabständige L^* -Graustufen; PS-Operator: *rgb/cmy0*

Prüfvorlage TG77; ME16(ISO 9241-306), 3(ISO/IEC 15775)
Achromatische Prüfvorlage N, 3D=0, de=0, *cmyk*

Eingabe: *rgb/cmyk* -> *rgb_d*
Ausgabe: Transfer nach *cmyk_d*

Umfeldstufe
Hex-Code

0
7
E
2
8
F



Landoltringe W-N

1 Ringstufe
Hex-Code

0-1
7-8
E-F
2-0
8-6
F-D

Code: Umfeld-Ring

TG771-1, Bild C4Wd: Element D: Landoltringe W-N; PS-Operator: *rgb/cmy0*

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	

Rasterweite in lpi

TG771-3, Bild C5Wd: Element E: Linienraster unter 45° (oder 135°); PS-Operator: *rgb/cmy0*

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	

Rasterweite in lpi

TG771-5, Bild C6Wd: Element F: Linienraster unter 90° (oder 0°); PS-Operator: *rgb/cmy0*

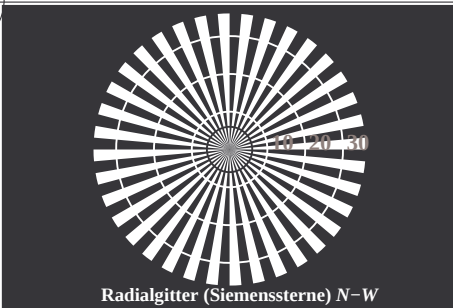
Siehe ähnliche Dateien: <http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT> /PS
Technische Information: <http://www.ps.bam.de> oder <http://130.149.60.45/~farbmetrik>



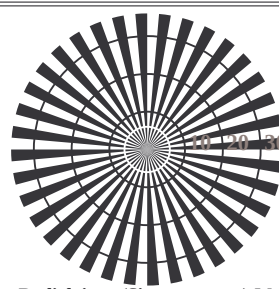
<http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT> /PS; Transfer Ausgabe
N: Keine 3D-Linearisierung (OL) in Datei (F) oder PS-Startup (S), Seite 4/22



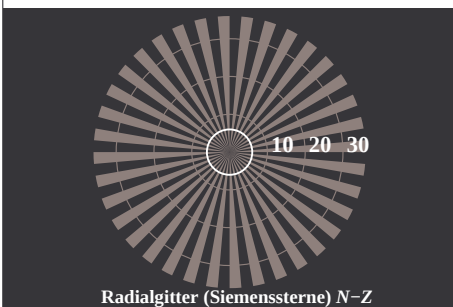
TUB-Registrierung: 20150901-TG77/TG77L0NA.TXT /PS TUB-Material: Code=rh4ta
Anwendung für Messung von Offsetdruck-Ausgabe, Separation cmyk6 (CMY0)



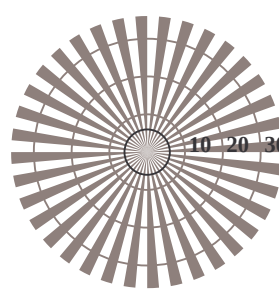
Radialgitter (Siemenssterne) N-W



Radialgitter (Siemenssterne) W-N



Radialgitter (Siemenssterne) N-Z



Radialgitter (Siemenssterne) W-Z

TG770-3, Bild C1Wd: Element A: Radialgitter N-W, W-N, N-Z und W-Z; PS-Operator: *rgb/cmy0*

L^*/Y_{Eingabe} (absolut)	18.0/2.5	37.3/9.7	56.7/24.6	76.1/49.9	95.4/88.6	N_0 (min.)	W_1 (max.)
$w^*=l^*_{\text{CIELAB}, r}$ (relativ)						N_0 (min.)	W_1 (max.)
w^*_{Eingabe}	0,000	0,250	0,500	0,750	1,000	N_0 (min.)	W_1 (max.)

TG770-5, Bild C2Wd: Element B: 5 visuell gleichabständige L^* -Graustufen + N_0 + W_1 ; PS-Operator: *rgb/cmy0*

L^*/Y_{Eingabe} (absolut)	18.0/2.5	23.2/3.8	28.3/5.6	33.5/7.8	38.6/10.5	43.8/13.7	49.0/17.6	54.1/22.1	59.3/27.3	64.4/33.3	69.6/40.2	74.8/47.9	79.9/56.5	85.1/66.2	90.2/76.8	95.4/88.6
Nr. und Hex-Code	00;F	01;E	02;D	03;C	04;B	05;A	06;9	07;8	08;7	09;6	10;5	11;4	12;3	13;2	14;1	15;0
$w^*=l^*_{\text{CIELAB}, r}$ (relativ)																
w^*_{Eingabe}	0,000	0,067	0,133	0,200	0,267	0,333	0,400	0,467	0,533	0,600	0,667	0,733	0,800	0,867	0,933	1,000

TG770-7, Bild C3Wd: Element C: 16 visuell gleichabständige L^* -Graustufen; PS-Operator: *rgb/cmy0*



Prüfvorlage TG77; ME16(ISO 9241-306), 3(ISO/IEC 15775)
Achromatische Prüfvorlage N, 3D=0, de=0, cmyk

Eingabe: *rgb/cmyk* -> *rgb_d*
Ausgabe: Transfer nach *cmyk_d*



Umfeldstufe Hex-Code	0	7	E	2	8	F
Landoltringe W-N						

Landoltringe W-N

Ringstufe Hex-Code	0-1	7-8	E-F	2-0	8-6	F-D
Code: Umfeld-Ring						

Code: Umfeld-Ring

TG771-1, Bild C4Wd: Element D: Landoltringe W-N; PS-Operator: *rgb/cmy0*

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	

Rasterweite in lpi

TG771-3, Bild C5Wd: Element E: Linienraster unter 45° (oder 135°); PS-Operator: *rgb/cmy0*

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	

Rasterweite in lpi

TG771-5, Bild C6Wd: Element F: Linienraster unter 90° (oder 0°); PS-Operator: *rgb/cmy0*

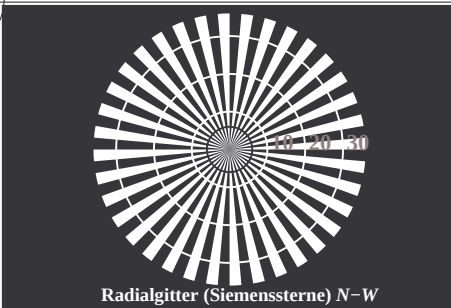
Siehe ähnliche Dateien: <http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT> /PS
Technische Information: <http://www.ps.bam.de> oder <http://130.149.60.45/~farbmetrik>



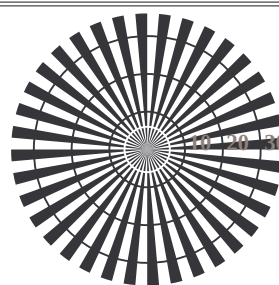
<http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT> /PS; Transfer Ausgabe
N: Keine 3D-Linearisierung (OL) in Datei (F) oder PS-Startup (S), Seite 5/22



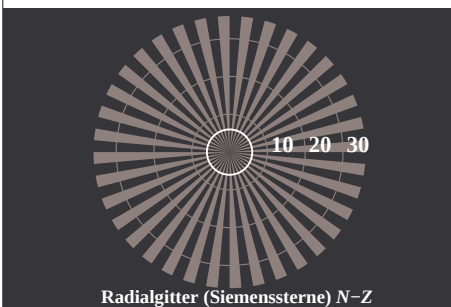
TUB-Registrierung: 20150901-TG77/TG77L0NA.TXT /PS TUB-Material: Code=rh4ta
Anwendung für Messung von Offsetdruck-Ausgabe, Separation cmyk0 (CMY0)



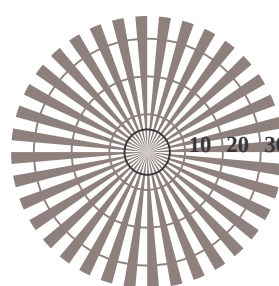
Radialgitter (Siemenssterne) N-W



Radialgitter (Siemenssterne) W-N



Radialgitter (Siemenssterne) N-Z



Radialgitter (Siemenssterne) W-Z

TG770-3, Bild C1Wd: Element A: Radialgitter N-W, W-N, N-Z und W-Z; PS-Operator: *rgb/cmy0*

L^*/Y_{Eingabe} (absolut)	18.0/2.5	37.3/9.7	56.7/24.6	76.1/49.9	95.4/88.6	N_0 (min.)	W_1 (max.)
$w^*=l^*_{\text{CIE LAB}, r}$ (relativ)						N_0 (min.)	W_1 (max.)
w^*_{Eingabe}	0,000	0,250	0,500	0,750	1,000	N_0 (min.)	W_1 (max.)

TG770-5, Bild C2Wd: Element B: 5 visuell gleichabständige L^* -Graustufen + N_0 + W_1 ; PS-Operator: *rgb/cmy0*

L^*/Y_{Eingabe} (absolut)	18.0/2.5	23.2/3.8	28.3/5.6	33.5/7.8	38.6/10.5	43.8/13.7	49.0/17.6	54.1/22.1	59.3/27.3	64.4/33.3	69.6/40.2	74.8/47.9	79.9/56.5	85.1/66.2	90.2/76.8	95.4/88.6
Nr. und Hex-Code	00;F	01;E	02;D	03;C	04;B	05;A	06;9	07;8	08;7	09;6	10;5	11;4	12;3	13;2	14;1	15;0
$w^*=l^*_{\text{CIE LAB}, r}$ (relativ)																
w^*_{Eingabe}	0,000	0,067	0,133	0,200	0,267	0,333	0,400	0,467	0,533	0,600	0,667	0,733	0,800	0,867	0,933	1,000

TG770-7, Bild C3Wd: Element C: 16 visuell gleichabständige L^* -Graustufen; PS-Operator: *rgb/cmy0*



Prüfvorlage TG77; ME16(ISO 9241-306), 3(ISO/IEC 15775)
Achromatische Prüfvorlage N, 3D=0, de=0, *cmyk*

Eingabe: *rgb/cmyk* -> *rgb_d*
Ausgabe: Transfer nach *cmyk_d*



Umfeldstufe Hex-Code	0		1	Ringstufe Hex-Code	0-1
	7		8		7-8
	E		F		E-F
	2		0		2-0
	8		6		8-6
	F		D		F-D
Landoltringe W-N					Code: Umfeld-Ring

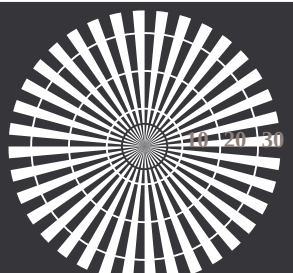
TG771-1, Bild C4Wd: Element D: Landoltringe W-N; PS-Operator: *rgb/cmy0*

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	
Rasterweite in lpi																	

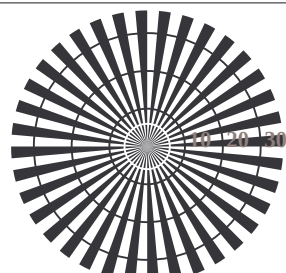
TG771-3, Bild C5Wd: Element E: Linienraster unter 45° (oder 135°); PS-Operator: *rgb/cmy0*

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	
Rasterweite in lpi																	

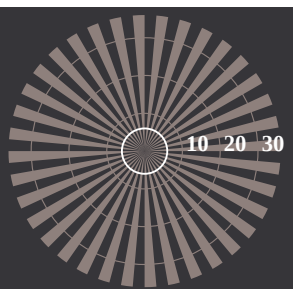
TG771-5, Bild C6Wd: Element F: Linienraster unter 90° (oder 0°); PS-Operator: *rgb/cmy0*



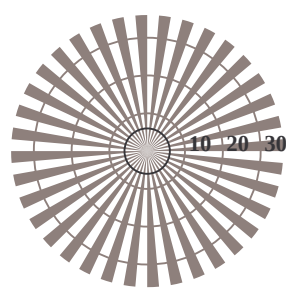
Radialgitter (Siemenssterne) N-W



Radialgitter (Siemenssterne) W-N



Radialgitter (Siemenssterne) N-Z



Radialgitter (Siemenssterne) W-Z

TG770-3, Bild C1Wd: Element A: Radialgitter N-W, W-N, N-Z und W-Z; PS-Operator: *rgb/cmy0*

L^*/Y_{Eingabe} (absolut)	18.0/2.5	37.3/9.7	56.7/24.6	76.1/49.9	95.4/88.6	N_0 (min.)	W_1 (max.)
$w^*=l^*_{\text{CIE LAB}, r}$ (relativ)							
w^*_{Eingabe}	0,000	0,250	0,500	0,750	1,000	N_0 (min.)	W_1 (max.)

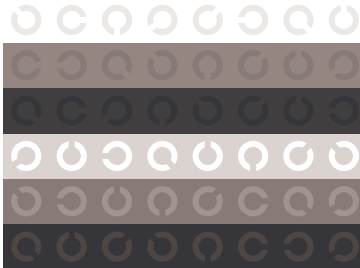
TG770-5, Bild C2Wd: Element B: 5 visuell gleichabständige L^* -Graustufen + N_0 + W_1 ; PS-Operator: *rgb/cmy0*

L^*/Y_{Eingabe} (absolut)	18.0/2.5	23.2/3.8	28.3/5.6	33.5/7.8	38.6/10.5	43.8/13.7	49.0/17.6	54.1/22.1	59.3/27.3	64.4/33.3	69.6/40.2	74.8/47.9	79.9/56.5	85.1/66.2	90.2/76.8	95.4/88.6
Nr. und Hex-Code	00;F	01;E	02;D	03;C	04;B	05;A	06;9	07;8	08;7	09;6	10;5	11;4	12;3	13;2	14;1	15;0
$w^*=l^*_{\text{CIE LAB}, r}$ (relativ)																
w^*_{Eingabe}	0,000	0,067	0,133	0,200	0,267	0,333	0,400	0,467	0,533	0,600	0,667	0,733	0,800	0,867	0,933	1,000

TG770-7, Bild C3Wd: Element C: 16 visuell gleichabständige L^* -Graustufen; PS-Operator: *rgb/cmy0*

Umfeldstufe
Hex-Code

0
7
E
2
8
F



Landoltringe W-N

1 Ringstufe
Hex-Code

0-1
7-8
E-F
2-0
8-6
F-D

Code: Umfeld-Ring

TG771-1, Bild C4Wd: Element D: Landoltringe W-N; PS-Operator: *rgb/cmy0*

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	

Rasterweite in lpi

TG771-3, Bild C5Wd: Element E: Linienraster unter 45° (oder 135°); PS-Operator: *rgb/cmy0*

	120	128	136	144	152	160	168	176	184	192	200	208	216	224	232	240	
120 (+8)																	240
60 (+4)																	120
30 (+2)																	60
15 (+1)																	30
	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	

Rasterweite in lpi

TG771-5, Bild C6Wd: Element F: Linienraster unter 90° (oder 0°); PS-Operator: *rgb/cmy0*

Siehe ähnliche Dateien: <http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT> /PS
Technische Information: <http://www.ps.bam.de> oder <http://130.149.60.45/~farbmetrik>

TUB-Registrierung: 20150901-TG77/TG77L0NA.TXT /PS
Anwendung für Messung von Offsetdruck-Ausgabe, Separation cmyk6 (CMY0)

TUB-Material: Code=rh4ta

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_100d	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	0.0 389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
1/657	R13Y_100_100d	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	48.6 63.3 49.1	80.2 37.7	1.0 0.125 0.0	48.9 62.8 49.4	79.9 38.1	0.6 36	1.0 0.116 0.0	48.6 63.3 49.1	80.2 37.7
2/666	R25Y_100_100d	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7	1.0 0.25 0.0	53.6 51.9 55.5	76.0 46.8	1.7 42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7
3/675	R38Y_100_100d	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	58.8 41.1 61.7	74.1 56.3	1.0 0.375 0.0	59.1 40.3 62.0	74.0 56.9	0.9 51	1.0 0.366 0.0	58.8 41.1 61.7	74.1 56.3
4/684	R50Y_100_100d	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1	0.0 59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
5/693	R63Y_100_100d	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	72.5 14.8 77.6	79.0 79.1	1.0 0.625 0.0	72.1 15.4 77.1	78.6 78.6	0.8 68	1.0 0.633 0.0	72.5 14.8 77.6	79.0 79.1
6/702	R75Y_100_100d	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	78.6 4.3 84.7	84.8 87.0	1.0 0.75 0.0	77.9 5.4 83.8	84.0 86.2	1.6 77	1.0 0.766 0.0	78.6 4.3 84.7	84.8 87.0
7/711	R88Y_100_100d	1.0 0.875 0.0	1.0 1.0 0.5	83	1.0 0.883 0.0	83.7 -3.8 90.5	90.6 92.4	1.0 0.875 0.0	83.4 -3.4 90.2	90.2 92.1	0.6 83	1.0 0.883 0.0	83.7 -3.8 90.5	90.6 92.4
8/720	Y00G_100_100d	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1	0.0 89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
9/639	Y13G_100_100d	0.875 1.0 0.0	1.0 1.0 0.5	97	0.883 1.0 0.0	84.5 -13.6 89.7	90.7 98.6	0.875 1.0 0.0	84.3 -13.9 89.2	90.3 98.8	0.5 96	0.883 1.0 0.0	84.5 -13.6 89.7	90.7 98.6
10/558	Y25G_100_100d	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	81.2 -17.0 84.3	86.0 101.4	0.75 1.0 0.0	80.7 -17.5 83.5	85.3 101.8	1.0 102	0.766 1.0 0.0	81.2 -17.0 84.3	86.0 101.4
11/477	Y38G_100_100d	0.625 1.0 0.0	1.0 1.0 0.5	112	0.633 1.0 0.0	75.6 -23.6 76.2	79.8 107.2	0.625 1.0 0.0	75.3 -24.0 75.7	79.4 107.6	0.7 111	0.633 1.0 0.0	75.6 -23.6 76.2	79.8 107.2
12/396	Y50G_100_100d	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0	0.0 119	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0
13/315	Y63G_100_100d	0.375 1.0 0.0	1.0 1.0 0.5	128	0.366 1.0 0.0	65.2 -36.4 57.6	68.2 122.3	0.375 1.0 0.0	65.7 -35.6 58.3	68.3 121.4	1.2 128	0.366 1.0 0.0	65.2 -36.4 57.6	68.2 122.3
14/234	Y75G_100_100d	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	57.9 -48.3 45.8	66.5 136.5	0.25 1.0 0.0	58.4 -47.3 46.8	66.6 135.3	1.4 137	0.233 1.0 0.0	57.9 -48.3 45.8	66.5 136.5
15/153	Y88G_100_100d	0.125 1.0 0.0	1.0 1.0 0.5	143	0.116 1.0 0.0	54.4 -54.7 38.0	66.6 145.1	0.125 1.0 0.0	54.7 -53.9 38.5	66.3 144.4	0.9 143	0.116 1.0 0.0	54.4 -54.7 38.0	66.6 145.1
16/72	G00C_100_100d	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5	0.0 149	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5
17/73	G13C_100_100d	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.116	50.5 -62.9 22.4	66.8 160.4	0.0 1.0 0.125	50.5 -62.8 21.9	66.5 160.7	0.5 156	0.0 1.0 0.116	50.5 -62.9 22.4	66.8 160.4
18/74	G25C_100_100d	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.233	51.1 -59.5 13.9	61.1 166.8	0.0 1.0 0.25	51.2 -58.9 12.7	60.3 167.7	1.2 162	0.0 1.0 0.233	51.1 -59.5 13.9	61.1 166.8
19/75	G38C_100_100d	0.0 1.0 0.375	1.0 1.0 0.5	172	0.0 1.0 0.366	51.9 -54.9 3.7	55.0 176.1	0.0 1.0 0.375	52.0 -54.5 3.1	54.6 176.7	0.6 171	0.0 1.0 0.366	51.9 -54.9 3.7	55.0 176.1
20/76	G50C_100_100d	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	52.9 -48.6 -8.0	49.3 189.3	0.0 1.0 0.5	52.9 -48.6 -8.0	49.3 189.3	0.0 180	0.0 1.0 0.5	52.9 -48.6 -8.0	49.3 189.3
21/77	G63C_100_100d	0.0 1.0 0.625	1.0 1.0 0.5	188	0.0 1.0 0.633	54.1 -42.0 -18.8	46.0 204.1	0.0 1.0 0.625	54.0 -42.3 -18.1	46.1 203.2	0.7 188	0.0 1.0 0.633	54.1 -42.0 -18.8	46.0 204.1
22/78	G75C_100_100d	0.0 1.0 0.75	1.0 1.0 0.5	196	0.0 1.0 0.766	55.1 -35.4 -28.4	45.4 218.7	0.0 1.0 0.75	55.0 -36.0 -27.4	45.3 217.2	1.1 197	0.0 1.0 0.766	55.1 -35.4 -28.4	45.4 218.7
23/79	G88C_100_100d	0.0 1.0 0.875	1.0 1.0 0.5	203	0.0 1.0 0.883	55.9 -30.4 -35.0	46.3 229.0	0.0 1.0 0.875	55.8 -30.7 -34.5	46.2 228.3	0.5 203	0.0 1.0 0.883	55.9 -30.4 -35.0	46.3 229.0
24/80	C00B_100_100d	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4	0.0 210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4
25/71	C13B_100_100d	0.0 0.875 1.0	1.0 1.0 0.5	217	0.0 0.883 1.0	54.3 -21.4 -41.4	46.6 242.6	0.0 0.875 1.0	54.1 -21.1 -41.3	46.4 242.9	0.3 216	0.0 0.883 1.0	54.3 -21.4 -41.4	46.6 242.6
26/62	C25B_100_100d	0.0 0.75 1.0	1.0 1.0 0.5	224	0.0 0.766 1.0	50.9 -16.2 -41.2	44.2 248.4	0.0 0.75 1.0	50.4 -15.5 -41.1	43.9 249.3	0.8 222	0.0 0.766 1.0	50.9 -16.2 -41.2	44.2 248.4
27/53	C38B_100_100d	0.0 0.625 1.0	1.0 1.0 0.5	232	0.0 0.633 1.0	46.8 -9.8 -40.9	42.1 256.4	0.0 0.625 1.0	46.5 -9.4 -40.8	41.9 256.9	0.4 231	0.0 0.633 1.0	46.8 -9.8 -40.9	42.1 256.4
28/44	C50B_100_100d	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	41.7 -1.2 -40.6	40.6 268.2	0.0 0.5 1.0	41.7 -1.2 -40.6	40.6 268.2	0.0 240	0.0 0.5 1.0	41.7 -1.2 -40.6	40.6 268.2
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 6.6 -40.2	40.8 279.3	0.0 0.375 1.0	37.3 6.1 -40.2	40.7 278.6	0.6 248	0.0 0.366 1.0	37.0 6.6 -40.2	40.8 279.3
30/26	C75B_100_100d	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.233 1.0	32.2 15.3 -40.3	43.1 290.8	0.0 0.25 1.0	32.8 14.3 -40.2	42.7 289.6	1.1 257	0.0 0.233 1.0	32.2 15.3 -40.3	43.1 290.8
31/17	C88B_100_100d	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.116 1.0	28.4 22.8 -40.3	46.3 299.5	0.0 0.125 1.0	28.6 22.4 -40.2	46.1 299.0	0.5 263	0.0 0.116 1.0	28.4 22.8 -40.3	46.3 299.5
32/8	B00M_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
33/89	B13M_100_100d	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	27.7 35.6 -36.7	51.1 314.1	0.125 0.0 1.0	27.9 36.0 -36.4	51.2 314.7	0.5 276	0.116 0.0 1.0	27.7 35.6 -36.7	51.1 314.1
34/170	B25M_100_100d	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1	0.25 0.0 1.0	28.8 41.9 -32.5	53.1 322.1	0.9 282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
35/251	B38M_100_100d	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	32.5 51.2 -26.5	57.7 332.6	0.375 0.0 1.0	32.7 51.8 -26.0	58.0 333.3	0.8 291	0.366 0.0 1.0	32.5 51.2 -26.5	57.7 332.6
36/332	B50M_100_100d	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5	0.0 300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
37/413	B63M_100_100d	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	38.3 65.8 -13.7	67.2 348.2	0.625 0.0 1.0	38.1 65.4 -14.0	66.9 347.9	0.5 308	0.633 0.0 1.0	38.3 65.8 -13.7	67.2 348.2
38/494	B75M_100_100d	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0	0.75 0.0 1.0	41.8 71.0 -9.2	71.6 352.5	0.8 317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0
39/575	B88M_100_100d	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	44.3 75.4 -4.7	75.6 356.3	0.875 0.0 1.0	44.2 75.2 -5.0	75.3 356.1	0.4 323	0.883 0.0 1.0	44.3 75.4 -4.7	75.6 356.3
40/656	M00R_100_100d	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	0.0 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
41/655	M13R_100_100d	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	45.9 78.3 3.8	78.4 2.8	1.0 0.0 0.875	45.9 78.2 4.1	78.3 363.0	0.2 336	1.0 0.0 0.883	45.9 78.3 3.8	78.4 2.8
42/654	M25R_100_100d	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9	1.0 0.0 0.75	45.9 77.1 8.6	77.6 366.4	0.6 342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
43/653	M38R_100_100d	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	46.0 75.7 14.4	77.1 10.8	1.0 0.0 0.625	46.0 75.6 14.8	77.0 371.1	0.4 351	1.0 0.0 0.633	46.0 75.7 14.4	77.1 10.8
44/652	M50R_100_100d	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.0									

Siehe ähnliche Dateien: <http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT> /PS
Technische Information: <http://www.ps.bam.de> oder <http://130.149.60.45/~farbmetrik>

TUB-Registrierung: 20150901-TG77/TG77L0NA.TXT /PS TUB-Material: Code=rha4ta
Anwendung für Messung von Offsetdruck-Ausgabe, Separation cmy6 (CMY0)

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	45.4 70.9 44.8 83.9 32.3	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	0.0 389		1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3
1/666	R25Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.0 53.4 54.8 76.5 45.7	1.0 0.25 0.0	53.6 51.9 55.5 76.0 46.8 1.7	42		1.0 0.233 0.0	53.0 53.4 54.8 76.5 45.7
2/684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	64.9 28.9 68.6 74.5 67.1	1.0 0.5 0.0	64.9 28.9 68.6 74.5 67.1	0.0 59		1.0 0.5 0.0	64.9 28.9 68.6 74.5 67.1
3/702	R75Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	78.6 4.3 84.7 84.8 87.0	1.0 0.75 0.0	77.9 5.4 83.8 84.0 86.2 1.6	77		1.0 0.766 0.0	78.6 4.3 84.7 84.8 87.0
4/720	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	87.8 -10.2 95.4 96.0 96.1	1.0 1.0 0.0	87.8 -10.2 95.4 96.0 96.1	0.0 89		1.0 1.0 0.0	87.8 -10.2 95.4 96.0 96.1
5/558	Y25G_100_100a	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	81.2 -17.0 84.3 86.0 101.4	0.75 1.0 0.0	80.7 -17.5 83.5 85.3 101.8 1.0	102	0.766 1.0 0.0	81.2 -17.0 84.3 86.0 101.4	
6/396	Y50G_100_100a	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	70.6 -29.7 66.5 72.8 114.0	0.5 1.0 0.0	70.6 -29.7 66.5 72.8 114.0	0.0 119	0.5 1.0 0.0	70.6 -29.7 66.5 72.8 114.0	
7/234	Y75G_100_100a	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	57.9 -48.3 45.8 66.5 136.5	0.25 1.0 0.0	58.4 -47.3 46.8 66.6 135.3 1.4	137	0.233 1.0 0.0	57.9 -48.3 45.8 66.5 136.5	
8/72	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	0.0 149	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	
9/72	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	0.0 149	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	
10/76	G25B_100_100a	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	52.9 -48.6 -8.0 49.3 189.3	0.0 1.0 0.5	52.9 -48.6 -8.0 49.3 189.3	0.0 180	0.0 1.0 0.5	52.9 -48.6 -8.0 49.3 189.3	
11/80	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5 48.7 238.4	0.0 1.0 1.0	56.8 -25.5 -41.5 48.7 238.4	0.0 210	0.0 1.0 1.0	56.8 -25.5 -41.5 48.7 238.4	
12/44	G75B_100_100a	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	41.7 -1.2 -40.6 40.6 268.2	0.0 0.5 1.0	41.7 -1.2 -40.6 40.6 268.2	0.0 240	0.0 0.5 1.0	41.7 -1.2 -40.6 40.6 268.2	
13/8	B00M_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	25.0 29.5 -40.4 50.0 306.2	0.0 0.0 1.0	25.0 29.5 -40.4 50.0 306.2	0.0 270	0.0 0.0 1.0	25.0 29.5 -40.4 50.0 306.2	
14/332	B25R_100_100a	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	35.6 58.6 -20.7 62.1 340.5	0.5 0.0 1.0	35.6 58.6 -20.7 62.1 340.5	0.0 300	0.5 0.0 1.0	35.6 58.6 -20.7 62.1 340.5	
15/656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2 79.3 359.8	1.0 0.0 1.0	46.1 79.3 -0.2 79.3 359.8	0.0 330	1.0 0.0 1.0	46.1 79.3 -0.2 79.3 359.8	
16/652	B75R_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	45.9 74.2 21.1 77.1 15.9	1.0 0.0 0.5	45.9 74.2 21.1 77.1 15.9	0.0 360	1.0 0.0 0.5	45.9 74.2 21.1 77.1 15.9	
17/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	0.0 389	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	
18/688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	70.5 35.4 22.4 41.9 32.3	1.0 0.5 0.5	68.0 29.9 28.7 41.5 43.8 8.7	389	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	
19/706	R50Y_100_050a	1.0 0.75 0.5	1.0 0.5 0.75	60	1.0 0.75 0.5	80.2 14.4 34.3 37.2 67.1	1.0 0.75 0.5	80.4 9.0 35.3 36.5 75.5 5.4	59	1.0 0.5 0.0	64.9 28.9 68.6 74.5 67.1	
20/724	Y00G_100_050a	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	91.7 -5.1 47.7 48.0 96.1	1.0 1.0 0.5	91.4 -7.7 42.5 43.2 100.3 5.8	89	1.0 1.0 0.0	87.8 -10.2 95.4 96.0 96.1	
21/562	Y50G_100_050a	0.75 1.0 0.5	1.0 0.5 0.75	120	0.75 1.0 0.5	83.1 -14.8 33.2 36.4 114.0	0.75 1.0 0.5	84.2 -14.1 31.5 34.5 114.0 2.1	119	0.5 1.0 0.0	70.6 -29.7 66.5 72.8 114.0	
22/400	G00B_100_050a	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	72.8 -32.5 14.8 35.7 155.5	0.5 1.0 0.5	73.9 -23.7 19.9 31.0 140.0 10.1	149	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	
23/404	G50B_100_050a	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	76.2 -12.7 -20.7 24.3 238.4	0.5 1.0 1.0	78.7 -11.6 -18.3 21.7 237.6 3.6	210	0.0 1.0 1.0	56.8 -25.5 -41.5 48.7 238.4	
24/368	B00R_100_050a	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	60.3 14.7 -20.2 25.0 306.2	0.5 0.5 1.0	57.9 18.3 -20.7 27.7 311.4 4.3	270	0.0 0.0 1.0	25.0 29.5 -40.4 50.0 306.2	
25/692	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	70.8 39.6 -0.1 39.6 359.8	1.0 0.5 1.0	70.7 35.2 -3.7 35.4 353.9 5.7	330	1.0 0.0 1.0	46.1 79.3 -0.2 79.3 359.8	
26/688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	70.5 35.4 22.4 41.9 32.3	1.0 0.5 0.5	68.0 29.9 28.7 41.5 43.8 8.7	389	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	
27/506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	52.7 35.4 22.4 41.9 32.3	0.75 0.25 0.25	50.4 39.4 31.9 50.7 38.9 10.5	389	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	
28/524	R50Y_075_050a	0.75 0.5 0.25	0.75 0.5 0.5	60	0.75 0.5 0.25	62.4 14.4 34.3 37.2 67.1	0.75 0.5 0.25	61.2 18.1 39.5 43.4 65.3 6.4	59	1.0 0.5 0.0	64.9 28.9 68.6 74.5 67.1	
29/542	Y00G_075_050a	0.75 0.75 0.25	0.75 0.5 0.5	90	0.75 0.75 0.25	73.9 -5.1 47.7 48.0 96.1	0.75 0.75 0.25	72.4 -1.4 48.0 48.0 91.7 3.9	89	1.0 1.0 0.0	87.8 -10.2 95.4 96.0 96.1	
30/380	Y50G_075_050a	0.5 0.75 0.25	0.75 0.5 0.5	120	0.5 0.75 0.25	65.3 -14.8 33.2 36.4 114.0	0.5 0.75 0.25	63.2 -12.6 35.5 37.7 109.6 3.7	119	0.5 1.0 0.0	70.6 -29.7 66.5 72.8 114.0	
31/218	G00B_075_050a	0.25 0.75 0.25	0.75 0.5 0.5	150	0.25 0.75 0.25	55.0 -32.5 14.8 35.7 155.5	0.25 0.75 0.25	53.0 -27.9 21.7 35.3 142.0 8.5	149	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	
32/222	G50B_075_050a	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.75	58.4 -12.7 -20.7 24.3 238.4	0.25 0.75 0.75	55.9 -14.3 -16.3 21.7 228.6 5.3	210	0.0 1.0 1.0	56.8 -25.5 -41.5 48.7 238.4	
33/186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	42.5 14.7 -20.2 25.0 306.2	0.25 0.25 0.75	37.5 18.9 -20.4 27.9 312.8 6.5	270	0.0 0.0 1.0	25.0 29.5 -40.4 50.0 306.2	
34/510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	53.0 39.6 -0.1 39.6 359.8	0.75 0.25 0.75	52.4 44.4 0.5 44.4 0.6 4.8	330	1.0 0.0 1.0	46.1 79.3 -0.2 79.3 359.8	
35/506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	52.7 35.4 22.4 41.9 32.3	0.75 0.25 0.25	50.4 39.4 31.9 50.7 38.9 10.5	389	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	
36/324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.0	34.9 35.4 22.4 41.9 32.3	0.5 0.0 0.0	34.8 44.7 22.4 50.0 26.6 9.2	389	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	
37/342	R50Y_050_050a	0.5 0.25 0.0	0.5 0.5 0.25	60	0.5 0.25 0.0	44.6 14.4 34.3 37.2 67.1	0.5 0.25 0.0	43.4 24.2 33.3 41.2 53.9 9.9	59	1.0 0.5 0.0	64.9 28.9 68.6 74.5 67.1	
38/360	Y00G_050_050a	0.5 0.5 0.0	0.5 0.5 0.25	90	0.5 0.5 0.0	56.1 -5.1 47.7 48.0 96.1	0.5 0.5 0.0	52.6 3.9 44.2 44.3 84.8 10.3	89	1.0 1.0 0.0	87.8 -10.2 95.4 96.0 96.1	
39/198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	47.4 -14.8 33.2 36.4 114.0	0.25 0.5 0.0	43.1 -14.1 28.4 31.7 116.4 6.5	119	0.5 1.0 0.0	70.6 -29.7 66.5 72.8 114.0	
40/36	G00B_050_050a	0.0 0.5 0.0	0.5 0.5 0.25	150	0.0 0.5 0.0	37.2 -32.5 14.8 35.7 155.5	0.0 0.5 0.0	37.3 -36.4 15.2 39.5 157.2 3.9	149	0.0 1.0 0.0	50.0 -65.0 29.6 71.4 155.5	
41/40	G50B_050_050a	0.0 0.5 0.5	0.5 0.5 0.25	210	0.0 0.5 0.5	40.5 -12.7 -20.7 24.3 238.4	0.0 0.5 0.5	39.1 -21.5 -13.3 25.3 211.8 11.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5 48.7 238.4	
42/4	B00R_050_050a	0.0 0.0 0.5	0.5 0.5 0.25	270	0.0 0.0 0.5	24.7 14.7 -20.2 25.0 306.2	0.0 0.0 0.5	24.3 11.6 -18.9 22.1 301.5 3.4	270	0.0 0.0 1.0	25.0 29.5 -40.4 50.0 306.2	
43/328	B50R_050_050a	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	35.2 39.6 -0.1 39.6 359.8	0.5 0.0 0.5	35.0 49.8 0.6 49.8 0.7 10.2	330	1.0 0.0 1.0	46.1 79.3 -0.2 79.3 359.8	
44/324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.0	34.9 35.4 22.4 41.9 32.3	0.5 0.0 0.0	34.8 44.7 22.4 50.0 26.6 9.2	389	1.0 0.0 0.0	45.4 70.9 44.8 83.9 32.3	
45/0	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	24.3 0.0 0.0 0.0 0.0	0.0 0.0 0.0	24.3 0.0 0.0 0.0 0.0	360	1.0 1.0 1.0	95.6 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	33.2 0.0 0.0 0.0 0.0	0.125 0.125 0.125	29.8 7.2 3.6 8.1 26.3 8.7	360	1.0 1.0 1.0	95.6 0.0 0.0 0.0 0.0	
47/182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	42.1 0.0 0.0 0.0 0.0	0.25 0.25 0.25	35.7 7.5 7.1 10.4 43.4 12.2	36			

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Technische Information: <http://www.ps.bam.de> oder <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	NW_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
1	B00R_012_012a	0.0	0.0	0.125	0.125	0.125	0.062	0.062	2.3	2.1	2.70	2.70
2	B00R_025_025a	0.0	0.0	0.25	0.25	0.25	0.125	0.125	4.8	-8.0	9.4	300.8
3	B00R_037_037a	0.0	0.0	0.375	0.375	0.375	0.187	0.187	6.9	-12.1	13.9	299.8
4	B00R_050_050a	0.0	0.0	0.5	0.5	0.5	0.25	0.25	11.6	-18.9	22.1	301.5
5	B00R_062_062a	0.0	0.0	0.625	0.625	0.625	0.312	0.312	15.8	-24.6	29.2	307.2
6	B00R_075_075a	0.0	0.0	0.75	0.75	0.75	0.375	0.375	20.7	-30.7	37.0	303.9
7	B00R_087_087a	0.0	0.0	0.875	0.875	0.875	0.437	0.437	24.8	-35.9	44.0	305.3
8	B00R_100_100a	0.0	0.0	1.0	1.0	1.0	0.5	0.5	29.5	-40.4	50.0	306.2
9	G00B_012_012a	0.0	0.125	0.0	0.125	0.125	0.062	0.062	2.9	8.7	160.0	0.8
10	G50B_012_012a	0.0	0.125	0.125	0.125	0.125	0.062	0.062	5.9	16.0	190.5	5.2
11	G75B_025_025a	0.0	0.125	0.25	0.25	0.25	0.125	0.125	6.7	-3.6	5.7	6.8
12	G84B_037_037a	0.0	0.125	0.375	0.375	0.375	0.187	0.187	10.2	-10.8	10.8	26.8
13	G88B_050_050a	0.0	0.125	0.5	0.5	0.5	0.25	0.25	13.4	-17.8	18.3	28.4
14	G90B_062_062a	0.0	0.125	0.625	0.625	0.625	0.312	0.312	17.8	-24.2	25.7	28.9
15	G92B_075_075a	0.0	0.125	0.75	0.75	0.75	0.375	0.375	21.1	-30.2	33.0	29.3
16	G93B_087_087a	0.0	0.125	0.875	0.875	0.875	0.437	0.437	26.2	-35.6	39.9	29.6
17	G94B_100_100a	0.0	0.125	1.0	1.0	1.0	0.5	0.5	28.6	-40.2	46.1	29.9
18	G00B_025_025a	0.0	0.25	0.0	0.25	0.25	0.125	0.125	3.0	15.7	2.2	149
19	G25B_025_025a	0.0	0.25	0.125	0.25	0.25	0.125	0.125	6.5	16.9	6.5	189
20	G50B_025_025a	0.0	0.25	0.25	0.25	0.25	0.125	0.125	10.6	21.0	10.0	1.0
21	G65B_037_037a	0.0	0.25	0.375	0.375	0.375	0.187	0.187	13.7	-11.0	8.3	13.7
22	G75B_050_050a	0.0	0.25	0.5	0.5	0.5	0.25	0.25	18.1	-15.7	16.7	25.0
23	G80B_062_062a	0.0	0.25	0.625	0.625	0.625	0.312	0.312	22.5	-22.5	22.5	26.8
24	G84B_075_075a	0.0	0.25	0.75	0.75	0.75	0.375	0.375	25.4	-29.1	29.5	27.4
25	G86B_087_087a	0.0	0.25	0.875	0.875	0.875	0.437	0.437	28.4	-34.9	36.3	28.5
26	G88B_100_100a	0.0	0.25	1.0	1.0	1.0	0.5	0.5	32.2	-40.2	42.7	28.6
27	G00B_037_037a	0.0	0.375	0.0	0.375	0.375	0.187	0.187	3.9	27.6	11.4	29.8
28	G15B_037_037a	0.0	0.375	0.125	0.375	0.375	0.187	0.187	6.6	26.4	16.4	16.4
29	G34B_037_037a	0.0	0.375	0.25	0.375	0.375	0.187	0.187	10.8	-22.1	-0.5	22.1
30	G50B_037_037a	0.0	0.375	0.375	0.375	0.375	0.187	0.187	12.6	-28.4	-6.6	19.6
31	G61B_050_050a	0.0	0.375	0.5	0.5	0.5	0.25	0.25	16.1	-14.1	-14.6	20.3
32	G69B_062_062a	0.0	0.375	0.625	0.625	0.625	0.312	0.312	20.0	-21.5	23.7	24.0
33	G75B_075_075a	0.0	0.375	0.75	0.75	0.75	0.375	0.375	24.0	-28.4	28.6	26.1
34	G79B_087_087a	0.0	0.375	0.875	0.875	0.875	0.437	0.437	28.4	-34.7	34.8	27.3
35	G81B_100_100a	0.0	0.375	1.0	1.0	1.0	0.5	0.5	32.2	-40.2	40.7	27.8
36	G00B_050_050a	0.0	0.5	0.0	0.5	0.5	0.25	0.25	3.9	15.7	2.2	149
37	G11B_050_050a	0.0	0.5	0.125	0.5	0.5	0.25	0.25	6.5	16.9	6.5	189
38	G25B_050_050a	0.0	0.5	0.25	0.5	0.5	0.25	0.25	10.6	21.0	10.0	1.0
39	G38B_050_050a	0.0	0.5	0.375	0.5	0.5	0.25	0.25	13.7	-11.0	8.3	13.7
40	G50B_050_050a	0.0	0.5	0.5	0.5	0.5	0.25	0.25	18.1	-15.7	16.7	25.0
41	G59B_062_062a	0.0	0.5	0.625	0.625	0.625	0.312	0.312	22.5	-22.5	22.5	26.8
42	G65B_075_075a	0.0	0.5	0.75	0.75	0.75	0.375	0.375	25.4	-29.1	29.5	27.4
43	G70B_087_087a	0.0	0.5	0.875	0.875	0.875	0.437	0.437	28.4	-34.9	36.3	28.5
44	G75B_100_100a	0.0	0.5	1.0	1.0	1.0	0.5	0.5	32.2	-40.2	40.7	27.8
45	G00B_062_062a	0.0	0.625	0.0	0.625	0.625	0.312	0.312	3.9	27.6	11.4	29.8
46	G09B_062_062a	0.0	0.625	0.125	0.625	0.625	0.312	0.312	6.6	26.4	16.4	16.4
47	G19B_062_062a	0.0	0.625	0.25	0.625	0.625	0.312	0.312	10.8	-22.1	-0.5	22.1
48	G30B_062_062a	0.0	0.625	0.375	0.625	0.625	0.312	0.312	12.6	-28.4	-6.6	19.6
49	G40B_062_062a	0.0	0.625	0.5	0.625	0.625	0.312	0.312	16.1	-14.1	-14.6	20.3
50	G50B_062_062a	0.0	0.625	0.625	0.625	0.625	0.312	0.312	20.0	-21.5	23.7	24.0
51	G57B_075_075a	0.0	0.625	0.75	0.75	0.75	0.375	0.375	24.0	-28.4	28.6	26.1
52	G63B_087_087a	0.0	0.625	0.875	0.875	0.875	0.437	0.437	28.4	-34.7	34.8	27.3
53	G68B_100_100a	0.0	0.625	1.0	1.0	1.0	0.5	0.5	32.2	-40.2	40.7	27.8
54	G00B_075_075a	0.0	0.75	0.0	0.75	0.75	0.375	0.375	3.9	27.6	11.4	29.8
55	G07B_075_075a	0.0	0.75	0.125	0.75	0.75	0.375	0.375	6.6	26.4	16.4	16.4
56	G15B_075_075a	0.0	0.75	0.25	0.75	0.75	0.375	0.375	10.8	-22.1	-0.5	22.1
57	G25B_075_075a	0.0	0.75	0.375	0.75	0.75	0.375	0.375	12.6	-28.4	-6.6	19.6
58	G34B_075_075a	0.0	0.75	0.5	0.75	0.75	0.375	0.375	16.1	-14.1	-14.6	20.3
59	G42B_075_075a	0.0	0.75	0.625	0.75	0.75	0.375	0.375	20.0	-21.5	23.7	24.0
60	G50B_075_075a	0.0	0.75	0.75	0.75	0.75	0.375	0.375	24.0	-28.4	28.6	26.1
61	G56B_087_087a	0.0	0.75	0.875	0.875	0.875	0.437	0.437	28.4	-34.7	34.8	27.3
62	G61B_100_100a	0.0	0.75	1.0	1.0	1.0	0.5	0.5	32.2	-40.2	40.7	27.8
63	G00B_087_087a	0.0	0.875	0.0	0.875	0.875	0.437	0.437	3.9	27.6	11.4	29.8
64	G06B_087_087a	0.0	0.875	0.125	0.875	0.875	0.437	0.437	6.6	26.4	16.4	16.4
65	G13B_087_087a	0.0	0.875	0.25	0.875	0.875	0.437	0.437	10.8	-22.1	-0.5	22.1
66	G20B_087_087a	0.0	0.875	0.375	0.875	0.875	0.437	0.437	12.6	-28.4	-6.6	19.6
67	G29B_087_087a	0.0	0.875	0.5	0.875	0.875	0.437	0.437	16.1	-14.1	-14.6	20.3
68	G36B_087_087a	0.0	0.875	0.625	0.875	0.875	0.437	0.437	20.0	-21.5	23.7	24.0
69	G43B_087_087a	0.0	0.875	0.75	0.875	0.875	0.437	0.437	24.0	-28.4	28.6	26.1
70	G50B_087_087a	0.0	0.875	0.875	0.875	0.875	0.437	0.437	28.4	-34.7	34.8	27.3
71	G55B_100_100a	0.0	0.875	1.0	1.0	1.0	0.5	0.5	32.2	-40.2	40.7	27.8
72	G00B_100_100a	0.0	1.0	0.0	1.0	1.0	0.5	0.5	3.9	27.6	11.4	29.8
73	G05B_100_100a	0.0	1.0	0.125	1.0	1.0	0.5	0.5	6.6	26.4	16.4	16.4
74	G11B_100_100a	0.0	1.0	0.25	1.0	1.0	0.5	0.5	10.6	21.0	10.0	1.0
75	G18B_100_100a	0.0	1.0	0.375	1.0	1.0	0.5	0.5	13.7	-11.0	8.3	13.7
76	G25B_100_100a	0.0	1.0	0.5	1.0	1.0	0.5	0.5	18.1	-15.7	16.7	25.0
77	G31B_100_100a	0.0	1.0	0.625	1.0	1.0	0.5	0.5	22.5	-22.5	22.5	26.8
78	G38B_100_100a	0.0	1.0	0.75	1.0	1.0	0.5	0.5	25.4	-29.1	29.5	27.4
79	G44B_100_100a	0.0	1.0	0.875	1.0	1.0	0.5	0.5	28.4	-34.9	36.3	28.5
80	G50B_100_100a	0.0	1.0	1.0	1.0	1.0	0.5	0.5	32.2	-40.2	40.7	27.8

delta E* = 4.2

0-003831-F0

TG770-7N, Seite 9/22-F

Prüfvorlage TG77; ME16(ISO 9241-306), 3(ISO/IEC 15775)
Farben und Farbabstände, ΔE^* , 3D=0, de=0, *cmlyk*

Eingabe: *rgb/cmyk* -> *rgb*_d
Ausgabe: Transfer nach *cmlyk*_d

0-003831-F0

TUB-Registrierung: 20150901-TG77/TG77L0NA.TXT /PS
Anwendung für Messung von Offsetdruck-Ausgabe, Separation *cmyn6* (CMY0)
TUB-Material: Code=rha4ta

Siehe ähnliche Dateien: <http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT> /PS
Technische Information: <http://www.ps.bam.de> oder <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_Md	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	27.0 8.8 5.6	10.4 32.3	0.125 0.0 0.0	26.6 14.6 4.2	15.2 16.1 5.9	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	
82	B50R_012_012a	0.125 0.0 0.125	0.125 0.125 0.062	330	0.125 0.0 0.125	27.0 9.9 0.0	9.9 359.8	0.125 0.0 0.125	26.7 15.8 0.3	15.8 1.1 5.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	
83	B25R_025_025a	0.125 0.0 0.25	0.25 0.25 0.125	300	0.125 0.0 0.25	27.1 14.6 -5.1	15.5 340.5	0.125 0.0 0.25	26.9 17.8 -4.5	18.4 345.8 3.2	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5	
84	B15R_037_037a	0.125 0.0 0.375	0.375 0.375 0.187	289	0.118 0.0 0.375	26.8 17.7 -11.0	20.9 328.1	0.125 0.0 0.375	26.6 19.3 -9.3	21.5 334.2 2.3	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1	
85	B11R_050_050a	0.125 0.0 0.5	0.5 0.5 0.25	284	0.116 0.0 0.5	26.5 20.6 -16.5	26.4 321.1	0.125 0.0 0.5	27.0 21.7 -15.4	26.6 324.6 1.7	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1	
86	B09R_062_062a	0.125 0.0 0.625	0.625 0.625 0.312	281	0.114 0.0 0.625	26.8 24.2 -21.7	32.5 318.2	0.125 0.0 0.625	27.1 25.2 -21.3	33.1 319.7 1.0	279	0.183 0.0 1.0	28.3 38.8 -34.7	52.1 318.2	
87	B07R_075_075a	0.125 0.0 0.75	0.75 0.75 0.375	279	0.112 0.0 0.75	27.1 27.9 -26.8	38.7 316.2	0.125 0.0 0.75	27.4 29.1 -26.9	39.7 317.2 1.2	278	0.15 0.0 1.0	28.1 37.2 -35.7	51.6 316.2	
88	B06R_087_087a	0.125 0.0 0.875	0.875 0.875 0.437	278	0.116 0.0 0.875	27.5 31.9 -31.6	44.9 315.2	0.125 0.0 0.875	27.4 33.0 -32.0	46.0 315.8 1.1	277	0.133 0.0 1.0	27.9 36.4 -36.2	51.3 315.2	
89	B05R_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	27.7 35.6 -36.7	51.1 314.1	0.125 0.0 1.0	27.9 36.0 -36.4	51.2 314.7 0.5	276	0.116 0.0 1.0	27.7 35.6 -36.7	51.1 314.1	
90	Y00G_012_012a	0.125 0.125 0.0	0.125 0.125 0.062	90	0.125 0.125 0.0	32.3 -1.2 11.9	12.0 96.1	0.125 0.125 0.0	29.6 5.9 7.7	9.7 52.8 8.6	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0	96.1
91	NW_012a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	33.2 0.0 0.0	0.0 0.0	0.0 0.125 0.125 0.125	29.8 7.2 3.6	8.1 26.3 8.7	360	1.0 1.0 1.0	95.6 0.0 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	33.3 3.6 -5.0	6.2 306.2	0.125 0.125 0.25	30.0 8.9 -1.7	9.1 349.1 7.0	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
93	B00R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.124 0.375	33.4 7.3 -10.1	12.5 306.2	0.125 0.125 0.375	30.4 11.8 -7.5	14.0 327.5 5.9	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
94	B00R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.124 0.5	33.5 11.0 -15.1	18.7 306.2	0.125 0.125 0.5	30.5 14.5 -14.1	20.3 315.8 4.7	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
95	B00R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	33.6 14.7 -20.2	25.0 306.2	0.125 0.125 0.625	30.9 17.9 -20.2	27.0 311.4 4.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
96	B00R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	33.7 18.4 -25.2	31.3 306.2	0.125 0.125 0.75	31.5 21.1 -26.2	33.7 308.7 3.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
97	B00R_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	33.8 22.1 -30.3	37.5 306.2	0.125 0.125 0.875	31.5 25.0 -31.5	40.2 308.4 3.8	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
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 25.8 -35.3	43.8 306.2	0.125 0.125 1.0	32.0 28.2 -36.3	46.0 307.8 3.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	
99	Y50G_025_025a	0.125 0.25 0.0	0.25 0.25 0.125	120	0.125 0.25 0.0	35.9 -7.4 16.6	18.2 114.0	0.125 0.25 0.0	33.7 -4.5 12.9	13.6 109.2 5.2	119	0.5 1.0 0.0	70.6 -29.7	66.5 72.8	114.0
100	G00B_025_012a	0.125 0.25 0.125	0.25 0.125 0.187	150	0.124 0.25 0.124	36.4 -8.1 3.7	8.9 155.5	0.125 0.25 0.125	33.9 -3.6 8.3	9.1 113.6 6.9	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4	155.5
101	G50B_025_012a	0.125 0.25 0.25	0.25 0.125 0.187	210	0.124 0.25 0.25	37.3 -3.1 -5.1	6.0 238.4	0.125 0.25 0.25	34.4 -1.1 1.6	2.0 124.6 7.7	210	0.0 1.0 1.0	56.8 -25.5	-41.5 48.7	238.4
102	G75B_037_025a	0.125 0.25 0.375	0.375 0.25 0.25	240	0.124 0.25 0.375	37.6 -0.3 -10.1	10.1 268.2	0.125 0.25 0.375	34.7 1.3 -4.5	4.7 286.1 6.5	240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6	268.2
103	G84B_050_037a	0.125 0.25 0.5	0.5 0.375 0.312	251	0.124 0.243 0.5	37.3 3.7 -15.1	15.6 283.7	0.125 0.25 0.5	35.0 4.5 -11.8	12.7 291.0 4.1	251	0.0 0.316 1.0	35.2 9.9	-40.4 41.6	283.7
104	G88B_062_050a	0.125 0.25 0.625	0.625 0.5 0.375	256	0.125 0.241 0.625	37.2 7.6 -20.1	21.5 290.8	0.125 0.25 0.625	35.2 8.5 -18.0	20.0 295.3 2.9	257	0.0 0.233 1.0	32.2 15.3	-40.3 43.1	290.8
105	G90B_075_062a	0.125 0.25 0.75	0.75 0.625 0.437	259	0.125 0.239 0.75	37.1 11.6 -25.2	27.8 294.6	0.125 0.25 0.75	35.7 12.5 -24.8	27.8 296.7 1.7	260	0.0 0.183 1.0	30.6 18.5	-40.4 44.5	294.6
106	G92B_087_075a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.237 0.875	37.1 15.5 -30.3	34.0 297.1	0.125 0.25 0.875	36.1 16.4 -30.6	34.8 298.2 1.3	262	0.0 0.15 1.0	29.5 20.7	-40.4 45.4	297.1
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.241 1.0	37.2 19.1 -35.2	40.1 298.4	0.125 0.25 1.0	36.4 19.7 -35.8	40.8 298.8 1.1	262	0.0 0.133 1.0	28.9 21.8	-40.3 45.8	298.4
108	Y68G_037_037a	0.125 0.375 0.0	0.375 0.375 0.187	131	0.118 0.375 0.0	38.6 -15.5 19.9	25.3 127.8	0.125 0.375 0.0	37.4 -15.0 17.0	22.7 131.3 3.1	131	0.316 1.0 0.0	62.3 -41.4	53.2 67.5	127.8
109	G00B_037_025a	0.125 0.375 0.125	0.375 0.25 0.25	150	0.124 0.375 0.124	39.7 -16.2 7.4	17.8 155.5	0.125 0.375 0.125	37.6 -12.8 11.7	17.3 137.3 5.9	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4	155.5
110	G25B_037_025a	0.125 0.375 0.25	0.375 0.25 0.25	180	0.124 0.375 0.25	40.4 -12.1 -2.0	12.3 189.3	0.125 0.375 0.25	38.4 -10.8 5.2	12.0 154.3 7.6	180	0.0 1.0 0.5	52.9 -48.6	-8.0 49.3	189.3
111	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	41.3 -6.3 -10.3	12.1 238.4	0.125 0.375 0.375	38.8 -7.8 -2.3	8.2 196.2 8.6	210	0.0 1.0 1.0	56.8 -25.5	-41.5 48.7	238.4
112	G65B_050_037a	0.125 0.375 0.5	0.5 0.375 0.312	229	0.124 0.381 0.5	42.2 -4.6 -15.4	16.0 253.3	0.125 0.375 0.5	39.7 -5.2 -9.5	10.8 241.1 6.4	228	0.0 0.683 1.0	48.3 -12.2	-41.1 42.9	253.3
113	G75B_062_050a	0.125 0.375 0.625	0.625 0.5 0.375	240	0.125 0.375 0.625	41.9 -0.6 -20.3	20.3 268.2	0.125 0.375 0.625	39.7 -0.9 -16.6	16.6 266.8 4.2	240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6	268.2
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.364 0.75	41.5 3.5 -25.1	25.4 277.9	0.125 0.375 0.75	39.8 4.0 -24.0	24.4 279.5 2.0	247	0.0 0.383 1.0	37.6 5.6	-40.3 40.7	277.9
115	G84B_087_075a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.362 0.875	41.4 7.4 -30.3	31.2 283.7	0.125 0.375 0.875	40.3 8.1 -30.2	31.3 285.1 1.3	251	0.0 0.316 1.0	35.2 9.9	-40.4 41.6	283.7
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 11.5 -35.2	37.1 288.1	0.125 0.375 1.0	40.4 12.6 -35.8	37.9 289.4 1.4	255	0.0 0.266 1.0	33.4 13.2	-40.3 42.4	288.1
117	Y76G_050_050a	0.125 0.5 0.0	0.5 0.5 0.25	136	0.116 0.5 0.0	41.1 -24.1 22.9	33.2 136.5	0.125 0.5 0.0	41.0 -23.7 21.5	32.0 137.7 1.4	137	0.233 1.0 0.0	57.9 -48.3	45.8 66.5	136.5
118	G00B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.124	42.9 -24.3 11.1	26.7 155.5	0.125 0.5 0.125	41.5 -19.6 15.4	26.6 144.4 5.3	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4	155.5
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.243	43.5 -21.3 2.7	21.4 172.5	0.125 0.5 0.25	42.1 -21.2 8.0	20.8 157.3 5.8	168	0.0 1.0 0.316	51.6 -56.8	7.4 57.3	172.5
120	G34B_050_037a	0.125 0.5 0.375	0.5 0.3												

Siehe ähnliche Dateien: <http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT> /PS
Technische Information: <http://www.ps.bam.de> oder <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		
162	R00Y_025_025a	0.25 0.0 0.0	0.25 0.25 0.125	390	0.25 0.0 0.0	29.6 17.7 11.2	20.9 32.3	0.25 0.0 0.0	28.1 24.0 7.8	25.2 18.0 7.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
163	R00Y_025_025a	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.125	29.7 18.5 5.2	19.2 15.9	0.25 0.0 0.125	28.1 25.5 4.4	25.9 9.7 7.2	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
164	B50R_025_025a	0.25 0.0 0.25	0.25 0.25 0.125	330	0.25 0.0 0.25	29.8 19.8 0.0	19.8 359.8	0.25 0.0 0.25	28.3 27.3 -0.1	27.3 359.7 7.6	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
165	B34R_037_037a	0.25 0.0 0.375	0.375 0.375 0.187	311	0.256 0.0 0.375	30.1 25.5 -4.4	25.9 350.0	0.25 0.0 0.375	28.5 29.3 -4.3	29.6 351.6 4.1	311	1.0 0.683 0.0	39.8 68.1 -11.9	69.1 350.0
166	B25R_050_050a	0.25 0.0 0.5	0.5 0.5 0.25	300	0.25 0.0 0.5	29.9 29.3 -10.3	31.0 340.5	0.25 0.0 0.5	28.5 30.6 -10.4	32.3 341.1 1.9	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
167	B19R_062_062a	0.25 0.0 0.625	0.625 0.625 0.312	293	0.239 0.0 0.625	29.7 32.7 -16.0	36.4 333.8	0.25 0.0 0.625	28.5 32.6 -17.0	36.8 332.3 1.5	292	0.383 0.0 1.0	32.9 52.3 -25.7	58.3 333.8
168	B15R_075_075a	0.25 0.0 0.75	0.75 0.75 0.375	289	0.237 0.0 0.75	29.3 35.5 -22.0	41.8 328.1	0.25 0.0 0.75	28.7 36.0 -23.1	42.8 327.3 1.3	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
169	B13R_087_087a	0.25 0.0 0.875	0.875 0.875 0.437	286	0.233 0.0 0.875	28.7 37.9 -27.8	47.0 323.6	0.25 0.0 0.875	28.6 39.2 -28.1	48.2 324.3 1.3	284	0.266 0.0 1.0	29.4 43.3 -31.8	53.8 323.6
170	B11R_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1	0.25 0.0 1.0	28.8 41.9 -32.5	53.1 322.1 0.9	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
171	R50Y_025_025a	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.125 0.0	34.5 7.2 17.1	18.6 67.1	0.25 0.125 0.0	31.0 16.0 11.8	19.9 36.4 10.8	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
172	R00Y_025_012a	0.25 0.125 0.125	0.25 0.125 0.187	390	0.25 0.124 0.124	35.9 8.8 5.6	10.4 32.3	0.25 0.125 0.125	31.0 16.8 8.0	18.6 25.3 9.7	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
173	B50R_025_012a	0.25 0.125 0.25	0.25 0.125 0.187	330	0.25 0.124 0.25	36.0 9.9 0.0	9.9 359.8	0.25 0.125 0.25	31.5 18.6 3.2	18.9 9.7 10.3	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
174	B25R_037_025a	0.25 0.125 0.375	0.375 0.25 0.25	300	0.25 0.124 0.375	36.0 14.6 -5.1	15.5 340.5	0.25 0.125 0.375	31.7 20.5 -2.2	20.6 353.7 7.9	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
175	B15R_050_037a	0.25 0.125 0.5	0.5 0.375 0.312	289	0.243 0.124 0.5	35.7 17.7 -11.0	20.9 328.1	0.25 0.125 0.5	31.9 22.3 -9.9	24.4 359.6 6.0	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
176	B11R_062_050a	0.25 0.125 0.625	0.625 0.5 0.375	284	0.241 0.125 0.625	35.4 20.6 -16.5	26.4 321.1	0.25 0.125 0.625	32.1 24.7 -16.1	29.5 326.9 5.3	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
177	B09R_075_062a	0.25 0.125 0.75	0.75 0.625 0.437	281	0.239 0.125 0.75	35.7 24.2 -21.7	32.5 318.2	0.25 0.125 0.75	32.4 28.2 -22.2	35.9 321.8 5.1	279	0.183 0.0 1.0	28.3 38.8 -34.7	52.1 318.2
178	B07R_087_062a	0.25 0.125 0.875	0.875 0.75 0.5	279	0.237 0.125 0.875	36.0 27.9 -26.8	38.7 316.2	0.25 0.125 0.875	32.8 30.9 -27.8	41.6 318.0 4.5	278	0.15 0.0 1.0	28.1 37.2 -35.7	51.6 316.2
179	B06R_100_087a	0.25 0.125 1.0	1.0 0.875 0.562	278	0.241 0.125 1.0	36.4 31.9 -31.6	44.9 315.2	0.25 0.125 1.0	33.1 33.9 -32.5	47.0 316.2 3.9	277	0.133 0.0 1.0	27.9 36.4 -36.2	51.3 315.2
180	Y00G_025_025a	0.25 0.25 0.0	0.25 0.25 0.125	90	0.25 0.25 0.0	40.2 -2.5 23.8	24.0 96.1	0.25 0.25 0.0	35.1 5.0 17.5	18.2 74.0 11.1	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
181	Y00G_025_012a	0.25 0.25 0.125	0.25 0.125 0.187	90	0.25 0.25 0.124	41.2 -1.2 11.9	12.0 96.1	0.25 0.25 0.125	35.4 5.7 13.1	14.3 66.3 9.1	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	42.1 0.0 0.0	0.0 0.0	0.25 0.25 0.25	35.7 7.5 7.1	10.4 43.4 12.2	360	1.0 1.0 1.0	95.6 0.0 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	42.2 3.6 -5.0	6.2 306.2	0.25 0.25 0.375	36.3 9.3 0.7	9.4 4.5 10.0	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
184	B00R_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.249 0.5	42.3 7.3 -10.1	12.5 306.2	0.25 0.25 0.5	36.4 12.3 -6.8	14.1 330.9 8.3	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
185	B00R_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	42.4 11.0 -15.1	18.7 306.2	0.25 0.25 0.625	37.0 15.5 -14.0	20.9 318.0 7.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	42.5 14.7 -20.2	25.0 306.2	0.25 0.25 0.75	37.5 18.9 -20.4	27.9 312.8 6.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
187	B00R_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.6 18.4 -25.2	31.3 306.2	0.25 0.25 0.875	38.0 22.0 -26.5	34.4 309.7 5.9	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
188	B00R_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	42.7 22.1 -30.3	37.5 306.2	0.25 0.25 1.0	38.2 25.3 -31.6	40.5 308.7 5.6	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
189	Y31G_037_037a	0.25 0.375 0.0	0.375 0.375 0.187	109	0.256 0.375 0.0	44.4 -7.9 29.8	30.8 104.9	0.25 0.375 0.0	39.4 -4.2 23.2	23.6 100.4 9.0	108	0.683 1.0 0.0	77.8 -21.1 79.4	82.2 104.9
190	Y50G_037_025a	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	44.8 -7.4 16.6	18.2 114.0	0.25 0.375 0.125	39.4 -3.5 17.7	18.1 101.1 6.7	119	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0
191	G00B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	45.4 -8.1 3.7	8.9 155.5	0.25 0.375 0.25	40.0 -1.6 10.9	11.0 98.5 11.0	149	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5
192	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	46.2 -3.1 -5.1	6.0 238.4	0.25 0.375 0.375	40.5 0.2 4.0	4.0 87.0 11.3	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4
193	G75B_050_025a	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.375 0.5	46.5 -0.3 -10.1	10.1 268.2	0.25 0.375 0.5	40.8 3.5 -4.5	5.8 307.8 8.8	240	0.0 0.5 1.0	41.7 -1.2 -40.6	40.6 268.2
194	G84B_062_037a	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.368 0.625	46.2 3.7 -15.1	15.6 283.7	0.25 0.375 0.625	40.7 7.2 -11.9	13.9 301.0 7.2	251	0.0 0.316 1.0	35.2 9.9 -40.4	41.6 283.7
195	G88B_075_050a	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.366 0.75	46.1 7.6 -20.1	21.5 290.8	0.25 0.375 0.75	41.7 10.7 -19.4	22.2 299.0 5.4	257	0.0 0.233 1.0	32.2 15.3 -40.3	43.1 290.8
196	G90B_087_062a	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.364 0.875	46.0 11.6 -25.2	27.8 294.6	0.25 0.375 0.875	41.7 14.8 -26.1	30.1 299.6 5.5	260	0.0 0.183 1.0	30.6 18.5 -40.4	44.5 294.6
197	G92B_100_075a	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.362 1.0	46.0 15.5 -30.3	34.0 297.1	0.25 0.375 1.0	42.1 18.3 -31.8	36.7 299.9 5.0	262	0.0 0.15 1.0	29.5 20.7 -40.4	45.4 297.1
198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	47.4 -14.8 33.2	36.4 114.0	0.25 0.5 0.0	43.1 -14.1 28.4	31.7 116.4 6.5	119	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0
199	Y68G_050_037a	0.25 0.5 0.125	0.5 0.375 0.312	131	0.243 0.5 0.124	47.5 -15.5 19.9	25.3 127.8	0.25 0.5 0.125	43.2 -13.5 21.7	25.6 121.9 4.9	131	0.316 1.0 0.0	62.3 -41.4 53.2	67.5 127.8
200	G00B_050_025a	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.249	48.6 -16.2 7.4	17.8 155.5	0.25 0.5 0.25	44.2 -10.9 14.5	18.2 126.7 9.9	149	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5
201	G25B_050_025a	0.25 0.5 0.375	0.5 0.25 0.375	180	0.249 0.5 0.375	49.3 -12.1 -2.0	12.3 189.3	0.25 0.5 0.375	44.9 -8.5 6.8	10.9 141.0 10.5	180	0.0 1.0 0.5	52.9 -48.6 -8.0	49.3 189.3
202	G50B_050_025a	0.25 0.5 0												

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Technische Information: <http://www.ps.bam.de> oder <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_Md	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	32.2 26.6 16.8	31.4 32.3	0.375 0.0 0.0	31.7 36.2 17.7	40.3 26.1 9.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
244		R18Y_037_037a	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.118	32.3 27.2 11.7	29.6 23.2	0.375 0.0 0.125	31.6 36.7 13.2	39.0 19.8 9.6	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
245		B65R_037_037a	0.375 0.0 0.25	0.375 0.375 0.187	349	0.375 0.0 0.256	32.4 28.6 4.4	29.0 8.9	0.375 0.0 0.25	31.7 38.5 8.1	39.3 11.9 10.5	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
246		B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	32.5 29.7 0.0	29.7 359.8	0.375 0.0 0.375	31.7 39.8 3.0	39.9 4.3 10.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
247		B38R_050_050a	0.375 0.0 0.5	0.5 0.5 0.25	316	0.383 0.0 0.5	33.2 35.8 -4.3	36.0 353.0	0.375 0.0 0.5	32.2 42.9 -3.3	43.0 355.5 7.3	317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0
248		B30R_062_062a	0.375 0.0 0.625	0.625 0.625 0.312	307	0.385 0.0 0.625	32.8 40.6 -9.0	41.6 347.4	0.375 0.0 0.625	32.4 45.1 -9.5	46.1 348.0 4.5	307	0.616 0.0 1.0	37.9 65.0 -14.5	66.6 347.4
249		B25R_075_075a	0.375 0.0 0.75	0.75 0.75 0.375	300	0.375 0.0 0.75	32.7 43.9 -15.5	46.6 340.5	0.375 0.0 0.75	32.5 47.1 -15.8	49.6 341.4 3.1	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
250		B20R_087_087a	0.375 0.0 0.875	0.875 0.875 0.437	295	0.364 0.0 0.875	32.5 47.4 -21.3	51.9 335.7	0.375 0.0 0.875	32.6 49.3 -21.4	53.8 336.5 1.9	294	0.416 0.0 1.0	33.7 54.1 -24.4	59.4 335.7
251		B18R_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	32.5 51.2 -26.5	57.7 332.6	0.375 0.0 1.0	32.7 51.8 -26.0	58.0 333.3 0.8	291	0.366 0.0 1.0	32.5 51.2 -26.5	57.7 332.6
252		R31Y_037_037a	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.118 0.0	36.4 17.1 22.2	28.1 52.2	0.375 0.125 0.0	34.8 28.0 21.3	35.2 37.3 10.9	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
253		R00Y_037_025a	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.124	38.5 17.7 11.2	20.9 32.3	0.375 0.125 0.125	35.1 28.3 16.7	32.9 30.6 12.4	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
254		R00Y_037_025a	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.25	38.6 18.5 5.2	19.2 15.9	0.375 0.125 0.25	35.3 29.6 10.7	31.5 19.8 12.7	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 52.3
255		B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.124 0.375	38.7 19.8 0.0	19.8 359.8	0.375 0.125 0.375	35.5 31.2 5.0	31.6 9.2 12.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
256		B34R_050_037a	0.375 0.125 0.5	0.5 0.375 0.312	311	0.381 0.124 0.5	39.0 25.5 -4.4	25.9 350.0	0.375 0.125 0.5	36.2 33.7 -2.3	33.7 355.9 8.9	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
257		B25R_062_050a	0.375 0.125 0.625	0.625 0.5 0.375	300	0.375 0.125 0.625	38.8 29.3 -10.3	31.0 340.5	0.375 0.125 0.625	36.2 35.2 -9.0	36.3 345.6 6.6	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
258		B19R_075_062a	0.375 0.125 0.75	0.75 0.625 0.437	293	0.364 0.125 0.75	38.6 32.7 -16.0	36.4 333.8	0.375 0.125 0.75	36.6 37.1 -15.7	40.3 337.0 4.8	292	0.383 0.0 1.0	32.9 52.3 -25.7	58.3 333.8
259		B15R_087_075a	0.375 0.125 0.875	0.875 0.75 0.5	289	0.362 0.125 0.875	38.2 35.5 -22.0	41.8 328.1	0.375 0.125 0.875	36.9 39.8 -21.4	45.2 331.6 4.5	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
260		B13R_100_087a	0.375 0.125 1.0	1.0 0.875 0.562	286	0.358 0.125 1.0	37.6 37.9 -27.8	47.0 323.6	0.375 0.125 1.0	36.8 42.2 -26.6	49.9 327.7 4.5	284	0.266 0.0 1.0	29.4 43.3 -31.8	53.8 323.6
261		R68Y_037_037a	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.256 0.0	43.2 4.1 30.1	30.4 82.1	0.375 0.25 0.0	39.9 16.0 27.6	31.9 59.7 12.6	71	1.0 0.683 0.0	74.8 11.0 80.4	81.1 82.1
262		R50Y_037_025a	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.25 0.124	43.4 7.2 17.1	18.6 67.1	0.375 0.25 0.125	39.9 17.1 21.7	27.7 51.6 11.5	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
263		R00Y_037_012a	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.249	44.8 8.8 5.6	10.4 32.3	0.375 0.25 0.25	40.0 18.4 15.1	23.9 39.3 14.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
264		B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.249 0.375	44.9 9.9 0.0	9.9 359.8	0.375 0.25 0.375	40.7 19.7 8.1	21.3 22.2 13.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
265		B25R_050_025a	0.375 0.25 0.5	0.5 0.25 0.375	300	0.375 0.249 0.5	44.9 14.6 -5.1	15.5 340.5	0.375 0.25 0.5	41.2 22.1 -0.1	22.1 359.5 9.7	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
266		B15R_062_037a	0.375 0.25 0.625	0.625 0.375 0.437	289	0.368 0.25 0.625	44.6 17.7 -11.0	20.9 328.1	0.375 0.25 0.625	41.6 23.9 -7.1	25.0 343.2 7.9	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
267		B11R_075_050a	0.375 0.25 0.75	0.75 0.5 0.5	284	0.366 0.25 0.75	44.3 20.6 -16.5	26.4 321.1	0.375 0.25 0.75	42.1 26.2 -14.0	29.7 331.7 6.5	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
268		B09R_087_062a	0.375 0.25 0.875	0.875 0.625 0.562	281	0.364 0.25 0.875	44.6 24.2 -17.7	32.5 318.2	0.375 0.25 0.875	42.9 28.9 -20.3	35.3 324.8 5.1	279	0.183 0.0 1.0	28.3 38.8 -34.7	52.1 318.2
269		B07R_100_075a	0.375 0.25 1.0	1.0 0.75 0.625	279	0.362 0.25 1.0	44.9 27.9 -26.8	38.7 316.2	0.375 0.25 1.0	43.1 31.3 -26.0	40.7 320.3 3.9	278	0.15 0.0 1.0	28.1 37.2 -35.7	51.6 316.2
270		Y00G_037_037a	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.375 0.0	48.1 -3.8 35.8	36.0 96.1	0.375 0.375 0.0	44.1 6.7 33.2	33.8 78.5 11.5	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
271		Y00G_037_025a	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.375 0.124	49.1 -2.5 23.8	24.0 96.1	0.375 0.375 0.125	44.5 7.0 26.3	27.2 75.0 10.9	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
272		Y00G_037_012a	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.375 0.249	50.1 -1.2 11.9	12.0 96.1	0.375 0.375 0.25	44.7 8.5 18.5	20.4 65.3 12.9	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
273		NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	51.0 0.0 0.0	0.0 0.0	0.375 0.375 0.375	45.3 10.0 11.0	14.9 47.8 16.0	360	1.0 1.0 1.0	95.6 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	51.1 3.6 -5.0	6.2 306.2	0.375 0.375 0.5	46.1 12.2 2.1	12.3 10.0 12.2	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
275		B00R_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	51.2 7.3 -10.1	12.5 306.2	0.375 0.375 0.625	46.7 14.8 -5.3	15.7 340.2 9.9	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
276		B00R_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	51.3 11.0 -15.1	18.7 306.2	0.375 0.375 0.75	47.4 17.2 -12.5	21.3 323.8 7.7	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
277		B00R_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	51.4 14.7 -20.2	25.0 306.2	0.375 0.375 0.875	48.1 19.9 -19.3	27.7 315.9 6.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
278		B00R_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	51.5 18.4 -25.2	31.3 306.2	0.375 0.375 1.0	48.4 23.0 -25.3	34.2 312.3 5.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
279		Y23G_050_050a	0.375 0.5 0.0	0.5 0.5 0.25	104	0.383 0.5 0.0	52.8 -8.5 42.1	43.0 101.4	0.375 0.5 0.0	49.1 -2.0 38.9	38.9 9.9 8.1	102	0.766 1.0 0.0	81.2 -17.0	84.3 86.0 101.4
280		Y31G_050_037a	0.375 0.5 0.125	0.5 0.375 0.312	109	0.381 0.5 0.124	53.3 -7.9 29.8	30.8 104.9	0.375 0.5 0.125	49.5 -1.7 31.0	31.0 93.2 7.3	108	0.683 1.0 0.0	77.8 -21.1	79.4 82.2 104.9
281		Y50G_050_025a	0.375 0.5 0.25	0.5 0.25 0.375	120	0.375 0.5 0.249	53.7 -7.4 16.6	18.2 114.0	0.375 0.5 0.25	49.7 -0.9 22.3	22.3 92.5 9.4	119	0.5 1.0 0.0	70.6 -29.7	66.5 72.8 114.0
282		G00B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437											

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Technische Information: <http://www.ps.bam.de> oder <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	34.9	35.4	22.4	41.9	32.3
325	R26Y_050_050a	0.5	0.0	0.125	0.5	0.5	0.0	0.116	35.0	36.0	17.6	40.1
326	R00Y_050_050a	0.5	0.0	0.25	0.5	0.5	0.0	0.25	35.1	37.1	10.5	38.5
327	B61R_050_050a	0.5	0.0	0.375	0.5	0.5	0.0	0.383	35.1	38.6	4.0	38.8
328	B50R_050_050a	0.5	0.0	0.5	0.5	0.5	0.0	0.5	35.2	39.6	-0.1	39.6
329	B40R_062_062a	0.5	0.0	0.625	0.625	0.625	0.312	319	0.51	0.0	0.625	36.0
330	B34R_075_075a	0.5	0.0	0.75	0.75	0.75	0.375	311	0.512	0.0	0.75	35.9
331	B29R_087_087a	0.5	0.0	0.875	0.875	0.875	0.437	305	0.51	0.0	0.875	35.6
332	B25R_100_100a	0.5	0.0	1.0	1.0	1.0	0.5	300	0.5	0.0	1.0	35.6
333	R23Y_050_050a	0.5	0.125	0.0	0.5	0.5	0.25	44	0.5	0.116	0.0	38.7
334	R00Y_050_037a	0.5	0.125	0.125	0.5	0.375	0.312	390	0.5	0.124	0.124	41.1
335	R18Y_050_037a	0.5	0.125	0.25	0.5	0.375	0.312	371	0.5	0.124	0.243	41.2
336	B65R_050_037a	0.5	0.125	0.375	0.5	0.375	0.312	349	0.5	0.124	0.381	41.3
337	B50R_050_037a	0.5	0.125	0.5	0.5	0.375	0.312	330	0.5	0.124	0.5	41.4
338	B38R_062_050a	0.5	0.125	0.625	0.625	0.5	0.375	316	0.508	0.125	0.625	42.1
339	B30R_075_062a	0.5	0.125	0.75	0.75	0.625	0.437	307	0.51	0.125	0.75	41.7
340	B25R_087_075a	0.5	0.125	0.875	0.875	0.75	0.5	300	0.5	0.125	0.875	41.7
341	B20R_100_087a	0.5	0.125	1.0	1.0	0.875	0.562	295	0.489	0.125	1.0	41.4
342	R50Y_050_050a	0.5	0.25	0.0	0.5	0.5	0.25	60	0.5	0.25	0.0	44.6
343	R31Y_050_037a	0.5	0.25	0.125	0.5	0.375	0.312	49	0.5	0.243	0.124	45.3
344	R00Y_050_025a	0.5	0.25	0.25	0.5	0.25	0.375	390	0.5	0.249	0.249	47.4
345	R00Y_050_025a	0.5	0.25	0.375	0.5	0.25	0.375	360	0.5	0.249	0.375	47.5
346	B50R_050_025a	0.5	0.25	0.5	0.5	0.25	0.375	330	0.5	0.249	0.5	47.6
347	B34R_062_037a	0.5	0.25	0.625	0.625	0.375	0.437	311	0.506	0.25	0.625	47.9
348	B25R_075_050a	0.5	0.25	0.75	0.75	0.5	0.5	300	0.5	0.25	0.75	47.8
349	B19R_087_062a	0.5	0.25	0.875	0.875	0.625	0.562	293	0.489	0.25	0.875	47.5
350	B15R_100_075a	0.5	0.25	1.0	1.0	0.75	0.625	289	0.487	0.25	1.0	47.1
351	R76Y_050_050a	0.5	0.375	0.0	0.5	0.5	0.25	76	0.5	0.383	0.0	51.5
352	R68Y_050_037a	0.5	0.375	0.125	0.5	0.375	0.312	71	0.5	0.381	0.124	52.2
353	R50Y_050_025a	0.5	0.375	0.25	0.5	0.25	0.375	60	0.5	0.375	0.249	52.3
354	R00Y_050_012a	0.5	0.375	0.375	0.5	0.125	0.437	390	0.5	0.375	0.375	53.7
355	B50R_050_012a	0.5	0.375	0.5	0.5	0.125	0.437	330	0.5	0.375	0.5	53.8
356	B25R_062_025a	0.5	0.375	0.625	0.625	0.25	0.5	300	0.5	0.375	0.625	53.9
357	B15R_075_037a	0.5	0.375	0.75	0.75	0.375	0.562	289	0.493	0.375	0.75	53.5
358	B11R_087_050a	0.5	0.375	0.875	0.875	0.5	0.625	284	0.491	0.375	0.875	53.2
359	B09R_100_062a	0.5	0.375	1.0	1.0	0.625	0.687	281	0.489	0.375	1.0	53.5
360	Y00G_050_050a	0.5	0.5	0.0	0.5	0.5	0.25	90	0.5	0.5	0.0	56.1
361	Y00G_050_037a	0.5	0.5	0.125	0.5	0.375	0.312	90	0.5	0.5	0.124	57.0
362	Y00G_050_025a	0.5	0.5	0.25	0.5	0.25	0.375	90	0.5	0.5	0.249	58.0
363	Y00G_050_012a	0.5	0.5	0.375	0.5	0.125	0.437	90	0.5	0.5	0.375	59.0
364	NW_050a	0.5	0.5	0.5	0.5	0.0	0.5	360	0.5	0.5	0.5	60.0
365	B00R_062_012a	0.5	0.5	0.625	0.625	0.125	0.562	270	0.5	0.5	0.625	60.0
366	B00R_075_025a	0.5	0.5	0.75	0.75	0.25	0.625	270	0.5	0.5	0.75	60.1
367	B00R_087_037a	0.5	0.5	0.875	0.875	0.375	0.687	270	0.5	0.5	0.875	60.2
368	B00R_100_050a	0.5	0.5	1.0	1.0	0.5	0.75	270	0.5	0.5	1.0	60.3
369	Y18G_062_062a	0.5	0.625	0.0	0.625	0.625	0.312	101	0.51	0.625	0.0	60.8
370	Y23G_062_050a	0.5	0.625	0.125	0.625	0.5	0.375	104	0.508	0.625	0.125	61.7
371	Y31G_062_037a	0.5	0.625	0.25	0.625	0.375	0.437	109	0.506	0.625	0.25	62.2
372	Y50G_062_025a	0.5	0.625	0.375	0.625	0.25	0.5	120	0.5	0.625	0.375	62.6
373	G00B_062_012a	0.5	0.625	0.5	0.625	0.125	0.562	150	0.5	0.625	0.5	63.2
374	G50B_062_012a	0.5	0.625	0.625	0.625	0.125	0.562	210	0.5	0.625	0.625	64.0
375	G75B_075_025a	0.5	0.625	0.75	0.75	0.25	0.625	240	0.5	0.625	0.75	64.3
376	G84B_087_037a	0.5	0.625	0.875	0.875	0.375	0.687	251	0.5	0.618	0.875	64.0
377	G88B_100_050a	0.5	0.625	1.0	1.0	0.5	0.75	256	0.5	0.616	1.0	63.9
378	Y31G_075_075a	0.5	0.75	0.0	0.75	0.75	0.375	109	0.512	0.75	0.0	64.4
379	Y38G_075_062a	0.5	0.75	0.125	0.75	0.625	0.437	113	0.51	0.75	0.125	64.9
380	Y50G_075_050a	0.5	0.75	0.25	0.75	0.5	0.5	120	0.5	0.75	0.25	65.3
381	Y68G_075_037a	0.5	0.75	0.375	0.75	0.375	0.562	131	0.493	0.75	0.375	65.3
382	G00B_075_025a	0.5	0.75	0.5	0.75	0.25	0.625	150	0.5	0.75	0.5	66.4
383	G25B_075_025a	0.5	0.75	0.625	0.75	0.25	0.625	180	0.5	0.75	0.625	67.1
384	G50B_075_025a	0.5	0.75	0.75	0.75	0.25	0.625	210	0.5	0.75	0.75	68.1
385	G65B_087_037a	0.5	0.75	0.875	0.875	0.375	0.687	229	0.5	0.756	0.875	68.9
386	G75B_100_050a	0.5	0.75	1.0	1.0	0.5	0.75	240	0.5	0.75	1.0	68.6
387	Y41G_087_087a	0.5	0.875	0.0	0.875	0.875	0.437	115	0.51	0.875	0.0	67.5
388	Y50G_087_075a	0.5	0.875	0.125	0.875	0.75	0.5	120	0.5	0.875	0.125	67.9
389	Y61G_087_062a	0.5	0.875	0.25	0.875	0.625	0.562	127	0.489	0.875	0.25	68.2
390	Y76G_087_050a	0.5	0.875	0.375	0.875	0.5	0.625	136	0.491	0.875	0.375	67.8
391	G00B_087_037a	0.5	0.875	0.5	0.875	0.375	0.687	150	0.5	0.875	0.5	69.6
392	G15B_087_037a	0.5	0.875	0.625	0.875	0.375	0.687	169	0.5	0.875	0.625	70.2
393	G34B_087_037a	0.5	0.875	0.75	0.875	0.375	0.687	191	0.5	0.875	0.75	71.3
394	G50B_087_037a	0.5	0.875	0.875	0.875	0.375	0.687	210	0.5	0.875	0.875	72.1
395	G61B_100_050a	0.5	0.875	1.0	1.0	0.5	0.75	224	0.5	0.883	1.0	73.2
396	Y50G_100_100a	0.5	1.0	0.0	1.0	1.0	0.5	120	0.5	1.0	0.0	70.6
397	Y58G_100_087a	0.5	1.0	0.125	1.0	0.875	0.562	125	0.489	1.0	0.125	70.8
398	Y68G_100_075a	0.5	1.0	0.25	1.0	0.75	0.625	131	0.487	1.0	0.25	71.4
399	Y81G_100_062a	0.5	1.0	0.375	1.0	0.625	0.687	139	0.489	1.0	0.375	72.3
400	G00B_100_050a	0.5	1.0	0.5	1.0	0.5	0.75	150	0.5	1.0	0.5	72.8
401	G11B_100_050a	0.5	1.0	0.625	1.0	0.5	0.75	164	0.5	1.0	0.616	73.3
402	G25B_100_050a	0.5	1.0									

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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		
405	R00Y_062_062a	0.625 0.0 0.0	0.625 0.625 0.312	390	0.625 0.0 0.0	37.5 44.3 28.0	52.4 32.3	0.625 0.0 0.0	37.2 53.3 28.6	60.5 28.2 9.0	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
406	R31Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.114	37.6 44.9 23.4	50.6 27.5	0.625 0.0 0.125	37.4 54.0 24.4	59.3 24.3 9.2	380	1.0 0.0 0.183	45.5 71.8 37.5	81.0 27.5
407	R11Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.239	37.7 45.6 17.4	48.8 20.8	0.625 0.0 0.25	37.3 54.8 19.5	58.2 19.6 9.4	367	1.0 0.0 0.383	45.8 73.0 27.8	78.2 20.8
408	B69R_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.385	37.8 47.2 9.5	48.1 11.4	0.625 0.0 0.375	37.4 56.1 13.0	57.6 13.0 9.5	352	1.0 0.0 0.616	46.0 75.5 15.2	77.1 11.4
409	B59R_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.625 0.0 0.51	37.8 48.6 3.9	48.7 4.6	0.625 0.0 0.5	37.4 57.9 6.5	58.2 6.4 9.6	339	1.0 0.0 0.816	45.9 77.7 6.2	78.0 4.6
410	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	37.9 49.5 -0.1	49.5 359.8	0.625 0.0 0.625	37.4 59.3 1.1	59.3 1.0 9.8	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
411	B42R_075_075a	0.625 0.0 0.75	0.75 0.75 0.375	321	0.637 0.0 0.75	38.9 55.7 -4.4	55.9 355.4	0.625 0.0 0.75	37.9 61.6 -4.2	61.8 356.0 5.9	322	0.85 0.0 1.0	43.7 74.3 -5.9	74.6 355.4
412	B36R_087_087a	0.625 0.0 0.875	0.875 0.875 0.437	314	0.641 0.0 0.875	39.2 61.5 -8.7	62.1 351.9	0.625 0.0 0.875	38.3 64.0 -9.1	64.6 351.8 2.6	315	0.733 0.0 1.0	41.3 70.3 -9.9	71.0 351.9
413	B31R_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	38.3 65.8 -13.7	67.2 348.2	0.625 0.0 1.0	38.1 65.4 -14.0	66.9 347.9 0.5	308	0.633 0.0 1.0	38.3 65.8 -13.7	67.2 348.2
414	R18Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	41	0.625 0.114 0.0	41.1 36.1 32.8	48.8 42.2	0.625 0.125 0.0	40.5 45.1 32.7	55.7 35.9 9.0	39	1.0 0.183 0.0	51.1 57.8 52.5	78.1 42.2
415	R00Y_062_050a	0.625 0.125 0.125	0.625 0.5 0.375	390	0.625 0.125 0.125	43.8 35.4 22.4	41.9 32.3	0.625 0.125 0.125	41.0 44.9 28.0	53.0 31.9 11.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
416	R26Y_062_050a	0.625 0.125 0.25	0.625 0.5 0.375	376	0.625 0.125 0.241	43.9 36.0 17.6	40.1 26.1	0.625 0.125 0.25	41.0 45.8 22.3	51.0 25.9 11.2	377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1
417	R00Y_062_050a	0.625 0.125 0.375	0.625 0.5 0.375	360	0.625 0.125 0.375	44.0 37.1 10.5	38.5 15.9	0.625 0.125 0.375	41.1 47.2 15.5	49.7 18.2 11.6	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
418	B61R_062_050a	0.625 0.125 0.5	0.625 0.5 0.375	344	0.625 0.125 0.508	44.0 38.6 4.0	38.8 5.9	0.625 0.125 0.5	41.4 48.6 7.7	49.3 9.0 11.0	342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
419	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	44.1 39.6 -0.1	39.6 359.8	0.625 0.125 0.625	41.7 50.4 1.6	50.4 1.8 11.1	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
420	B40R_075_062a	0.625 0.125 0.75	0.75 0.625 0.437	319	0.635 0.125 0.75	44.9 45.8 -4.4	46.0 354.4	0.625 0.125 0.75	42.7 52.1 -4.3	52.3 355.2 6.7	320	0.816 0.0 1.0	43.1 73.2 -7.0	73.6 354.4
421	B34R_087_075a	0.625 0.125 0.875	0.875 0.75 0.5	311	0.637 0.125 0.875	44.8 51.0 -8.9	51.8 350.0	0.625 0.125 0.875	42.7 54.6 -10.3	55.5 349.2 4.3	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
422	B29R_100_087a	0.625 0.125 1.0	1.0 0.875 0.562	305	0.635 0.125 1.0	44.5 55.3 -14.3	57.1 345.4	0.625 0.125 1.0	43.0 56.2 -15.1	58.2 344.9 1.9	305	0.583 0.0 1.0	37.2 63.2 -16.4	65.3 345.4
423	R38Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	53	0.625 0.239 0.0	46.3 24.7 39.1	46.2 57.6	0.625 0.25 0.0	45.1 34.1 38.7	51.6 48.5 9.5	52	1.0 0.383 0.0	59.5 39.5 62.5	74.0 57.6
424	R23Y_062_050a	0.625 0.25 0.125	0.625 0.5 0.375	44	0.625 0.241 0.125	47.6 26.7 27.4	38.2 45.7	0.625 0.25 0.125	45.7 34.0 33.2	47.6 44.3 9.6	42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7
425	R00Y_062_037a	0.625 0.25 0.25	0.625 0.375 0.437	390	0.625 0.25 0.25	50.1 26.6 16.8	31.4 32.3	0.625 0.25 0.25	46.1 34.0 26.2	43.0 37.6 12.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
426	R18Y_062_037a	0.625 0.25 0.375	0.625 0.375 0.437	371	0.625 0.25 0.368	50.2 27.2 11.7	29.6 23.2	0.625 0.25 0.375	46.5 35.2 19.1	40.1 28.4 11.4	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
427	B65R_062_037a	0.625 0.25 0.5	0.625 0.375 0.437	349	0.625 0.25 0.506	50.2 28.6 4.4	29.0 8.9	0.625 0.25 0.5	46.9 37.0 10.1	38.4 15.3 10.6	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
428	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	50.3 29.7 0.0	29.7 359.8	0.625 0.25 0.625	47.5 38.1 3.1	38.3 4.7 9.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
429	B38R_075_050a	0.625 0.25 0.75	0.75 0.5 0.5	316	0.633 0.25 0.75	51.0 35.8 -4.3	36.0 353.0	0.625 0.25 0.75	48.6 39.4 -3.6	39.6 354.7 4.4	317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0
430	B30R_087_062a	0.625 0.25 0.875	0.875 0.625 0.562	307	0.635 0.25 0.875	50.6 40.6 -9.0	41.6 347.4	0.625 0.25 0.875	49.0 42.1 -9.7	43.2 346.9 2.2	307	0.616 0.0 1.0	37.9 65.0 -14.5	66.6 347.4
431	B25R_100_075a	0.625 0.25 1.0	1.0 0.75 0.625	300	0.635 0.25 1.0	50.6 49.9 -15.5	46.6 340.5	0.625 0.25 1.0	49.1 43.7 -15.5	46.4 340.3 1.5	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
432	R61Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	67	0.625 0.385 0.0	53.9 10.2 47.9	49.0 77.8	0.625 0.375 0.0	50.8 21.2 46.0	50.6 65.2 11.5	67	1.0 0.616 0.0	71.6 16.4 76.6	78.4 77.8
433	R50Y_062_050a	0.625 0.375 0.125	0.625 0.5 0.375	60	0.625 0.375 0.125	53.5 14.4 34.3	37.2 6.7	0.625 0.375 0.125	50.7 22.7 38.2	44.5 59.2 9.5	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
434	R31Y_062_037a	0.625 0.375 0.25	0.625 0.375 0.437	49	0.625 0.368 0.25	54.2 17.1 22.2	28.1 52.2	0.625 0.375 0.25	50.9 23.8 30.0	38.3 51.5 10.8	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
435	R00Y_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	56.3 17.7 11.2	20.9 32.3	0.625 0.375 0.375	51.6 24.4 22.1	33.0 42.1 13.7	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
436	R00Y_062_025a	0.625 0.375 0.5	0.625 0.25 0.5	360	0.625 0.375 0.5	56.4 18.5 5.2	19.2 15.9	0.625 0.375 0.5	52.0 26.1 13.2	29.2 26.9 11.8	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
437	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	56.5 19.8 0.0	19.8 359.8	0.625 0.375 0.625	52.6 27.8 4.7	28.2 9.6 10.1	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
438	B34R_075_037a	0.625 0.375 0.75	0.75 0.375 0.562	311	0.631 0.375 0.75	56.8 25.5 -4.4	25.9 350.0	0.625 0.375 0.75	53.8 29.6 -2.9	29.8 354.2 5.3	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
439	B25R_087_050a	0.625 0.375 0.875	0.875 0.5 0.625	300	0.625 0.375 0.875	56.7 29.3 -10.3	31.0 340.5	0.625 0.375 0.875	54.2 31.4 -9.8	32.9 342.6 3.3	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
440	B19R_100_062a	0.625 0.375 1.0	1.0 0.625 0.687	293	0.614 0.375 1.0	56.4 32.7 -16.0	36.4 333.8	0.625 0.375 1.0	54.3 32.9 -16.3	36.8 333.5 2.1	292	0.383 0.0 1.0	32.9 52.3 -25.7	58.3 333.8
441	R81Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	79	0.625 0.51 0.0	59.7 0.5 54.6	54.6 89.4	0.625 0.5 0.0	55.7 11.1 52.4	53.6 77.9 11.5	80	1.0 0.816 0.0	80.8 0.8 87.3	87.3 89.4
442	R76Y_062_050a	0.625 0.5 0.125	0.625 0.5 0.375	76	0.625 0.508 0.125	60.4 2.1 42.3	42.4 87.0	0.625 0.5 0.125	56.2 11.5 43.8	45.3 75.3 10.3	77	1.0 0.766 0.0	78.6 4.3 84.7	84.8 87.0
443	R68Y_062_037a	0.625 0.5 0.25	0.625 0.375 0.437	71	0.625 0.506 0.25	61.1 4.1 30.1	30.4 82.1	0.625 0.5 0.25	56.7 12.5 34.7	36.9 70.0 10.5	71	1.0 0.683 0.0	74.8 11.0 80.4	81.1 82.1
444	R50Y_062_025a	0.625 0.5 0.375	0.625 0.25 0.5	60	0.625 0.5 0.375	61.2 7.2 17.1	18.6 67.1	0.625 0.5 0.375	57.0 14.3 25.0	28.8 60.2 11.4	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
445	R00Y_062_012a													

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Technische Information: <http://www.ps.bam.de> oder <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_Md	rgb*Md	LabCh*Md																						
486		R00Y_075_075a	0.75	0.0	0.0	0.75	0.375	390	0.75	0.0	0.0	40.7	59.2	36.3	69.4	31.5	6.6	389	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3									
487		R35Y_075_075a	0.75	0.0	0.125	0.75	0.375	381	0.75	0.0	0.112	40.2	53.7	29.2	61.1	28.5	0.75	0.0	0.125	40.6	60.2	31.6	68.0	27.7	6.9	382	1.0	0.0	0.15	45.5	71.6	39.0	81.5	28.5	
488		R18Y_075_075a	0.75	0.0	0.25	0.75	0.375	371	0.75	0.0	0.237	40.4	54.5	23.4	59.3	23.2	0.75	0.0	0.25	40.9	61.1	25.5	66.2	22.6	6.9	371	1.0	0.0	0.316	45.7	72.6	31.2	79.1	23.2	
489		R00Y_075_075a	0.75	0.0	0.375	0.75	0.375	360	0.75	0.0	0.375	40.5	55.6	15.8	57.8	15.9	0.75	0.0	0.375	41.0	62.2	19.2	65.1	17.1	7.4	360	1.0	0.0	0.5	45.9	74.2	21.1	77.1	15.9	
490		B65R_075_075a	0.75	0.0	0.5	0.75	0.375	349	0.75	0.0	0.512	40.5	57.3	8.9	58.0	8.9	0.75	0.0	0.5	40.9	64.0	11.4	65.1	10.1	7.2	348	1.0	0.0	0.683	45.9	76.4	11.9	77.3	8.9	
491		B57R_075_075a	0.75	0.0	0.625	0.75	0.375	339	0.75	0.0	0.637	40.5	58.5	3.7	58.6	3.7	0.75	0.0	0.625	41.1	65.4	5.1	65.6	4.4	7.0	337	1.0	0.0	0.85	45.9	78.0	5.0	78.2	3.7	
492		B50R_075_075a	0.75	0.0	0.75	0.75	0.375	330	0.75	0.0	0.75	40.6	59.4	-0.1	59.4	359.8	0.75	0.0	0.75	41.1	66.9	0.0	66.9	0.0	7.4	330	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8	
493		B43R_087_087a	0.75	0.0	0.875	0.875	0.375	322	0.758	0.0	0.875	41.6	65.5	-4.6	65.7	355.9	0.75	0.0	0.875	41.4	69.0	-4.7	69.2	356.0	3.4	322	0.866	0.0	1.0	44.0	74.9	-5.3	75.1	355.9	
494		B38R_100_100a	0.75	0.0	1.0	1.0	0.5	316	0.766	0.0	1.0	42.1	71.6	-8.7	72.1	353.0	0.75	0.0	1.0	41.8	71.0	-9.2	71.6	352.5	0.8	317	0.766	0.0	1.0	42.1	71.6	-8.7	72.1	353.0	
495		R15Y_075_075a	0.75	0.125	0.0	0.75	0.375	39	0.75	0.112	0.0	43.4	45.5	38.0	59.3	39.9	0.75	0.125	0.0	43.9	51.3	40.0	65.1	37.8	6.1	37	1.0	0.15	0.0	49.8	60.7	50.7	79.1	39.9	
496		R00Y_075_062a	0.75	0.125	0.125	0.75	0.625	437	0.75	0.125	0.125	46.4	44.3	28.0	52.4	32.3	0.75	0.125	0.125	44.5	50.6	34.5	61.3	34.3	9.2	389	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3	
497		R31Y_075_062a	0.75	0.125	0.25	0.75	0.625	437	0.75	0.125	0.239	46.5	44.9	23.4	50.6	27.5	0.75	0.125	0.25	44.8	51.4	28.4	58.8	28.9	8.3	380	1.0	0.0	0.183	45.5	71.8	37.5	81.0	27.5	
498		R11Y_075_062a	0.75	0.125	0.375	0.75	0.625	437	0.75	0.125	0.364	46.6	45.6	17.4	48.8	20.8	0.75	0.125	0.375	45.0	52.4	21.2	56.5	22.0	7.9	367	1.0	0.0	0.383	45.8	73.0	27.8	78.2	20.8	
499		B69R_075_062a	0.75	0.125	0.5	0.75	0.625	437	0.75	0.125	0.51	46.8	47.2	9.5	48.1	11.4	0.75	0.125	0.5	45.4	54.0	12.4	55.4	12.9	7.5	352	1.0	0.0	0.616	46.0	75.5	15.2	77.1	11.4	
500		B59R_075_062a	0.75	0.125	0.625	0.75	0.625	437	0.75	0.125	0.635	46.7	48.6	3.9	48.7	4.6	0.75	0.125	0.625	45.8	55.0	5.5	55.3	5.7	6.6	339	1.0	0.0	0.816	45.9	77.7	6.2	78.0	4.6	
501		B50R_075_062a	0.75	0.125	0.75	0.75	0.625	437	0.75	0.125	0.75	46.8	49.5	-0.1	49.5	359.8	0.75	0.125	0.75	45.9	56.5	-0.2	56.5	359.7	7.0	330	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8	
502		B42R_087_075a	0.75	0.125	0.875	0.875	0.75	0.5	321	0.762	0.125	0.875	47.8	55.7	-4.4	55.9	355.4	0.75	0.125	0.875	46.6	58.6	-5.6	58.9	354.5	3.3	322	0.85	0.0	1.0	43.7	74.3	-5.9	74.6	355.4
503		B36R_100_087a	0.75	0.125	1.0	1.0	0.875	314	0.766	0.125	1.0	48.1	61.5	-8.7	62.1	351.9	0.75	0.125	1.0	47.0	60.4	-10.4	61.3	350.2	2.2	315	0.733	0.0	1.0	41.3	70.3	-9.9	71.0	351.9	
504		R31Y_075_075a	0.75	0.25	0.0	0.75	0.375	49	0.75	0.237	0.0	48.5	34.3	44.4	56.2	52.2	0.75	0.25	0.0	48.9	39.7	46.7	61.3	49.6	5.8	48	1.0	0.316	0.0	0.616	45.8	59.2	74.9	52.2	
505		R18Y_075_062a	0.75	0.25	0.125	0.75	0.625	437	0.75	0.239	0.125	50.0	36.1	32.8	48.8	42.2	0.75	0.25	0.125	49.3	39.8	39.4	56.1	44.7	7.6	39	1.0	0.183	0.0	51.1	57.8	52.5	78.1	42.2	
506		R00Y_075_050a	0.75	0.25	0.25	0.75	0.5	0.5	390	0.75	0.25	0.25	52.7	35.4	22.4	41.9	32.3	0.75	0.25	0.25	50.4	39.4	31.9	50.7	38.9	10.5	389	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3
507		R26Y_075_050a	0.75	0.25	0.375	0.75	0.5	0.5	376	0.75	0.25	0.366	52.8	36.0	17.6	40.1	26.1	0.75	0.25	0.375	51.0	39.9	24.4	46.8	31.4	8.0	377	1.0	0.0	0.233	45.6	72.1	35.3	80.3	26.1
508		R00Y_075_050a	0.75	0.25	0.5	0.75	0.5	0.5	360	0.75	0.25	0.5	52.9	37.1	10.5	38.5	15.9	0.75	0.25	0.5	51.3	41.4	15.2	44.1	20.2	6.5	360	1.0	0.0	0.5	45.9	74.2	21.1	77.1	15.9
509		B61R_075_050a	0.75	0.25	0.625	0.75	0.5	0.5	344	0.75	0.25	0.633	52.9	38.6	4.0	38.8	5.9	0.75	0.25	0.625	52.0	42.7	7.1	43.3	9.4	5.1	342	1.0	0.0	0.766	45.9	77.3	8.0	77.7	5.9
510		B50R_075_050a	0.75	0.25	0.75	0.75	0.5	0.5	330	0.75	0.25	0.75	53.0	39.6	-0.1	39.6	359.8	0.75	0.25	0.75	52.4	44.4	0.5	44.4	0.6	4.8	330	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8
511		B40R_087_062a	0.75	0.25	0.875	0.875	0.625	319	0.76	0.25	0.875	53.9	45.8	-4.4	46.0	354.4	0.75	0.25	0.875	53.4	46.0	-5.4	46.3	353.2	1.1	320	0.816	0.0	1.0	43.1	73.2	-7.0	73.6	354.4	
512		B34R_100_075a	0.75	0.25	1.0	1.0	0.75	0.625	311	0.762	0.25	1.0	53.7	51.0	-8.9	51.8	350.0	0.75	0.25	1.0	53.7	47.7	-10.9	48.9	347.1	3.9	311	0.683	0.0	1.0	39.8	68.1	-11.9	69.1	350.0
513		R50Y_075_075a	0.75	0.375	0.0	0.75	0.375	60	0.75	0.375	0.0	54.7	21.6	51.5	55.9	67.1	0.75	0.375	0.0	54.3	28.1	53.1	60.1	62.1	6.6	59	1.0	0.5	0.0	64.9	28.9	68.6	74.5	67.1	
514		R38Y_075_062a	0.75	0.375	0.125	0.75	0.625	437	0.75	0.364	0.125	55.2	24.7	39.1	46.2	57.6	0.75	0.375	0.125	54.7	28.8	44.2	52.8	56.8	6.6	52	1.0	0.383	0.0	59.5	39.5	62.5	74.0	57.6	
515		R23Y_075_050a	0.75	0.375	0.25	0.75	0.5	0.5	44	0.75	0.366	0.25	56.5	26.7	27.4	38.2	45.7	0.75	0.375	0.25	55.2	29.4	35.2	45.9	50.0	8.3	42	1.0	0.233	0.0	53.0	53.4	54.8	76.5	45.7
516		R00Y_075_037a	0.75	0.375	0.375	0.75	0.375	562	0.75	0.375	0.375	59.0	26.6	16.8	31.4	32.3	0.75	0.375	0.375	56.5	29.0	26.5	39.3	42.3	10.3	389	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3	
517		R18Y_075_037a	0.75	0.375	0.5	0.75	0.375	562	0.75	0.375	0.493</																								

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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		
567	R00Y_087_087a	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.0	42.8 62.0 39.2	73.4 32.3	0.875 0.0 0.0	43.2 65.4 40.5	76.9 31.8	3.6 389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
568	R36Y_087_087a	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.116	42.9 62.5 34.7	71.6 29.0	0.875 0.0 0.125	43.3 66.0 35.3	74.9 28.1 3.5	382	1.0 0.0 0.133	45.5 71.5 39.7	81.8 29.0
569	R23Y_087_087a	0.875 0.0 0.25	0.875 0.875 0.437	374	0.875 0.0 0.233	43.0 63.2 29.5	69.8 25.0	0.875 0.0 0.25	43.6 66.5 29.6	72.8 23.9 3.3	375	1.0 0.0 0.266	45.6 72.3 33.8	79.8 25.0
570	R08Y_087_087a	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.364	43.1 64.2 22.7	68.1 19.4	0.875 0.0 0.375	43.6 67.7 23.3	71.6 19.0 3.5	365	1.0 0.0 0.416	45.8 73.4 25.9	77.9 19.4
571	B70R_087_087a	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.51	43.2 65.8 14.8	67.4 12.7	0.875 0.0 0.5	43.7 69.3 16.0	71.2 13.0 3.7	354	1.0 0.0 0.583	45.9 75.2 16.9	77.1 12.7
572	B63R_087_087a	0.875 0.0 0.625	0.875 0.875 0.437	346	0.875 0.0 0.641	43.2 67.3 8.3	67.8 7.0	0.875 0.0 0.625	43.8 70.8 9.3	71.4 7.5 3.6	344	1.0 0.0 0.733	45.9 77.0 9.4	77.5 7.0
573	B56R_087_087a	0.875 0.0 0.75	0.875 0.875 0.437	338	0.875 0.0 0.758	43.2 68.4 3.8	68.5 3.2	0.875 0.0 0.75	43.8 72.3 4.2	72.5 3.3 4.0	337	1.0 0.0 0.866	45.9 78.1 4.4	78.3 3.2
574	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	43.4 69.4 -0.1	69.4 359.8	0.875 0.0 0.875	44.0 73.5 -0.8	73.5 359.3 4.2	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
575	B44R_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	44.3 75.4 -4.7	75.6 356.3	0.875 0.0 1.0	44.2 75.2 -5.0	75.3 356.1 0.4	323	0.883 0.0 1.0	44.3 75.4 -4.7	75.6 356.3
576	R13Y_087_087a	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.116 0.0	46.1 54.3 43.6	69.7 38.7	0.875 0.125 0.0	47.3 56.4 44.0	71.5 38.0 2.4	37	1.0 0.133 0.0	49.2 62.1 49.8	79.6 38.7
577	R00Y_087_075a	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.125	49.1 53.2 33.6	62.9 32.3	0.875 0.125 0.125	47.6 56.0 38.5	67.9 34.5 5.8	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
578	R35Y_087_075a	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.237	49.1 53.7 29.2	61.1 28.5	0.875 0.125 0.25	47.9 56.7 32.6	65.4 29.8 4.6	382	1.0 0.0 0.15	45.5 71.6 39.0	81.5 28.5
579	R18Y_087_075a	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.362	49.3 54.5 23.4	59.3 23.2	0.875 0.125 0.375	48.2 57.5 25.3	62.8 23.7 3.7	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
580	R00Y_087_075a	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.5	49.4 55.6 15.8	57.8 15.9	0.875 0.125 0.5	48.4 59.1 16.9	61.5 15.9 3.7	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
581	B65R_087_075a	0.875 0.125 0.625	0.875 0.75 0.5	349	0.875 0.125 0.637	49.4 57.3 8.9	58.0 8.9	0.875 0.125 0.625	48.8 60.3 9.3	61.0 8.8 3.1	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
582	B57R_087_075a	0.875 0.125 0.75	0.875 0.75 0.5	339	0.875 0.125 0.762	49.4 58.5 3.7	58.6 3.7	0.875 0.125 0.75	48.9 62.0 2.9	62.0 2.7 3.6	337	1.0 0.0 0.85	45.9 78.0 5.0	78.2 3.7
583	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	49.5 59.4 -0.1	59.4 359.8	0.875 0.125 0.875	49.3 62.9 -2.0	62.9 358.1 3.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
584	B43R_100_087a	0.875 0.125 1.0	1.0 0.875 0.562	322	0.883 0.125 1.0	50.5 65.5 -4.6	65.7 355.9	0.875 0.125 1.0	49.6 64.5 -6.6	64.9 354.1 2.3	322	0.866 0.0 1.0	44.0 74.9 -5.3	75.1 355.9
585	R26Y_087_087a	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.233 0.0	50.6 44.1 49.4	66.2 48.2	0.875 0.25 0.0	51.7 45.6 50.7	68.2 48.0 2.3	44	1.0 0.266 0.0	54.4 50.4 56.5	75.7 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	52.4 45.5 38.0	59.3 39.9	0.875 0.25 0.125	52.6 45.0 43.6	62.7 44.1 5.6	37	1.0 0.15 0.0	49.8 60.7 50.7	79.1 39.9
587	R00Y_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	55.3 44.3 28.0	52.4 32.3	0.875 0.25 0.25	53.7 44.1 35.9	56.8 39.1 8.0	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
588	R31Y_087_062a	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.364	55.4 44.9 23.4	50.6 27.5	0.875 0.25 0.375	54.3 44.5 28.2	52.7 32.3 4.8	380	1.0 0.0 0.183	45.5 71.8 37.5	81.0 27.5
589	R11Y_087_062a	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.489	55.6 45.6 17.4	48.8 20.8	0.875 0.25 0.5	54.5 45.9 19.9	50.0 23.4 2.7	367	1.0 0.0 0.383	45.8 73.0 27.8	78.2 20.8
590	B69R_087_062a	0.875 0.25 0.625	0.875 0.625 0.562	353	0.875 0.25 0.635	55.7 47.2 9.5	48.1 11.4	0.875 0.25 0.625	55.1 47.5 10.8	48.7 12.8 1.4	352	1.0 0.0 0.616	46.0 75.5 15.2	77.1 11.4
591	B59R_087_062a	0.875 0.25 0.75	0.875 0.625 0.562	341	0.875 0.25 0.76	55.6 48.6 3.9	48.7 4.6	0.875 0.25 0.75	55.4 48.8 4.0	49.0 4.6 0.3	339	1.0 0.0 0.816	45.9 77.7 6.2	78.0 4.6
592	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	55.7 49.5 -0.1	49.5 359.8	0.875 0.25 0.875	56.0 49.9 -1.8	49.9 357.9 1.7	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
593	B42R_100_075a	0.875 0.25 1.0	1.0 0.75 0.625	321	0.887 0.25 1.0	56.7 55.7 -4.4	55.9 355.4	0.875 0.25 1.0	56.7 51.9 -6.8	52.3 352.4 4.5	322	0.85 0.0 1.0	43.7 74.3 -5.9	74.6 355.4
594	R41Y_087_087a	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.364 0.0	56.5 32.0 56.4	64.9 60.3	0.875 0.375 0.0	57.5 33.5 57.7	66.8 59.8 2.2	54	1.0 0.416 0.0	61.0 36.6 64.5	74.1 60.3
595	R31Y_087_075a	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.362 0.125	57.4 34.3 44.4	56.2 52.2	0.875 0.375 0.125	57.9 33.6 48.9	59.4 55.5 4.6	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
596	R18Y_087_062a	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.364 0.25	58.9 36.1 32.8	48.8 42.2	0.875 0.375 0.25	58.6 34.1 39.3	52.1 49.0 6.8	39	1.0 0.183 0.0	51.1 57.8 52.5	78.1 42.2
597	R00Y_087_050a	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	61.6 35.4 22.4	41.9 32.3	0.875 0.375 0.375	59.7 33.8 30.7	45.6 42.2 8.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
598	R26Y_087_050a	0.875 0.375 0.5	0.875 0.5 0.625	376	0.875 0.375 0.491	61.7 36.0 17.6	40.1 26.1	0.875 0.375 0.5	60.3 34.8 21.9	41.1 32.1 4.6	377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1
599	R00Y_087_050a	0.875 0.375 0.625	0.875 0.5 0.625	360	0.875 0.375 0.625	61.8 37.1 10.5	38.5 15.9	0.875 0.375 0.625	61.1 36.1 12.9	38.3 19.7 2.6	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
600	B61R_087_050a	0.875 0.375 0.75	0.875 0.5 0.625	344	0.875 0.375 0.758	61.8 38.6 4.0	38.8 5.9	0.875 0.375 0.75	61.4 37.8 4.6	38.1 7.0 1.0	342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
601	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	61.9 39.6 -0.1	39.6 359.8	0.875 0.375 0.875	62.3 38.7 -1.4	38.7 357.8 1.6	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
602	B40R_100_062a	0.875 0.375 1.0	1.0 0.625 0.687	319	0.885 0.375 1.0	62.8 45.8 -4.4	46.0 354.4	0.875 0.375 1.0	63.0 40.3 -7.2	40.9 349.7 6.1	320	0.816 0.0 1.0	43.1 73.2 -7.0	73.6 354.4
603	R58Y_087_087a	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.51 0.0	64.0 17.7 65.2	67.6 74.8	0.875 0.5 0.0	63.7 21.0 64.7	68.1 72.0 3.3	65	1.0 0.583 0.0	69.7 20.2 74.6	77.3 74.8
604	R50Y_087_075a	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.5 0.125	63.6 21.6 51.5	55.9 67.1	0.875 0.5 0.125	63.9 22.1 53.8	58.2 67.6 2.4	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
605	R38Y_087_062a	0.875 0.5 0.25	0.875 0.625 0.562	53	0.875 0.489 0.25	64.1 24.7 39.1	46.2 57.6	0.875 0.5 0.25	64.0 23.7 43.4	49.4 61.3 4.4	52	1.0 0.383 0.0	59.5 39.5 62.5	74.0 57.6
606	R23Y_087_050a	0.875 0.5 0.375	0.875 0.5 0.625	44	0.875 0.491 0.375	65.4 26.7 27.4	38.2 45.7	0.875 0.5 0.375	64.9 24.1 3					

n	HIC*Fd	rgb_Fd	ict_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb**Fd	LabCh**Fd	DE**Fd	hsiMd	rgb**Md	LabCh**Md	
648	R00Y_100_100a	1.0	0.0	0.0	1.0	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3
649	R38Y_100_100a	1.0	0.0	0.125	1.0	1.0	0.0	0.116	45.5	71.4	40.1	81.9	29.5
650	R26Y_100_100a	1.0	0.0	0.25	1.0	1.0	0.0	0.233	45.6	72.1	34.6	80.0	26.1
651	R13Y_100_100a	1.0	0.0	0.375	1.0	1.0	0.0	0.366	45.8	72.9	28.3	78.3	21.5
652	R00Y_100_100a	1.0	0.0	0.5	1.0	1.0	0.0	0.5	45.9	74.2	21.1	77.1	15.9
653	B68R_100_100a	1.0	0.0	0.625	1.0	1.0	0.0	0.633	46.0	75.6	14.8	77.0	11.1
654	B61R_100_100a	1.0	0.0	0.75	1.0	1.0	0.0	0.766	45.9	77.3	8.0	77.7	5.9
655	B55R_100_100a	1.0	0.0	0.875	1.0	1.0	0.0	0.883	45.9	78.3	3.8	78.4	2.8
656	B50R_100_100a	1.0	0.0	1.0	1.0	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8
657	R11Y_100_100a	1.0	0.125	0.0	1.0	1.0	0.116	0.0	48.6	63.3	49.1	80.2	37.7
658	R00Y_100_087a	1.0	0.125	0.125	1.0	0.875	0.125	0.125	51.7	62.0	39.2	73.4	32.3
659	R36Y_100_087a	1.0	0.125	0.25	1.0	0.875	0.125	0.241	51.8	62.5	34.7	71.6	29.0
660	R23Y_100_087a	1.0	0.125	0.375	1.0	0.875	0.125	0.358	51.9	63.2	29.5	69.8	25.0
661	R08Y_100_087a	1.0	0.125	0.5	1.0	0.875	0.125	0.489	52.0	64.2	22.7	68.1	19.4
662	B70R_100_087a	1.0	0.125	0.625	1.0	0.875	0.125	0.635	52.1	65.8	14.8	67.4	12.7
663	B63R_100_087a	1.0	0.125	0.75	1.0	0.875	0.125	0.766	52.1	67.3	8.3	67.8	7.0
664	B56R_100_087a	1.0	0.125	0.875	1.0	0.875	0.125	0.883	52.1	68.4	3.8	68.5	3.2
665	B50R_100_087a	1.0	0.125	1.0	1.0	0.875	0.125	1.0	52.3	69.4	-0.1	69.4	359.8
666	R23Y_100_100a	1.0	0.25	0.0	1.0	1.0	0.233	0.0	53.0	53.4	54.8	76.5	45.7
667	R13Y_100_087a	1.0	0.25	0.125	1.0	0.875	0.241	0.125	55.0	54.3	43.6	69.7	38.7
668	R00Y_100_075a	1.0	0.25	0.25	1.0	0.75	0.25	0.25	58.0	53.2	33.6	62.9	32.3
669	R35Y_100_075a	1.0	0.25	0.375	1.0	0.75	0.25	0.362	58.0	53.7	29.2	61.1	28.5
670	R18Y_100_075a	1.0	0.25	0.5	1.0	0.75	0.25	0.487	58.2	54.5	23.4	59.3	23.2
671	R00Y_100_075a	1.0	0.25	0.625	1.0	0.75	0.25	0.625	58.3	55.6	15.8	57.8	15.9
672	B65R_100_075a	1.0	0.25	0.75	1.0	0.75	0.25	0.762	58.3	57.3	8.9	58.0	8.9
673	B57R_100_075a	1.0	0.25	0.875	1.0	0.75	0.25	0.887	58.3	58.5	3.7	58.6	3.7
674	B50R_100_075a	1.0	0.25	1.0	1.0	0.75	0.25	1.0	58.4	59.4	-0.1	59.4	359.8
675	R36Y_100_100a	1.0	0.375	0.0	1.0	1.0	0.366	0.0	58.8	41.1	61.7	74.1	56.3
676	R26Y_100_087a	1.0	0.375	0.125	1.0	0.875	0.358	0.125	59.2	41.2	53.0	67.1	46.4
677	R15Y_100_075a	1.0	0.375	0.25	1.0	0.75	0.362	0.25	61.3	45.5	38.0	59.3	39.9
678	R00Y_100_062a	1.0	0.375	0.375	1.0	0.625	0.375	0.375	64.2	44.3	28.0	52.4	32.3
679	R31Y_100_062a	1.0	0.375	0.5	1.0	0.625	0.375	0.489	64.3	44.9	23.4	50.6	27.5
680	R11Y_100_062a	1.0	0.375	0.625	1.0	0.625	0.375	0.614	64.5	45.6	17.4	48.8	20.8
681	B69R_100_062a	1.0	0.375	0.75	1.0	0.625	0.375	0.76	64.6	47.2	9.5	48.1	11.4
682	B59R_100_062a	1.0	0.375	0.875	1.0	0.625	0.375	0.885	64.5	48.6	3.9	48.7	4.6
683	B50R_100_062a	1.0	0.375	1.0	1.0	0.625	0.375	1.0	64.6	49.5	-0.1	49.5	359.8
684	R50Y_100_100a	1.0	0.5	0.0	1.0	1.0	0.5	0.0	64.9	28.9	68.6	74.5	67.1
685	R41Y_100_087a	1.0	0.5	0.125	1.0	0.875	0.489	0.125	65.4	32.0	56.4	64.9	60.3
686	R31Y_100_075a	1.0	0.5	0.25	1.0	0.75	0.487	0.25	66.3	34.3	44.4	56.2	52.2
687	R18Y_100_062a	1.0	0.5	0.375	1.0	0.625	0.489	0.375	67.8	36.1	32.8	48.8	42.2
688	R00Y_100_050a	1.0	0.5	0.5	1.0	0.5	0.5	0.5	70.5	35.4	22.4	41.9	32.3
689	R26Y_100_050a	1.0	0.5	0.625	1.0	0.5	0.5	0.616	70.6	36.0	17.6	40.1	26.1
690	R00Y_100_050a	1.0	0.5	0.75	1.0	0.5	0.5	0.75	70.7	37.1	10.5	38.5	15.9
691	B61R_100_050a	1.0	0.5	0.875	1.0	0.5	0.5	0.883	70.7	38.6	4.0	38.8	5.9
692	B50R_100_050a	1.0	0.5	1.0	1.0	0.5	0.5	1.0	70.8	39.6	-0.1	39.6	359.8
693	R63Y_100_100a	1.0	0.625	0.0	1.0	1.0	0.633	0.0	72.5	14.8	77.6	79.0	79.1
694	R58Y_100_087a	1.0	0.625	0.125	1.0	0.875	0.635	0.125	72.9	17.7	65.2	67.6	74.8
695	R50Y_100_075a	1.0	0.625	0.25	1.0	0.75	0.625	0.25	72.5	21.6	51.5	55.9	67.1
696	R38Y_100_062a	1.0	0.625	0.375	1.0	0.625	0.614	0.375	73.0	24.7	39.1	46.2	57.6
697	R23Y_100_050a	1.0	0.625	0.5	1.0	0.5	0.616	0.5	74.3	26.7	27.4	38.2	45.7
698	R00Y_100_037a	1.0	0.625	0.625	1.0	0.375	0.625	0.625	76.8	26.6	16.8	31.4	32.3
699	R18Y_100_037a	1.0	0.625	0.75	1.0	0.375	0.625	0.743	76.9	27.2	11.7	29.6	23.2
700	B65R_100_037a	1.0	0.625	0.875	1.0	0.375	0.625	0.881	77.0	28.6	4.4	29.0	8.9
701	B50R_100_037a	1.0	0.625	1.0	1.0	0.375	0.625	1.0	77.0	29.7	0.0	29.7	359.8
702	R76Y_100_100a	1.0	0.75	0.0	1.0	1.0	0.766	0.0	78.6	4.3	84.7	84.8	87.0
703	R73Y_100_087a	1.0	0.75	0.125	1.0	0.875	0.766	0.125	79.4	6.0	72.6	72.9	85.2
704	R68Y_100_075a	1.0	0.75	0.25	1.0	0.75	0.762	0.25	80.0	8.2	60.3	60.8	82.1
705	R61Y_100_062a	1.0	0.75	0.375	1.0	0.625	0.76	0.375	80.6	10.2	47.9	49.0	77.8
706	R50Y_100_050a	1.0	0.75	0.5	1.0	0.5	0.75	0.5	80.2	14.4	34.3	37.2	67.1
707	R31Y_100_037a	1.0	0.75	0.625	1.0	0.375	0.743	0.625	80.9	17.1	22.2	28.1	52.2
708	R00Y_100_025a	1.0	0.75	0.75	1.0	0.25	0.75	0.75	83.0	17.7	11.2	20.9	32.3
709	R00Y_100_025a	1.0	0.75	0.875	1.0	0.25	0.75	0.875	83.1	18.5	5.2	19.2	15.9
710	B50R_100_025a	1.0	0.75	1.0	1.0	0.25	0.75	1.0	83.2	19.8	0.0	19.8	359.8
711	R88Y_100_100a	1.0	0.875	0.0	1.0	1.0	0.883	0.0	83.7	-3.8	90.5	90.6	92.4
712	R86Y_100_087a	1.0	0.875	0.125	1.0	0.875	0.883	0.125	84.6	-2.4	78.6	78.6	91.7
713	R85Y_100_075a	1.0	0.875	0.25	1.0	0.75	0.887	0.25	85.6	-1.1	66.7	66.7	91.0
714	R81Y_100_062a	1.0	0.875	0.375	1.0	0.625	0.885	0.375	86.4	0.5	54.6	54.6	89.4
715	R76Y_100_050a	1.0	0.875	0.5	1.0	0.5	0.883	0.5	87.1	2.1	42.3	42.4	87.0
716	R68Y_100_037a	1.0	0.875	0.625	1.0	0.375	0.881	0.625	87.8	4.1	30.1	30.4	82.1
717	R50Y_100_025a	1.0	0.875	0.75	1.0	0.25	0.875	0.75	87.9	7.2	17.1	18.6	67.1
718	R00Y_100_012a	1.0	0.875	0.875	1.0	0.125	0.875	0.875	89.3	8.8	5.6	10.4	32.3
719	B50R_100_012a	1.0	0.875	1.0	1.0	0.125	0.875	1.0	89.4	9.9	0.0	9.9	359.8
720	Y00G_100_100a	1.0	1.0	0.0	1.0	1.0	1.0	0.0	87.8	-10.2	95.4	96.0	96.1
721	Y00G_100_087a	1.0	1.0	0.125	1.0	0.875	1.0	0.125	88.8	-8.9	83.5	84.0	96.1
722	Y00G_100_075a	1.0	1.0	0.25	1.0	0.75	1.0	0.25	89.7	-7.6	71.6	72.0	96.1
723	Y00G_100_062a	1.0	1.0	0.375	1.0	0.625	1.0	0.375	90.7	-6.3	59.6	60.0	96.1
724	Y00G_100_050a	1.0	1.0	0.5	1.0	0.5	1.0	0.5	91.7	-5.1	47.7	48.0	96.1
725	Y00G_100_037a	1.0	1.0	0.625	1.0	0.375	1.0	0.625	92.6	-3.8	35.8	36.0	96.1
726	Y00G_100_025a	1.0	1.0	0.75	1.0	0.25	0.875	0.75	93.6	-2.5	23.8	24.0	96.1
727	Y00G_100_012a	1.0	1.0	0.875	1.0	0.125	0.875	0.875	94.6	-1.2	11.9	12.0	96.1
728	NW_100a	1.0	1.0	1.0	1.0	1.0	1.0	1.0	95.6	0.0	0.0	0.0	0.0

delta E* = 3.7

Siehe ähnliche Dateien: <http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT> /PS
Technische Information: <http://www.ps.bam.de> oder <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			
729	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.5 0.0 0.0	0.1 112.0 0.1	360	1.0 1.0 1.0	95.6 0.0 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	90.7 -3.1 -5.1	6.0 238.4	0.875 1.0 1.0	91.9 -2.9 -4.1	5.0 234.3 1.6	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
731	G50B_100_025d	0.75 1.0 1.0	1.0 0.25 0.875	210	0.75 1.0 1.0	85.9 -6.3 -10.3	12.1 238.4	0.75 1.0 1.0	87.8 -5.7 -8.6	10.3 236.4 2.7	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
732	G50B_100_037d	0.625 1.0 1.0	1.0 0.375 0.812	210	0.625 1.0 1.0	81.0 -9.5 -15.5	18.2 238.4	0.625 1.0 1.0	83.2 -8.6 -13.4	15.9 237.2 3.2	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
733	G50B_100_050d	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	76.2 -12.7 -20.7	24.3 238.4	0.5 1.0 1.0	77.6 -12.2 -19.4	22.9 237.6 2.0	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
734	G50B_100_062d	0.375 1.0 1.0	1.0 0.625 0.687	210	0.375 1.0 1.0	71.3 -15.9 -25.9	30.4 238.4	0.375 1.0 1.0	72.3 -15.5 -24.9	29.4 238.1 1.4	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
735	G50B_100_075d	0.25 1.0 1.0	1.0 0.75 0.625	210	0.25 1.0 1.0	66.5 -19.1 -31.1	36.5 238.4	0.25 1.0 1.0	66.5 -19.1 -31.2	36.6 238.4 0.0	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
736	G50B_100_087d	0.125 1.0 1.0	1.0 0.875 0.562	210	0.125 1.0 1.0	61.6 -22.3 -36.3	42.6 238.4	0.125 1.0 1.0	61.2 -21.8 -36.5	42.5 239.0 0.6	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
737	G50B_100_100d	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4	0.0 1.0 1.0	55.3 -24.7 -42.3	49.0 239.6 1.7	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
738	ROOY_100_012d	1.0 0.875 0.875	1.0 1.125 0.937	390	1.0 0.875 0.875	89.3 8.8 5.6	10.4 32.3	1.0 0.875 0.875	89.7 4.4 7.8	9.0 60.1 4.9	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3
739	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0	0.875 0.875 0.875	86.1 1.2 3.6	3.8 70.9 3.8	360	1.0 1.0 1.0	95.6 0.0 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 -3.1 -5.1	6.0 238.4	0.75 0.875 0.875	82.2 -1.9 -0.8	2.1 204.3 4.4	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
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 -6.3 -10.3	12.1 238.4	0.625 0.875 0.875	77.9 -5.4 -5.5	7.8 225.6 4.9	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
742	G50B_087_037d	0.5 0.875 0.875	0.875 0.375 0.687	210	0.5 0.875 0.875	72.1 -9.5 -15.5	18.2 238.4	0.5 0.875 0.875	72.8 -9.5 -11.3	14.8 229.9 4.2	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
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 -12.7 -20.7	24.3 238.4	0.375 0.875 0.875	67.6 -13.7 -16.9	21.8 230.9 3.9	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
744	G50B_087_062d	0.25 0.875 0.875	0.875 0.625 0.562	210	0.25 0.875 0.875	62.4 -15.9 -25.9	30.4 238.4	0.25 0.875 0.875	62.2 -18.3 -23.4	29.8 231.9 3.4	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
745	G50B_087_075d	0.125 0.875 0.875	0.875 0.75 0.5	210	0.125 0.875 0.875	57.6 -19.1 -31.1	36.5 238.4	0.125 0.875 0.875	57.2 -22.1 -28.6	36.1 232.2 3.9	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
746	G50B_087_087d	0.0 0.875 0.875	0.875 0.875 0.437	210	0.0 0.875 0.875	52.7 -22.3 -36.3	42.6 238.4	0.0 0.875 0.875	51.9 -26.3 -34.9	43.7 232.9 4.3	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
747	ROOY_100_025d	1.0 0.75 0.75	1.0 0.25 0.875	390	1.0 0.75 0.75	83.0 17.7 11.2	20.9 32.3	1.0 0.75 0.75	82.3 11.7 15.1	19.1 52.1 7.1	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3
748	ROOY_087_012d	0.875 0.75 0.75	0.875 0.125 0.812	390	0.875 0.75 0.75	80.4 8.8 5.6	10.4 32.3	0.875 0.75 0.75	79.1 8.0 10.9	13.6 53.6 5.5	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3
749	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0	0.75 0.75 0.75	75.6 4.4 6.7	8.0 56.1 8.3	360	1.0 1.0 1.0	95.6 0.0 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	72.9 -3.1 -5.1	6.0 238.4	0.625 0.75 0.75	71.2 0.3 1.9	2.0 79.0 8.2	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
751	G50B_075_025d	0.5 0.75 0.75	0.75 0.25 0.625	210	0.5 0.75 0.75	68.1 -6.3 -10.3	12.1 238.4	0.5 0.75 0.75	66.4 -4.7 -3.8	6.1 219.4 6.9	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
752	G50B_075_037d	0.375 0.75 0.75	0.75 0.375 0.562	210	0.375 0.75 0.75	63.2 -9.5 -15.5	18.2 238.4	0.375 0.75 0.75	61.8 -9.3 -9.6	13.4 225.8 6.0	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
753	G50B_075_050d	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.75	58.4 -12.7 -20.7	24.3 238.4	0.25 0.75 0.75	56.5 -15.2 -16.0	22.1 226.3 5.6	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
754	G50B_075_062d	0.125 0.75 0.75	0.75 0.625 0.437	210	0.125 0.75 0.75	53.5 -15.9 -25.9	30.4 238.4	0.125 0.75 0.75	52.2 -19.8 -21.1	28.9 226.8 6.3	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
755	G50B_075_075d	0.0 0.75 0.75	0.75 0.75 0.375	210	0.0 0.75 0.75	48.7 -19.1 -31.1	36.5 238.4	0.0 0.75 0.75	47.3 -25.7 -27.2	37.5 226.6 7.8	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
756	ROOY_100_037d	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	76.8 26.6 16.8	31.4 32.3	1.0 0.625 0.625	76.1 18.3 22.9	29.3 51.3 10.2	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3
757	ROOY_087_025d	0.875 0.625 0.625	0.875 0.25 0.75	390	0.875 0.625 0.625	74.1 17.7 11.2	20.9 32.3	0.875 0.625 0.625	73.0 14.4 18.5	23.5 52.0 8.0	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3
758	ROOY_075_012d	0.75 0.625 0.625	0.75 0.125 0.687	390	0.75 0.625 0.625	71.5 8.8 5.6	10.4 32.3	0.75 0.625 0.625	69.8 10.1 14.0	17.3 54.0 8.6	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3
759	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0	0.625 0.625 0.625	65.4 5.8 9.1	10.9 57.3 11.4	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	
760	G50B_062_012d	0.5 0.625 0.625	0.625 0.125 0.562	210	0.5 0.625 0.625	64.0 -3.1 -5.1	6.0 238.4	0.5 0.625 0.625	61.0 0.4 3.7	3.7 83.2 10.1	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
761	G50B_062_025d	0.375 0.625 0.625	0.625 0.25 0.5	210	0.375 0.625 0.625	59.2 -6.3 -10.3	12.1 238.4	0.375 0.625 0.625	56.7 -5.3 -2.1	5.7 201.6 8.6	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
762	G50B_062_037d	0.25 0.625 0.625	0.625 0.375 0.437	210	0.25 0.625 0.625	54.3 -9.5 -15.5	18.2 238.4	0.25 0.625 0.625	51.9 -12.3 -8.5	14.9 214.7 7.9	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
763	G50B_062_050d	0.125 0.625 0.625	0.625 0.5 0.375	210	0.125 0.625 0.625	49.4 -12.7 -20.7	24.3 238.4	0.125 0.625 0.625	48.0 -18.0 -13.9	22.8 217.6 8.7	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
764	G50B_062_062d	0.0 0.625 0.625	0.625 0.625 0.312	210	0.0 0.625 0.625	44.6 -15.9 -25.9	30.4 238.4	0.0 0.625 0.625	43.3 -25.1 -20.1	32.1 218.6 10.9	210	0.0 1.0 1.0	95.6 -25.5 -41.5	48.7 238.4	
765	ROOY_100_050d	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	70.5 35.4 22.4	41.9 32.3	1.0 0.5 0.5	68.2 29.0 29.0	41.1 45.0 9.5	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3
766	ROOY_087_037d	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.5	67.9 26.6 16.8	31.4 32.3	0.875 0.5 0.5	65.3 24.5 25.2	35.1 45.7 9.0	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3
767	ROOY_075_025d	0.75 0.5 0.5	0.75 0.25 0.625	390	0.75 0.5 0.5	65.2 17.7 11.2	20.9 32.3	0.75 0.5 0.5	62.2 20.1 20.1	28.5 45.0 9.7	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3
768	ROOY_062_012d	0.625 0.5 0.5	0.625 0.125 0.562	390	0.625 0.5 0.5	62.6 8.8 5.6	10.4 32.3	0.625 0.5 0.5	58.7 14.9 15.6	21.6 46.3 12.3	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3
769	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	60.0 0.0 0.0	0.0 0.0	0.5 0.5 0.5	54.3 8.9 10.1	13.5 48.5 14.6	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	
770	G50B_050_012d	0.375 0.5 0.5	0.5 0.125 0.4												

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Technische Information: <http://www.ps.bam.de> oder <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		
810	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.6 0.0 0.1	0.1 116.7 0.1	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
811	BOOR_100_012d	0.875 0.875 1.0	1.0 0.125 0.937	270	0.875 0.875 1.0	86.8 3.6 -5.0	6.2 306.2	0.875 0.875 1.0	87.2 3.8 -5.3	6.6 305.3 0.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
812	BOOR_100_025d	0.75 0.75 1.0	1.0 0.25 0.875	270	0.75 0.75 1.0	77.9 7.3 -10.1	12.5 306.2	0.75 0.75 1.0	76.6 9.6 -10.6	14.3 312.1 2.6	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
813	BOOR_100_037d	0.625 0.625 1.0	1.0 0.375 0.812	270	0.625 0.625 1.0	69.1 11.0 -15.1	18.7 306.2	0.625 0.625 1.0	67.2 13.6 -15.6	20.8 311.0 3.2	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
814	BOOR_100_050d	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	60.3 14.7 -20.2	25.0 306.2	0.5 0.5 1.0	55.8 19.6 -21.4	29.1 312.4 6.7	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
815	BOOR_100_062d	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	51.5 18.4 -25.2	31.3 306.2	0.375 0.375 1.0	45.8 24.1 -26.3	35.7 312.5 8.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
816	BOOR_100_075d	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	42.7 22.1 -30.3	37.5 306.2	0.25 0.25 1.0	37.4 26.6 -31.6	41.3 310.1 6.9	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
817	BOOR_100_087d	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	33.9 25.8 -35.3	43.8 306.2	0.125 0.125 1.0	28.7 31.4 -36.1	47.8 311.0 7.6	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
818	BOOR_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 0.0 1.0	23.4 30.6 -39.6	50.1 307.6 2.0	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
819	Y00G_100_012d	1.0 1.0 0.875	1.0 0.125 0.937	90	1.0 1.0 0.875	94.6 -1.2 11.9	12.0 96.1	1.0 1.0 0.875	94.6 -2.5 9.9	10.2 104.1 2.3	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
820	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0	0.875 0.875 0.875	86.3 1.2 3.7	3.9 71.1 3.9	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
821	BOOR_087_012d	0.75 0.75 0.875	0.875 0.125 0.812	270	0.75 0.75 0.875	77.9 3.6 -5.0	6.2 306.2	0.75 0.75 0.875	76.0 6.9 -2.3	7.3 341.0 4.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
822	BOOR_087_025d	0.625 0.625 0.875	0.875 0.25 0.75	270	0.625 0.625 0.875	69.0 7.3 -10.1	12.5 306.2	0.625 0.625 0.875	66.7 11.0 -8.0	13.6 323.8 4.7	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
823	BOOR_087_037d	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.5 0.875	60.2 10.0 -15.1	18.7 306.2	0.5 0.5 0.875	55.5 16.6 -14.6	22.1 318.6 7.2	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
824	BOOR_087_050d	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	51.4 14.7 -20.2	25.0 306.2	0.375 0.375 0.875	45.6 21.0 -20.4	29.2 315.8 8.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
825	BOOR_087_062d	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.6 18.4 -25.2	31.3 306.2	0.25 0.25 0.875	37.1 23.2 -26.2	35.0 311.5 7.3	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
826	BOOR_087_075d	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	33.8 22.1 -30.3	37.5 306.2	0.125 0.125 0.875	29.0 26.9 -31.2	41.2 310.8 6.8	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
827	BOOR_087_087d	0.0 0.0 0.875	0.875 0.875 0.437	270	0.0 0.0 0.875	24.9 25.8 -35.3	43.8 306.2	0.0 0.0 0.875	23.4 26.1 -35.1	43.8 306.6 1.6	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
828	Y00G_100_025d	1.0 1.0 0.75	1.0 0.25 0.875	90	1.0 1.0 0.75	93.6 -2.5 23.8	24.0 96.1	1.0 1.0 0.75	93.5 -4.4 20.0	20.4 102.4 4.2	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
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 -1.2 11.9	12.0 96.1	0.875 0.875 0.75	85.2 -0.7 13.0	13.1 93.4 1.3	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
830	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0	0.75 0.75 0.75	75.1 4.6 6.6	6.1 54.7 8.5	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
831	BOOR_075_012d	0.625 0.625 0.75	0.75 0.125 0.687	270	0.625 0.625 0.75	68.9 3.6 -5.0	6.2 306.2	0.625 0.625 0.75	66.1 8.4 0.2	8.4 1.7 7.7	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
832	BOOR_075_025d	0.5 0.5 0.75	0.75 0.25 0.625	270	0.5 0.5 0.75	60.1 7.3 -10.1	12.5 306.2	0.5 0.5 0.75	54.8 13.8 -6.8	15.4 333.6 8.9	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
833	BOOR_075_037d	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	51.3 10.0 -15.1	18.7 306.2	0.375 0.375 0.75	45.6 17.2 -13.3	21.7 322.1 8.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
834	BOOR_075_050d	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	42.5 14.7 -20.2	25.0 306.2	0.25 0.25 0.75	37.2 19.3 -19.7	27.6 314.5 7.0	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
835	BOOR_075_062d	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	33.7 18.4 -25.2	31.3 306.2	0.125 0.125 0.75	29.3 22.6 -25.7	34.2 311.4 6.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
836	BOOR_075_075d	0.0 0.0 0.75	0.75 0.75 0.375	270	0.0 0.0 0.75	24.9 22.1 -30.3	37.5 306.2	0.0 0.0 0.75	23.6 21.0 -30.2	36.9 304.8 1.6	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
837	Y00G_100_037d	1.0 1.0 0.625	1.0 0.375 0.812	90	1.0 1.0 0.625	92.6 -3.8 35.8	36.0 96.1	1.0 1.0 0.625	92.4 -6.1 30.9	31.6 101.2 5.3	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
838	Y00G_087_025d	0.875 0.875 0.625	0.875 0.25 0.75	90	0.875 0.875 0.625	84.7 -2.5 23.8	24.0 96.1	0.875 0.875 0.625	84.2 -2.8 23.6	23.8 96.7 0.5	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
839	Y00G_075_012d	0.75 0.75 0.625	0.75 0.125 0.687	90	0.75 0.75 0.625	76.8 -1.2 11.9	12.0 96.1	0.75 0.75 0.625	74.4 2.4 16.3	16.5 81.4 6.2	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
840	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0	0.625 0.625 0.625	65.5 5.9 9.4	11.1 57.6 11.6	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
841	BOOR_062_012d	0.5 0.5 0.625	0.625 0.125 0.562	270	0.5 0.5 0.625	60.0 3.6 -5.0	6.2 306.2	0.5 0.5 0.625	54.5 11.4 1.1	11.4 5.8 11.3	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
842	BOOR_062_025d	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	51.2 7.3 -10.1	12.5 306.2	0.375 0.375 0.625	45.2 14.8 -6.0	16.0 337.7 10.3	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
843	BOOR_062_037d	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	42.4 10.0 -15.1	18.7 306.2	0.25 0.25 0.625	36.9 16.3 -13.2	21.0 320.9 7.8	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
844	BOOR_062_050d	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	33.6 14.7 -20.2	25.0 306.2	0.125 0.125 0.625	29.1 19.3 -19.9	27.7 314.1 6.3	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
845	BOOR_062_062d	0.0 0.0 0.625	0.625 0.625 0.312	270	0.0 0.0 0.625	24.8 18.4 -25.2	31.3 306.2	0.0 0.0 0.625	23.5 16.8 -24.9	30.0 304.0 2.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
846	Y00G_100_050d	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	91.7 -5.1 47.7	48.0 96.1	1.0 1.0 0.5	91.2 -7.6 43.4	44.1 100.0 5.0	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
847	Y00G_087_037d	0.875 0.875 0.5	0.875 0.375 0.687	90	0.875 0.875 0.5	83.7 -3.8 35.8	36.0 96.1	0.875 0.875 0.5	83.1 -4.5 35.6	35.8 97.2 0.9	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
848	Y00G_075_025d	0.75 0.75 0.5	0.75 0.25 0.625	90	0.75 0.75 0.5	75.8 -2.5 23.8	24.0 96.1	0.75 0.75 0.5	73.6 0.4 27.0	27.0 88.9 4.9	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
849	Y00G_062_012d	0.625 0.625 0.5	0.625 0.125 0.562	90	0.625 0.625 0.5	67.9 -1.2 11.9	12.0 96.1	0.625 0.625 0.5	64.7 3.9 19.0	19.4 78.1 9.3	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
850	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	60.0 0.0 0.0	0.0 0.0	0.5 0.5 0.5	54.3 9.1 9.8	13.4 47.1 14.5	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
851	BOOR_050_012d	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375									

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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
891	NW_100d	1.0 1.0 1.0	1.0 0.0 0.0	1.0 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	111.4 0.1 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	89.4 9.9 0.0	9.9 359.8	1.0 0.875 1.0	348.2 3.6 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
893	B50R_100_025d	1.0 0.75 1.0	1.0 0.25 0.875	330	1.0 0.75 1.0	83.2 19.8 0.0	19.8 359.8	1.0 0.75 1.0	351.1 4.9 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
894	B50R_100_037d	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	77.0 29.7 0.0	29.7 359.8	1.0 0.625 1.0	352.2 7.0 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
895	B50R_100_050d	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	70.8 39.6 -0.1	39.6 359.8	1.0 0.5 1.0	353.8 5.5 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
896	B50R_100_062d	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	64.6 49.5 -0.1	49.5 359.8	1.0 0.375 1.0	355.3 4.7 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
897	B50R_100_075d	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	58.4 59.4 -0.1	59.4 359.8	1.0 0.25 1.0	357.1 3.4 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
898	B50R_100_087d	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	52.3 69.4 -0.1	69.4 359.8	1.0 0.125 1.0	358.6 2.6 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
899	B50R_100_100d	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	1.0 0.0 1.0	359.5 0.0 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
900	GO0B_100_012d	0.875 1.0 0.875	1.0 0.125 0.937	150	0.875 1.0 0.875	89.9 -81.1 3.7	8.9 155.5	0.875 1.0 0.875	90.9 -5.6 5.6	7.9 135.3 3.2	149	50.0 -65.0 29.6 71.4 155.5
901	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	86.2 1.2 3.6	3.8 71.0 3.8	360	1.0 1.0 1.0 95.6 0.0 0.0 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	80.5 9.9 0.0	9.9 359.8	0.875 0.75 0.875	80.1 10.0 2.1	10.2 11.8 2.1	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
903	B50R_087_025d	0.875 0.625 0.875	0.875 0.25 0.75	330	0.875 0.625 0.875	74.3 19.8 0.0	19.8 359.8	0.875 0.625 0.875	74.6 18.0 0.9	18.1 2.9 2.0	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
904	B50R_087_037d	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	68.1 29.7 0.0	29.7 359.8	0.875 0.5 0.875	66.7 30.6 -0.6	30.6 358.7 1.7	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
905	B50R_087_050d	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	61.9 39.6 -0.1	39.6 359.8	0.875 0.375 0.875	60.5 40.8 -1.0	40.8 358.5 2.0	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
906	B50R_087_062d	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	55.7 49.5 -0.1	49.5 359.8	0.875 0.25 0.875	54.0 52.3 -1.0	52.3 358.7 3.3	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
907	B50R_087_075d	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	49.5 59.4 -0.1	59.4 359.8	0.875 0.125 0.875	47.7 64.4 -0.5	64.4 359.4 5.3	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
908	B50R_087_087d	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	43.4 69.4 -0.1	69.4 359.8	0.875 0.0 0.875	42.9 73.7 1.1	73.7 0.8 4.5	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
909	GO0B_100_025d	0.75 1.0 0.75	1.0 0.25 0.875	150	0.75 1.0 0.75	84.2 -16.2 7.4	17.8 155.5	0.75 1.0 0.75	85.6 -11.0 10.4	15.2 136.5 6.2	149	50.0 -65.0 29.6 71.4 155.5
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 -8.1 3.7	8.9 155.5	0.75 0.875 0.75	81.1 -4.3 8.3	9.4 117.5 5.9	149	50.0 -65.0 29.6 71.4 155.5
911	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	75.6 4.3 6.4	7.8 56.1 8.1	360	1.0 1.0 1.0 95.6 0.0 0.0 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	71.6 9.9 0.0	9.9 359.8	0.75 0.625 0.75	70.5 12.2 4.7	13.1 21.4 5.4	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
913	B50R_075_025d	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	65.4 19.8 0.0	19.8 359.8	0.75 0.5 0.75	63.2 23.9 2.7	24.1 6.6 5.4	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
914	B50R_075_037d	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	59.2 29.7 0.0	29.7 359.8	0.75 0.375 0.75	57.3 34.4 1.7	34.4 2.9 5.4	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
915	B50R_075_050d	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	53.0 39.6 -0.1	39.6 359.8	0.75 0.25 0.75	50.7 45.7 0.7	45.8 0.9 6.6	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
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 49.5 -0.1	49.5 359.8	0.75 0.125 0.75	44.9 57.7 0.1	57.7 0.1 8.4	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
917	B50R_075_075d	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	40.6 59.4 -0.1	59.4 359.8	0.75 0.0 0.75	40.3 67.0 1.0	67.0 0.8 7.6	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
918	GO0B_100_037d	0.625 1.0 0.625	1.0 0.375 0.812	150	0.625 1.0 0.625	78.5 -24.3 11.1	26.7 155.5	0.625 1.0 0.625	79.8 -17.2 15.5	23.2 137.8 8.5	149	50.0 -65.0 29.6 71.4 155.5
919	GO0B_087_025d	0.625 0.875 0.625	0.875 0.25 0.75	150	0.625 0.875 0.625	75.3 -16.2 7.4	17.8 155.5	0.625 0.875 0.625	76.0 -10.5 12.9	16.7 129.1 8.0	149	50.0 -65.0 29.6 71.4 155.5
920	GO0B_075_012d	0.625 0.75 0.625	0.75 0.125 0.687	150	0.625 0.75 0.625	72.1 -8.1 3.7	8.9 155.5	0.625 0.75 0.625	70.7 -2.0 10.9	11.1 100.3 9.5	149	50.0 -65.0 29.6 71.4 155.5
921	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	66.0 5.6 8.9	10.5 57.5 10.9	360	1.0 1.0 1.0 95.6 0.0 0.0 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	62.7 9.9 0.0	9.9 359.8	0.625 0.5 0.625	59.5 17.0 6.1	18.1 19.9 9.9	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
923	B50R_062_025d	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	56.5 19.8 0.0	19.8 359.8	0.625 0.375 0.625	53.7 26.9 4.3	27.3 9.1 8.8	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
924	B50R_062_037d	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	50.3 29.7 0.0	29.7 359.8	0.625 0.25 0.625	47.9 38.2 2.9	38.3 4.3 9.3	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
925	B50R_062_050d	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	44.1 39.6 -0.1	39.6 359.8	0.625 0.125 0.625	42.0 50.1 1.3	50.1 1.5 10.7	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
926	B50R_062_062d	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	37.9 49.5 -0.1	49.5 359.8	0.625 0.0 0.625	37.5 59.5 0.8	59.5 0.7 10.0	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
927	GO0B_100_050d	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	72.8 -32.5 14.8	35.7 155.5	0.5 1.0 0.5	73.8 -24.0 19.6	31.0 140.7 9.7	149	50.0 -65.0 29.6 71.4 155.5
928	GO0B_087_037d	0.5 0.875 0.5	0.875 0.375 0.687	150	0.5 0.875 0.5	69.6 -24.3 11.1	26.7 155.5	0.5 0.875 0.5	70.0 -18.0 17.2	24.9 136.3 8.8	149	50.0 -65.0 29.6 71.4 155.5
929	GO0B_075_025d	0.5 0.75 0.5	0.75 0.25 0.625	150	0.5 0.75 0.5	66.4 -16.2 7.4	17.8 155.5	0.5 0.75 0.5	65.3 -9.6 14.9	17.7 122.9 10.0	149	50.0 -65.0 29.6 71.4 155.5
930	GO0B_062_012d	0.5 0.625 0.5	0.625 0.125 0.562	150	0.5 0.625 0.5	63.2 -8.1 3.7	8.9 155.5	0.5 0.625 0.5	61.0 -2.3 12.4	12.6 100.7 10.6	149	50.0 -65.0 29.6 71.4 155.5
931	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	60.0 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	54.8 8.7 9.3	12.7 40.7 13.7	360	1.0 1.0 1.0 95.6 0.0 0.0 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	53.8 9.9 0.0	9.9 359.8	0.5 0.375 0.5	49.6 18.6 6.7	19.8 19.7 11.8	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
933	B50R_050_025d	0.5 0.25 0.5	0.5 0.25 0.375	330	0.5 0.25 0.5	47.6 19.8 0.0	19.8 359.8	0.5 0.25 0.5	44.1 29.4 4.1	29.7 7.9 11.0	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
934	B50R_050_037d	0.5 0.125 0.5	0.5 0.375 0.312	330	0.5 0.125 0.5	41.4 29.7 0.0	29.7 359.8	0.5 0.125 0.5	38.7 41.2 1.8	41.3 2.5 12.0	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
935	B50R_050_050d	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	35.2 39.6 -0.1	39.6 359.8	0.5 0.0 0.5	34.5 50.1 0.7	50.1 0.8 10.5	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
936	GO0B_100_062d	0.375 1.0 0.375	1.0 0.625 0.687	150	0.375 1.0 0.375	67.1 -40.6 18.5	44.6 155.5	0.375 1.0 0.375	67.5 -31.6 23.8	39.6 143.0 10.4	149	50.0 -65.0 29.6 71.4 155.5
937	GO0B_087_050d	0.375 0.875 0.375	0.875 0.5 0.625	150	0.375 0.875 0.375	63.9 -32.5 14.8	35.7 155.5	0.375 0.875 0.375	64.2 -26.0 21.1	33.5 140.9 9.0	149	50.0 -65.0 29.6 71.4 155.5
938	GO0B_075_037d	0.375 0.75 0.375	0.75 0.375 0.562	150	0.375 0.75 0.375	60.7 -24.3 11.1	26.7 155.5	0.375 0.75 0.375	60.0 -17.7 18.6	25.7 133.4 10.1	149	50.0 -65.0 29.6 71.4 155.5
939	GO0B_062_025d	0.375 0.625 0.375	0.625 0.25 0.5	150	0.375 0.625 0.375	57.5 -16.2 7.4	17.8 155.5	0.375 0.625 0.375	56.3 -10.4 16.0	19.1 123.1 10.4	149	50.0 -65.0 29.6 71.4 155.5
940	GO0B_050_012d	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	54.3 -8.1 3.7	8.9 155.5	0.375 0.5 0.375	50.7 0.3 12.7	12.7 88.6 12.8	149	50.0 -65.0 29.6 71.4 155.5
941	NW_037d	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	51.0 0.0 0.0	0.0 0.0 0.0	0.375 0.375 0.375	45.5 9.7 9.2	13.4 43.5 14.5	360	1.0 1.0 1.0 95.6 0.0 0.0 0.0 0.0
942	B50R_037_012d	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.25 0.375	44.9 9.9 0.0	9.9 359.8	0.375 0.25 0.375	40.5 20.4 6.1	21.3 16.7 12.9	330	1.0 0.0 1.0 46.1 79.3 -0.2 79.3 359.8
943	B50R_037_025d	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.125 0.375	38.7 19.8 0.0	19.8 359.8	0.375 0.125 0.375	35.5 31.3 3.0	31.5 5.4 12.3	330	1.0 0.0 1.0 46.1 79.

Siehe ähnliche Dateien: <http://130.149.60.45/~farbmetrik/TG77/TG77L0NA.TXT> /PS
Technische Information: <http://www.ps.bam.de> oder <http://130.149.60.45/~farbmetrik>

		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_000a	0.0	0.0	0.0	0.0	0.0	0.0	0.0	360	0.0	0.0	0.0	24.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0

Siehe ähnliche Dateien: <http://130.149.60.45/~farbmetrik/TG77/TG77.HTM>
Technische Information: <http://www.ps.bam.de> oder <http://130.149.60.45/~farbmetrik>

TUB-Registrierung: 20150901-TG77/TG77L0NA.TXT /.PS TUB-Material: Code=rh4ta
Anwendung für Messung von Offsetdruck-Ausgabe, Separation cmyñ6 (CMY0)

	HIC*Fd	rgb_Fd			ict_Fd			hsi_Fd			rgb*Fd			LabCh*Fd			rgb**Fd			LabCh**Fd			DE*Fd			hsiM,d	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	86.0	0.0	0.0	0.0	0.0	0.866	0.866	0.866	86.1	1.2	3.4	3.7	69.9	3.7	360	1.0	1.0	1.0	95.6	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	90.8	0.0	0.0	0.0	0.0	0.933	0.933	0.933	90.8	0.4	1.4	1.5	71.6	1.5	360	1.0	1.0	1.0	95.6	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	95.6	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.6	0.0	0.1	0.1	114.3	0.1	360	1.0	1.0	1.0	95.6	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	24.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	23.0	0.7	-0.9	1.1	308.5	1.7	360	1.0	1.0	1.0	95.6	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	29.0	0.0	0.0	0.0	0.0	0.066	0.066	0.066	25.6	5.5	0.6	5.5	6.7	6.5	360	1.0	1.0	1.0	95.6	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	33.8	0.0	0.0	0.0	0.0	0.133	0.133	0.133	28.2	8.3	3.4	9.0	22.4	10.6	360	1.0	1.0	1.0	95.6	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	38.6	0.0	0.0	0.0	0.0	0.2	0.2	0.2	32.0	10.0	5.8	11.6	30.4	13.3	360	1.0	1.0	1.0	95.6	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	43.3	0.0	0.0	0.0	0.0	0.266	0.266	0.266	36.7	8.8	8.7	12.4	44.7	14.0	360	1.0	1.0	1.0	95.6	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	48.1	0.0	0.0	0.0	0.0	0.333	0.333	0.333	40.7	10.4	8.9	13.7	40.4	15.5	360	1.0	1.0	1.0	95.6	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	52.8	0.0	0.0	0.0	0.0	0.4	0.4	0.4	46.8	8.7	10.2	13.4	49.7	14.7	360	1.0	1.0	1.0	95.6	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	57.5	0.0	0.0	0.0	0.0	0.466	0.466	0.466	51.8	8.8	9.9	13.3	48.4	14.5	360	1.0	1.0	1.0	95.6	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	62.3	0.0	0.0	0.0	0.0	0.533	0.533	0.533	57.5	7.3	9.2	11.8	51.6	12.7	360	1.0	1.0	1.0	95.6	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	67.1	0.0	0.0	0.0	0.0	0.6	0.6	0.6	63.6	6.0	9.2	11.0	56.7	11.5	360	1.0	1.0	1.0	95.6	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	71.8	0.0	0.0	0.0	0.0	0.666	0.666	0.666	69.3	5.2	8.3	9.8	57.5	10.1	360	1.0	1.0	1.0	95.6	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	76.6	0.0	0.0	0.0	0.0	0.734	0.734	0.734	74.5	4.8	6.5	8.1	53.5	8.3	360	1.0	1.0	1.0	95.6	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	81.3	0.0	0.0	0.0	0.0	0.8	0.8	0.8	80.5	2.7	5.2	5.9	62.0	5.9	360	1.0	1.0	1.0	95.6	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	86.0	0.0	0.0	0.0	0.0	0.866	0.866	0.866	86.1	1.2	3.4	3.6	69.4	3.6	360	1.0	1.0	1.0	95.6	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	90.8	0.0	0.0	0.0	0.0	0.933	0.933	0.933	90.7	0.4	1.4	1.5	71.7	1.5	360	1.0	1.0	1.0	95.6	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	95.6	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.7	0.0	0.0	0.1	118.4	0.1	360	1.0	1.0	1.0	95.6	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	24.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	23.3	1.3	-2.4	2.8	299.2	2.9	360	1.0	1.0	1.0	95.6	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	95.6	0.0	0.0	0.0	0.0	1.0	1.0	1.0	95.7	0.0	0.0	0.0	138.7	0.0	360	1.0	1.0	1.0	95.6	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	45.4	70.9	44.8	83.9	32.3	1.0	0.0	0.0	45.4	70.5	45.5	83.9	32.8	0.7	389	1.0	0.0	0.0	45.4	70.9	44.8	83.9	32.3
1075	G50B_100_100d	0.0	1.0	1.0	1.0	1.0	0.5	210	0.0	1.0	1.0	56.8	-25.5	-41.5	48.7	238.4	0.0	1.0	1.0	56.4	-25.2	-41.8	48.8	238.9	0.5	210	0.0	1.0	1.0	56.8	-25.5	-41.5	48.7	238.4
1076	Y00G_100_100d	1.0	1.0	0.0	1.0	1.0	0.5	90	1.0	1.0	0.0	87.8	-10.2	95.4	96.0	96.1	1.0	1.0	0.0	87.5	-10.0	95.1	95.7	96.0	0.4	89	1.0	1.0	0.0	87.8	-10.2	95.4	96.0	96.1
1077	B00R_100_100d	0.0	0.0	1.0	1.0	1.0	0.5	270	0.0	0.0	1.0	25.0	29.5	-40.4	50.0	306.2	0.0	0.0	1.0	24.7	29.8	-40.1	49.9	306.6	0.5	270	0.0	0.0	1.0	25.0	29.5	-40.4	50.0	306.2
1078	G00B_100_100d	0.0	1.0	0.0	1.0	1.0	0.5	150	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5	0.0	1.0	0.0	49.2	-65.4	28.0	71.2	156.7	1.8	149	0.0	1.0	0.0	50.0	-65.0	29.6	71.4	155.5
1079	B50R_100_100d	1.0	0.0	1.0	1.0	1.0	0.5	330	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8	1.0	0.0	1.0	45.8	79.2	-0.2	79.2	359.8	0.2	330	1.0	0.0	1.0	46.1	79.3	-0.2	79.3	359.8

delta E* = 5.8

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TG770-7N, Seite 22/22-F

Prüfvorlage TG77; ME16(ISO 9241-306), 3(ISO/IEC 15775)
Farben und Farbabstände, ΔE^* , 3D=0, de=0, cmyk

Eingabe: *rgb/cmyk* -> *rgb_d*
Ausgabe: Transfer nach *cmyk_d*

0-0032131-F0