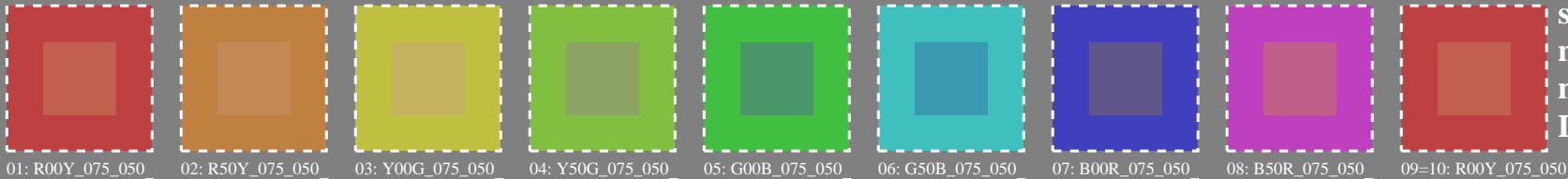


test no 2 pour un rendu de couleurs – couleurs métamères pour D65 et D50; écran standard (sRGB)



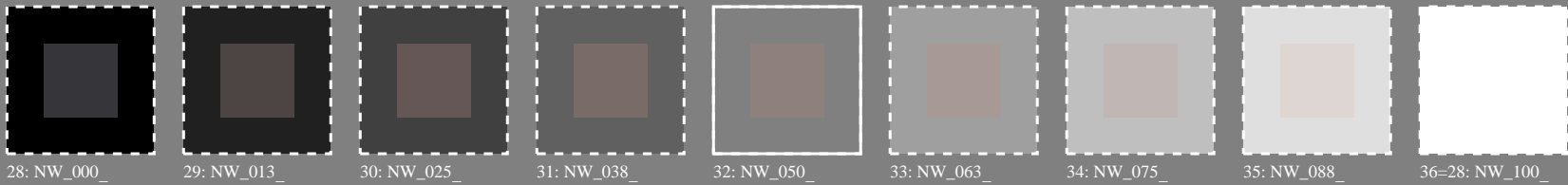
series :
métamère
m
D65



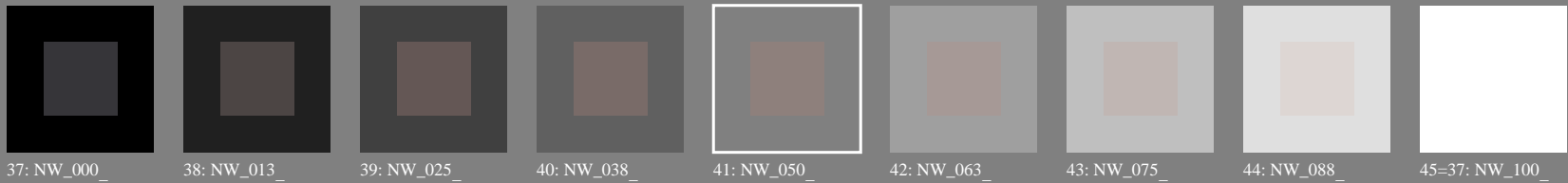
central
z
D65/D50

couleurs métamères uniquement possibles pour impression offset
et les imprimantes avec du mois quatre couleurs, *CMYK* ou *CMY0*

métamère
m
D50



métamère
m
D65



gris
g
D65/D50

couleurs métamères uniquement possibles pour impression offset
et les imprimantes avec du mois quatre couleurs, *CMYK* ou *CMY0*

métamère
m
D50

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF21/PF21.HTM>
informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

TUB enregistrement: 20130201 -PF21/PF21L0NP.PDF /.PS
application pour la mesure de sortie sur écran
TUB matériel: code=rh4ta

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF21/PF21.HTM>
informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

TUB enregistrement: 20130201 -PF21/PF21L0NP.PDF /.PS
application pour la mesure de sortie sur écran, aucune séparation
TUB matériel: code=rh4ta

test no 2 pour un rendu de couleurs – couleurs métamères pour D65 et D50; écran standard (sRGB); *rgb*→*rgb*d*



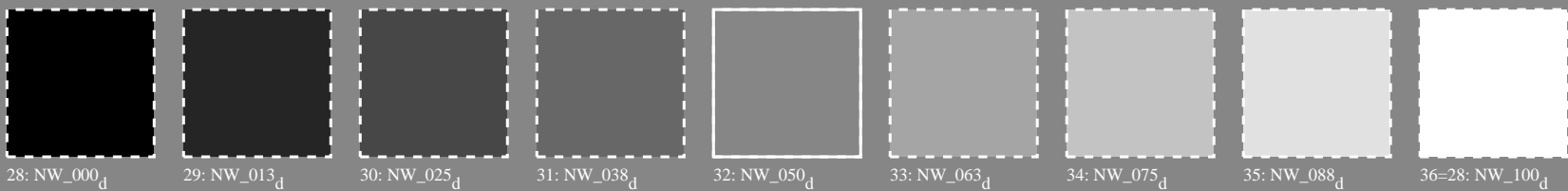
series :
métamère
m
D65



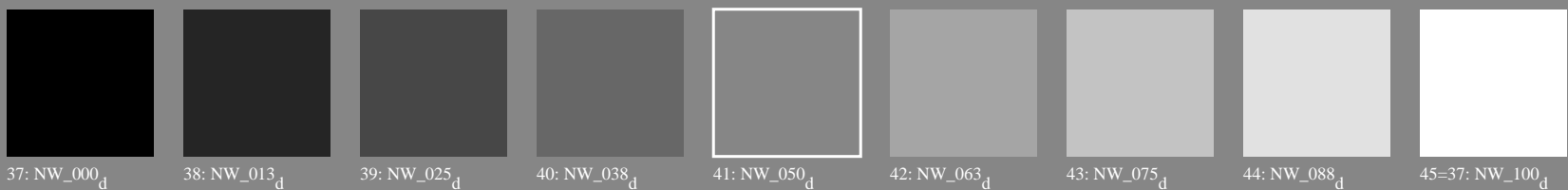
central
z
D65/D50

couleurs métamères uniquement possibles pour impression offset
et les imprimantes avec du mois quatre couleurs, *CMYK* ou *CMY0*

métamère
m
D50



métamère
m
D65
Lab*W=95.4, 0.0, 0.0



gris
g
D65/D50

couleurs métamères uniquement possibles pour impression offset
et les imprimantes avec du mois quatre couleurs, *CMYK* ou *CMY0*

métamère
m
D50

nj	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Ma	LabCh*Ma
0/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5 100.4 39.9 0.0	389	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0	
1/657	R13Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	51.4 74.1 64.9 98.5 41.2	1.0 0.125 0.0	51.5 73.9 64.9 98.3 41.3 0.2	36	1.0 0.116 0.0	51.4 74.1 64.9 98.5 41.2	
2/666	R25Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.7 67.6 65.8 94.4 44.2	1.0 0.25 0.0	54.0 66.7 65.9 93.8 44.6 1.0	42	1.0 0.233 0.0	53.7 67.6 65.8 94.4 44.2	
3/675	R38Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	57.9 56.2 67.9 88.1 50.3	1.0 0.375 0.0	58.2 55.4 67.9 87.7 50.7 0.7	51	1.0 0.366 0.0	57.9 56.2 67.9 88.1 50.3	
4/684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	63.6 41.3 71.0 82.2 59.7	1.0 0.5 0.0	63.6 41.3 71.0 82.2 59.7 0.0	59	1.0 0.5 0.0	63.6 41.3 71.0 82.2 59.7	
5/693	R63Y_100_100a	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	70.5 24.7 75.4 79.4 71.8	1.0 0.625 0.0	70.1 25.8 75.0 79.3 71.0 1.2	68	1.0 0.633 0.0	70.5 24.7 75.4 79.4 71.8	
6/702	R75Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	78.2 7.8 80.6 81.0 84.4	1.0 0.75 0.0	77.2 9.8 79.7 80.3 82.9 2.3	77	1.0 0.766 0.0	78.2 7.8 80.6 81.0 84.4	
7/711	R88Y_100_100a	1.0 0.875 0.0	1.0 1.0 0.5	83	1.0 0.883 0.0	85.3 -6.7 85.5 85.8 94.4	1.0 0.875 0.0	84.8 -5.7 85.0 85.2 93.8 1.1	83	1.0 0.883 0.0	85.3 -6.7 85.5 85.8 94.4	
8/720	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	92.6 -20.7 90.7 93.0 102.8	1.0 1.0 0.0	92.6 -20.6 90.7 93.0 102.8 0.0	89	1.0 1.0 0.0	92.6 -20.7 90.7 93.0 102.8	
9/639	Y13G_100_100a	0.875 1.0 0.0	1.0 1.0 0.5	97	0.883 1.0 0.0	90.5 -32.2 88.3 94.0 110.0	0.875 1.0 0.0	90.4 -33.0 88.1 94.1 110.5 0.8	96	0.883 1.0 0.0	90.5 -32.2 88.3 94.0 110.0	
10/558	Y25G_100_100a	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	88.7 -43.3 86.2 96.5 116.6	0.75 1.0 0.0	88.5 -44.9 85.8 96.8 117.6 1.6	102	0.766 1.0 0.0	88.7 -43.3 86.2 96.5 116.6	
11/477	Y38G_100_100a	0.625 1.0 0.0	1.0 1.0 0.5	112	0.633 1.0 0.0	87.0 -55.0 84.1 105.1 123.2	0.625 1.0 0.0	86.9 -55.7 83.9 105.0 123.6 0.7	111	0.633 1.0 0.0	87.0 -55.0 84.1 105.1 123.2	
12/396	Y50G_100_100a	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	85.7 -65.2 82.4 100.5 128.3	0.5 1.0 0.0	85.7 -65.2 82.4 100.7 128.3 0.0	119	0.5 1.0 0.0	85.7 -65.2 82.4 100.5 128.3	
13/315	Y63G_100_100a	0.375 1.0 0.0	1.0 1.0 0.5	128	0.366 1.0 0.0	84.7 -73.2 81.2 109.3 132.0	0.375 1.0 0.0	84.7 -72.8 81.2 109.1 131.8 0.3	128	0.366 1.0 0.0	84.7 -73.2 81.2 109.3 132.0	
14/234	Y75G_100_100a	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	84.0 -78.7 80.4 112.5 134.3	0.25 1.0 0.0	84.1 -78.2 80.4 112.2 134.1 0.4	137	0.233 1.0 0.0	84.0 -78.7 80.4 112.5 134.3	
15/153	Y88G_100_100a	0.125 1.0 0.0	1.0 1.0 0.5	143	0.116 1.0 0.0	83.7 -81.5 80.0 114.2 135.5	0.125 1.0 0.0	83.7 -81.4 80.0 114.2 135.5 0.1	143	0.116 1.0 0.0	83.7 -81.5 80.0 114.2 135.5	
16/72	G00C_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	83.6 -82.7 79.8 115.0 136.0	0.0 1.0 0.0	83.6 -82.7 79.8 115.0 136.0 0.0	149	0.0 1.0 0.0	83.6 -82.7 79.8 115.0 136.0	
17/73	G13C_100_100a	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.116	83.6 -82.1 76.8 112.5 136.9	0.0 1.0 0.125	83.6 -82.1 76.5 112.3 137.0 0.2	156	0.0 1.0 0.116	83.6 -82.1 76.8 112.5 136.9	
18/74	G25C_100_100a	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.233	83.7 -80.8 70.1 106.9 139.0	0.0 1.0 0.25	83.8 -80.5 69.1 106.1 139.3 1.0	162	0.0 1.0 0.233	83.7 -80.8 70.1 106.9 139.0	
19/75	G38C_100_100a	0.0 1.0 0.375	1.0 1.0 0.5	172	0.0 1.0 0.366	84.0 -78.0 58.8 97.7 142.9	0.0 1.0 0.375	84.0 -77.7 58.1 97.1 143.2 0.7	171	0.0 1.0 0.366	84.0 -78.0 58.8 97.7 142.9	
20/76	G50C_100_100a	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	84.3 -73.7 44.9 86.4 148.6	0.0 1.0 0.5	84.3 -73.7 44.9 86.3 148.6 0.0	180	0.0 1.0 0.5	84.3 -73.7 44.9 86.4 148.6	
21/77	G63C_100_100a	0.0 1.0 0.625	1.0 1.0 0.5	188	0.0 1.0 0.633	84.8 -68.1 29.5 74.3 156.5	0.0 1.0 0.625	84.7 -68.5 30.6 75.0 155.9 1.1	188	0.0 1.0 0.633	84.8 -68.1 29.5 74.3 156.5	
22/78	G75C_100_100a	0.0 1.0 0.75	1.0 1.0 0.5	196	0.0 1.0 0.766	85.4 -61.2 13.7 62.8 167.3	0.0 1.0 0.75	85.3 -62.0 15.8 64.0 165.6 2.3	197	0.0 1.0 0.766	85.4 -61.2 13.7 62.8 167.3	
23/79	G88C_100_100a	0.0 1.0 0.875	1.0 1.0 0.5	203	0.0 1.0 0.883	86.1 -54.1 0.0 54.1 180.0	0.0 1.0 0.875	86.0 -54.5 1.0 54.5 178.8 1.1	203	0.0 1.0 0.883	86.1 -54.1 0.0 54.1 180.0	
24/80	C00B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	86.8 -46.1 -13.5 48.1 196.3	0.0 1.0 1.0	86.8 -46.1 -13.5 48.1 196.3 0.0	210	0.0 1.0 1.0	86.8 -46.1 -13.5 48.1 196.3	
25/71	C13B_100_100a	0.0 0.875 1.0	1.0 1.0 0.5	217	0.0 0.883 1.0	78.5 -33.4 -26.3 42.5 218.2	0.0 0.875 1.0	77.9 -32.3 -27.0 42.1 219.8 1.3	216	0.0 0.883 1.0	78.5 -33.4 -26.3 42.5 218.2	
26/62	C25B_100_100a	0.0 0.75 1.0	1.0 1.0 0.5	224	0.0 0.766 1.0	70.2 -19.5 -39.3 43.9 243.6	0.0 0.75 1.0	69.1 -17.0 -40.7 44.1 247.2 3.0	222	0.0 0.766 1.0	70.2 -19.5 -39.3 43.9 243.6	
27/53	C38B_100_100a	0.0 0.625 1.0	1.0 1.0 0.5	232	0.0 0.633 1.0	60.9 -1.5 -53.9 53.9 268.3	0.0 0.625 1.0	60.3 -0.1 -54.6 54.6 269.8 1.7	231	0.0 0.633 1.0	60.9 -1.5 -53.9 53.9 268.3	
28/44	C50B_100_100a	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	51.7 18.3 -68.3 70.7 285.0	0.0 0.5 1.0	51.7 18.3 -68.3 70.7 285.0 0.0	240	0.0 0.5 1.0	51.7 18.3 -68.3 70.7 285.0	
29/35	C63B_100_100a	0.0 0.375 1.0	1.0 1.0 0.5	248	0.0 0.366 1.0	43.4 38.7 -82.0 90.7 295.3	0.0 0.375 1.0	43.8 37.6 -81.2 89.5 294.8 1.4	248	0.0 0.366 1.0	43.4 38.7 -82.0 90.7 295.3	
30/26	C75B_100_100a	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.233 1.0	36.5 57.6 -93.4 109.7 301.6	0.0 0.25 1.0	37.1 55.9 -92.3 107.9 301.1 2.1	257	0.0 0.233 1.0	36.5 57.6 -93.4 109.7 301.6	
31/17	C88B_100_100a	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.116 1.0	32.3 70.0 -100.3 122.3 304.9	0.0 0.125 1.0	32.4 69.6 -100.0 121.9 304.8 0.5	263	0.0 0.116 1.0	32.3 70.0 -100.3 122.3 304.9	
32/8	B00M_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2 0.0	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2	
33/89	B13M_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	30.9 76.2 -102.5 127.8 306.6	0.125 0.0 1.0	31.0 76.2 -102.5 127.7 306.6 0.0	276	0.116 0.0 1.0	30.9 76.2 -102.5 127.8 306.6	
34/170	B25M_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	32.3 76.7 -100.1 126.2 307.4	0.25 0.0 1.0	32.6 76.8 -99.8 125.9 307.5 0.4	282	0.233 0.0 1.0	32.3 76.7 -100.1 126.2 307.4	
35/251	B38M_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	34.9 77.9 -95.7 123.4 309.1	0.375 0.0 1.0	35.1 77.9 -95.5 123.3 309.2 0.3	291	0.366 0.0 1.0	34.9 77.9 -95.7 123.4 309.1	
36/332	B50M_100_100a	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	38.5 79.8 -89.7 120.0 311.6	0.5 0.0 1.0	38.5 79.8 -89.7 120.1 311.6 0.0	300	0.5 0.0 1.0	38.5 79.8 -89.7 120.0 311.6	
37/413	B63M_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	43.0 82.7 -82.2 116.6 315.1	0.625 0.0 1.0	42.7 82.5 -82.8 116.8 314.8 0.6	308	0.633 0.0 1.0	43.0 82.7 -82.2 116.6 315.1	
38/494	B75M_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	47.9 86.4 -74.0 113.8 319.4	0.75 0.0 1.0	47.2 85.8 -75.1 114.1 318.8 1.3	317	0.766 0.0 1.0	47.9 86.4 -74.0 113.8 319.4	
39/575	B88M_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	52.5 90.1 -66.3 111.9 323.6	0.875 0.0 1.0	52.1 89.8 -66.9 112.0 323.3 0.7	323	0.883 0.0 1.0	52.5 90.1 -66.3 111.9 323.6	
40/656	M00R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	57.2 94.3 -58.4 110.9 328.2	1.0 0.0 1.0	57.2 94.3 -58.4 111.0 328.2 0.0	330	1.0 0.0 1.0	57.2 94.3 -58.4 110.9 328.2	
41/655	M13R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	55.7 90.6 -44.8 101.1 333.6	1.0 0.0 0.875	55.6 90.3 -43.9 100.4 334.0 0.9	336	1.0 0.0 0.883	55.7 90.6 -44.8 101.1 333.6	
42/654	M25R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	54.4 87.3 -30.6 92.5 340.6	1.0 0.0 0.75	54.2 86.7 -28.6 91.3 341.6 2.0	342	1.0 0.0 0.766	54.4 87.3 -30.6 92.5 340.6	
43/653	M38R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	53.0 83.9 -13.6 85.0 350.7	1.0 0.0 0.625	53.0 83.6 -12.6 84.6 351.4 1.0	351	1.0 0.0 0.633	53.0 83.9 -13.6 85.0 350.7	
44/652	M50R_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	52.0 81.1 4.1 81.2 2.9	1.0 0.0 0.5	52.0 81.1 4.1 81.2 2.9 0.0	360	1.0 0.0 0.5	52.0 81.1 4.1 81.2 2.9	
45/651	M63R_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	51.3 79.3 22.7 82.5 16.0	1.0 0.0 0.375	51.3 79.2 21.6 82.1 15.2 1.1	368	1.0 0.0 0.366	51.3 79.3 22.7 82.5 16.0	
46/650	M75R_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	50.8 78.0 41.2 88.2 27.8	1.0 0.0 0.25	50.8 77.9 39.2 87.2 26.6 2.0	377	1.0 0.0 0.233	50.8 78.0 41.2 88.2 27.8	
47/649	M88R_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	50.5 77.2 55.6 95.1 35.7	1.0 0.0 0.125	50.6 77.2 54.9 94.8 35.4 0.6	383	1.0 0.0 0.116	50.5 77.2 55.6 95.1 35.7	
48/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5 100.4 39.9 0.0	389	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0	

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF21/PF21.HTM>
 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

nj	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Ma	LabCh*Ma		
0/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5	100.4 39.9 0.0	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
1/666	R25Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.7 67.6 65.8	94.4 44.2	1.0 0.25 0.0	54.0 66.7 65.9	93.8 44.6 1.0	42	1.0 0.233 0.0	53.7 67.6 65.8	94.4 44.2
2/684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	63.6 61.3 71.0	82.2 59.7	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7 0.0	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
3/702	R75Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	78.2 78.8 80.6	81.0 84.4	1.0 0.75 0.0	77.2 9.8 79.7	80.3 82.9 2.3	77	1.0 0.766 0.0	78.2 78.8 80.6	81.0 84.4
4/720	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8	1.0 1.0 0.0	92.6 -20.6 90.7	93.0 102.8 0.0	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
5/558	Y25G_100_100a	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	88.7 -43.3 86.2	96.5 116.6	0.75 1.0 0.0	88.5 -44.9 85.8	96.8 117.6 1.6	102	0.766 1.0 0.0	88.7 -43.3 86.2	96.5 116.6
6/396	Y50G_100_100a	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3 0.0	119	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
7/234	Y75G_100_100a	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	84.0 -78.7 80.4	112.5 134.3	0.25 1.0 0.0	84.1 -78.2 80.4	112.3 134.1 0.4	137	0.233 1.0 0.0	84.0 -78.7 80.4	112.5 134.3
8/72	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0 0.0	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
9/72	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0 0.0	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
10/76	G25B_100_100a	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	84.3 -73.7 44.9	86.4 148.6	0.0 1.0 0.5	84.3 -73.7 44.9	86.3 148.6 0.0	180	0.0 1.0 0.5	84.3 -73.7 44.9	86.4 148.6
11/80	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3 0.0	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
12/44	G75B_100_100a	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	81.7 18.3 -68.3	70.7 285.0	0.0 0.5 1.0	81.7 18.3 -68.3	70.7 285.0 0.0	240	0.0 0.5 1.0	81.7 18.3 -68.3	70.7 285.0
13/8	B00M_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2 0.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
14/332	B25R_100_100a	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6	0.5 0.0 1.0	38.5 79.8 -89.7	120.1 311.6 0.0	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
15/656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	1.0 0.0 1.0	57.2 94.3 -58.4	111.0 328.2 0.0	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
16/652	B75R_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9 0.0	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
17/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
18/688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	72.9 38.4 32.2	50.2 40.0	1.0 0.5 0.5	64.7 46.4 21.9	51.3 25.2 15.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
19/706	R50Y_100_050a	1.0 0.75 0.5	1.0 0.5 0.75	60	1.0 0.75 0.5	79.5 20.6 35.5	41.1 59.7	1.0 0.75 0.5	78.0 15.0 39.2	42.0 69.0 6.9	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
20/724	Y00G_100_050a	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	94.0 -10.3 45.3	46.5 102.8	1.0 1.0 0.5	93.2 -15.9 57.8	59.9 105.3 13.6	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
21/562	Y50G_100_050a	0.75 1.0 0.5	1.0 0.5 0.75	120	0.75 1.0 0.5	90.5 -32.6 41.2	52.5 128.3	0.75 1.0 0.5	89.1 -38.7 51.9	64.8 126.7 12.4	119	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
22/400	G00B_100_050a	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	89.5 -41.3 39.9	57.5 136.0	0.5 1.0 0.5	86.3 -57.6 47.9	75.0 140.2 18.4	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
23/404	G50B_100_050a	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	91.1 -23.0 -6.7	24.0 196.3	0.5 1.0 1.0	88.8 -33.9 -10.4	35.4 197.1 11.6	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
24/368	B00R_100_050a	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	62.8 38.0 -51.7	64.2 306.2	0.5 0.5 1.0	56.0 31.9 -61.1	69.0 297.5 13.0	270	0.0 1.0 1.0	30.3 76.0 -103.5	128.5 306.2
25/692	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	76.3 47.1 -29.2	55.4 328.2	1.0 0.5 1.0	68.6 62.6 -40.5	74.6 327.0 20.6	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
26/688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	72.9 38.4 32.2	50.2 40.0	1.0 0.5 0.5	64.7 46.4 21.9	51.3 25.2 15.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
27/506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	49.0 38.4 32.2	50.2 40.0	0.75 0.25 0.25	43.3 48.9 27.4	56.0 29.2 12.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
28/524	R50Y_075_050a	0.75 0.5 0.25	0.75 0.5 0.5	60	0.75 0.5 0.25	55.6 20.6 35.5	41.1 59.7	0.75 0.5 0.25	55.8 17.8 42.0	45.6 66.9 7.1	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
29/542	Y00G_075_050a	0.75 0.75 0.25	0.75 0.5 0.5	90	0.75 0.75 0.25	70.1 -10.3 45.3	46.5 102.8	0.75 0.75 0.25	71.7 -14.8 58.9	60.8 104.1 14.4	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
30/380	Y50G_075_050a	0.5 0.75 0.25	0.75 0.5 0.5	120	0.5 0.75 0.25	66.7 -32.6 41.2	52.5 128.3	0.5 0.75 0.25	67.6 -39.2 53.4	66.3 126.3 13.9	119	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
31/218	G00B_075_050a	0.25 0.75 0.25	0.75 0.5 0.5	150	0.25 0.75 0.25	65.6 -41.3 39.9	57.5 136.0	0.25 0.75 0.25	65.2 -56.7 50.2	75.8 138.5 18.5	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
32/222	G50B_075_050a	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.75	67.2 -23.0 -6.7	24.0 196.3	0.25 0.75 0.75	67.5 -32.5 -9.7	33.9 196.7 9.8	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
33/186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	39.0 38.0 -51.7	64.2 306.2	0.25 0.25 0.75	32.9 38.5 -64.1	74.8 301.0 13.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
34/510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	52.5 47.1 -29.2	55.4 328.2	0.75 0.25 0.75	47.5 63.1 -39.9	74.6 327.6 19.8	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
35/506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	49.0 38.4 32.2	50.2 40.0	0.75 0.25 0.25	43.3 48.9 27.4	56.0 29.2 12.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
36/324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.0	25.2 38.4 32.2	50.2 40.0	0.5 0.0 0.0	23.7 46.0 35.7	58.2 37.8 8.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
37/342	R50Y_050_050a	0.5 0.25 0.0	0.5 0.5 0.25	60	0.5 0.25 0.0	31.8 20.6 35.5	41.1 59.7	0.5 0.25 0.0	32.3 22.9 42.9	48.6 61.8 7.7	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
38/360	Y00G_050_050a	0.5 0.5 0.0	0.5 0.5 0.25	90	0.5 0.5 0.0	46.3 -10.3 45.3	46.5 102.8	0.5 0.5 0.0	48.9 -12.3 54.2	55.6 102.8 9.5	89	1.0 1.0 0.0	92.6 -20.7 90.7	93.0 102.8
39/198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	42.8 -32.6 41.2	52.5 128.3	0.25 0.5 0.0	44.9 -37.9 49.4	62.3 127.5 10.0	119	0.5 1.0 0.0	85.7 -65.2 82.4	105.1 128.3
40/36	G00B_050_050a	0.0 0.5 0.0	0.5 0.5 0.25	150	0.0 0.5 0.0	41.8 -41.3 39.9	57.5 136.0	0.0 0.5 0.0	43.5 -49.5 47.7	68.8 136.0 11.4	149	0.0 1.0 0.0	83.6 -82.7 79.8	115.0 136.0
41/40	G50B_050_050a	0.0 0.5 0.5	0.5 0.5 0.25	210	0.0 0.5 0.5	43.4 -23.0 -6.7	24.0 196.3	0.0 0.5 0.5	45.5 -27.6 -8.1	28.7 196.3 5.1	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3
42/4	B00R_050_050a	0.0 0.0 0.5	0.5 0.5 0.25	270	0.0 0.0 0.5	15.1 38.0 -51.7	64.2 306.2	0.0 0.0 0.5	11.7 45.5 -61.9	76.8 306.2 13.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
43/328	B50R_050_050a	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	28.6 47.1 -29.2	55.4 328.2	0.5 0.0 0.5	27.8 56.4 -34.9	66.3 328.2 10.9	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
44/324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.0	25.2 38.4 32.2	50.2 40.0	0.5 0.0 0.0	23.7 46.0 35.7	58.2 37.8 8.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
45/0	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0
46/91	NW_013a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	11.9 0.0 0.0	0.0 0.0	0.125 0.125 0.125	11.0 0.0 0.0	0.0 0.0 325.7	0.8	360	1.0 1.0 1.0	95.4 0.

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n=j	HIC*Fa	rgb_Fa	icr_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	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 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0
1	BOOR_012_012a	0.0 0.0 0.125	0.125 0.125 0.125	0.062 0.062 0.062	0.0 0.0 0.125	3.7 9.5	-12.9 16.0 306.2	0.0 0.0 0.125	0.8 5.8	-15.5 16.6	290.4 5.4 270	30.3 76.0
2	BOOR_025_025a	0.0 0.0 0.25	0.25 0.25 0.25	0.125 0.125 0.125	0.0 0.0 0.25	7.5 19.0	-25.8 32.1 306.2	0.0 0.0 0.25	2.9 20.6	-35.3 40.9	300.2 10.6 270	30.3 76.0
3	BOOR_037_037a	0.0 0.0 0.375	0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.0 0.375	11.3 28.5	-38.8 48.1 306.2	0.0 0.0 0.375	6.7 36.7	-50.3 62.3	306.1 14.9 270	30.3 76.0
4	BOOR_050_050a	0.0 0.0 0.5	0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.0 0.5	15.1 38.0	-51.7 64.2 306.2	0.0 0.0 0.5	11.7 45.5	-61.9 76.8	306.2 13.0 270	30.3 76.0
5	BOOR_062_062a	0.0 0.0 0.625	0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.0 0.625	18.9 47.5	-64.7 80.3 306.2	0.0 0.0 0.625	16.6 55.5	-72.9 90.4	306.2 10.3 270	30.3 76.0
6	BOOR_075_075a	0.0 0.0 0.75	0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.0 0.75	22.7 57.0	-77.6 96.3 306.2	0.0 0.0 0.75	21.3 61.2	-83.4 103.5	306.2 7.2 270	30.3 76.0
7	BOOR_087_087a	0.0 0.0 0.875	0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.0 0.875	26.5 66.5	-90.6 112.4 306.2	0.0 0.0 0.875	25.9 68.7	-93.6 116.1	306.2 3.7 270	30.3 76.0
8	BOOR_100_100a	0.0 0.0 1.0	1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.0 1.0	30.3 76.0	-103.5 128.5 306.2	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2 0.0 270	30.3 76.0
9	GOOB_012_012a	0.0 0.125	0.125 0.125 0.125	0.062 0.062 0.062	0.0 0.125	10.4	-10.3 9.9	14.3 136.0	0.0 0.125	0.0 8.2	-16.7 11.9	20.6 144.4
10	G50B_012_012a	0.0 0.125	0.125 0.125 0.125	0.062 0.062 0.062	0.0 0.125	10.8	-5.7 -1.6	6.0 196.3	0.0 0.125	0.125 8.9	-10.7 -3.3	11.2 197.0
11	G75B_025_025a	0.0 0.125	0.25 0.25 0.25	0.125 0.125 0.125	0.0 0.125	12.9	4.5 -17.0	17.6 285.0	0.0 0.125	0.25 10.8	3.0 -22.5	22.7 277.6
12	G84B_037_037a	0.0 0.125	0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.118	0.375	15.2 17.1	-32.5 36.7 297.8	0.0 0.125	0.375 13.5	17.5 -39.0	42.8 299.1
13	G88B_050_050a	0.0 0.125	0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.116	0.5	18.2 28.8	-46.7 54.8 301.6	0.0 0.125	0.5 16.8	30.3 -53.4	61.4 294.7
14	G90B_062_062a	0.0 0.125	0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.114	0.625	21.6 39.4	-60.3 72.1 303.1	0.0 0.125	0.625 20.5	41.6 -66.3	78.3 302.1
15	G92B_075_075a	0.0 0.125	0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.112	0.75	25.0 50.0	-73.9 89.3 304.0	0.0 0.125	0.75 24.4	51.7 -78.2	93.8 303.5
16	G93B_087_087a	0.0 0.125	0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.116	0.875	28.7 60.0	-87.1 105.8 304.5	0.0 0.125	0.875 28.4	61.0 -89.4	108.2 304.3
17	G94B_100_100a	0.0 0.125	1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.116	1.0	32.3 70.0	-100.3 122.3 304.9	0.0 0.125	1.0 32.4	69.6 -100.0	121.9 304.8
18	GOOB_025_025a	0.0 0.25	0.0 0.25 0.25 0.25	0.125 0.125 0.125	0.0 0.25	0.0	20.9 -18.6	19.9 28.7 136.0	0.0 0.25	0.0 20.9	-30.6 28.3	41.7 137.2
19	G25B_025_025a	0.0 0.25	0.125 0.125 0.125 0.125	0.062 0.062 0.062	0.0 0.25	0.125	21.0 -20.4	11.2 21.6 148.6	0.0 0.25	0.125 21.2	-26.3 13.8	29.7 152.3
20	G50B_025_025a	0.0 0.25	0.25 0.25 0.25 0.25	0.125 0.125 0.125	0.0 0.25	0.25	21.7 -11.5	-3.3 12.0 196.3	0.0 0.25	0.25 22.1	-1.7 -5.0	17.8 196.3
21	G65B_037_037a	0.0 0.25	0.375 0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.256	0.375	24.1 -3.4	-18.3 18.6 259.3	0.0 0.25	0.375 23.5	-4.6 -22.9	23.4 258.4
22	G75B_050_050a	0.0 0.25	0.5 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.25	0.5	25.8 9.1	-34.1 35.3 285.0	0.0 0.25	0.5 25.5	8.7 -39.2	40.2 282.5
23	G80B_062_062a	0.0 0.25	0.625 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.239	0.625	27.7 22.6	-50.3 55.1 294.2	0.0 0.25	0.625 27.9	21.8 -54.1	58.3 291.9
24	G84B_075_075a	0.0 0.25	0.75 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.237	0.75	30.3 34.3	-65.0 73.5 297.8	0.0 0.25	0.75 30.7	34.0 -67.7	75.8 296.6
25	G88B_087_087a	0.0 0.25	0.875 0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.233	0.875	33.2 46.6	-79.6 92.3 300.3	0.0 0.25	0.875 30.8	45.4 -80.4	92.3 299.4
26	G88B_100_100a	0.0 0.25	1.0 1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.233	1.0	36.5 57.6	-93.9 109.7 301.6	0.0 0.25	1.0 37.1	55.9 -92.3	107.9 301.1
27	GOOB_037_037a	0.0 0.375	0.0 0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.375	0.0	31.3 -31.0	29.9 43.1 136.0	0.0 0.375	0.0 32.5	-40.3 38.9	56.1 136.0
28	G15B_037_037a	0.0 0.375	0.125 0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.375	0.118	31.4 -29.7	23.6 38.0 141.4	0.0 0.375	0.125 32.7	-37.7 27.7	46.9 143.6
29	G34B_037_037a	0.0 0.375	0.25 0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.375	0.256	31.8 -24.7	8.7 26.2 160.4	0.0 0.375	0.25 33.2	-31.7 11.0	33.6 160.8
30	G50B_037_037a	0.0 0.375	0.375 0.375 0.375 0.375	0.187 0.187 0.187	0.0 0.375	0.375	32.5 -17.3	-5.0 18.0 196.3	0.0 0.375	0.375 34.1	-22.5 -6.6	23.4 196.3
31	G61B_050_050a	0.0 0.375	0.5 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.383	0.5	35.1 -9.7	-19.6 21.9 243.6	0.0 0.375	0.5 35.4	-11.1 -23.5	26.0 244.6
32	G69B_062_062a	0.0 0.375	0.625 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.385	0.625	37.3 0.5	-34.8 34.8 270.8	0.0 0.375	0.625 37.0	1.1 -39.4	39.4 271.7
33	G75B_075_075a	0.0 0.375	0.75 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.375	0.75	38.8 13.7	-51.2 53.0 285.0	0.0 0.375	0.75 39.0	13.7 -54.2	56.0 284.1
34	G79B_087_087a	0.0 0.375	0.875 0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.364	0.875	40.6 26.8	-67.7 72.8 291.5	0.0 0.375	0.875 41.3	25.9 -68.1	72.9 290.8
35	G81B_100_100a	0.0 0.375	1.0 1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.366	1.0	43.4 38.7	-82.0 90.7 295.3	0.0 0.375	1.0 43.8	37.6 -81.2	89.5 294.8
36	GOOB_050_050a	0.0 0.5	0.0 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.5	0.0	41.8 -41.3	39.9 57.5 136.0	0.0 0.5	0.0 43.5	-49.5 47.7	68.8 136.0
37	G11B_050_050a	0.0 0.5	0.125 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.5	0.116	41.8 -40.4	35.0 53.4 139.0	0.0 0.5	0.125 43.7	-47.7 39.5	62.0 140.3
38	G25B_050_050a	0.0 0.5	0.25 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.5	0.25	42.1 -36.8	22.4 43.2 148.6	0.0 0.5	0.25 44.0	-43.5 25.2	50.3 149.9
39	G38B_050_050a	0.0 0.5	0.375 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.5	0.383	42.7 -36.0	6.8 31.4 167.3	0.0 0.5	0.375 44.6	-36.7 8.6	37.7 166.7
40	G50B_050_050a	0.0 0.5	0.5 0.5 0.5 0.5	0.25 0.25 0.25	0.0 0.5	0.5	43.4 -23.0	-6.7 24.0 196.3	0.0 0.5	0.5 45.5	-27.6 -8.1	28.7 196.3
41	G59B_062_062a	0.0 0.5	0.625 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.51	0.625	46.1 -16.3	-21.4 26.9 232.6	0.0 0.5	0.625 46.6	-16.9 -24.3	29.6 235.0
42	G65B_075_075a	0.0 0.5	0.75 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.512	0.75	48.3 -6.9	-36.6 37.3 259.3	0.0 0.5	0.75 48.1	-5.4 -39.7	40.1 262.2
43	G70B_087_087a	0.0 0.5	0.875 0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.51	0.875	50.4 4.4	-52.3 52.4 274.9	0.0 0.5	0.875 49.8	6.4 -54.4	54.8 276.7
44	G75B_100_100a	0.0 0.5	1.0 1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.5	1.0	51.7 18.3	-68.3 70.7 285.0	0.0 0.5	1.0 51.7	18.3 -68.3	70.7 285.0
45	GOOB_062_062a	0.0 0.625	0.0 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.0	52.2 -51.7	49.9 71.9 136.0	0.0 0.625	0.0 54.1	-58.2 56.2	80.9 136.0
46	G09B_062_062a	0.0 0.625	0.125 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.114	52.3 -50.9	45.6 68.4 138.1	0.0 0.625	0.125 54.2	-56.9 49.9	75.7 138.7
47	G19B_062_062a	0.0 0.625	0.25 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.239	52.5 -48.5	35.7 60.2 143.5	0.0 0.625	0.25 54.4	-53.8 37.8	65.8 144.9
48	G30B_062_062a	0.0 0.625	0.375 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.385	52.9 -44.0	19.7 47.3 155.4	0.0 0.625	0.375 54.8	-48.6 22.6	63.6 155.0
49	G40B_062_062a	0.0 0.625	0.5 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.51	53.5 -36.5	4.6 36.8 172.6	0.0 0.625	0.5 55.5	-41.3 6.5	41.8 171.0
50	G50B_062_062a	0.0 0.625	0.625 0.625 0.625 0.625	0.312 0.312 0.312	0.0 0.625	0.625	54.2 -28.8	-8.4 30.0 196.3	0.0 0.625	0.625 56.3	-32.4 -9.5	33.8 196.3
51	G57B_075_075a	0.0 0.625	0.75 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.637	0.75	57.1 -22.4	-22.6 31.9 225.3	0.0 0.625	0.75 57.4	-22.3 -25.1	33.6 228.3
52	G63B_087_087a	0.0 0.625	0.875 0.875 0.875 0.875	0.437 0.437 0.437	0.0 0.641	0.875	59.4 -13.4	-37.5 39.8 250.2	0.0 0.625	0.875 58.7	-11.4 -40.2	41.8 254.0
53	G68B_100_100a	0.0 0.625	1.0 1.0 1.0 1.0	0.5 0.5 0.5	0.0 0.633	1.0	60.9 -1.5	-53.9 53.9 268.3	0.0 0.625	1.0 60.3	-0.1 -54.6	54.6 269.8
54	GOOB_075_075a	0.0 0.75	0.0 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.75	0.0	62.7 -62.0	59.9 86.2 136.0	0.0 0.75	0.0 64.2	-66.6 64.3	92.6 136.0
55	G07B_075_075a	0.0 0.75	0.125 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.75	0.112	62.9 -61.4	56.3 83.3 137.4	0.0 0.75	0.125 64.3	-65.6 59.4	88.5 137.8
56	G15B_075_075a	0.0 0.75	0.25 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.75	0.237	62.9 -59.4	47.3 76.0 141.4	0.0 0.75	0.25 64.5	-63.2 49.1	80.1 142.1
57	G25B_075_075a	0.0 0.75	0.375 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.75	0.375	63.2 -55.3	33.7 64.8 148.6	0.0 0.75	0.375 64.8	-59.1 35.4	68.9 149.0
58	G34B_075_075a	0.0 0.75	0.5 0.75 0.75 0.75	0.375 0.375 0.375	0.0 0.75	0.512	63.7 -49.4	17.5 52.4 160.4	0.0 0.75	0.5 65.3	-53.3 20.1	57.0 159.2
59	G42B_075_075a	0.0 0.7										

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 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	ief_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	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	6.3 9.6 8.0	12.5 40.0	0.125 0.0 0.0	2.4 10.9 3.8	11.6 19.4 5.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
82	B50R_012_012a	0.125 0.0 0.125	0.125 0.125 0.062	330	0.125 0.0 0.125	7.1 11.7 -7.3	13.8 328.2	0.125 0.0 0.125	3.2 16.7 -11.6	20.4 325.1 7.6	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
83	B25R_025_025a	0.125 0.0 0.25	0.25 0.25 0.125	300	0.125 0.0 0.25	9.6 19.9 -22.4	30.0 311.6	0.125 0.0 0.25	5.3 28.5 -31.2	42.3 312.3 13.0	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
84	B15R_037_037a	0.125 0.0 0.375	0.375 0.375 0.187	289	0.118 0.0 0.375	12.7 29.0 -36.5	46.7 308.4	0.125 0.0 0.375	9.0 38.1 -46.3	60.0 309.4 13.8	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
85	B11R_050_050a	0.125 0.0 0.5	0.5 0.5 0.25	284	0.116 0.0 0.5	16.1 38.3 -50.0	63.1 307.4	0.125 0.0 0.5	13.4 46.1 -59.0	74.9 307.9 12.1	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
86	B09R_062_062a	0.125 0.0 0.625	0.625 0.625 0.312	281	0.114 0.0 0.625	19.8 47.8 -63.2	79.3 307.0	0.125 0.0 0.625	17.9 53.9 -70.7	88.9 307.3 9.8	279	0.183 0.0 1.0	31.7 76.5 -101.2	126.9 307.0
87	B07R_075_075a	0.125 0.0 0.75	0.75 0.75 0.375	279	0.112 0.0 0.75	23.5 57.2 -76.4	95.5 306.8	0.125 0.0 0.75	22.3 61.5 -81.7	102.3 306.9 6.9	278	0.15 0.0 1.0	31.3 76.3 -101.9	127.4 306.8
88	B06R_087_087a	0.125 0.0 0.875	0.875 0.875 0.437	278	0.110 0.0 0.875	27.2 66.7 -89.5	111.6 306.7	0.125 0.0 0.875	26.7 69.0 -92.3	115.2 306.7 3.6	277	0.133 0.0 1.0	31.1 76.3 -102.3	127.6 306.7
89	B05R_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	30.9 76.2 -102.5	127.8 306.6	0.125 0.0 1.0	31.0 76.2 -102.5	127.7 306.6 0.0	276	0.116 0.0 1.0	30.9 76.2 -102.5	127.8 306.6
90	Y00G_012_012a	0.125 0.125 0.0	0.125 0.125 0.062	90	0.125 0.125 0.0	11.5 -25.5 11.3	11.6 102.8	0.125 0.125 0.0	10.4 -5.0 15.4	16.2 108.0 4.8	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8
91	NW_012a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	11.9 0.0 0.0	0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 325.7 0.8	360	1.0 1.0 1.0	95.4 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	15.7 9.5 -12.9	16.0 306.2	0.125 0.125 0.25	12.6 9.6 -19.5	21.8 296.2 7.3	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
93	B00R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.124 0.375	19.5 19.0 -25.8	32.1 306.2	0.125 0.125 0.375	15.0 21.1 -36.5	42.1 300.0 11.6	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
94	B00R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.124 0.5	23.3 28.5 -38.8	48.1 306.2	0.125 0.125 0.5	18.1 32.4 -51.3	60.6 302.2 14.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
95	B00R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	27.1 38.0 -51.7	64.2 306.2	0.125 0.125 0.625	21.6 42.8 -64.6	77.5 303.5 14.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
96	B00R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	30.9 47.5 -64.7	80.3 306.2	0.125 0.125 0.75	25.3 52.5 -76.8	93.0 304.3 14.2	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
97	B00R_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	34.7 57.0 -77.6	96.3 306.2	0.125 0.125 0.875	29.1 61.5 -88.2	107.5 304.8 12.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
98	B00R_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	38.5 66.5 -90.6	112.4 306.2	0.125 0.125 1.0	33.0 69.9 -99.0	121.3 305.2 10.6	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
99	Y50G_025_012a	0.125 0.25 0.0	0.25 0.25 0.125	120	0.125 0.25 0.0	21.4 -16.3 20.6	26.2 128.3	0.125 0.25 0.0	21.9 -22.3 29.7	37.2 126.9 10.9	119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3
100	G00B_025_012a	0.125 0.25 0.125	0.25 0.125 0.187	150	0.124 0.25 0.124	22.3 -10.3 9.9	14.3 136.0	0.125 0.25 0.125	22.2 -18.8 15.2	24.2 141.0 10.0	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
101	G50B_025_012a	0.125 0.25 0.25	0.25 0.125 0.187	210	0.124 0.25 0.25	22.7 -5.7 -1.6	6.0 196.3	0.125 0.25 0.25	23.0 -11.2 -3.5	11.7 197.3 5.7	210	0.0 1.0 0.0	86.8 -46.1	-13.5 48.1 196.3
102	G75B_037_025a	0.125 0.25 0.375	0.375 0.25 0.25	240	0.124 0.25 0.375	24.8 4.5 -17.0	17.6 285.0	0.125 0.25 0.375	24.4 -0.5 -21.5	21.5 268.6 6.7	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
103	G84B_050_037a	0.125 0.25 0.5	0.5 0.375 0.312	251	0.124 0.243 0.5	27.2 17.1 -32.5	36.7 297.8	0.125 0.25 0.5	26.3 11.5 -37.9	39.6 286.9 7.8	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
104	G88B_062_050a	0.125 0.25 0.625	0.625 0.5 0.375	256	0.125 0.241 0.625	30.1 28.8 -46.7	54.8 301.6	0.125 0.25 0.625	28.7 23.7 -52.9	59.0 294.1 8.1	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6
105	G90B_075_062a	0.125 0.25 0.75	0.75 0.625 0.437	259	0.125 0.239 0.75	33.5 39.4 -60.3	72.1 303.1	0.125 0.25 0.75	31.4 35.4 -66.7	75.5 297.7 7.7	260	0.0 0.183 1.0	34.6 63.0 -96.6	115.3 303.1
106	G92B_087_075a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.237 0.875	36.9 50.0 -73.9	89.3 304.0	0.125 0.25 0.875	34.4 46.3 -79.5	92.0 300.2 7.1	262	0.0 0.15 1.0	33.4 66.7 -98.6	119.1 304.0
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.241 1.0	40.6 60.0 -87.1	105.8 304.5	0.125 0.25 1.0	37.6 56.5 -91.4	107.5 301.7 6.3	262	0.0 0.133 1.0	32.8 68.6 -99.6	120.9 304.5
108	Y68G_037_037a	0.125 0.375 0.0	0.375 0.375 0.187	131	0.118 0.375 0.0	31.6 -28.2 30.3	41.4 132.9	0.125 0.375 0.0	33.1 -35.2 39.6	53.0 131.5 11.7	131	0.316 1.0 0.0	84.4 -75.3	80.9 110.6 132.9
109	G00B_037_025a	0.125 0.375 0.125	0.375 0.25 0.25	150	0.124 0.375 0.124	32.8 -20.6 19.9	28.7 136.0	0.125 0.375 0.125	33.3 -22.9 28.6	43.6 138.9 14.9	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
110	G25B_037_025a	0.125 0.375 0.25	0.375 0.25 0.25	180	0.124 0.375 0.25	33.0 -18.4 11.2	21.6 148.6	0.125 0.375 0.25	33.8 -27.4 11.9	29.9 156.5 9.0	180	0.0 1.0 0.5	84.3 -73.7	44.9 86.4 148.6
111	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	33.6 -11.5 3.3	12.0 196.3	0.125 0.375 0.375	34.7 -18.9 -5.7	19.8 196.8 7.8	210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3
112	G65B_050_037a	0.125 0.375 0.5	0.5 0.375 0.312	229	0.124 0.381 0.5	36.0 -3.4 -18.3	18.6 259.3	0.125 0.375 0.5	35.9 -8.3 -22.7	24.1 249.7 6.5	228	0.0 0.683 1.0	64.4 -9.2	-48.8 49.7 259.3
113	G75B_062_050a	0.125 0.375 0.625	0.625 0.5 0.375	240	0.125 0.375 0.625	37.8 9.1 -31.1	35.3 285.0	0.125 0.375 0.625	37.5 3.3 -38.6	38.7 274.9 7.3	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.364 0.75	39.6 22.6 -50.3	55.1 294.2	0.125 0.375 0.75	39.5 15.3 -53.5	55.6 285.9 7.9	247	0.0 0.383 1.0	44.3 36.2 -80.5	88.2 294.2
115	G84B_087_075a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.362 0.875	42.4 34.3 -60.5	73.5 297.8	0.125 0.375 0.875	41.7 27.1 -67.4	72.7 291.9 7.5	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
116	G86B_100_087a	0.125 0.375 1.0	1.0 0.875 0.562	254	0.125 0.358 1.0	45.2 46.6 -79.6	92.2 300.3	0.125 0.375 1.0	44.2 38.6 -80.5	89.3 295.5 8.1	255	0.0 0.266 1.0	38.0 53.3 -91.0	105.4 300.3
117	Y76G_050_050a	0.125 0.5 0.0	0.5 0.5 0.25	136	0.116 0.5 0.0	42.0 -39.3 40.2	56.2 134.3	0.125 0.5 0.0	43.9 -45.9 48.2	66.6 133.6 10.6	137	0.233 1.0 0.0	84.0 -78.7	80.4 112.5 134.3
118	G00B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.124	43.2 -31.0 29.9	43.1 136.0	0.125 0.5 0.125	44.1 -44.3 40.1	59.8 137.8 16.7	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.243	43.3 -29.1 23.6	38.0 141.4	0.125 0.5 0.25	44.4 -40.3 25.7	47.9 147.4 10.9	168	0.0 1.0 0.316	83.9 -79.2	63.1 101.3 141.4
120	G34B_050_037a	0.125 0.5 0.375	0.5 0.375 0.312	191	0.124 0.5 0.381	43.8 -24.7 8.7	26.2 160.4	0.125 0.5 0.375	45.0 -33.8 9.2	35.1 164.7 9.2	191	0.0 1.0 0.683	85.0 -65.8	23.4 69.9 160.4
121	G50B_050_037a	0.125 0.5 0.5	0.5 0.375 0.312	210	0.124 0.5 0.5	44.5 -17.3 -5.0	18.0 196.3	0.125 0.5 0.5	45.9 -25.2 -7.5	26.3 196.6 8.3	210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3
122	G61B_062_050a	0.125 0.5 0.625	0.625 0.5 0.375	224	0.125 0.508 0.625	47.0 -9.7 -19.6	21.9 243.6	0.125 0.5 0.625	47.0 -14.9 -23.7	28.0 237.7 6.6	222	0.0 0.766 1.0	70.2 -19.5	-39.3 43.9 243.6
123	G69B_075_062a	0.125 0.5 0.75	0.75 0.625 0.437	233	0.125 0.51 0.75	49.2 0.5 -34.8	34.8 270.8	0.125 0.5 0.75	48.4 -3.8 -39.2	39.3 264.4 6.2	232	0.0 0.616 1.0	59.7 70.8 -55.6	55.7 270.8
124	G75B_087_075a	0.125 0.5 0.875	0.875 0.75 0.5	240	0.125 0.5 0.875	50.7 13.7 -51.2	53.0 285.0	0.125 0.5 0.875	50.1 7.7 -53.8	54.4 278.2 6.5	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
125	G79B_100_087a	0.125 0.5 1.0	1.0 0.875 0.562	245	0.125 0.489 1.0	52.6 26.8 -67.7	72.8 291.5	0.125 0.5 1.0	52.0 19.4 -67.8	70.5 285.9 7.4	245	0.0 0.416 1.0	46.5 30.6 -77.4	83.2 291.5
126	Y81G_062_062a	0.125 0.625 0.0	0.625 0.625 0.312	139	0.114 0.625 0.0	52.4 -49.9 50.1	70.8 134.8	0.125 0.625 0.0	54.3 -55.6 56.5	79.3 134.5 8.7	149	0.183 1.0 0.0	83.9 -79.9	80.2 113.3 134.8
127	G00B_062_050a	0.125 0.625 0.125	0.625 0.5 0.375	150	0.125 0.625 0.125	53.7 -41.3 39.9	57.5 136.0	0.125 0.625 0.125	54.4 -54.4 50.3	74.1 137.2 16.7	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
128	G11B_062_050a	0.125 0.625 0.25	0.625 0.5 0.375	164	0.125 0.625 0.241	53.8 -40.4 35.0	53.4 139.0	0.125 0.625 0.25	54.7 -51.4 38.2	61.1				

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n	HIC*Fa	rgb_Fa	icr_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Ma	LabCh*Ma		
162	R00Y_025_025a	0.25 0.0 0.0	0.25 0.25 0.125	390	0.25 0.0 0.0	12.6 19.2 16.1	25.1 25.1 4.0	0.25 0.0 0.0	8.6 28.5 13.6	31.6 25.5 10.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
163	R00Y_025_025a	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.125	13.0 20.2 1.0	20.3 2.9 2.0	0.25 0.0 0.125	9.4 30.5 -1.8	30.6 35.6 11.2	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 29.0
164	B50R_025_025a	0.25 0.0 0.25	0.25 0.25 0.125	330	0.25 0.0 0.25	14.3 23.5 -14.6	27.7 328.2	0.25 0.0 0.25	11.1 34.9 -21.6	41.1 32.8 13.7	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
165	B34R_037_037a	0.25 0.0 0.375	0.25 0.375 0.187	311	0.256 0.0 0.375	16.8 31.5 -29.7	43.3 316.7	0.25 0.0 0.375	13.8 41.1 -38.3	56.2 316.9 13.2	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7
166	B25R_050_050a	0.25 0.0 0.5	0.5 0.5 0.25	300	0.25 0.0 0.5	19.2 39.9 -44.8	60.0 311.6	0.25 0.0 0.5	17.1 48.0 -52.8	71.4 312.2 11.6	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
167	B19R_062_062a	0.25 0.0 0.625	0.625 0.625 0.312	293	0.239 0.0 0.625	22.1 48.8 -59.9	76.9 309.3	0.25 0.0 0.625	20.7 55.2 -65.9	86.0 309.9 9.2	292	0.383 0.0 1.0	35.3 78.1 -95.1	123.0 309.3
168	B15R_075_075a	0.25 0.0 0.75	0.75 0.75 0.375	289	0.237 0.0 0.75	25.4 58.1 -73.1	93.4 308.4	0.25 0.0 0.75	24.6 62.5 -77.8	99.8 308.7 6.5	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
169	B13R_087_087a	0.25 0.0 0.875	0.875 0.875 0.437	286	0.233 0.0 0.875	28.8 67.3 -86.8	109.9 307.8	0.25 0.0 0.875	28.6 69.7 -89.1	113.1 308.0 3.2	284	0.266 0.0 1.0	32.9 77.0 -99.2	125.6 307.8
170	B11R_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4	0.25 0.0 1.0	32.6 76.8 -99.8	125.9 307.5 0.4	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
171	R50Y_025_025a	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.125 0.0	15.9 10.3 17.7	20.5 59.7	0.25 0.125 0.0	14.7 12.2 22.0	25.2 60.9 4.8	59	1.0 0.5 0.0	63.6 46.3 71.0	82.2 59.7
172	R00Y_025_012a	0.25 0.125 0.125	0.25 0.125 0.187	390	0.25 0.124 0.124	18.2 9.6 8.0	12.5 40.0	0.25 0.125 0.125	15.2 14.7 6.5	16.1 23.9 6.1	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
173	B50R_025_012a	0.25 0.125 0.25	0.25 0.125 0.187	330	0.25 0.124 0.25	19.0 11.7 -7.3	13.8 328.2	0.25 0.125 0.25	16.4 20.2 -13.2	24.2 326.7 10.6	330	1.0 0.0 0.0	57.2 94.3 -58.4	110.9 328.2
174	B25R_037_025a	0.25 0.125 0.375	0.375 0.25 0.25	300	0.25 0.124 0.375	21.5 19.9 -22.4	30.0 311.6	0.25 0.125 0.375	18.4 28.0 -30.9	41.7 312.1 12.1	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
175	B15R_050_037a	0.25 0.125 0.5	0.5 0.375 0.312	289	0.243 0.124 0.5	24.6 29.0 -36.5	46.7 308.4	0.25 0.125 0.5	20.9 36.7 -46.5	59.3 308.3 13.1	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
176	B11R_062_050a	0.25 0.125 0.625	0.625 0.5 0.375	284	0.241 0.125 0.625	28.1 38.3 -50.0	63.1 307.4	0.25 0.125 0.625	23.9 45.7 -60.5	75.9 307.0 13.4	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
177	B09R_075_062a	0.25 0.125 0.75	0.75 0.625 0.437	281	0.239 0.125 0.75	31.7 47.8 -63.2	79.3 307.0	0.25 0.125 0.75	27.3 54.4 -73.4	91.4 306.5 12.9	279	0.233 0.0 1.0	31.7 76.5 -101.2	126.9 307.0
178	B07R_087_075a	0.25 0.125 0.875	0.875 0.75 0.5	279	0.237 0.125 0.875	35.4 57.2 -76.4	95.5 306.8	0.25 0.125 0.875	30.8 62.8 -85.3	106.0 306.3 11.4	278	0.15 0.0 1.0	31.3 76.3 -101.9	127.4 306.8
179	B06R_100_087a	0.25 0.125 1.0	1.0 0.875 0.562	278	0.231 0.125 1.0	39.1 66.7 -89.5	111.6 306.2	0.25 0.125 1.0	34.5 70.9 -96.6	119.8 306.2 9.4	277	0.133 0.0 1.0	31.1 76.3 -102.3	127.6 306.7
180	Y00G_025_025a	0.25 0.25 0.0	0.25 0.25 0.125	90	0.25 0.25 0.0	23.1 -5.1 22.6	23.2 102.8	0.25 0.25 0.0	24.2 -7.6 32.9	33.7 103.1 10.5	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8
181	Y00G_025_012a	0.25 0.25 0.125	0.25 0.125 0.187	90	0.25 0.25 0.124	23.5 -2.5 11.3	11.6 102.8	0.25 0.25 0.125	24.5 -5.3 18.6	19.4 105.9 7.8	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8
182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	23.8 0.0 0.0	0.0 0.0	0.25 0.25 0.25	25.2 0.0 0.0	0.0 32.5 1.4	360	1.0 1.0 1.0	95.4 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	27.6 9.5 -12.9	16.0 306.2	0.25 0.25 0.375	26.5 8.0 -18.0	19.8 294.0 5.4	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
184	B00R_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.249 0.5	31.4 19.0 -25.8	32.1 306.2	0.25 0.25 0.5	28.2 17.7 -34.7	39.0 297.0 9.5	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
185	B00R_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.245 0.25 0.625	35.2 28.5 -38.8	48.1 306.2	0.25 0.25 0.625	30.4 28.1 -50.0	57.4 299.3 12.2	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	39.0 38.0 -51.7	64.2 306.2	0.25 0.25 0.75	32.9 38.5 -64.1	74.8 301.0 13.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
187	B00R_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.8 47.5 -64.7	80.3 306.2	0.25 0.25 0.875	35.8 48.6 -77.1	91.2 302.1 14.3	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
188	B00R_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	46.6 57.0 -77.6	96.3 306.2	0.25 0.25 1.0	38.8 58.2 -89.4	106.7 303.0 14.1	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2
189	Y31G_037_037a	0.25 0.375 0.0	0.375 0.375 0.187	109	0.256 0.375 0.0	32.8 -19.0 31.8	37.1 120.8	0.25 0.375 0.0	34.6 -24.3 41.4	48.0 120.4 11.0	108	0.683 1.0 0.0	87.6 -50.7	84.9 98.9 120.8
190	Y50G_037_025a	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	33.3 -16.3 20.6	26.2 128.3	0.25 0.375 0.125	34.8 -22.5 30.5	38.0 126.3 11.8	119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3
191	G00B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	34.3 -10.3 9.9	14.3 136.0	0.25 0.375 0.25	35.2 -18.1 14.0	22.9 142.2 8.8	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
192	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	34.7 -5.7 -1.6	6.0 196.3	0.25 0.375 0.375	36.0 -11.0 -3.5	11.6 197.8 5.8	210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3
193	G75B_050_025a	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.375 0.5	36.7 4.5 -17.0	17.6 285.0	0.25 0.375 0.5	37.2 -2.0 -20.5	20.6 264.3 7.4	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
194	G84B_062_037a	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.368 0.625	39.1 17.1 -32.5	36.7 297.8	0.25 0.375 0.625	38.7 8.2 -36.6	37.5 282.7 9.7	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
195	G88B_075_050a	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.366 0.75	42.1 28.8 -46.7	54.8 301.6	0.25 0.375 0.75	40.6 19.1 -51.6	55.0 290.3 10.9	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6
196	G90B_087_062a	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.364 0.875	45.5 39.4 -60.3	72.1 303.1	0.25 0.375 0.875	42.8 30.1 -65.7	72.2 294.6 11.0	260	0.0 0.183 1.0	34.6 63.0 -96.6	115.3 303.1
197	G92B_100_075a	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.362 1.0	48.9 50.0 -73.9	89.3 304.0	0.25 0.375 1.0	45.2 40.8 -78.9	88.9 297.3 11.1	262	0.0 0.15 1.0	33.4 66.7	-98.6 119.1 304.0
198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.25 0.125	120	0.25 0.5 0.0	42.8 -32.6 41.2	52.5 128.3	0.25 0.5 0.0	44.9 -37.9 49.4	62.3 127.5 10.0	119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3
199	Y68G_050_037a	0.25 0.5 0.125	0.5 0.375 0.312	131	0.243 0.5 0.124	43.6 -28.2 30.3	41.4 132.9	0.25 0.5 0.125	45.0 -36.5 41.4	55.2 131.4 13.9	131	0.316 1.0 0.0	84.4 -75.3	80.9 110.6 132.9
200	G00B_050_025a	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.249	44.7 -20.6 19.9	28.7 136.0	0.25 0.5 0.25	45.4 -33.0 27.2	42.8 140.5 14.3	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
201	G25B_050_025a	0.25 0.5 0.375	0.5 0.25 0.375	180	0.249 0.5 0.375	44.9 -18.4 11.2	21.6 148.6	0.25 0.5 0.375	45.9 -19.3 10.6	29.3 158.6 8.9	180	0.0 1.0 0.5	84.3 -73.7	44.9 86.4 148.6
202	G50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	45.5 -11.5 -3.3	12.0 196.3	0.25 0.5 0.5	46.8 -27.5 -6.0	20.4 197.2 8.5	210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3
203	G65B_062_037a	0.25 0.5 0.625	0.625 0.375 0.437	229	0.25 0.506 0.625	48.0 -3.4 -18.3	18.6 259.3	0.25 0.5 0.625	47.9 -10.2 -22.3	24.5 245.3 7.8	228	0.0 0.683 1.0	64.4 -9.2	-48.8 49.7 259.3
204	G75B_075_050a	0.25 0.5 0.75	0.75 0.5 0.5	240	0.25 0.5 0.75	49.7 9.1 -34.1	35.3 285.0	0.25 0.5 0.75	49.3 0.1 -37.8	37.8 270.1 9.7	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
205	G80B_087_062a	0.25 0.5 0.875	0.875 0.625 0.562	247	0.25 0.489 0.875	51.5 22.6 -50.3	55.1 294.2	0.25 0.5 0.875	50.9 10.9 -52.5	53.6 281.7 11.9	247	0.0 0.383 1.0	44.3 36.2 -80.5	88.2 294.2
206	G84B_100_075a	0.25 0.5 1.0	1.0 0.75 0.625	251	0.25 0.487 1.0	54.4 34.3 -60.5	73.5 297.8	0.25 0.5 1.0	52.8 21.9 -66.5	70.0 288.2 12.5	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8
207	Y61G_062_062a	0.25 0.625 0.0	0.625 0.625 0.312	127	0.239 0.625 0.0	53.0 -45.2 50.8	68.0 131.6	0.25 0.625 0.0	55.1 -49.5 57.4	75.8 130.7 8.1	127	0.383 1.0 0.0	84.8 -72.3	81.3 108.8 131.6
208	Y76G_062_050a	0.25 0.625 0.125	0.625 0.5 0.375	136	0.241 0.625 0.125	53.9 -39.3 40.2	56.2 134.3	0.25 0.625 0.125	55.2 -48.4 51.2	70.5 133.3 14.3	137	0.233 1.0 0.0	84.0 -78.7	80.4 112.5 134.3
209	G00B_062_037a	0.25 0.625 0.25	0.625 0.375 0.437	150	0.25 0.625 0.25	55.2 -31.0 29.9	43.1 136.0	0.25 0.625 0.25	55.4 -45.7 39.2	60.2 139.3 17.3	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
210	G15B_062													

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informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md			
243	R00Y_037_037a	0.375 0.0 0.0	0.375 0.375 0.187	390	0.375 0.0 0.0	18.9 28.8 24.2	37.6 40.0	0.375 0.0 0.0	16.4 37.5 25.4	45.3 34.1 9.1	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
244	R18Y_037_037a	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.118	19.1 29.6 11.1	31.7 20.6	0.375 0.0 0.125	16.8 38.7 9.7	39.9 14.1 9.4	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6	
245	B65R_037_037a	0.375 0.0 0.25	0.375 0.375 0.187	349	0.375 0.0 0.256	20.0 32.0 -7.4	32.9 346.8	0.375 0.0 0.25	17.9 41.5 -10.4	42.8 345.8 10.2	348	1.0 0.0 0.683	53.5 85.4 -19.9	87.7 346.8	
246	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	21.4 35.5 -21.9	41.6 328.2	0.375 0.0 0.375	19.7 46.0 -28.5	54.1 328.2 12.6	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	
247	B38R_050_050a	0.375 0.0 0.5	0.5 0.5 0.25	316	0.383 0.0 0.5	23.9 43.2 -37.0	56.9 319.4	0.375 0.0 0.5	22.1 51.5 -44.4	68.1 319.2 11.3	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4	
248	B30R_062_062a	0.375 0.0 0.625	0.625 0.625 0.312	307	0.385 0.0 0.625	26.5 51.4 -52.0	73.1 314.6	0.375 0.0 0.625	24.9 57.8 -58.7	82.4 314.5 9.4	307	0.616 0.0 1.0	42.4 82.3 -83.2	117.0 314.6	
249	B25R_075_075a	0.375 0.0 0.75	0.75 0.75 0.375	300	0.375 0.0 0.75	28.9 59.8 -67.2	90.0 311.6	0.375 0.0 0.75	28.1 64.4 -71.9	96.5 311.8 6.5	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6	
250	B20R_087_087a	0.375 0.0 0.875	0.875 0.875 0.437	295	0.364 0.0 0.875	31.7 68.8 -81.8	106.9 310.0	0.375 0.0 0.875	31.6 71.2 -84.0	110.1 310.2 3.2	294	0.416 0.0 1.0	36.3 78.6 -93.5	122.2 310.0	
251	B18R_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	34.9 77.9 -95.7	123.4 309.1	0.375 0.0 1.0	35.1 77.9 -95.5	123.3 309.2 0.3	291	0.366 0.0 1.0	34.9 77.9 -95.7	123.4 309.1	
252	R31Y_037_037a	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.118 0.0	21.1 22.7 25.2	33.9 47.9	0.375 0.125 0.0	20.4 26.4 30.1	40.1 48.7 6.2	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9	
253	R00Y_037_025a	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.124	24.5 19.2 16.1	25.1 40.0	0.375 0.125 0.125	20.7 27.8 14.8	31.5 28.0 9.5	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
254	R00Y_037_025a	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.25	24.9 20.2 1.0	20.3 2.9	0.375 0.125 0.25	21.6 31.1 -4.9	31.5 25.0 12.8	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	
255	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.124 0.375	26.2 23.5 -14.6	27.7 328.2	0.375 0.125 0.375	23.1 36.3 -23.1	43.0 327.5 15.6	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	
256	B34R_050_037a	0.375 0.125 0.5	0.5 0.5 0.375	312	0.381 0.124 0.5	28.7 31.5 -29.7	43.3 316.7	0.375 0.125 0.5	25.1 42.8 -39.5	58.3 317.2 15.3	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7	
257	B25R_062_050a	0.375 0.125 0.625	0.625 0.5 0.375	300	0.375 0.125 0.625	31.2 39.9 -44.8	60.0 311.6	0.375 0.125 0.625	27.6 50.0 -54.4	73.9 312.5 14.4	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6	
258	B19R_075_062a	0.375 0.125 0.75	0.75 0.625 0.437	293	0.364 0.125 0.75	34.0 48.8 -59.4	76.9 309.7	0.375 0.125 0.75	30.4 57.5 -68.1	89.1 310.2 12.8	292	0.383 0.0 1.0	35.3 78.1 -95.1	123.0 309.3	
259	B15R_087_075a	0.375 0.125 0.875	0.875 0.75 0.5	289	0.362 0.125 0.875	37.4 58.1 -73.1	93.4 308.4	0.375 0.125 0.875	33.6 65.1 -80.7	103.7 308.9 11.0	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4	
260	B13R_100_087a	0.375 0.125 1.0	1.0 0.875 0.562	286	0.358 0.125 1.0	40.7 67.3 -86.8	109.9 307.8	0.375 0.125 1.0	36.9 72.6 -92.6	117.7 308.1 8.7	284	0.266 0.0 1.0	32.9 77.0 -99.2	125.6 307.8	
261	R68Y_037_037a	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.256 0.0	27.5 6.9 29.1	29.9 76.5	0.375 0.25 0.0	27.8 8.3 37.5	38.4 77.4 8.5	71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5	
262	R50Y_037_025a	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.25 0.124	27.8 10.3 17.7	20.5 59.7	0.375 0.25 0.125	28.1 9.8 23.7	25.7 67.5 6.0	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7	
263	R00Y_037_012a	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.249	30.1 9.6 8.0	12.5 40.0	0.375 0.25 0.25	28.7 13.3 5.4	14.4 22.0 4.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
264	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.249 0.375	31.0 11.7 -7.3	13.8 328.2	0.375 0.25 0.375	29.7 19.0 -12.7	22.9 326.1 9.1	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	
265	B25R_050_025a	0.375 0.25 0.5	0.5 0.25 0.375	300	0.375 0.249 0.5	33.5 19.9 -22.4	30.0 311.6	0.375 0.25 0.5	31.2 26.3 -29.7	39.7 311.5 9.9	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6	
266	B15R_062_037a	0.375 0.25 0.625	0.625 0.375 0.437	289	0.368 0.25 0.625	36.5 29.0 -36.5	46.7 308.4	0.375 0.25 0.625	33.2 34.6 -45.4	57.0 307.3 10.9	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4	
267	B11R_075_050a	0.375 0.25 0.75	0.75 0.5 0.5	284	0.366 0.25 0.75	40.0 38.3 -50.0	63.1 307.4	0.375 0.25 0.75	35.4 43.3 -59.8	73.9 305.9 11.8	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4	
268	B09R_087_062a	0.375 0.25 0.875	0.875 0.625 0.562	281	0.364 0.25 0.875	43.7 47.8 -63.2	79.3 307.0	0.375 0.25 0.875	38.0 52.2 -73.3	90.0 305.4 12.3	279	0.183 0.0 1.0	31.7 76.5 -101.2	126.9 307.0	
269	B07R_100_075a	0.375 0.25 1.0	1.0 0.75 0.625	279	0.362 0.25 1.0	47.3 -76.4	95.5 306.8	0.375 0.25 1.0	40.9 60.9 -86.0	105.4 305.3 12.0	278	0.15 0.0 1.0	31.3 76.3 -101.9	127.4 306.8	
270	Y00G_037_037a	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.375 0.0	34.7 -7.7	34.0 34.9	0.375 0.375 0.0	36.9 -10.0	44.2 45.3 10.2	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8	
271	Y00G_037_025a	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.375 0.124	35.0 -5.1	22.6 23.2	0.375 0.375 0.125	37.1 -8.7	33.8 34.9 10.4	11.8 89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8	
272	Y00G_037_012a	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.375 0.249	35.4 -2.5	11.3 11.6	0.375 0.375 0.25	37.5 -5.4	17.5 18.3 10.7	7.1 89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0 102.8	
273	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	35.7 0.0 0.0	0.0 0.0	0.375 0.375 0.375	38.3 0.0 0.0	0.0 0.0 0.0	360	1.0 1.0 1.0	95.4 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	39.5 9.5 -12.9	16.0 306.2	0.375 0.375 0.5	39.4 7.2 -17.0	18.5 292.9 4.7	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
275	B00R_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	43.3 19.0 -25.8	32.1 306.2	0.375 0.375 0.625	40.8 15.7 -33.2	36.8 295.4 8.4	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
276	B00R_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	47.1 28.5 -38.8	48.1 306.2	0.375 0.375 0.75	42.5 25.1 -48.4	54.5 297.4 11.1	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
277	B00R_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	50.9 38.0 -51.7	64.2 306.2	0.375 0.375 0.875	44.6 34.8 -62.7	71.7 299.0 13.0	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
278	B00R_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	54.7 47.5 -64.7	80.3 306.2	0.375 0.375 1.0	46.8 44.5 -76.1	88.2 300.3 14.2	270	0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2	
279	Y23G_050_050a	0.375 0.5 0.0	0.5 0.5 0.25	104	0.383 0.5 0.0	44.3 -21.6	43.1 48.2	0.375 0.5 0.0	46.6 -26.1	51.4 57.7	116.9 9.7	102	0.766 1.0 0.0	88.7 -43.3	86.2 96.5 116.6
280	Y31G_050_037a	0.375 0.5 0.125	0.5 0.375 0.312	109	0.381 0.5 0.124	44.8 -19.0	31.8 37.1	0.375 0.5 0.125	46.7 -25.0	43.6 50.2	119.8 13.3	108	0.683 1.0 0.0	87.6 -50.7	84.9 98.9 120.8
281	Y50G_050_025a	0.375 0.5 0.25	0.5 0.25 0.375	120	0.375 0.5 0.249	45.2 -16.3	20.6 26.2	0.375 0.5 0.25	47.0 -22.1	29.6 36.9	126.8 10.8	119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3
282	G00B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	46.2 -10.3	9.9 14.3	0.375 0.5 0.375	47.6 -17.3	13.1 21.8	142.8 7.8	149	0.0 1.0 0.0	83.6 -82.7	79.8 115.0 136.0
283	G50B_050_012a	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.5	46.6 -5.7	-1.6 6.0	0.375 0.5 0.5	48.4 -10.7	-3.5 11.3	198.2 5.6	210	0.0 1.0 1.0	86.8 -46.1	-13.5 48.1 196.3
284	G75B_062_025a	0.375 0.5 0.625	0.625 0.25 0.5	240	0.375 0.5 0.625	48.7 4.5 -17.0	17.6 285.0	0.375 0.5 0.625	49.4 -2.7	-19.8 20.0	262.1 7.8	240	0.0 0.5 1.0	51.7 18.3 -68.3	70.7 285.0
285	G84B_075_037a	0.375 0.5 0.75	0.75 0.375 0.562	251	0.375 0.493 0.75	51.0 17.1 -32.5	36.7 297.8	0.375 0.5 0.75	50.7 6.3 -35.4	35.9 280.2 11.1	251	0.0 0.316 1.0	40.7 45.8 -86.7	98.1 297.8	
286	G88B_087_050a	0.375 0.5 0.875	0.875 0.5 0.625	256	0.375 0.491 0.875	54.0 28.8 -46.7	54.8 301.6	0.375 0.5 0.875	52.3 16.1 -50.2	52.7 287.8 13.2	257	0.0 0.233 1.0	36.5 57.6 -93.4	109.7 301.6	
287	G90B_100_062a	0.375 0.5 1.0	1.0 0.625 0.687	259	0.375 0.489 1.0	57.4 39.4 -60.3	72.1 303.1	0.375 0.5 1.0	54.1 26.2 -64.3	69.4 292.1 14.1	260	0.0 0.183 1.0	34.6 63.0 -96.6	115.3 303.1	
288	Y38G_062_062a	0.375 0.625 0.0	0.625 0.625 0.312	113	0.385 0.625 0.0	54.2 -35.2	52.4 63.1	0.375 0.625 0.0	56.3 -39.9	58.9 71.2	124.1 8.3	112	0.616 1.0 0.0	86.8 -56.4	83.8 101.0 123.9
289	Y50G_062_050a	0.375 0.625 0.125	0.625 0.5 0.375	120	0.375 0.625 0.125	54.7 -32.6	41.2 52.5	0.375 0.625 0.125	56.4 -39.0	52.8 65.7	126.4 13.4	119	0.5 1.0 0.0	85.7 -65.2	82.4 105.1 128.3
290	Y68G_062_037a	0.375 0.625 0.25	0.625 0.375 0.437	131	0.368 0.625 0.25	55.5 -28.2	30.3 41.4	0.375 0.625 0.25	56.6 -36.6	40.9 54.9	131.8 13.5	131	0.316 1.0 0.0	84.4 -75.3	80.9 110.6 132.9
291	G00B_062_025a	0.375 0.625													

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF21/PF21.HTM>
 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md			
324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.5	0.25 0.25 0.25	390	0.5 0.0 0.0	25.2 38.4 32.2	50.2 40.0 27.8	0.5 0.0 0.0	23.7 46.0 35.7	58.2 37.8 8.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
325	R26Y_050_050a	0.5 0.0 0.125	0.5 0.5 0.5	0.25 0.25 0.25	376	0.5 0.0 0.116	25.4 39.0 20.6	44.1 27.0 40.0	0.5 0.0 0.125	24.0 46.8 20.3	51.0 23.5 7.9	377	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8
326	R00Y_050_050a	0.5 0.0 0.25	0.5 0.5 0.5	0.25 0.25 0.25	360	0.5 0.0 0.25	26.0 40.5 2.0	40.6 2.9 0.5	0.5 0.0 0.25	24.8 48.8 0.4	48.8 0.5 8.4	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
327	B61R_050_050a	0.5 0.0 0.375	0.5 0.5 0.5	0.25 0.25 0.25	344	0.5 0.0 0.383	27.2 43.6 -15.3	46.2 340.6 0.5	0.5 0.0 0.375	26.0 52.0 -18.0	55.1 340.8 8.9	342	1.0 0.0 0.766	54.4 87.3 -30.6	92.5 340.6
328	B50R_050_050a	0.5 0.0 0.5	0.5 0.5 0.5	0.25 0.25 0.25	330	0.5 0.0 0.5	28.6 47.1 -29.2	55.4 328.2 0.5	0.5 0.0 0.5	27.8 56.4 -34.9	66.3 328.2 10.9	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
329	B40R_062_062a	0.5 0.0 0.625	0.625 0.625 0.625	0.312 0.312 0.312	319	0.51 0.0 0.625	31.1 55.0 -44.2	70.6 321.2 0.5	0.5 0.0 0.625	30.0 61.6 -50.3	79.5 320.7 9.0	320	0.816 0.0 1.0	49.8 88.1 -70.7	113.0 321.2
330	B34R_075_075a	0.5 0.0 0.75	0.75 0.75 0.75	0.375 0.375 0.375	311	0.512 0.0 0.75	33.6 63.1 -59.4	86.6 316.7 0.5	0.5 0.0 0.75	32.6 67.4 -64.4	93.2 316.3 6.6	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7
331	B29R_087_087a	0.5 0.0 0.875	0.875 0.875 0.875	0.437 0.437 0.437	305	0.51 0.0 0.875	36.1 71.4 -74.4	103.2 317.8 0.5	0.5 0.0 0.875	35.5 73.5 -77.4	106.8 313.5 3.7	305	0.583 0.0 1.0	41.3 81.6 -85.1	117.9 313.8
332	B25R_100_100a	0.5 0.0 1.0	1.0 1.0 1.0	0.5 0.5 0.5	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6 0.5	0.5 0.0 1.0	38.5 79.8 -89.7	120.1 311.6 0.0	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
333	R23Y_050_050a	0.5 0.125 0.0	0.5 0.5 0.5	0.25 0.25 0.25	44	0.5 0.116 0.0	26.8 33.8 32.9	47.2 44.2 0.5	0.5 0.125 0.0	26.5 38.1 38.3	54.1 45.1 6.9	42	1.0 0.233 0.0	53.7 67.6 65.8	94.4 44.2
334	R00Y_050_037a	0.5 0.125 0.125	0.5 0.375 0.312	0.312 0.312 0.312	390	0.5 0.124 0.124	30.8 28.8 24.2	37.6 40.0 0.5	0.5 0.125 0.125	26.8 39.0 23.5	45.6 31.1 10.9	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
335	R18Y_050_037a	0.5 0.125 0.25	0.5 0.375 0.312	0.312 0.312 0.312	371	0.5 0.124 0.243	31.0 29.6 11.1	31.7 20.6 0.5	0.5 0.125 0.25	27.4 41.2 4.3	41.4 5.9 13.9	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6
336	B63R_050_037a	0.5 0.125 0.375	0.5 0.375 0.312	0.312 0.312 0.312	349	0.5 0.124 0.381	32.0 32.0 -7.4	32.9 346.8 0.5	0.5 0.125 0.375	28.5 44.8 -14.1	47.0 342.4 14.8	348	1.0 0.0 0.683	53.5 85.4 -19.9	87.7 346.8
337	B50R_050_037a	0.5 0.125 0.5	0.5 0.375 0.312	0.312 0.312 0.312	330	0.5 0.124 0.5	33.4 35.3 -21.9	41.6 328.2 0.5	0.5 0.125 0.5	30.1 49.6 -31.2	58.6 327.8 17.3	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
338	B38R_062_050a	0.5 0.125 0.625	0.625 0.5 0.375	0.312 0.312 0.312	316	0.508 0.125 0.625	35.8 43.2 -37.0	56.9 319.4 0.5	0.5 0.125 0.625	32.1 55.3 -46.8	72.5 319.7 16.0	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4
339	B30R_075_062a	0.5 0.125 0.75	0.75 0.625 0.437	0.307 0.307 0.307	305	0.51 0.125 0.75	38.4 51.4 -52.0	73.1 314.6 0.5	0.5 0.125 0.75	34.5 61.7 -61.2	82.9 315.2 14.3	307	0.616 0.0 1.0	42.4 82.3 -83.2	117.0 314.6
340	B25R_087_075a	0.5 0.125 0.875	0.875 0.75 0.5	0.300 0.300 0.300	300	0.5 0.125 0.875	40.8 59.8 -67.2	90.0 311.6 0.5	0.5 0.125 0.875	37.2 68.3 -74.6	101.2 312.4 11.8	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
341	B20R_100_087a	0.5 0.125 1.0	1.0 0.875 0.562	0.295 0.295 0.295	295	0.489 0.125 1.0	43.6 68.8 -81.8	106.9 310.0 0.5	0.5 0.125 1.0	40.1 75.2 -87.1	115.1 310.7 9.0	294	0.416 0.0 1.0	36.3 78.6 -93.5	122.2 310.0
342	R50Y_050_050a	0.5 0.25 0.0	0.5 0.5 0.5	0.25 0.25 0.25	60	0.5 0.25 0.0	31.8 20.6 35.5	41.1 59.7 0.5	0.5 0.25 0.0	32.3 22.9 42.9	48.6 61.8 7.7	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
343	R31Y_050_037a	0.5 0.25 0.125	0.5 0.375 0.312	0.312 0.312 0.312	49	0.5 0.243 0.124	33.0 22.7 25.2	33.9 47.9 0.5	0.5 0.25 0.125	32.5 23.9 30.0	38.4 51.4 4.9	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9
344	R00Y_050_025a	0.5 0.25 0.25	0.5 0.25 0.375	0.390 0.390 0.390	390	0.5 0.249 0.249	36.4 19.2 16.1	25.1 40.0 0.5	0.5 0.25 0.25	33.0 26.3 12.1	29.0 24.7 8.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
345	R00Y_050_025a	0.5 0.25 0.375	0.5 0.25 0.375	0.360 0.360 0.360	360	0.5 0.249 0.375	36.8 20.2 1.0	20.3 2.9 0.5	0.5 0.25 0.375	33.9 30.3 -6.0	30.9 348.7 12.6	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
346	B50R_050_025a	0.5 0.25 0.5	0.5 0.25 0.375	0.330 0.330 0.330	330	0.5 0.249 0.5	38.1 23.5 -14.6	27.7 328.2 0.5	0.5 0.25 0.5	35.2 35.7 -23.2	42.6 326.9 15.2	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
347	B34R_062_037a	0.5 0.25 0.625	0.625 0.375 0.437	0.311 0.311 0.311	311	0.506 0.25 0.625	40.6 31.1 -29.7	43.3 316.7 0.5	0.5 0.25 0.625	36.8 42.2 -39.2	57.6 317.0 14.8	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7
348	B25R_075_050a	0.5 0.25 0.75	0.75 0.5 0.300 0.300	0.5 0.25 0.75	43.1 39.9 -44.8	60.0 311.6 0.5	0.25 0.75	38.8 49.3 -54.2	73.3 312.3 13.9	300 0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6			
349	B19R_087_062a	0.5 0.25 0.875	0.875 0.625 0.293 0.293	0.489 0.489 0.489	489	0.489 0.25 0.875	45.9 48.8 -59.4	76.9 309.3 0.5	0.25 0.875	41.1 56.9 -68.1	88.8 309.8 12.8	292	0.383 0.0 1.0	35.3 78.1 -95.1	123.0 309.3
350	B15R_100_075a	0.5 0.25 1.0	1.0 0.75 0.625	0.289 0.289 0.289	289	0.487 0.25 1.0	49.3 58.1 -73.1	93.4 308.4 0.5	0.25 1.0	43.7 64.7 -81.2	103.8 310.5 11.8	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
351	R76Y_050_050a	0.5 0.375 0.0	0.5 0.5 0.5	0.25 0.25 0.25	76	0.5 0.383 0.0	39.1 3.9 40.3	40.5 84.4 0.5	0.375 0.0	40.2 4.9 48.0	48.2 84.1 7.8	77	1.0 0.766 0.0	78.2 7.8 80.6	81.0 84.4
352	R68Y_050_037a	0.5 0.375 0.125	0.5 0.375 0.312	0.312 0.312 0.312	71	0.5 0.381 0.124	39.4 6.9 29.1	29.9 76.5 0.5	0.375 0.125	40.3 5.9 38.1	38.6 81.1 9.1	71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5
353	R50Y_050_025a	0.5 0.375 0.25	0.5 0.25 0.375	0.60 0.60 0.60	390	0.5 0.375 0.249	39.7 10.3 17.7	20.5 59.7 0.5	0.375 0.25	40.7 8.3 22.2	23.7 69.3 5.0	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
354	R00Y_050_012a	0.5 0.375 0.375	0.5 0.125 0.437	0.390 0.390 0.390	390	0.5 0.375 0.375	42.0 9.6 8.0	12.5 40.0 0.5	0.375 0.375	41.4 12.4 4.8	13.3 21.2 4.3	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
355	B50R_050_012a	0.5 0.375 0.5	0.5 0.125 0.437	0.330 0.330 0.330	330	0.5 0.375 0.5	42.9 11.7 -7.3	13.8 328.2 0.5	0.375 0.5	42.3 18.0 -12.2	21.8 325.7 8.0	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
356	B25R_062_025a	0.5 0.375 0.625	0.625 0.25 0.5	0.300 0.300 0.300	300	0.5 0.375 0.625	45.4 19.9 -22.4	30.0 311.6 0.5	0.375 0.625	43.6 24.8 -28.6	37.9 311.0 8.0	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
357	B15R_075_037a	0.5 0.375 0.75	0.75 0.375 0.562	0.289 0.289 0.289	289	0.493 0.375 0.75	48.5 29.0 -36.5	46.7 308.4 0.5	0.375 0.75	45.2 32.6 -44.0	54.7 306.5 8.8	288	0.316 0.0 1.0	33.9 77.4 -97.5	124.5 308.4
358	B11R_087_050a	0.5 0.375 0.875	0.875 0.5 0.625	0.284 0.284 0.284	284	0.491 0.375 0.875	51.9 38.3 -50.0	63.1 307.4 0.5	0.375 0.875	47.1 40.9 -58.5	71.4 304.9 10.0	282	0.233 0.0 1.0	32.3 76.7 -100.1	126.2 307.4
359	B09R_100_062a	0.5 0.375 1.0	1.0 0.625 0.687	0.281 0.281 0.281	281	0.489 0.375 1.0	55.6 47.8 -63.2	79.3 307.0 0.5	0.375 1.0	49.2 49.5 -72.2	87.6 304.4 11.1	279	0.183 0.0 1.0	31.7 76.5 -101.2	126.9 307.0
360	Y00G_050_050a	0.5 0.5 0.0	0.5 0.5 0.5	0.25 0.25 0.25	90	0.5 0.5 0.0	46.3 -10.3 45.3	46.5 102.8 0.5	0.5 0.0	48.9 -12.3	54.2 55.6 102.8	9.5 8.9 1.0	1.0 0.0 1.0	92.6 -20.7	90.7 93.0
361	Y00G_050_037a	0.5 0.5 0.125	0.5 0.375 0.312	0.30 0.30 0.30	90	0.5 0.5 0.124	46.6 -7.7 34.0	34.9 102.8 0.5	0.5 0.125 0.91	-11.4 46.7	48.0 103.7 13.4	89 1.0 1.0 0.0	92.6 -20.7	90.7 93.0	
362	Y00G_050_025a	0.5 0.5 0.25	0.5 0.25 0.375	0.90 0.90 0.90	390	0.5 0.5 0.249	47.0 -5.1 22.6	23.2 102.8 0.5	0.5 0.25 49.3	-9.2 32.9	34.2 105.6 11.3	89 1.0 1.0 0.0	92.6 -20.7	90.7 93.0	
363	Y00G_050_012a	0.5 0.5 0.375	0.5 0.125 0.437	0.90 0.90 0.90	390	0.5 0.5 0.375	47.3 -2.5 11.3	11.6 102.8 0.5	0.5 0.375 49.8	-5.3 16.6	17.5 107.8 6.5	89 1.0 1.0 0.0	92.6 -20.7	90.7 93.0	
364	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	0.360 0.360 0.360	360	0.5 0.5 0.5	47.7 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	49.6 0.0 0.0	0.0 325.3 2.9	360 1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	
365	B00R_062_012a	0.5 0.625 0.625	0.125 0.625 0.270	0.5 0.625 0.625	270	0.5 0.5 0.625	51.5 9.5 -12.9	16.0 306.2 0.5	0.5 0.625 51.6	6.7 -16.3 17.6	292.4 4.3 270 0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2		
366	B00R_075_025a	0.5 0.5 0.75	0.75 0.25 0.625	0.270 0.270 0.270	270	0.5 0.5 0.75	55.3 19.0 -25.8	32.1 306.2 0.5	0.5 0.75 52.8	14.4 -31.9 35.1	294.3 7.9 270 0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2		
367	B00R_087_037a	0.5 0.5 0.875	0.875 0.375 0.687	0.270 0.270 0.270	270	0.5 0.5 0.875	59.1 28.5 -38.8	48.1 306.2 0.5	0.5 0.875 54.3	23.0 -46.9 52.2	296.5 10.8 270 0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2		
368	B00R_100_050a	0.5 0.5 1.0	1.0 0.5 0.75	0.270 0.270 0.270	270	0.5 0.5 1.0	62.8 38.0 -51.7	64.2 306.2 0.5	0.5 1.0 56.0	31.9 -61.1 69.0	297.5 13.0 270 0.0 0.0 1.0	30.3 76.0 -103.5	128.5 306.2		
369	Y18G_062_062a	0.5 0.625 0.0	0.625 0.625 0.312												

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF21/PF21.L0NP.PDF>
 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	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	31.5 48.0 40.3	62.7 40.0	0.625 0.0 0.0	30.7 54.1 44.5	70.1 39.4 7.4	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
406	R31Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.114	31.7 48.7 29.7	57.0 31.3	0.625 0.1125 0.114	31.0 54.7 30.0	62.4 28.7 6.0	380	1.0 0.0 0.183	50.7 77.9 47.5	91.2 31.3
407	R11Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.239	32.1 49.6 12.8	51.3 14.4	0.625 0.0 0.25	31.5 56.2 10.9	57.2 11.0 6.7	367	1.0 0.0 0.383	51.4 79.5 20.4	82.1 14.4
408	B69R_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.385	33.0 52.2	-7.1 52.7 35.2	0.625 0.0 0.375	32.4 58.6	-7.7 59.1 35.2	6.4 352	1.0 0.0 0.616	52.9 83.6	-11.4 84.3 35.2
409	B59R_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.625 0.0 0.51	34.3 55.5	-22.8 60.1 337.6	0.625 0.0 0.5	33.8 62.1	-25.0 67.0 338.0	6.9 339	1.0 0.0 0.816	54.9 88.9	-36.6 96.2 337.6
410	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	35.8 58.9	-36.5 69.3 328.2	0.625 0.0 0.625	35.5 66.4	-41.1 78.1 328.2	8.7 330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
411	B42R_075_075a	0.625 0.0 0.75	0.75 0.75 0.375	321	0.637 0.0 0.75	38.4 66.8	-51.4 84.3 322.4	0.625 0.0 0.75	37.6 71.3	-55.9 90.6 321.8	6.4 322	0.85 0.0 1.0	51.2 89.1	-68.5 112.4 322.4
412	B36R_087_087a	0.625 0.0 0.875	0.875 0.875 0.437	314	0.641 0.0 0.875	40.8 74.7	-66.6 100.1 318.3	0.625 0.0 0.875	40.0 76.7	-69.8 103.7 317.7	3.8 315	0.733 0.0 1.0	46.6 85.4	-76.1 114.4 318.3
413	B31R_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	43.0 82.7	-82.2 116.6 315.1	0.625 0.0 1.0	42.7 82.5	-82.8 116.8 314.8	0.6 308	0.633 0.0 1.0	43.0 82.7	-82.2 116.6 315.1
414	R18Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	41	0.625 0.114 0.0	32.9 44.0	40.9 60.1 42.8	0.625 0.125 0.0	32.8 48.2	45.9 66.6 6.5 39	399	1.0 0.183 0.0	52.7 70.5	65.5 96.2 42.8
415	R00Y_062_050a	0.625 0.125 0.125	0.625 0.5 0.375	390	0.625 0.125 0.125	37.1 38.4 32.2	50.2 40.0	0.625 0.125 0.125	33.0 48.8	32.2 58.5 33.3 11.1	389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0
416	R26Y_062_050a	0.625 0.125 0.25	0.625 0.5 0.375	376	0.625 0.125 0.241	37.3 39.0	20.6 44.1 27.8	0.625 0.125 0.25	33.5 50.4	13.6 52.2 15.1 13.9	377	1.0 0.0 0.233	50.8 78.0	41.2 88.2 27.8
417	R00Y_062_050a	0.625 0.125 0.375	0.625 0.5 0.375	360	0.625 0.125 0.375	37.9 40.5	2.0 40.6 2.9	0.625 0.125 0.375	34.4 53.1	-4.8 53.3 354.8	14.7 360	1.0 0.0 0.5	52.0 81.1	4.1 81.2 2.9
418	B61R_062_050a	0.625 0.125 0.5	0.625 0.5 0.375	344	0.625 0.125 0.508	39.1 43.6	-15.3 46.2 340.6	0.625 0.125 0.5	35.6 56.7	-22.2 60.9 338.6	15.2 342	1.0 0.0 0.766	54.4 87.3	-30.6 92.5 340.6
419	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	40.5 47.1	-29.2 55.4 328.2	0.625 0.125 0.625	37.3 61.3	-38.3 72.3 327.9	17.2 330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
420	B40R_075_062a	0.625 0.125 0.75	0.75 0.625 0.437	319	0.635 0.125 0.75	43.1 55.0	-44.2 70.6 321.2	0.625 0.125 0.75	39.2 66.6	-53.4 85.3 321.2	15.2 320	0.816 0.0 1.0	49.8 88.1	-70.7 113.0 321.2
421	B34R_087_075a	0.625 0.125 0.875	0.875 0.75 0.5	311	0.637 0.125 0.875	45.5 63.1	-59.4 86.6 316.7	0.625 0.125 0.875	41.5 72.3	-67.4 98.9 317.0	12.9 311	0.683 0.0 1.0	44.8 84.1	-79.2 115.5 316.7
422	B29R_100_087a	0.625 0.125 1.0	1.0 0.875 0.562	305	0.635 0.125 1.0	48.0 71.4	-74.4 103.2 313.8	0.625 0.125 1.0	44.0 78.4	-80.5 112.4 314.2	10.1 305	0.583 0.0 1.0	41.3 81.6	-85.1 117.9 313.8
423	R38Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	53	0.625 0.239 0.0	36.6 34.0	42.6 54.6 51.3	0.625 0.25 0.0	37.4 35.7	48.5 60.2 53.5 6.1	52	1.0 0.383 0.0	58.5 54.5	68.2 87.3 51.3
424	R23Y_062_050a	0.625 0.25 0.125	0.625 0.5 0.375	44	0.625 0.241 0.125	38.8 33.8	32.9 47.2 44.2	0.625 0.25 0.125	37.6 36.4	36.8 51.8 45.2 4.8	42	1.0 0.233 0.0	57.7 67.6	65.8 94.4 44.2
425	R00Y_062_037a	0.625 0.25 0.25	0.625 0.375 0.437	390	0.625 0.25 0.25	42.7 28.8	24.2 37.6 40.0	0.625 0.25 0.25	38.0 38.2	19.6 42.9 27.1 11.4	389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0
426	R18Y_062_037a	0.625 0.25 0.375	0.625 0.375 0.437	371	0.625 0.25 0.368	43.0 29.6	11.1 31.7 20.6	0.625 0.25 0.375	38.7 41.1	1.5 41.1 21.1 15.5	371	1.0 0.0 0.316	51.1 79.1	29.7 84.5 20.6
427	B65R_062_037a	0.625 0.25 0.5	0.625 0.375 0.437	349	0.625 0.25 0.506	43.9 32.0	-7.4 32.9 346.8	0.625 0.25 0.5	39.8 45.1	-15.7 47.8 340.7	16.0 349	1.0 0.0 0.683	53.5 85.4	-19.9 87.7 346.8
428	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	45.3 35.3	-21.9 41.6 328.2	0.625 0.25 0.625	41.2 50.2	-32.1 59.6 327.4	18.5 330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
429	B38R_075_050a	0.625 0.25 0.75	0.75 0.5 0.5	316	0.633 0.25 0.75	47.8 43.2	-37.0 56.9 319.4	0.625 0.25 0.75	42.9 56.0	-47.4 73.4 319.7	17.2 317	0.766 0.0 1.0	47.9 86.4	-74.0 113.8 319.4
430	B30R_087_062a	0.625 0.25 0.875	0.875 0.625 0.562	307	0.635 0.25 0.875	50.3 51.4	-52.0 73.1 314.6	0.625 0.25 0.875	44.9 62.4	-61.8 87.9 315.2	15.7 307	0.616 0.0 1.0	42.4 82.3	-83.2 117.0 314.6
431	B25R_100_075a	0.625 0.25 1.0	1.0 0.75 0.625	300	0.625 0.25 1.0	52.8 59.8	-67.2 90.0 311.6	0.625 0.25 1.0	47.2 69.2	-75.4 102.3 312.5	13.5 300	0.5 0.0 1.0	38.5 79.8	-89.7 120.0 311.6
432	R61Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	67	0.625 0.385 0.0	43.5 16.7	46.8 49.7 70.2	0.625 0.375 0.0	44.1 19.3	52.4 55.9 69.7 6.2	67	1.0 0.616 0.0	69.6 26.8	74.8 79.5 70.2
433	R50Y_062_050a	0.625 0.375 0.125	0.625 0.5 0.375	60	0.625 0.375 0.125	43.7 20.6	35.5 41.1 59.7	0.625 0.375 0.125	44.2 20.0	43.2 47.6 65.1 7.7	59	1.0 0.5 0.0	63.6 41.3	71.0 82.2 59.7
434	R31Y_062_037a	0.625 0.375 0.25	0.625 0.375 0.437	49	0.625 0.368 0.25	44.9 22.7	25.2 33.9 47.9	0.625 0.375 0.25	44.5 21.8	27.8 35.4 51.9 2.8	48	1.0 0.316 0.0	56.2 60.6	67.2 90.5 47.9
435	R00Y_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	48.4 19.2	16.1 25.1 40.0	0.625 0.375 0.375	45.1 24.9	10.6 27.0 23.1 8.5	389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0
436	R00Y_062_025a	0.625 0.375 0.5	0.625 0.25 0.5	360	0.625 0.375 0.5	48.7 20.2	1.0 20.3 2.9	0.625 0.375 0.5	46.0 29.2	-6.4 29.9 347.5 11.9	360	1.0 0.0 0.5	52.0 81.1	4.1 81.2 2.9
437	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	50.1 23.5	-14.6 27.7 328.2	0.625 0.375 0.625	47.1 34.6	-22.9 41.5 326.5 14.1	330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
438	B34R_075_037a	0.625 0.375 0.75	0.75 0.375 0.562	311	0.631 0.375 0.75	52.5 31.5	-29.7 43.3 316.7	0.625 0.375 0.75	48.5 40.9	-38.5 56.2 316.7 13.5	311	0.683 0.0 1.0	44.8 84.1	-79.2 115.5 316.7
439	B25R_087_050a	0.625 0.375 0.875	0.875 0.5 0.625	300	0.625 0.375 0.875	55.0 39.9	-44.8 60.0 311.6	0.625 0.375 0.875	50.2 47.9	-53.3 71.7 311.9 12.6	300	0.5 0.0 1.0	38.5 79.8	-89.7 120.0 311.6
440	B19R_100_062a	0.625 0.375 1.0	1.0 0.625 0.687	293	0.614 0.375 1.0	57.8 48.8	-59.4 76.9 309.3	0.625 0.375 1.0	52.1 55.3	-67.3 87.1 309.4 11.7	292	0.383 0.0 1.0	35.3 78.1	-95.1 123.0 309.3
441	R81Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	79	0.625 0.51 0.0	50.8 1.0	51.8 51.8 88.7	0.625 0.5 0.0	51.9 1.9	57.7 57.8 88.0 6.0	80	1.0 0.816 0.0	81.2 1.7	82.9 83.0 88.7
442	R76Y_062_050a	0.625 0.5 0.125	0.625 0.5 0.375	76	0.625 0.508 0.125	51.0 3.9	40.3 40.5 84.4	0.625 0.5 0.125	52.0 2.6	50.5 50.6 86.9 10.3	77	1.0 0.766 0.0	73.2 78.8	80.6 81.0 84.4
443	R68Y_062_037a	0.625 0.5 0.25	0.625 0.375 0.437	71	0.625 0.506 0.25	51.3 6.9	29.1 29.9 76.5	0.625 0.5 0.25	52.3 4.4	37.1 37.4 83.2 8.4	71	1.0 0.683 0.0	73.4 18.5	77.6 79.8 76.5
444	R50Y_062_025a	0.625 0.5 0.375	0.625 0.25 0.5	60	0.625 0.5 0.375	51.6 10.3	17.7 20.5 59.7	0.625 0.5 0.375	52.8 7.4	21.1 22.3 70.5 4.5	59	1.0 0.5 0.0	63.6 41.3	71.0 82.2 59.7
445	R00Y_062_012a	0.625 0.5 0.5	0.625 0.125 0.562	390	0.625 0.5 0.5	54.0 9.6	8.0 12.5 40.0	0.625 0.5 0.5	53.4 11.7	4.4 12.6 20.7 4.2	389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0
446	B50R_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	54.8 11.7	-7.3 13.8 328.2	0.625 0.5 0.625	54.4 17.2	-11.8 20.9 325.5 7.1	330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
447	B25R_075_025a	0.625 0.5 0.75	0.75 0.25 0.625	300	0.625 0.5 0.75	57.3 19.9	-22.4 30.0 311.6	0.625 0.5 0.75	55.5 23.7	-27.6 36.4 310.7 6.6	300	0.5 0.0 1.0	38.5 79.8	-89.7 120.0 311.6
448	B15R_087_037a	0.625 0.5 0.875	0.875 0.375 0.687	289	0.618 0.5 0.875	60.4 29.0	-36.5 46.7 308.4	0.625 0.5 0.875	56.9 31.0	-42.7 52.8 305.9 7.3	288	0.316 0.0 1.0	33.9 77.4	-97.5 124.5 308.4
449	B11R_100_050a	0.625 0.5 1.0	1.0 0.5 0.75 284	284	0.616 0.5 1.0	63.9 38.3	-50.0 63.1 307.4	0.625 0.5 1.0	58.5 38.8	-57.1 69.0 304.2 8.8	282	0.233 0.0 1.0	32.3 76.7	-100.1 126.2 307.4
450	Y00G_062_062a	0.625 0.625 0.0	0.625 0.625 0.312	90	0.625 0.625 0.0	57.9	-12.9 56.7 58.1 102.8	0.625 0.625 0.0	60.4	-14.5 63.8 65.4 102.8 7.7	89	1.0 1.0 0.0	92.6	-20.7 90.7 93.0 102.8
451	Y00G_062_050a	0.625 0.625 0.125	0.625 0.5 0.375	90	0.625 0.625 0.125	58.2	-10.3 45.3 46.5 102.8	0.625 0.625 0.125	60.5	-13.9 58.1 59.7 103.4 13.4	89	1.0 1.0 0.0	92.6	-20.7 90.7 93.0 102.8
452	Y00G_062_037a	0.625 0.625 0.25	0.625 0.375 0.437	90	0.625 0.625 0.25	58.5	-7.7 34.0 34.9 102.8	0.625 0.625 0.25	60.7	-12.2 46.6 48.2 104.7 13.5</				

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Ma	LabCh*Ma		
486	R00Y_075_075a	0.75 0.0 0.0	0.75 0.75 0.375	390	0.75 0.0 0.0	37.8 57.7 48.4	75.3 40.0	0.75 0.0 0.0	37.5 61.9 51.9	80.8 39.9 5.5	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
487	R35Y_075_075a	0.75 0.0 0.125	0.75 0.75 0.375	381	0.75 0.0 0.112	37.9 58.2 38.8	69.9 33.6	0.75 0.0 0.125	37.7 62.4 38.9	73.5 31.9 4.2	382	1.0 0.0 0.15	50.6 77.6 51.7	93.3 33.6
488	R18Y_075_075a	0.75 0.0 0.25	0.75 0.75 0.375	371	0.75 0.0 0.237	38.3 59.3 22.3	63.4 20.6	0.75 0.0 0.25	38.1 63.5 20.8	66.9 18.1 4.4	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6
489	R00Y_075_075a	0.75 0.0 0.375	0.75 0.75 0.375	360	0.75 0.0 0.375	39.0 60.8 3.1	60.9 2.9	0.75 0.0 0.375	38.8 65.5 2.4	65.5 2.1 4.6	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
490	B65R_075_075a	0.75 0.0 0.5	0.75 0.75 0.375	349	0.75 0.0 0.512	40.1 64.1 -14.9	65.8 346.8	0.75 0.0 0.5	39.9 68.2 -15.1	69.9 347.4 4.1	348	1.0 0.0 0.683	53.5 85.4 -19.9	87.7 346.8
491	B57R_075_075a	0.75 0.0 0.625	0.75 0.75 0.375	339	0.75 0.0 0.637	41.5 67.3 -30.5	73.9 335.5	0.75 0.0 0.625	41.3 71.8 -31.6	78.4 336.2 4.5	337	1.0 0.0 0.85	55.3 89.8 -40.7	98.6 335.5
492	B50R_075_075a	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	42.9 70.7 -43.8	83.2 328.2	0.75 0.0 0.75	43.0 76.0 -47.0	89.4 328.2 6.1	330	1.0 0.0 1.0	57.2 84.3 -58.4	110.9 328.2
493	B43R_087_087a	0.75 0.0 0.875	0.875 0.875 0.375	322	0.758 0.0 0.875	45.3 78.4 -59.0	98.1 323.0	0.75 0.0 0.875	45.0 80.7 -61.5	101.5 326.3 3.4	322	0.866 0.0 1.0	51.8 94.6 -67.4	112.1 323.0
494	B38R_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4	0.75 0.0 1.0	47.2 85.8 -75.1	114.1 318.8 1.3	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4
495	R15Y_075_075a	0.75 0.125 0.0	0.75 0.75 0.375	39	0.75 0.112 0.0	39.0 54.3 48.9	73.1 41.9	0.75 0.125 0.0	39.1 57.3 52.5	77.8 42.5 4.7	37	1.0 0.15 0.0	52.0 72.4 65.2	97.4 41.9
496	R00Y_075_062a	0.75 0.125 0.125	0.75 0.625 0.437	390	0.75 0.125 0.125	43.4 48.0 40.3	62.7 40.0	0.75 0.125 0.125	39.3 57.8 40.4	70.6 34.9 10.6	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
497	R31Y_075_062a	0.75 0.125 0.25	0.75 0.625 0.437	379	0.75 0.125 0.239	43.6 48.7 29.7	57.0 31.3	0.75 0.125 0.25	39.7 59.0 22.8	63.3 21.1 12.9	380	1.0 0.0 0.183	50.7 77.9 47.5	91.2 31.3
498	R11Y_075_062a	0.75 0.125 0.375	0.75 0.625 0.437	367	0.75 0.125 0.364	44.0 49.6 12.8	51.3 14.4	0.75 0.125 0.375	40.3 61.0 4.6	61.2 4.3 14.5	367	1.0 0.0 0.383	51.4 79.5 20.4	82.1 14.4
499	B69R_075_062a	0.75 0.125 0.5	0.75 0.625 0.437	353	0.75 0.125 0.51	45.0 52.2 -7.1	52.7 352.1	0.75 0.125 0.5	41.4 64.0 -12.9	65.2 348.5 13.5	352	1.0 0.0 0.616	52.9 83.6 -11.4	84.3 352.1
500	B59R_075_062a	0.75 0.125 0.625	0.75 0.625 0.437	341	0.75 0.125 0.635	46.2 55.5 -22.8	60.1 337.6	0.75 0.125 0.625	42.7 67.7 -29.4	73.8 336.4 14.2	339	1.0 0.0 0.816	54.9 88.9 -36.6	96.2 337.6
501	B50R_075_062a	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	47.7 58.9 -32.6	69.3 328.2	0.75 0.125 0.75	44.3 72.1 -44.9	84.9 328.0 15.9	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
502	B42R_087_075a	0.75 0.125 0.875	0.875 0.75 0.5	321	0.762 0.125 0.875	50.3 66.8 -51.4	84.3 322.4	0.75 0.125 0.875	46.2 77.0 -59.5	97.3 322.3 13.6	322	0.85 0.0 1.0	51.2 89.1 -68.5	112.4 322.4
503	B36R_100_087a	0.75 0.125 1.0	1.0 0.875 0.562	314	0.766 0.125 1.0	52.7 74.7 -66.6	100.1 318.3	0.75 0.125 1.0	48.4 82.4 -73.2	110.2 318.3 10.9	315	0.733 0.0 1.0	46.6 85.4 -76.1	114.4 318.3
504	R31Y_075_075a	0.75 0.25 0.0	0.75 0.75 0.375	49	0.75 0.237 0.0	42.2 45.5 50.4	67.9 47.9	0.75 0.25 0.0	42.8 47.1 54.2	71.8 49.0 4.2	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9
505	R18Y_075_062a	0.75 0.25 0.125	0.75 0.625 0.437	41	0.75 0.239 0.125	44.8 44.0 40.9	60.1 42.8	0.75 0.25 0.125	42.9 47.6 43.8	64.7 42.6 4.9	39	1.0 0.183 0.0	52.7 70.5 65.5	96.2 42.8
506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	49.0 38.4 32.2	50.2 40.0	0.75 0.25 0.25	43.3 48.9 27.4	56.0 29.2 12.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
507	R26Y_075_050a	0.75 0.25 0.375	0.75 0.5 0.5	376	0.75 0.25 0.366	49.2 39.0 20.6	44.1 27.8	0.75 0.25 0.375	43.9 51.1 9.6	52.0 10.6 17.1	377	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8
508	R00Y_075_050a	0.75 0.25 0.5	0.75 0.5 0.5	360	0.75 0.25 0.5	49.8 40.5 2.0	40.6 2.9	0.75 0.25 0.5	44.8 54.3 -7.7	54.8 35.1 17.6	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
509	B61R_075_050a	0.75 0.25 0.625	0.75 0.5 0.5	344	0.75 0.25 0.633	51.0 43.6 -15.3	46.2 340.6	0.75 0.25 0.625	46.0 58.3 -24.3	63.1 337.3 17.9	342	1.0 0.0 0.766	54.4 87.3 -30.6	92.5 340.6
510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	52.5 47.1 -29.2	55.4 328.7	0.75 0.25 0.75	47.5 63.1 -30.9	74.6 327.6 19.8	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.7
511	B40R_087_062a	0.75 0.25 0.875	0.875 0.625 0.5	319	0.76 0.25 0.875	55.0 55.0 -44.2	70.6 321.2	0.75 0.25 0.875	49.2 68.4 -54.7	87.6 321.3 18.0	320	0.816 0.0 1.0	42.8 88.1 -70.7	113.0 321.2
512	B34R_100_075a	0.75 0.25 1.0	1.0 0.75 0.625	311	0.762 0.25 1.0	57.4 63.1 -59.4	86.6 317.7	0.75 0.25 1.0	51.2 74.3 -68.7	101.2 317.2 15.8	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7
513	R50Y_075_075a	0.75 0.375 0.0	0.75 0.75 0.375	60	0.75 0.375 0.0	47.7 31.0 53.2	61.6 51.7	0.75 0.375 0.0	48.5 32.5 57.4	65.9 60.4 4.4	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
514	R38Y_075_062a	0.75 0.375 0.125	0.75 0.625 0.437	53	0.75 0.364 0.125	48.5 34.0 42.6	54.6 51.3	0.75 0.375 0.125	48.6 33.0 48.8	59.0 55.9 6.2	52	1.0 0.383 0.0	58.5 54.5 68.2	87.3 51.3
515	R23Y_075_050a	0.75 0.375 0.25	0.75 0.5 0.5	44	0.75 0.366 0.25	50.7 33.8 32.9	47.2 44.2	0.75 0.375 0.25	48.9 34.4 34.1	48.4 44.7 2.2	42	1.0 0.233 0.0	53.7 67.6 65.8	94.4 44.2
516	R00Y_075_037a	0.75 0.375 0.375	0.75 0.375 0.562	390	0.75 0.375 0.375	54.7 28.8 24.2	37.6 40.0	0.75 0.375 0.375	49.4 36.7 17.1	40.5 25.0 11.8	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
517	R18Y_075_037a	0.75 0.375 0.5	0.75 0.375 0.562	371	0.75 0.375 0.493	54.9 29.6 11.1	31.7 20.6	0.75 0.375 0.5	50.1 40.1 0.1	40.1 0.1 15.9	371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6
518	B65R_075_037a	0.75 0.375 0.625	0.75 0.375 0.562	349	0.75 0.375 0.631	55.8 32.0 -7.4	32.9 346.8	0.75 0.375 0.625	51.1 44.4 -16.4	47.4 339.7 16.0	348	1.0 0.0 0.683	53.5 85.4 -19.9	87.7 346.8
519	B50R_075_037a	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	57.2 35.3 -21.9	41.6 328.2	0.75 0.375 0.75	52.4 49.6 -32.2	59.1 327.0 18.2	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
520	B38R_087_050a	0.75 0.375 0.875	0.875 0.5 0.625	316	0.758 0.375 0.875	59.7 43.2 -37.0	56.9 319.4	0.75 0.375 0.875	53.9 55.4 -47.2	72.8 319.5 16.9	317	0.766 0.0 1.0	47.9 86.4 -74.0	113.8 319.4
521	B30R_100_062a	0.75 0.375 1.0	1.0 0.625 0.687	307	0.76 0.375 1.0	62.2 51.4 -52.0	73.1 314.6	0.75 0.375 1.0	55.6 61.8 -61.5	87.2 315.1 15.5	307	0.616 0.0 1.0	42.4 82.3 -83.2	117.0 314.6
522	R68Y_075_075a	0.75 0.5 0.0	0.75 0.75 0.375	71	0.75 0.512 0.0	55.0 13.8 58.2	59.8 76.5	0.75 0.5 0.0	55.4 15.9 61.8	63.8 75.5 4.1	71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5
523	R61Y_075_062a	0.75 0.5 0.125	0.75 0.625 0.437	67	0.75 0.51 0.125	55.4 16.7 46.8	49.7 70.2	0.75 0.5 0.125	55.5 16.4 54.9	57.3 73.3 8.1	67	1.0 0.616 0.0	69.6 26.8 74.8	79.5 70.2
524	R50Y_075_050a	0.75 0.5 0.25	0.75 0.5 0.5	60	0.75 0.5 0.25	55.6 20.6 35.5	41.1 59.7	0.75 0.5 0.25	55.8 17.8 42.0	45.6 66.9 7.1	59	1.0 0.5 0.0	63.6 41.3 71.0	82.2 59.7
525	R31Y_075_037a	0.75 0.5 0.375	0.75 0.375 0.562	49	0.75 0.493 0.375	56.8 22.7 25.2	33.9 47.9	0.75 0.5 0.375	56.2 20.2 26.2	33.1 52.3 2.7	48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9
526	R00Y_075_025a	0.75 0.5 0.5	0.75 0.25 0.625	390	0.75 0.5 0.5	60.3 19.2 16.1	25.1 40.0	0.75 0.5 0.5	56.8 23.7 9.7	25.6 22.2 8.5	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
527	R00Y_075_025a	0.75 0.5 0.625	0.75 0.25 0.625	360	0.75 0.5 0.625	60.7 20.2 1.0	20.3 2.9	0.75 0.5 0.625	57.6 28.2 -6.6	28.9 346.7 11.4	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
528	B50R_075_025a	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	62.0 23.5 -14.6	27.7 328.2	0.75 0.5 0.75	58.7 33.5 -22.4	40.4 326.2 13.1	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2
529	B34R_087_037a	0.75 0.5 0.875	0.875 0.375 0.687	311	0.756 0.5 0.875	64.5 31.5 -29.7	43.3 316.7	0.75 0.5 0.875	60.0 39.7 -37.6	54.7 316.5 12.2	311	0.683 0.0 1.0	44.8 84.1 -79.2	115.5 316.7
530	B25R_100_050a	0.75 0.5 1.0	1.0 0.5 0.75	300	0.75 0.5 1.0	67.0 39.9 -44.8	60.0 311.6	0.75 0.5 1.0	61.4 46.4 -52.2	69.9 316.6 11.3	300	0.5 0.0 1.0	38.5 79.8 -89.7	120.0 311.6
531	R85Y_075_075a	0.75 0.625 0.0	0.75 0.75 0.375	81	0.75 0.637 0.0	62.4 -1.8 63.2	63.2 91.7	0.75 0.625 0.0	63.2 -0.7 67.1	67.1 90.6 4.1	81	1.0 0.85 0.0	83.3 -2.5 84.2	84.3 91.7
532	R81Y_075_062a	0.75 0.625 0.125	0.75 0.625 0.437	79	0.75 0.635 0.125	62.7 1.0 51.8	51.8 88.7	0.75 0.625 0.125	63.3 -0.2 61.6	61.6 90.2 9.8	80	1.0 0.816 0.0	81.2 17.7 82.9	83.0 88.7
533	R76Y_075_050a	0.75 0.625 0.25	0.75 0.5 0.5	76	0.75 0.633 0.25	62.9 3.9 40.3	40.5 84.4	0.75 0.625 0.25	63.5 1.0 50.4	50.5 88.7 10.5	77	1.0 0.766 0.0	78.2 7.8 80.6	81.0 84.4
534	R68Y_075_037a	0.75 0.625 0.375	0.75 0.375 0.562	71	0.75 0.631 0.375	63.3 6.9 29.1	29.9 76.5	0.75 0.625 0.375	63.8 3.4 35.9	36.1 84.5 7.7	71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5
535	R50Y_075_025a	0.75 0.625 0.5	0.75 0.25 0.625	60	0.75 0.625 0.5									

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF21/PF21L0NP.PDF> / .PS
 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	iet_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMa	rgb*Ma	LabCh*Ma			
567	R00Y_087_087a	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.0	44.1 67.3 56.4	87.8 40.0	0.875 0.0 0.0	44.1 69.5 58.3	90.8 39.9	2.9 389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
568	R36Y_087_087a	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.116	44.2 67.7 47.1	82.5 34.8	0.875 0.0 0.125	44.2 69.9 47.2	84.3 34.0	2.2 382	1.0 0.0 0.133	50.6 77.3 53.9	94.4 34.8	
569	R23Y_087_087a	0.875 0.0 0.25	0.875 0.875 0.437	374	0.875 0.0 0.233	44.5 68.5 32.2	75.7 25.1	0.875 0.0 0.25	44.5 70.8 30.2	77.0 23.1	2.9 375	1.0 0.0 0.266	50.9 78.3 36.8	86.6 25.1	
570	R08Y_087_087a	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.364	45.1 70.2 13.8	71.6 11.1	0.875 0.0 0.375	45.1 72.4 12.2	73.4 9.5	2.6 365	1.0 0.0 0.416	51.5 80.3 15.8	81.8 11.1	
571	B70R_087_087a	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.51	46.1 72.8 6.0	73.0 355.2	0.875 0.0 0.5	46.0 74.6 5.3	74.8 355.8	1.9 354	1.0 0.0 0.583	52.7 83.2 6.9	83.5 355.2	
572	B63R_087_087a	0.875 0.0 0.625	0.875 0.875 0.437	346	0.875 0.0 0.641	47.2 75.6 23.1	79.1 342.9	0.875 0.0 0.625	47.1 77.6 22.1	80.7 344.0	2.2 344	1.0 0.0 0.733	54.0 86.5 26.4	90.4 342.9	
573	B56R_087_087a	0.875 0.0 0.75	0.875 0.875 0.437	338	0.875 0.0 0.758	48.6 78.8 37.5	87.3 334.5	0.875 0.0 0.75	48.5 81.2 37.9	89.6 334.9	2.3 337	1.0 0.0 0.866	55.5 90.1 42.8	99.8 334.5	
574	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	50.1 82.5 31.1	97.1 328.2	0.875 0.0 0.875	50.2 85.3 32.8	100.3 328.2	3.2 330	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2	
575	B44R_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	52.5 90.1 66.3	111.9 323.6	0.875 0.0 1.0	52.1 89.8 66.9	112.0 323.3	0.7 323	0.883 0.0 1.0	52.5 90.1 66.3	111.9 323.6	
576	R13Y_087_087a	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.116 0.0	45.2 64.2 56.9	85.8 41.9	0.875 0.125 0.0	45.3 65.8 58.8	88.3 41.7	2.5 37	1.0 0.133 0.0	51.7 73.4 65.0	98.0 41.5	
577	R00Y_087_075a	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.125	49.7 57.7 48.4	75.3 40.0	0.875 0.125 0.125	45.5 66.2 48.3	81.9 36.0	9.5 389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
578	R35Y_087_075a	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.237	49.9 58.2 38.8	69.9 33.6	0.875 0.125 0.25	45.8 67.1 41.3	74.3 25.3	12.0 382	1.0 0.0 0.15	50.6 77.6 51.7	93.3 33.6	
579	R18Y_087_075a	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.362	50.2 59.3 22.3	63.4 20.6	0.875 0.125 0.375	46.4 68.8 13.9	70.2 11.4	13.2 371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6	
580	R00Y_087_075a	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.5	50.9 60.8 3.1	60.9 2.9	0.875 0.125 0.5	47.2 71.1 3.6	71.2 357.1	12.8 360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	
581	B65R_087_075a	0.875 0.125 0.625	0.875 0.75 0.5	349	0.875 0.125 0.637	52.1 64.1 14.9	65.8 346.8	0.875 0.125 0.625	48.3 74.2 20.3	76.9 344.6	12.0 348	1.0 0.0 0.683	53.5 85.4 19.9	87.7 346.8	
582	B57R_087_075a	0.875 0.125 0.75	0.875 0.75 0.5	339	0.875 0.125 0.762	53.4 67.3 30.5	73.9 335.5	0.875 0.125 0.75	49.6 77.9 36.1	85.9 335.1	12.5 337	1.0 0.0 0.85	55.3 89.8 40.7	98.6 335.5	
583	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	54.9 70.7 43.8	83.2 328.2	0.875 0.125 0.875	51.3 82.1 51.1	96.7 328.1	14.0 330	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2	
584	B43R_100_087a	0.875 0.125 1.0	1.0 1.0 0.875	322	0.883 0.125 1.0	57.3 78.4 59.0	98.1 323.0	0.875 0.125 1.0	53.1 86.9 65.3	106.7 323.0	11.3 322	0.866 0.0 1.0	51.8 89.6 67.4	112.1 323.0	
585	R26Y_087_087a	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.233 0.0	47.8 57.0 58.0	81.3 45.5	0.875 0.25 0.0	48.4 57.3 60.1	83.0 46.3	2.1 44	1.0 0.266 0.0	54.6 65.1 66.3	93.0 45.5	
586	R15Y_087_075a	0.875 0.25 0.125	0.875 0.75 0.5	39	0.875 0.237 0.125	50.9 54.3 48.9	73.1 41.9	0.875 0.25 0.125	48.5 57.7 50.8	76.9 41.3	4.5 37	1.0 0.15 0.0	52.0 72.4 65.2	97.4 41.9	
587	R00Y_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	55.4 48.0 40.3	62.7 40.0	0.875 0.25 0.25	48.8 58.7 35.3	68.5 31.0	13.4 389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
588	R31Y_087_062a	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.364	55.5 48.7 29.7	57.0 31.3	0.875 0.25 0.375	49.3 60.4 17.9	63.0 16.5	17.7 380	1.0 0.0 0.183	50.7 77.9 47.5	91.2 31.3	
589	R11Y_087_062a	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.489	55.9 49.6 12.8	51.3 14.4	0.875 0.25 0.5	50.1 63.0 0.6	63.0 0.5	18.9 367	1.0 0.0 0.383	51.4 79.5 20.4	82.1 14.4	
590	B69R_087_062a	0.875 0.25 0.625	0.875 0.625 0.562	353	0.875 0.25 0.635	56.9 52.2 7.1	52.7 352.1	0.875 0.25 0.625	51.1 66.2 16.0	68.1 346.3	17.5 352	1.0 0.0 0.616	52.9 83.6 11.4	84.3 352.1	
591	B59R_087_062a	0.875 0.25 0.75	0.875 0.625 0.562	341	0.875 0.25 0.76	58.2 55.5 22.8	60.1 337.6	0.875 0.25 0.75	52.3 70.2 31.9	77.1 335.5	18.1 339	1.0 0.0 0.816	54.9 88.9 36.6	96.2 337.6	
592	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	59.6 58.9 36.5	69.3 328.2	0.875 0.25 0.875	53.8 74.7 47.0	88.3 327.8	19.8 330	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2	
593	B42R_100_075a	0.875 0.25 1.0	1.0 1.0 0.75	325	0.887 0.25 1.0	62.2 66.8 51.4	84.1 324.4	0.875 0.25 1.0	55.6 79.8 67.3	100.7 324.2	17.6 322	0.85 0.0 1.0	51.2 89.1 68.5	112.4 322.4	
594	R41Y_087_087a	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.364 0.0	52.5 44.4 60.6	75.1 53.7	0.875 0.375 0.0	53.2 44.5 62.6	76.8 54.5	2.1 54	1.0 0.416 0.0	60.0 50.7 69.3	85.9 53.7	
595	R31Y_087_075a	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.362 0.125	54.1 45.5 50.4	67.9 47.9	0.875 0.375 0.125	53.3 44.9 54.7	70.8 50.6	4.4 48	1.0 0.316 0.0	56.2 60.6 67.2	90.5 47.9	
596	R18Y_087_062a	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.364 0.25	56.8 44.0 40.9	60.1 42.8	0.875 0.375 0.25	53.5 45.9 40.7	61.4 41.5	3.7 39	1.0 0.183 0.0	57.2 70.5 65.5	96.2 42.8	
597	R00Y_087_050a	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	61.0 38.4 32.2	50.2 40.0	0.875 0.375 0.375	54.0 47.8 24.1	53.6 26.8	14.2 389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
598	R26Y_087_050a	0.875 0.375 0.5	0.875 0.5 0.625	376	0.875 0.375 0.491	61.1 39.0 20.6	44.1 27.8	0.875 0.375 0.5	54.6 50.5 7.2	51.0 8.1	18.8 377	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8	
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 40.5 2.0	40.6 2.9	0.875 0.375 0.625	55.5 54.0 9.3	54.8 35.0	18.7 360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	
600	B61R_087_050a	0.875 0.375 0.75	0.875 0.5 0.625	344	0.875 0.375 0.758	62.9 43.6 15.3	46.2 340.6	0.875 0.375 0.75	56.6 58.3 25.2	63.5 336.5	18.7 342	1.0 0.0 0.766	54.4 87.3 30.6	92.5 340.6	
601	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	64.4 47.1 29.2	55.4 328.2	0.875 0.375 0.875	58.0 63.2 40.5	75.0 327.3	20.6 330	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2	
602	B40R_100_062a	0.875 0.375 1.0	1.0 1.0 0.625	319	0.885 0.375 1.0	66.9 55.0 44.2	70.6 321.2	0.875 0.375 1.0	59.5 68.6 55.0	88.0 321.2	18.8 320	0.816 0.0 1.0	49.8 88.1 70.7	113.0 321.2	
603	R58Y_087_087a	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.51 0.0	59.4 27.1 64.8	70.2 67.2	0.875 0.5 0.0	59.4 29.0 66.2	72.3 66.2	2.4 65	1.0 0.583 0.0	67.9 31.0 74.0	80.3 67.2	
604	R50Y_087_075a	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.5 0.125	59.6 31.0 53.2	61.6 59.7	0.875 0.5 0.125	59.4 29.5 59.8	66.7 63.7	6.7 52	1.0 0.5 0.0	63.6 41.3	71.0 82.2	59.7
605	R38Y_087_062a	0.875 0.5 0.25	0.875 0.625 0.562	53	0.875 0.489 0.25	60.4 34.0 42.6	54.6 51.3	0.875 0.5 0.25	59.7 30.6 47.4	56.4 57.1	5.9 59	1.0 0.383 0.0	58.5 54.5	68.2 87.3	51.3
606	R23Y_087_050a	0.875 0.5 0.375	0.875 0.5 0.625	44	0.875 0.491 0.375	62.6 33.8 32.9	47.2 44.2	0.875 0.5 0.375	60.0 32.5 31.9	45.6 44.4	3.0 42	1.0 0.233 0.0	53.7 67.6	65.8 94.4	44.2
607	R00Y_087_037a	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.5	66.6 28.8 24.2	37.6 40.0	0.875 0.5 0.5	60.6 35.3 15.5	38.6 23.7	12.3 389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	
608	R18Y_087_037a	0.875 0.5 0.625	0.875 0.375 0.687	371	0.875 0.5 0.618	66.8 29.6 11.1	31.7 20.6	0.875 0.5 0.625	61.3 39.0 0.7	39.0 358.9	16.0 371	1.0 0.0 0.316	51.1 79.1 29.7	84.5 20.6	
609	B63R_087_037a	0.875 0.5 0.75	0.875 0.375 0.687	349	0.875 0.5 0.756	67.7 32.0 7.4	32.9 346.8	0.875 0.5 0.75	62.3 43.5 16.6	46.5 339.0	15.6 348	1.0 0.0 0.683	53.5 85.4 19.9	87.7 346.8	
610	B50R_087_037a	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	69.1 35.3 21.9	41.6 328.2	0.875 0.5 0.875	63.5 48.6 31.9	58.2 326.7	17.6 330	1.0 0.0 1.0	57.2 94.3 58.4	110.9 328.2	
611	B38R_100_050a	0.875 0.5 1.0	1.0 0.5 0.75	316	0.883 0.5 1.0	71.6 43.2 37.0	56.9 319.4	0.875 0.5 1.0	64.8 54.4 46.6	71.7 319.3	16.3 317	0.766 0.0 1.0	47.9 86.4 74.0	113.8 319.4	
612	R73Y_087_087a	0.875 0.625 0.0	0.875 0.875 0.437	74	0.875 0.641 0.0	66.7 10.5 69.4	70.2 81.3	0.875 0.625 0.0	66.5 12.7 70.9	72.0 79.7	2.7 75	1.0 0.733 0.0	76.2 12.0 79.3	80.2 81.3	
613	R68Y_087_075a	0.875 0.625 0.125	0.875 0.75 0.5	71	0.875 0.637 0.125	67.0 13.8 58.2	59.8 76.5	0.875 0.625 0.125	66.5 13.2 65.6	66.9 78.6	7.4 71	1.0 0.683 0.0	73.4 18.5 77.6	79.8 76.5	
614	R61Y_087_062a	0.875 0.625 0.25	0.875 0.625 0.562	67	0.875 0.635 0.25	67.4 16.7 46.8	49.7 70.7	0.875 0.625 0.25	66.7 14.3 54.8	56.6 75.3	8.4 67	1			

n	HIC*Fa	rgb_Fa	icr_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Ma		
648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5	100.4 39.9 0.0	389	1.0 0.0 0.0	50.4 76.9 64.5	100.4 40.0
649	R38Y_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	50.5 77.2 55.6	95.1 35.7	1.0 0.0 0.125	50.6 77.2 54.9	94.8 35.4 0.6	383	1.0 0.0 0.116	50.5 77.2 55.6	95.1 35.7
650	R26Y_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8	1.0 0.0 0.25	50.8 77.9 39.2	87.2 26.6 2.0	377	1.0 0.0 0.233	50.8 78.0 41.2	88.2 27.8
651	R13Y_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	51.3 79.3 22.7	82.5 16.0	1.0 0.0 0.375	51.3 79.2 21.6	82.1 15.2 1.1	368	1.0 0.0 0.366	51.3 79.3 22.7	82.5 16.0
652	R00Y_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9 0.0	360	1.0 0.0 0.5	52.0 81.1 4.1	81.2 2.9
653	B68R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	53.0 83.9	-13.6 85.0 350.7	1.0 0.0 0.625	53.0 83.6	-12.6 84.6 351.4 1.0	351	1.0 0.0 0.633	53.0 83.9	-13.6 85.0 350.7
654	B61R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	54.4 87.3	-30.6 92.5 340.6	1.0 0.0 0.75	54.2 86.7	-28.9 91.3 341.6 2.0	342	1.0 0.0 0.766	54.4 87.3	-30.6 92.5 340.6
655	B55R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	55.7 90.6	-44.8 101.1 333.6	1.0 0.0 0.875	55.6 90.3	-43.9 100.4 334.0 0.9	336	1.0 0.0 0.883	55.7 90.6	-44.8 101.1 333.6
656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2	1.0 0.0 1.0	57.2 94.3	-58.4 111.0 328.2 0.0	330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
657	R11Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	370	1.0 0.116 0.0	51.4 74.1	64.9 98.5 41.2	1.0 0.125 0.0	51.5 73.9	64.9 98.3 41.3 0.2	36	1.0 0.116 0.0	51.4 74.1	64.9 98.5 41.2
658	R00Y_100_087a	1.0 0.125 0.125	1.0 0.875 0.562	390	1.0 0.125 0.125	56.1 67.3	56.4 87.8 40.0	1.0 0.125 0.125	51.6 74.2	55.7 92.8 36.9 8.2	389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0
659	R36Y_100_087a	1.0 0.125 0.25	1.0 0.875 0.562	382	1.0 0.125 0.241	56.2 67.7	47.1 82.5 34.8	1.0 0.125 0.25	51.9 74.9	40.3 85.1 28.3 10.8	382	1.0 0.0 0.133	50.6 77.3	63.9 100.4 34.8
660	R23Y_100_087a	1.0 0.125 0.375	1.0 0.875 0.562	374	1.0 0.125 0.358	56.4 68.5	32.2 75.7 25.1	1.0 0.125 0.375	52.3 76.3	23.0 79.7 16.7 12.7	375	1.0 0.0 0.266	50.9 78.3	36.8 86.6 25.1
661	R08Y_100_087a	1.0 0.125 0.5	1.0 0.875 0.562	365	1.0 0.125 0.489	57.0 70.2	13.2 71.6 11.1	1.0 0.125 0.5	53.0 78.2	5.6 78.4 4.1 12.1	365	1.0 0.0 0.416	51.5 80.3	15.8 81.8 11.1
662	B70R_100_087a	1.0 0.125 0.625	1.0 0.875 0.562	355	1.0 0.125 0.635	58.0 72.8	-6.0 73.0 355.2	1.0 0.125 0.625	53.9 80.8	-11.1 81.5 352.1 10.3	354	1.0 0.0 0.583	52.7 83.2	-6.9 83.5 355.2
663	B63R_100_087a	1.0 0.125 0.75	1.0 0.875 0.562	346	1.0 0.125 0.766	59.2 75.6	-23.1 79.1 342.9	1.0 0.125 0.75	55.1 83.9	-27.2 88.2 342.0 10.0	344	1.0 0.0 0.733	54.0 86.5	-26.4 90.4 342.9
664	B56R_100_087a	1.0 0.125 0.875	1.0 0.875 0.562	338	1.0 0.125 0.883	60.5 78.8	-37.5 87.3 334.5	1.0 0.125 0.875	56.5 87.6	-42.5 97.4 334.1 10.8	337	1.0 0.0 0.866	55.5 90.1	-42.8 99.8 334.5
665	B50R_100_087a	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	62.0 82.5	-51.1 97.1 328.2	1.0 0.125 1.0	58.1 91.8	-57.0 108.0 328.1 11.6	330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
666	R23Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.7 67.6	65.8 94.4 44.2	1.0 0.25 0.0	54.0 66.7	65.9 93.8 44.6 1.0	42	1.0 0.233 0.0	53.7 67.6	65.8 94.4 44.2
667	R13Y_100_087a	1.0 0.25 0.125	1.0 0.875 0.562	38	1.0 0.241 0.125	57.1 64.2	56.9 85.8 41.5	1.0 0.25 0.125	54.1 67.0	57.6 88.4 40.7 4.1	37	1.0 0.133 0.0	51.7 73.4	65.0 98.0 41.5
668	R00Y_100_075a	1.0 0.25 0.25	1.0 0.75 0.625	390	1.0 0.25 0.25	61.7 57.7	48.4 75.3 40.0	1.0 0.25 0.25	54.4 67.8	43.1 80.3 32.4 13.5	389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0
669	R35Y_100_075a	1.0 0.25 0.375	1.0 0.75 0.625	381	1.0 0.25 0.362	61.8 58.2	38.8 69.9 33.6	1.0 0.25 0.375	54.8 69.2	26.2 74.0 20.7 18.1	382	1.0 0.0 0.15	50.6 77.6	51.7 93.3 33.6
670	R18Y_100_075a	1.0 0.25 0.5	1.0 0.75 0.625	371	1.0 0.25 0.487	62.1 59.3	22.3 63.4 20.6	1.0 0.25 0.5	55.5 71.3	9.0 71.8 7.2 19.0	371	1.0 0.0 0.316	51.1 79.1	29.7 84.5 20.6
671	R00Y_100_075a	1.0 0.25 0.625	1.0 0.75 0.625	360	1.0 0.25 0.625	62.9 60.8	3.1 60.9 2.9	1.0 0.25 0.625	56.3 74.0	-7.6 74.4 354.1 18.1	360	1.0 0.0 0.5	52.0 81.1	4.1 81.2 2.9
672	B65R_100_075a	1.0 0.25 0.75	1.0 0.75 0.625	349	1.0 0.25 0.762	64.0 64.1	-14.9 65.8 346.8	1.0 0.25 0.75	57.4 77.7	-23.6 80.8 342.9 17.1	348	1.0 0.0 0.683	53.5 85.4	-19.9 87.7 346.8
673	B57R_100_075a	1.0 0.25 0.875	1.0 0.75 0.625	339	1.0 0.25 0.887	65.3 67.3	-20.5 73.9 335.5	1.0 0.25 0.875	59.7 81.2	-39.0 91.1 334.3 17.5	337	1.0 0.0 0.85	55.3 89.8	-40.7 98.6 335.5
674	B50R_100_075a	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	66.8 70.7	-43.8 83.2 328.2	1.0 0.25 1.0	60.2 85.6	-53.6 101.0 327.9 18.9	330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
675	R36Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	57.9 56.2	67.9 88.1 50.3	1.0 0.375 0.0	58.2 55.4	67.9 87.7 50.7 0.7	51	1.0 0.366 0.0	57.9 56.2	67.9 88.1 50.3
676	R26Y_100_087a	1.0 0.375 0.125	1.0 0.875 0.562	46	1.0 0.358 0.125	59.7 57.0	58.0 81.3 45.5	1.0 0.375 0.125	58.2 55.8	60.8 82.5 47.4 3.3	44	1.0 0.266 0.0	54.6 65.1	66.3 93.0 45.5
677	R15Y_100_075a	1.0 0.375 0.25	1.0 0.75 0.625	39	1.0 0.362 0.25	62.8 54.3	48.9 73.1 41.9	1.0 0.375 0.25	58.5 56.6	47.5 73.9 39.9 5.1	37	1.0 0.15 0.0	52.0 72.4	65.2 97.4 41.9
678	R00Y_100_062a	1.0 0.375 0.375	1.0 0.625 0.687	390	1.0 0.375 0.375	67.3 48.0	40.3 62.7 40.0	1.0 0.375 0.375	58.9 58.1	31.4 66.1 28.3 15.8	389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0
679	R31Y_100_062a	1.0 0.375 0.5	1.0 0.625 0.687	379	1.0 0.375 0.489	67.4 48.7	29.7 57.0 31.3	1.0 0.375 0.5	59.4 60.3	14.6 62.1 13.6 20.6	380	1.0 0.0 0.183	50.7 77.9	47.5 91.2 31.3
680	R11Y_100_062a	1.0 0.375 0.625	1.0 0.625 0.687	367	1.0 0.375 0.614	67.9 49.6	12.8 51.3 14.4	1.0 0.375 0.625	60.2 63.2	-1.8 63.2 358.3 21.3	367	1.0 0.0 0.383	51.4 79.5	20.4 82.1 14.4
681	B69R_100_062a	1.0 0.375 0.75	1.0 0.625 0.687	353	1.0 0.375 0.76	68.8 52.2	-7.1 52.7 352.1	1.0 0.375 0.75	61.2 66.7	-17.9 69.1 344.9 19.6	352	1.0 0.0 0.616	52.9 83.6	-11.4 84.3 352.1
682	B59R_100_062a	1.0 0.375 0.875	1.0 0.625 0.687	341	1.0 0.375 0.885	70.1 55.5	-22.8 60.1 337.6	1.0 0.375 0.875	62.4 70.9	-33.3 78.3 334.8 20.0	339	1.0 0.0 0.816	54.9 88.9	-36.6 96.2 337.6
683	B50R_100_062a	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	71.5 58.9	-36.5 69.3 328.2	1.0 0.375 1.0	63.8 75.6	-48.0 89.6 327.5 21.6	330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	63.6 41.3	71.0 82.2 59.7	1.0 0.5 0.0	63.6 41.3	71.0 82.2 59.7 0.0	59	1.0 0.5 0.0	63.6 41.3	71.0 82.2 59.7
685	R41Y_100_087a	1.0 0.5 0.125	1.0 0.875 0.562	55	1.0 0.489 0.125	64.4 44.4	60.6 75.1 53.7	1.0 0.5 0.125	63.7 41.7	65.0 77.2 57.3 5.1	54	1.0 0.416 0.0	60.0 50.7	69.3 85.9 53.7
686	R31Y_100_075a	1.0 0.5 0.25	1.0 0.75 0.625	49	1.0 0.487 0.25	66.0 45.5	50.4 67.9 47.9	1.0 0.5 0.25	63.9 42.6	53.1 68.1 51.2 4.5	48	1.0 0.316 0.0	56.2 60.6	67.2 90.5 47.9
687	R18Y_100_062a	1.0 0.5 0.375	1.0 0.625 0.687	41	1.0 0.489 0.375	68.7 44.0	40.9 60.1 42.8	1.0 0.5 0.375	64.2 44.1	38.0 58.3 40.7 5.3	39	1.0 0.183 0.0	52.7 70.5	65.5 96.2 42.8
688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	72.9 38.4	32.2 50.2 40.0	1.0 0.5 0.5	64.7 46.4	21.9 51.3 25.2 15.4	389	1.0 0.0 0.0	50.4 76.9	64.5 100.4 40.0
689	R26Y_100_050a	1.0 0.5 0.625	1.0 0.5 0.75	376	1.0 0.5 0.616	73.1 39.0	20.6 44.1 27.8	1.0 0.5 0.625	65.4 49.5	5.6 49.8 6.5 19.8	377	1.0 0.0 0.233	50.8 78.0	41.2 88.2 27.8
690	R00Y_100_050a	1.0 0.5 0.75	1.0 0.5 0.75	360	1.0 0.5 0.75	73.7 40.5	20.0 40.6 2.9	1.0 0.5 0.75	66.3 53.2	-10.2 54.2 349.1 19.1	360	1.0 0.0 0.5	52.0 81.1	4.1 81.2 2.9
691	B61R_100_050a	1.0 0.5 0.875	1.0 0.5 0.75	344	1.0 0.5 0.883	74.9 43.6	-15.3 46.2 340.6	1.0 0.5 0.875	67.3 57.6	-25.6 63.1 335.9 18.9	342	1.0 0.0 0.766	54.4 87.3	-30.6 92.5 340.6
692	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	76.3 47.1	-29.2 55.4 328.2	1.0 0.5 1.0	68.6 62.6	-40.5 74.6 327.0 20.6	330	1.0 0.0 1.0	57.2 94.3	-58.4 110.9 328.2
693	R63Y_100_100a	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	70.5 24.7	75.4 79.4 71.8	1.0 0.625 0.0	70.1 25.8	75.0 79.3 71.0 1.2	68	1.0 0.633 0.0	70.5 24.7	75.4 79.4 71.8
694	R58Y_100_087a	1.0 0.625 0.125	1.0 0.875 0.562	65	1.0 0.635 0.125	71.3 27.1	64.8 70.2 67.2	1.0 0.625 0.125	70.1 26.1	70.0 74.7 69.5 5.4	65	1.0 0.583 0.0	67.9 31.0	74.0 80.3 67.2
695	R50Y_100_075a	1.0 0.625 0.25	1.0 0.75 0.625	60	1.0 0.625 0.25	71.6 31.0	53.2 61.6 59.7	1.0 0.625 0.25	70.3 27.0	59.6 65.4 65.5 7.5	59	1.0 0.5 0.0	63.6 41.3	71.0 82.2 59.7
696	R38Y_100_062a	1.0 0.625 0.375	1.0 0.625 0.687	53	1.0 0.614 0.375	72.3 34.0	42.6 54.6 51.3	1.0 0.625 0.375	70.6 28.6	45.7 53.9 57.8 6.4	52	1.0 0.383 0.0	58.5 54.5	68.2 87.3 51.3
697	R23Y_100_050a	1.0 0.625 0.5	1.0 0.5 0.75	44	1.0 0.616 0.5	74.5 33.8	32.9 47.2 44.2	1.0 0.625 0.5	71.0 31.0	30.2 43.3 44.3 5.2	42	1.0 0.233 0.0	53.7 67.6	65.8 94.4 44.2
698	R00Y_100_037a	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	78.5 28.8	24.2 37.6 40.0	1.0 0.625 0.625	71.6					

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	rgb*Md	LabCh*Md	
729	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 1.0 1.0	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0
730	G50B_100_012a	0.875 1.0 1.0	1.0 1.0 0.125	0.937 210	0.875 1.0 1.0	94.3 -5.7 -1.6	6.0 196.3	0.875 1.0 1.0	93.3 -9.7 -3.3	10.3 198.8 4.4	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
731	G50B_100_025a	0.75 1.0 1.0	1.0 0.25 0.875	210	0.75 1.0 1.0	93.2 -11.3 -3.3	12.0 196.3	0.75 1.0 1.0	91.5 -18.9 -6.2	19.9 198.1 8.0	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
732	G50B_100_037a	0.625 1.0 1.0	1.0 0.375 0.812	210	0.625 1.0 1.0	92.2 -17.3 -5.0	18.0 196.3	0.625 1.0 1.0	90.0 -27.0 -8.5	28.3 197.6 10.5	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
733	G50B_100_050a	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	91.1 -23.0 -6.7	24.0 196.3	0.5 1.0 1.0	88.8 -33.9 -10.4	35.4 197.1 11.6	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
734	G50B_100_062a	0.375 1.0 1.0	1.0 0.625 0.687	210	0.375 1.0 1.0	90.0 -28.8 -8.4	30.0 196.3	0.375 1.0 1.0	87.9 -39.3 -11.8	41.0 196.8 11.1	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
735	G50B_100_075a	0.25 1.0 1.0	1.0 0.75 0.625	210	0.25 1.0 1.0	89.0 -34.6 -10.1	36.1 196.3	0.25 1.0 1.0	87.3 -43.0 -12.8	44.9 196.5 9.0	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
736	G50B_100_087a	0.125 1.0 1.0	1.0 0.875 0.562	210	0.125 1.0 1.0	87.9 -40.4 -11.8	42.1 196.3	0.125 1.0 1.0	87.0 -45.2 -13.3	47.2 196.4 5.1	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
737	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3	0.0 1.0 1.0	86.8 -46.1 -13.5	48.1 196.3 0.0	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
738	ROOY_100_012a	1.0 0.875 0.875	1.0 0.125 0.937	390	1.0 0.875 0.875	89.7 9.6 8.0	12.5 40.0	1.0 0.875 0.875	87.1 10.5 3.8	11.2 20.1 5.0	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
739	NW_087a	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	83.4 0.0 0.0	0.0 0.0	0.875 0.875 0.875	84.7 0.0 0.0	0.0 32.5 1.2	360 1.0 1.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
740	G50B_087_012a	0.75 0.875 0.875	0.875 0.125 0.812	210	0.75 0.875 0.875	82.4 -5.7 -1.6	6.0 196.3	0.75 0.875 0.875	82.5 -10.0 -3.3	10.5 198.7 4.5	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
741	G50B_087_025a	0.625 0.875 0.875	0.875 0.25 0.75	210	0.625 0.875 0.875	81.3 -11.5 -3.3	12.0 196.3	0.625 0.875 0.875	80.7 -19.1 -6.2	20.1 197.9 8.1	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
742	G50B_087_037a	0.5 0.875 0.875	0.875 0.375 0.687	210	0.5 0.875 0.875	80.2 -17.3 -5.0	18.0 196.3	0.5 0.875 0.875	79.3 -27.1 -8.5	28.4 197.4 10.4	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
743	G50B_087_050a	0.375 0.875 0.875	0.875 0.5 0.625	210	0.375 0.875 0.875	79.2 -23.0 -6.7	24.0 196.3	0.375 0.875 0.875	78.3 -33.4 -10.2	34.9 196.9 10.9	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
744	G50B_087_062a	0.25 0.875 0.875	0.875 0.625 0.562	210	0.25 0.875 0.875	78.1 -28.8 -8.4	30.0 196.3	0.25 0.875 0.875	77.5 -37.9 -11.3	39.6 196.6 9.5	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
745	G50B_087_075a	0.125 0.875 0.875	0.875 0.75 0.5	210	0.125 0.875 0.875	77.0 -34.6 -10.1	36.1 196.3	0.125 0.875 0.875	77.1 -40.6 -12.0	42.4 196.4 6.3	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
746	G50B_087_087a	0.0 0.875 0.875	0.875 0.875 0.437	210	0.0 0.875 0.875	76.0 -40.4 -11.8	42.1 196.3	0.0 0.875 0.875	77.0 -41.7 -12.2	43.5 196.3 1.7	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
747	ROOY_100_025a	1.0 0.75 0.75	1.0 0.25 0.875	390	1.0 0.75 0.75	84.1 19.2 16.1	25.1 40.0	1.0 0.75 0.75	79.2 21.9 8.5	23.5 21.3 9.4	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
748	ROOY_087_012a	0.875 0.75 0.75	0.875 0.125 0.812	390	0.875 0.75 0.75	77.8 9.6 8.0	12.5 40.0	0.875 0.75 0.75	76.2 10.8 4.0	11.6 20.3 4.5	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
749	NW_075a	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	71.5 0.0 0.0	0.0 0.0	0.75 0.75 0.75	73.7 0.0 0.0	0.0 32.5 2.1	360 1.0 1.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
750	G50B_075_012a	0.625 0.75 0.75	0.75 0.125 0.687	210	0.625 0.75 0.75	70.4 -5.7 -1.6	6.0 196.3	0.625 0.75 0.75	71.5 -10.2 -3.4	10.8 198.5 4.9	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
751	G50B_075_025a	0.5 0.75 0.75	0.75 0.25 0.625	210	0.5 0.75 0.75	69.4 -11.5 -3.3	12.0 196.3	0.5 0.75 0.75	69.8 -19.4 -6.2	20.3 197.8 8.3	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
752	G50B_075_037a	0.375 0.75 0.75	0.75 0.375 0.562	210	0.375 0.75 0.75	68.3 -17.3 -5.0	18.0 196.3	0.375 0.75 0.75	68.4 -26.9 -8.3	28.2 197.1 10.1	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
753	G50B_075_050a	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.75	67.2 -23.0 -6.7	24.0 196.3	0.25 0.75 0.75	67.5 -32.5 -9.7	33.9 196.7 9.8	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
754	G50B_075_062a	0.125 0.75 0.75	0.75 0.625 0.437	210	0.125 0.75 0.75	66.2 -28.8 -8.4	30.0 196.3	0.125 0.75 0.75	67.0 -35.8 -10.6	37.3 196.4 7.5	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
755	G50B_075_075a	0.0 0.75 0.75	0.75 0.75 0.375	210	0.0 0.75 0.75	65.1 -34.6 -10.1	36.1 196.3	0.0 0.75 0.75	66.8 -37.1 -10.9	38.7 196.3 3.1	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
756	ROOY_100_037a	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	78.5 28.8 24.2	37.6 40.0	1.0 0.625 0.625	71.6 34.1 14.4	37.0 22.9 13.0	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
757	ROOY_087_025a	0.875 0.625 0.625	0.875 0.25 0.75	390	0.875 0.625 0.625	72.2 19.2 16.1	25.1 40.0	0.875 0.625 0.625	68.1 22.7 9.0	24.5 21.7 8.8	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
758	ROOY_075_012a	0.75 0.625 0.625	0.75 0.125 0.687	390	0.75 0.625 0.625	65.9 9.6 8.0	12.5 40.0	0.75 0.625 0.625	65.0 11.2 4.2	12.0 20.4 4.2	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
759	NW_062a	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	59.6 0.0 0.0	0.0 0.0	0.625 0.625 0.625	62.4 0.0 0.0	0.0 32.5 2.7	360 1.0 1.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
760	G50B_062_012a	0.5 0.625 0.625	0.625 0.125 0.562	210	0.5 0.625 0.625	58.5 -5.7 -1.6	6.0 196.3	0.5 0.625 0.625	60.1 -10.5 -3.5	11.0 198.4 5.3	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
761	G50B_062_025a	0.375 0.625 0.625	0.625 0.25 0.5	210	0.375 0.625 0.625	57.4 -11.5 -3.3	12.0 196.3	0.375 0.625 0.625	58.5 -19.5 -6.1	20.5 197.5 8.5	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
762	G50B_062_037a	0.25 0.625 0.625	0.625 0.375 0.437	210	0.25 0.625 0.625	56.4 -17.3 -5.0	18.0 196.3	0.25 0.625 0.625	57.3 -26.4 -8.0	27.6 196.9 9.6	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
763	G50B_062_050a	0.125 0.625 0.625	0.625 0.5 0.375	210	0.125 0.625 0.625	55.3 -23.0 -6.7	24.0 196.3	0.125 0.625 0.625	56.6 -30.7 -9.1	32.0 196.5 8.0	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
764	G50B_062_062a	0.0 0.625 0.625	0.625 0.625 0.312	210	0.0 0.625 0.625	54.2 -28.8 -8.4	30.0 196.3	0.0 0.625 0.625	56.3 -32.4 -9.5	33.8 196.3 4.3	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
765	ROOY_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	72.9 38.4 32.2	50.2 40.0	1.0 0.5 0.5	64.7 46.4 21.9	51.3 25.2 15.4	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
766	ROOY_087_037a	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.5	66.6 28.8 24.2	37.6 40.0	0.875 0.5 0.5	60.6 35.3 15.5	38.6 23.7 12.3	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
767	ROOY_075_025a	0.75 0.5 0.5	0.75 0.25 0.625	390	0.75 0.5 0.5	60.3 19.2 16.1	25.1 40.0	0.75 0.5 0.5	56.8 23.7 9.7	25.6 22.2 8.5	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
768	ROOY_062_012a	0.625 0.5 0.5	0.625 0.125 0.562	390	0.625 0.5 0.5	54.0 9.6 8.0	12.5 40.0	0.625 0.5 0.5	53.4 11.7 4.4	12.6 20.7 4.2	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
769	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	47.7 0.0 0.0	0.0 0.0	0.5 0.5 0.5	50.6 0.0 0.0	0.0 32.5 2.9	360 1.0 1.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
770	G50B_050_012a	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.5	46.6 -5.7 -1.6	6.0 196.3	0.375 0.5 0.5	48.4 -10.7 -3.5	11.3 198.2 5.6	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
771	G50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.25 0.5 0.5	45.5 -11.5 -3.3	12.0 196.3	0.25 0.5 0.5	46.8 -19.5 -6.0	20.4 197.2 8.5	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
772	G50B_050_037a	0.125 0.5 0.5	0.5 0.375 0.312	210	0.125 0.5 0.5	44.5 -17.3 -5.0	18.0 196.3	0.125 0.5 0.5	45.9 -25.2 -7.5	26.3 196.6 8.3	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
773	G50B_050_050a	0.0 0.5 0.5	0.5 0.5 0.25	210	0.0 0.5 0.5	43.4 -23.0 -6.7	24.0 196.3	0.0 0.5 0.5	45.5 -27.6 -8.1	28.7 196.3 5.1	210 0.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
774	ROOY_100_062a	1.0 0.375 0.375	1.0 0.625 0.687	390	1.0 0.375 0.375	67.3 48.0 40.3	62.7 40.0	1.0 0.375 0.375	58.9 58.1 31.4	66.1 28.3 15.8	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
775	ROOY_087_050a	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	61.0 38.4 32.2	50.2 40.0	0.875 0.375 0.375	54.0 47.8 24.1	53.6 26.8 14.2	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
776	ROOY_075_037a	0.75 0.375 0.375	0.75 0.375 0.562	390	0.75 0.375 0.375	54.7 28.8 24.2	37.6 40.0	0.75 0.375 0.375	49.4 36.7 17.1	40.5 25.0 11.8	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
777	ROOY_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	48.4 19.2 16.1	25.1 40.0	0.625 0.375 0.375	45.1 24.9 10.6	27.0 23.1 8.5	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
778	ROOY_050_012a	0.5 0.375 0.375	0.5 0.125 0.437	390	0.5 0.375 0.375	42.0 9.6 8.0	12.5 40.0	0.5 0.375 0.375	41.4 12.4 4.8	13.3 21.2 4.3	389 1.0 0.0 0.0	0.0 0.0 0.0	95.4 0.0 0.0
779	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	35.7 0.0 0.0	0.0 0.0	0.375 0.375 0.375	38.3 0.0 0.0	0.0 32.5 2.5	360 1.0 1.0 1.0	1.0 1.0 1.0	86.8 -46.1 -13.5
780	G50B_037_012a	0.25 0.375 0.375	0.3										

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 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	rgb*Md	LabCh*Md				
810	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	1.0 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	0.0 325.2	0.0 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0		
811	BOOR_100_012a	0.875 0.875 1.0	1.0 0.125 0.937	270	0.875 0.875 1.0	87.2 9.5	-12.9 16.0 306.2	0.875 0.875 1.0	85.5 5.8	-14.8 15.9	291.5 4.4	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
812	BOOR_100_025a	0.75 0.75 1.0	1.0 0.25 0.875	270	0.75 0.75 1.0	79.1 19.0	-25.8 32.1 306.2	0.75 0.75 1.0	75.6 12.8	-30.0 32.7	293.1 8.2	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
813	BOOR_100_037a	0.625 0.625 1.0	1.0 0.375 0.812	270	0.625 0.625 1.0	71.0 28.5	-38.8 48.1 306.2	0.625 0.625 1.0	65.7 21.4	-45.6 50.4	295.1 11.1	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
814	BOOR_100_050a	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	62.8 38.0	-51.7 64.2 306.2	0.5 0.5 1.0	56.0 31.9	-61.1 69.0	297.5 13.0	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
815	BOOR_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	54.7 47.5	-64.7 80.3 306.2	0.375 0.375 1.0	46.8 44.9	-76.1 88.2	300.3 14.2	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
816	BOOR_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	46.6 57.0	-77.6 96.3 306.2	0.25 0.25 1.0	38.8 58.2	-89.4 106.7	303.0 14.1	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
817	BOOR_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	38.5 66.5	-90.6 112.4 306.2	0.125 0.125 1.0	33.0 69.9	-99.0 121.3	305.2 10.6	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
818	BOOR_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
819	Y00G_100_012a	1.0 1.0 0.875	1.0 1.0 0.937	90	1.0 1.0 0.875	95.0 -2.5	11.3 11.6 102.8	1.0 1.0 0.875	94.7 -5.0	14.6 15.4	108.9 4.1	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
820	NW_087a	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	83.4 0.0	0.0 0.0 0.0	0.875 0.875 0.875	84.7 0.0	0.0 0.0	0.0 325.2	1.2 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0	0.0
821	BOOR_087_012a	0.75 0.75 0.875	0.875 0.125 0.812	270	0.75 0.75 0.875	75.3 9.5	-12.9 16.0 306.2	0.75 0.75 0.875	74.6 6.0	-15.2 16.4	291.7 4.1	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
822	BOOR_087_025a	0.625 0.625 0.875	0.875 0.25 0.75	270	0.625 0.625 0.875	67.2 19.0	-25.8 32.1 306.2	0.625 0.625 0.875	64.4 13.5	-30.9 33.8	293.6 7.9	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
823	BOOR_087_037a	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.5 0.875	59.1 28.5	-38.8 48.1 306.2	0.5 0.5 0.875	54.3 23.0	-46.9 52.2	296.1 10.8	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
824	BOOR_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	50.9 38.0	-51.7 64.2 306.2	0.375 0.375 0.875	44.6 34.8	-62.7 71.7	299.0 13.0	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
825	BOOR_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.8 47.5	-64.7 80.3 306.2	0.25 0.25 0.875	35.8 48.6	-77.1 91.2	302.1 14.3	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
826	BOOR_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	34.7 57.0	-77.6 96.3 306.2	0.125 0.125 0.875	29.1 61.5	-88.2 107.5	304.8 12.7	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
827	BOOR_087_087a	0.0 0.0 0.875	0.875 0.875 0.437	270	0.0 0.0 0.875	26.5 66.5	-90.6 112.4 306.2	0.0 0.0 0.875	25.9 68.7	-93.6 116.1	306.2 3.7	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
828	Y00G_100_025a	1.0 1.0 0.75	1.0 0.25 0.875	90	1.0 1.0 0.75	94.7 -5.1	22.6 23.2 102.8	1.0 1.0 0.75	94.1 -9.3	29.3 30.8	107.7 7.9	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
829	Y00G_087_012a	0.875 0.875 0.75	0.875 0.125 0.812	90	0.875 0.875 0.75	83.1 -2.5	11.3 11.6 102.8	0.875 0.875 0.75	84.0 -5.1	15.0 15.8	108.7 4.5	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
830	NW_075a	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	71.5 0.0	0.0 0.0 0.0	0.75 0.75 0.75	73.7 0.0	0.0 0.0	325.2 2.1	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0	0.0
831	BOOR_075_012a	0.625 0.625 0.75	0.75 0.125 0.687	270	0.625 0.625 0.75	63.4 9.5	-12.9 16.0 306.2	0.625 0.625 0.75	63.3 6.3	-15.7 16.9	292.0 4.1	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
832	BOOR_075_025a	0.5 0.5 0.75	0.75 0.25 0.625	270	0.5 0.5 0.75	55.3 19.0	-25.8 32.1 306.2	0.5 0.5 0.75	52.8 14.4	-31.9 35.1	294.3 7.9	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
833	BOOR_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	47.1 28.5	-38.8 48.1 306.2	0.375 0.375 0.75	42.5 25.1	-48.4 54.5	297.4 11.1	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
834	BOOR_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	39.0 38.0	-51.7 64.2 306.2	0.25 0.25 0.75	32.9 38.5	-64.1 74.8	301.0 13.7	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
835	BOOR_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	30.9 47.5	-64.7 80.3 306.2	0.125 0.125 0.75	25.3 52.5	-76.8 93.0	304.3 14.2	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
836	BOOR_075_075a	0.0 0.0 0.75	0.5 0.75 0.375	270	0.0 0.0 0.75	22.7 57.0	-77.6 96.3 306.2	0.0 0.0 0.75	21.3 61.2	-83.4 103.5	306.2 7.2	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
837	Y00G_100_037a	1.0 1.0 0.625	1.0 0.375 0.812	90	1.0 1.0 0.625	94.3 -7.7	34.0 34.9 102.8	1.0 1.0 0.625	93.6 -13.0	43.8 45.7	106.5 11.1	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
838	Y00G_087_025a	0.875 0.875 0.625	0.875 0.25 0.75	90	0.875 0.875 0.625	82.7 -5.1	22.6 23.2 102.8	0.875 0.875 0.625	83.4 -9.4	30.0 31.5	107.3 8.5	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
839	Y00G_075_012a	0.75 0.75 0.625	0.75 0.125 0.687	90	0.75 0.75 0.625	71.2 -2.5	11.3 11.6 102.8	0.75 0.75 0.625	73.0 -5.1	15.4 16.3	108.5 5.2	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
840	NW_062a	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	59.6 0.0	0.0 0.0 0.0	0.625 0.625 0.625	62.4 0.0	0.0 0.0	325.2 2.7	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0	0.0
841	BOOR_062_012a	0.5 0.5 0.625	0.625 0.125 0.562	270	0.5 0.5 0.625	51.5 9.5	-12.9 16.0 306.2	0.5 0.5 0.625	51.6 6.7	-16.3 17.6	292.4 4.3	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
842	BOOR_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	43.3 19.0	-25.8 32.1 306.2	0.375 0.375 0.625	40.8 15.7	-33.2 36.8	295.4 8.4	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
843	BOOR_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.25 0.25 0.625	35.2 28.5	-38.8 48.1 306.2	0.25 0.25 0.625	30.4 28.1	-50.0 57.4	299.3 12.2	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
844	BOOR_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	27.1 38.0	-51.7 64.2 306.2	0.125 0.125 0.625	21.6 42.8	-64.6 77.5	303.5 14.7	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
845	BOOR_062_062a	0.0 0.0 0.625	0.625 0.625 0.312	270	0.0 0.0 0.625	18.9 47.5	-64.7 80.3 306.2	0.0 0.0 0.625	16.6 53.5	-72.9 90.4	306.2 10.3	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
846	Y00G_100_050a	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	94.0 -10.0	45.3 46.5 102.8	1.0 1.0 0.5	93.2 -15.9	57.8 59.9	105.3 13.6	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
847	Y00G_087_037a	0.875 0.875 0.5	0.875 0.375 0.687	90	0.875 0.875 0.5	82.4 -7.7	34.0 34.9 102.8	0.875 0.875 0.5	82.9 -12.9	44.8 46.6	106.0 11.9	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
848	Y00G_075_025a	0.75 0.75 0.5	0.75 0.25 0.625	90	0.75 0.75 0.5	70.8 -5.1	22.6 23.2 102.8	0.75 0.75 0.5	72.4 -9.4	30.9 32.3	106.9 9.4	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
849	Y00G_062_012a	0.625 0.625 0.5	0.625 0.125 0.562	90	0.625 0.625 0.5	59.2 -2.5	11.3 11.6 102.8	0.625 0.625 0.5	61.6 -5.2	16.0 16.8	108.2 5.8	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
850	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	47.7 0.0	0.0 0.0 0.0	0.5 0.5 0.5	50.6 0.0	0.0 0.0	325.3 2.9	360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0	0.0
851	BOOR_050_012a	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.375 0.5	39.5 9.5	-12.9 16.0 306.2	0.375 0.375 0.5	39.4 7.2	-17.0 18.5	292.9 4.7	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
852	BOOR_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.25 0.25 0.5	31.4 19.0	-25.8 32.1 306.2	0.25 0.25 0.5	28.2 17.7	-34.7 39.0	297.0 9.5	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
853	BOOR_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.125 0.125 0.5	23.3 28.5	-38.8 48.1 306.2	0.125 0.125 0.5	18.1 32.4	-51.3 60.6	302.2 14.0	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
854	BOOR_050_050a	0.0 0.0 0.5	0.5 0.5 0.25	270	0.0 0.0 0.5	15.1 38.0	-51.7 64.2 306.2	0.0 0.0 0.5	11.7 45.5	-61.9 76.8	306.2 13.0	270	0.0 0.0 1.0	30.3 76.0	-103.5 128.5	306.2
855	Y00G_100_062a	1.0 1.0 0.375	1.0 0.625 0.687	90	1.0 1.0 0.375	93.6 -12.9	56.7 58.1 102.8	1.0 1.0 0.375	92.9 -18.0	70.4 72.7	104.3 14.7	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
856	Y00G_087_050a	0.875 0.875 0.375	0.875 0.5 0.625	90	0.875 0.875 0.375	82.1 -10.0	45.3 46.5 102.8	0.875 0.875 0.375	82.6 -15.5	58.6 60.6	104.8 14.2	89	1.0 1.0 0.0	92.6 -20.7	90.7 93.0	102.8
857	Y00G_075_037a	0.75 0.75 0.375	0.7													

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 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	ief_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	rgb*Md	LabCh*Md
891	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	1.0 360	1.0 1.0 1.0	95.4 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	0.0 360	1.0 1.0 1.0	95.4 0.0 0.0
892	B50R_100_012a	1.0 0.875 1.0	1.0 0.125 0.937	330	1.0 0.875 1.0	90.6 11.7 -7.3	13.8 328.2	1.0 0.875 1.0	87.9 15.7 -10.9	19.1 325.1 5.9	330	1.0 0.875 1.0
893	B50R_100_025a	1.0 0.75 1.0	1.0 0.25 0.875	330	1.0 0.75 1.0	85.8 23.5 -14.6	27.7 328.2	1.0 0.75 1.0	80.9 31.7 -21.5	38.4 325.8 11.8	330	1.0 0.75 1.0
894	B50R_100_037a	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	81.1 35.3 -21.9	41.6 328.2	1.0 0.625 1.0	74.3 47.6 -31.5	57.1 326.4 17.0	330	1.0 0.625 1.0
895	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	76.3 47.1 -29.2	55.4 328.2	1.0 0.5 1.0	68.6 62.6 -40.5	74.6 327.0 20.6	330	1.0 0.5 1.0
896	B50R_100_062a	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	71.5 58.9 -36.5	69.3 328.2	1.0 0.375 1.0	63.8 75.6 -48.0	89.6 325.5 21.6	330	1.0 0.375 1.0
897	B50R_100_075a	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	66.8 70.7 -43.8	83.2 328.2	1.0 0.25 1.0	60.2 85.6 -53.6	101.0 327.9 18.9	330	1.0 0.25 1.0
898	B50R_100_087a	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	62.0 82.5 -51.1	97.1 328.2	1.0 0.125 1.0	58.1 91.8 -57.0	108.0 328.1 11.6	330	1.0 0.125 1.0
899	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	57.2 94.3 -58.4	110.9 328.2	1.0 0.0 1.0	57.2 94.3 -58.4	111.0 328.2 0.0	330	1.0 0.0 1.0
900	GO0B_100_012a	0.875 1.0 0.875	1.0 0.125 0.937	150	0.875 1.0 0.875	93.9 -10.3 9.9	14.3 136.0	0.875 1.0 0.875	92.5 -15.4 11.3	19.1 143.6 5.4	149	0.875 1.0 0.875
901	NW_087a	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	83.4 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	84.7 0.0 0.0	0.0 325.2 1.2	360	1.0 1.0 1.0
902	B50R_087_012a	0.875 0.75 0.875	0.875 0.125 0.812	330	0.875 0.75 0.875	78.7 11.7 -7.3	13.8 328.2	0.875 0.75 0.875	77.1 16.1 -11.2	19.6 325.2 6.0	330	1.0 0.75 1.0
903	B50R_087_025a	0.875 0.625 0.875	0.875 0.25 0.75	330	0.875 0.625 0.875	73.9 23.5 -14.6	27.7 328.2	0.875 0.625 0.875	69.9 32.6 -22.0	39.3 325.9 12.3	330	1.0 0.625 1.0
904	B50R_087_037a	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	69.1 35.3 -21.9	41.6 328.2	0.875 0.5 0.875	63.5 48.6 -31.9	58.2 326.7 17.6	330	1.0 0.5 1.0
905	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	64.4 47.1 -29.2	55.4 328.2	0.875 0.375 0.875	58.0 63.2 -40.5	75.0 327.3 20.6	330	1.0 0.375 1.0
906	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	59.6 58.9 -36.5	69.3 328.2	0.875 0.25 0.875	53.8 74.7 -47.0	88.3 327.8 19.8	330	1.0 0.25 1.0
907	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	54.9 70.7 -43.8	83.2 328.2	0.875 0.125 0.875	51.3 82.1 -51.1	96.7 328.1 14.0	330	1.0 0.125 1.0
908	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	50.1 82.5 -51.1	97.1 328.2	0.875 0.0 0.875	50.2 85.3 -52.8	100.3 328.2 3.2	330	1.0 0.0 1.0
909	GO0B_100_025a	0.75 1.0 0.75	1.0 0.25 0.875	150	0.75 1.0 0.75	92.4 -20.6 19.9	28.7 136.0	0.75 1.0 0.75	90.1 -20.5 23.2	38.3 142.7 10.6	149	0.75 1.0 0.75
910	GO0B_087_012a	0.75 0.875 0.75	0.875 0.125 0.812	150	0.75 0.875 0.75	82.0 -10.3 9.9	14.3 136.0	0.75 0.875 0.75	81.8 -15.7 11.6	19.6 143.5 5.6	149	0.75 0.875 0.75
911	NW_075a	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	71.5 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	73.7 0.0 0.0	0.0 325.2 2.1	360	1.0 1.0 1.0
912	B50R_075_012a	0.75 0.625 0.75	0.75 0.125 0.687	330	0.75 0.625 0.75	66.7 11.7 -7.3	13.8 328.2	0.75 0.625 0.75	65.9 16.6 -11.5	20.2 325.3 6.4	330	1.0 0.625 1.0
913	B50R_075_025a	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	62.0 23.5 -14.6	27.7 328.2	0.75 0.5 0.75	58.7 33.5 -22.4	40.4 326.2 13.1	330	1.0 0.5 1.0
914	B50R_075_037a	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	57.2 35.3 -21.9	41.6 328.2	0.75 0.375 0.75	52.4 49.6 -32.2	59.1 327.0 18.2	330	1.0 0.375 1.0
915	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	52.5 47.1 -29.2	55.4 328.2	0.75 0.25 0.75	47.5 63.1 -39.9	74.6 327.6 19.8	330	1.0 0.25 1.0
916	B50R_075_062a	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	47.7 58.9 -36.5	69.3 328.2	0.75 0.125 0.75	44.3 72.1 -44.9	84.9 328.0 15.9	330	1.0 0.125 1.0
917	B50R_075_075a	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	42.9 70.7 -43.8	83.2 328.2	0.75 0.0 0.75	40.3 75.0 76.0	-47.0 89.4 328.2	6.1 330	1.0 0.0 1.0
918	GO0B_100_037a	0.625 1.0 0.625	1.0 0.375 0.812	150	0.625 1.0 0.625	90.9 -31.0 29.9	43.1 136.0	0.625 1.0 0.625	88.0 -44.8 35.5	57.2 141.6 15.1	149	0.625 1.0 0.625
919	GO0B_087_025a	0.625 0.875 0.625	0.875 0.25 0.75	150	0.625 0.875 0.625	80.5 -20.6 19.9	28.7 136.0	0.625 0.875 0.625	79.3 -31.1 23.9	39.3 142.4 11.2	149	0.625 0.875 0.625
920	GO0B_075_012a	0.625 0.75 0.625	0.75 0.125 0.687	150	0.625 0.75 0.625	70.0 -10.3 9.9	14.3 136.0	0.625 0.75 0.625	70.8 -16.2 12.0	20.2 143.4 6.2	149	0.625 0.75 0.625
921	NW_062a	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	59.6 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	62.4 0.0 0.0	0.0 325.2 2.7	360	1.0 1.0 1.0
922	B50R_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	54.8 11.7 -7.3	13.8 328.2	0.625 0.5 0.625	54.4 17.2 -11.8	20.9 325.5 7.1	330	1.0 0.5 1.0
923	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	50.1 23.5 -14.6	27.7 328.2	0.625 0.375 0.625	47.1 34.6 -22.9	41.5 326.5 14.1	330	1.0 0.375 1.0
924	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	45.3 35.3 -21.9	41.6 328.2	0.625 0.25 0.625	41.2 50.2 -32.1	59.6 327.4 18.5	330	1.0 0.25 1.0
925	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	40.5 47.1 -29.2	55.4 328.2	0.625 0.125 0.625	37.3 61.3 -38.3	72.3 327.9 17.2	330	1.0 0.125 1.0
926	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	35.8 58.9 -36.5	69.3 328.2	0.625 0.0 0.625	35.5 66.4 -41.1	78.1 328.2 8.7	330	1.0 0.0 1.0
927	GO0B_100_050a	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	89.5 -41.3 39.9	57.5 136.0	0.5 1.0 0.5	86.3 -57.6 47.9	75.0 140.2 18.4	149	0.5 1.0 0.5
928	GO0B_087_037a	0.5 0.875 0.5	0.875 0.375 0.687	150	0.5 0.875 0.5	79.0 -31.0 29.9	43.1 136.0	0.5 0.875 0.5	77.4 -45.4 36.6	58.3 141.0 15.9	149	0.5 0.875 0.5
929	GO0B_075_025a	0.5 0.75 0.5	0.75 0.25 0.625	150	0.5 0.75 0.5	68.6 -20.6 19.9	28.7 136.0	0.5 0.75 0.5	68.3 -31.8 24.8	40.4 142.0 12.1	149	0.5 0.75 0.5
930	GO0B_062_012a	0.5 0.625 0.5	0.625 0.125 0.562	150	0.5 0.625 0.5	58.1 -10.3 9.9	14.3 136.0	0.5 0.625 0.5	59.4 -16.7 12.5	20.9 143.1 7.0	149	0.5 0.625 0.5
931	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	47.7 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	50.6 0.0 0.0	0.0 325.3 2.9	360	1.0 1.0 1.0
932	B50R_050_012a	0.5 0.375 0.5	0.5 0.125 0.437	330	0.5 0.375 0.5	42.9 11.7 -7.3	13.8 328.2	0.5 0.375 0.5	42.3 18.0 -12.2	21.8 325.7 8.0	330	1.0 0.375 1.0
933	B50R_050_025a	0.5 0.25 0.5	0.5 0.25 0.375	330	0.5 0.25 0.5	38.1 23.5 -14.6	27.7 328.2	0.5 0.25 0.5	35.7 -23.2 42.6	326.9 15.2 330	1.0 0.25 1.0	0.5 0.25 0.5
934	B50R_050_037a	0.5 0.125 0.5	0.5 0.375 0.312	330	0.5 0.125 0.5	33.4 35.3 -21.9	41.6 328.2	0.5 0.125 0.5	30.1 49.6 -31.2	58.6 327.8 17.3	330	1.0 0.125 1.0
935	B50R_050_050a	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	28.6 47.1 -29.2	55.4 328.2	0.5 0.0 0.5	27.8 56.4 -34.9	66.3 328.2 10.9	330	1.0 0.0 1.0
936	GO0B_100_062a	0.375 1.0 0.375	1.0 0.625 0.687	150	0.375 1.0 0.375	88.0 -51.7 49.9	71.9 136.0	0.375 1.0 0.375	85.1 -68.3 59.7	90.7 138.8 19.4	149	0.375 1.0 0.375
937	GO0B_087_050a	0.375 0.875 0.375	0.875 0.5 0.625	150	0.375 0.875 0.375	77.5 -41.3 39.9	57.5 136.0	0.375 0.875 0.375	75.9 -57.6 49.1	75.7 139.5 18.7	149	0.375 0.875 0.375
938	GO0B_075_037a	0.375 0.75 0.375	0.75 0.375 0.562	150	0.375 0.75 0.375	67.1 -31.0 29.9	43.1 136.0	0.375 0.75 0.375	66.5 -45.8 37.9	59.4 140.3 16.7	149	0.375 0.75 0.375
939	GO0B_062_025a	0.375 0.625 0.375	0.625 0.25 0.5	150	0.375 0.625 0.375	56.6 -20.6 19.9	28.7 136.0	0.375 0.625 0.375	57.0 -32.5 25.9	41.6 141.4 13.2	149	0.375 0.625 0.375
940	GO0B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	46.2 -10.3 9.9	14.3 136.0	0.375 0.5 0.375	47.6 -17.3 13.1	21.8 142.8 7.8	149	0.375 0.5 0.375
941	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	35.7 0.0 0.0	0.0 0.0 0.0	0.375 0.375 0.375	38.3 0.0 0.0	0.0 325.3 2.5	360	1.0 1.0 1.0
942	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.25 0.375	31.0 11.7 -7.3	13.8 328.2	0.375 0.25 0.375	29.7 19.0 -12.7	22.9 326.1 9.1	330	1.0 0.25 0.375
943	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.125 0.375	26.2 23.5 -14.6	27.7 328.2	0.375 0.125 0.375	23.1 36.3 -23.1	43.0 327.5 15.6	330	1.0 0.125 0.375
944	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	21.4 35.3 -21.9	41.6 328.2	0.375 0.0 0.375	19.7 46.0 -28.5	54.1 328.2 12.6	330	1.0 0.0 1.0
945	GO0B_100_075a	0.25 1.0 0.25	1.0 0.75 0.625	150	0.25 1.0 0.25	86.5 -60.0 59.9	86.2 136.0	0.25 1.0 0.25	84.2 -76.1 69.7	103.3 137.4 17.3	149	0.25 1.0 0.25
946	GO0B_087_062a	0.25 0.875 0.25	0.875 0.625 0.562	150	0.25 0.875 0.25	76.1 -51.7 49.9	71.9 136.0	0.25 0.875 0.25	74.8 -66.8			

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 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsi_Md	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	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0	
973	NW_012a	0.125 0.125	0.125 0.125	0.125 360	0.125 0.125	0.125 11.9	0.0 0.0 0.0	0.125 0.125	0.125 11.0	0.0 0.0 0.0	325.7 0.8	360 1.0 1.0 1.0	95.4 0.0 0.0
974	NW_025a	0.25 0.25 0.25	0.25 0.25 0.25	0.25 360	0.25 0.25 0.25	23.8	0.0 0.0 0.0	0.25 0.25 0.25	25.2 25.2	0.0 0.0 0.0	325.5 1.4	360 1.0 1.0 1.0	95.4 0.0 0.0
975	NW_037a	0.375 0.375	0.375 0.375	0.375 360	0.375 0.375	35.7	0.0 0.0 0.0	0.375 0.375	38.3 38.3	0.0 0.0 0.0	325.3 2.5	360 1.0 1.0 1.0	95.4 0.0 0.0
976	NW_050a	0.5 0.5 0.5	0.5 0.5 0.5	0.5 360	0.5 0.5 0.5	47.7	0.0 0.0 0.0	0.5 0.5 0.5	50.6 50.6	0.0 0.0 0.0	325.3 2.9	360 1.0 1.0 1.0	95.4 0.0 0.0
977	NW_062a	0.625 0.625	0.625 0.625	0.625 360	0.625 0.625	59.6	0.0 0.0 0.0	0.625 0.625	62.4 62.4	0.0 0.0 0.0	325.2 2.7	360 1.0 1.0 1.0	95.4 0.0 0.0
978	NW_075a	0.75 0.75 0.75	0.75 0.75 0.75	0.75 360	0.75 0.75 0.75	71.5	0.0 0.0 0.0	0.75 0.75 0.75	73.7 73.7	0.0 0.0 0.0	325.2 2.1	360 1.0 1.0 1.0	95.4 0.0 0.0
979	NW_087a	0.875 0.875	0.875 0.875	0.875 360	0.875 0.875	83.4	0.0 0.0 0.0	0.875 0.875	84.7 84.7	0.0 0.0 0.0	325.2 1.2	360 1.0 1.0 1.0	95.4 0.0 0.0
980	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 360	1.0 1.0 1.0	95.4	0.0 0.0 0.0	1.0 1.0 1.0	95.4 95.4	0.0 0.0 0.0	325.2 0.0	360 1.0 1.0 1.0	95.4 0.0 0.0
981	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	360 1.0 1.0 1.0	95.4 0.0 0.0
982	NW_012a	0.125 0.125	0.125 0.125	0.125 360	0.125 0.125	11.9	0.0 0.0 0.0	0.125 0.125	11.0 11.0	0.0 0.0 0.0	325.7 0.8	360 1.0 1.0 1.0	95.4 0.0 0.0
983	NW_025a	0.25 0.25 0.25	0.25 0.25 0.25	0.25 360	0.25 0.25 0.25	23.8	0.0 0.0 0.0	0.25 0.25 0.25	25.2 25.2	0.0 0.0 0.0	325.5 1.4	360 1.0 1.0 1.0	95.4 0.0 0.0
984	NW_037a	0.375 0.375	0.375 0.375	0.375 360	0.375 0.375	35.7	0.0 0.0 0.0	0.375 0.375	38.3 38.3	0.0 0.0 0.0	325.3 2.5	360 1.0 1.0 1.0	95.4 0.0 0.0
985	NW_050a	0.5 0.5 0.5	0.5 0.5 0.5	0.5 360	0.5 0.5 0.5	47.7	0.0 0.0 0.0	0.5 0.5 0.5	50.6 50.6	0.0 0.0 0.0	325.3 2.9	360 1.0 1.0 1.0	95.4 0.0 0.0
986	NW_062a	0.625 0.625	0.625 0.625	0.625 360	0.625 0.625	59.6	0.0 0.0 0.0	0.625 0.625	62.4 62.4	0.0 0.0 0.0	325.2 2.7	360 1.0 1.0 1.0	95.4 0.0 0.0
987	NW_075a	0.75 0.75 0.75	0.75 0.75 0.75	0.75 360	0.75 0.75 0.75	71.5	0.0 0.0 0.0	0.75 0.75 0.75	73.7 73.7	0.0 0.0 0.0	325.2 2.1	360 1.0 1.0 1.0	95.4 0.0 0.0
988	NW_087a	0.875 0.875	0.875 0.875	0.875 360	0.875 0.875	83.4	0.0 0.0 0.0	0.875 0.875	84.7 84.7	0.0 0.0 0.0	325.2 1.2	360 1.0 1.0 1.0	95.4 0.0 0.0
989	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 360	1.0 1.0 1.0	95.4	0.0 0.0 0.0	1.0 1.0 1.0	95.4 95.4	0.0 0.0 0.0	325.2 0.0	360 1.0 1.0 1.0	95.4 0.0 0.0
990	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	360 1.0 1.0 1.0	95.4 0.0 0.0
991	NW_012a	0.125 0.125	0.125 0.125	0.125 360	0.125 0.125	11.9	0.0 0.0 0.0	0.125 0.125	11.0 11.0	0.0 0.0 0.0	325.7 0.8	360 1.0 1.0 1.0	95.4 0.0 0.0
992	NW_025a	0.25 0.25 0.25	0.25 0.25 0.25	0.25 360	0.25 0.25 0.25	23.8	0.0 0.0 0.0	0.25 0.25 0.25	25.2 25.2	0.0 0.0 0.0	325.5 1.4	360 1.0 1.0 1.0	95.4 0.0 0.0
993	NW_037a	0.375 0.375	0.375 0.375	0.375 360	0.375 0.375	35.7	0.0 0.0 0.0	0.375 0.375	38.3 38.3	0.0 0.0 0.0	325.3 2.5	360 1.0 1.0 1.0	95.4 0.0 0.0
994	NW_050a	0.5 0.5 0.5	0.5 0.5 0.5	0.5 360	0.5 0.5 0.5	47.7	0.0 0.0 0.0	0.5 0.5 0.5	50.6 50.6	0.0 0.0 0.0	325.3 2.9	360 1.0 1.0 1.0	95.4 0.0 0.0
995	NW_062a	0.625 0.625	0.625 0.625	0.625 360	0.625 0.625	59.6	0.0 0.0 0.0	0.625 0.625	62.4 62.4	0.0 0.0 0.0	325.2 2.7	360 1.0 1.0 1.0	95.4 0.0 0.0
996	NW_075a	0.75 0.75 0.75	0.75 0.75 0.75	0.75 360	0.75 0.75 0.75	71.5	0.0 0.0 0.0	0.75 0.75 0.75	73.7 73.7	0.0 0.0 0.0	325.2 2.1	360 1.0 1.0 1.0	95.4 0.0 0.0
997	NW_087a	0.875 0.875	0.875 0.875	0.875 360	0.875 0.875	83.4	0.0 0.0 0.0	0.875 0.875	84.7 84.7	0.0 0.0 0.0	325.2 1.2	360 1.0 1.0 1.0	95.4 0.0 0.0
998	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 360	1.0 1.0 1.0	95.4	0.0 0.0 0.0	1.0 1.0 1.0	95.4 95.4	0.0 0.0 0.0	325.2 0.0	360 1.0 1.0 1.0	95.4 0.0 0.0
999	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	360 1.0 1.0 1.0	95.4 0.0 0.0
1000	NW_012a	0.125 0.125	0.125 0.125	0.125 360	0.125 0.125	11.9	0.0 0.0 0.0	0.125 0.125	11.0 11.0	0.0 0.0 0.0	325.7 0.8	360 1.0 1.0 1.0	95.4 0.0 0.0
1001	NW_025a	0.25 0.25 0.25	0.25 0.25 0.25	0.25 360	0.25 0.25 0.25	23.8	0.0 0.0 0.0	0.25 0.25 0.25	25.2 25.2	0.0 0.0 0.0	325.5 1.4	360 1.0 1.0 1.0	95.4 0.0 0.0
1002	NW_037a	0.375 0.375	0.375 0.375	0.375 360	0.375 0.375	35.7	0.0 0.0 0.0	0.375 0.375	38.3 38.3	0.0 0.0 0.0	325.3 2.5	360 1.0 1.0 1.0	95.4 0.0 0.0
1003	NW_050a	0.5 0.5 0.5	0.5 0.5 0.5	0.5 360	0.5 0.5 0.5	47.7	0.0 0.0 0.0	0.5 0.5 0.5	50.6 50.6	0.0 0.0 0.0	325.3 2.9	360 1.0 1.0 1.0	95.4 0.0 0.0
1004	NW_062a	0.625 0.625	0.625 0.625	0.625 360	0.625 0.625	59.6	0.0 0.0 0.0	0.625 0.625	62.4 62.4	0.0 0.0 0.0	325.2 2.7	360 1.0 1.0 1.0	95.4 0.0 0.0
1005	NW_075a	0.75 0.75 0.75	0.75 0.75 0.75	0.75 360	0.75 0.75 0.75	71.5	0.0 0.0 0.0	0.75 0.75 0.75	73.7 73.7	0.0 0.0 0.0	325.2 2.1	360 1.0 1.0 1.0	95.4 0.0 0.0
1006	NW_087a	0.875 0.875	0.875 0.875	0.875 360	0.875 0.875	83.4	0.0 0.0 0.0	0.875 0.875	84.7 84.7	0.0 0.0 0.0	325.2 1.2	360 1.0 1.0 1.0	95.4 0.0 0.0
1007	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 360	1.0 1.0 1.0	95.4	0.0 0.0 0.0	1.0 1.0 1.0	95.4 95.4	0.0 0.0 0.0	325.2 0.0	360 1.0 1.0 1.0	95.4 0.0 0.0
1008	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	360 1.0 1.0 1.0	95.4 0.0 0.0
1009	NW_006a	0.066 0.066	0.066 0.066	0.066 360	0.066 0.066	6.2	0.0 0.0 0.0	0.066 0.066	6.4 6.4	0.0 0.0 0.0	326.3 1.8	360 1.0 1.0 1.0	95.4 0.0 0.0
1010	NW_013a	0.133 0.133	0.133 0.133	0.133 360	0.133 0.133	12.6	0.0 0.0 0.0	0.133 0.133	12.0 12.0	0.0 0.0 0.0	325.6 0.6	360 1.0 1.0 1.0	95.4 0.0 0.0
1011	NW_020a	0.2 0.2 0.2	0.2 0.2 0.2	0.2 360	0.2 0.2 0.2	19.0	0.0 0.0 0.0	0.2 0.2 0.2	19.7 19.7	0.0 0.0 0.0	325.5 0.6	360 1.0 1.0 1.0	95.4 0.0 0.0
1012	NW_026a	0.266 0.266	0.266 0.266	0.266 360	0.266 0.266	25.3	0.0 0.0 0.0	0.266 0.266	27.0 27.0	0.0 0.0 0.0	325.4 1.6	360 1.0 1.0 1.0	95.4 0.0 0.0
1013	NW_033a	0.333 0.333	0.333 0.333	0.333 360	0.333 0.333	31.7	0.0 0.0 0.0	0.333 0.333	34.0 34.0	0.0 0.0 0.0	325.3 2.2	360 1.0 1.0 1.0	95.4 0.0 0.0
1014	NW_040a	0.4 0.4 0.4	0.4 0.4 0.4	0.4 360	0.4 0.4 0.4	38.1	0.0 0.0 0.0	0.4 0.4 0.4	40.8 40.8	0.0 0.0 0.0	325.3 2.6	360 1.0 1.0 1.0	95.4 0.0 0.0
1015	NW_046a	0.466 0.466	0.466 0.466	0.466 360	0.466 0.466	44.4	0.0 0.0 0.0	0.466 0.466	47.3 47.3	0.0 0.0 0.0	325.4 2.8	360 1.0 1.0 1.0	95.4 0.0 0.0
1016	NW_053a	0.533 0.533	0.533 0.533	0.533 360	0.533 0.533	50.8	0.0 0.0 0.0	0.533 0.533	53.7 53.7	0.0 0.0 0.0	325.3 2.9	360 1.0 1.0 1.0	95.4 0.0 0.0
1017	NW_060a	0.6 0.6 0.6	0.6 0.6 0.6	0.6 360	0.6 0.6 0.6	57.2	0.0 0.0 0.0	0.6 0.6 0.6	60.0 60.0	0.0 0.0 0.0	325.3 2.8	360 1.0 1.0 1.0	95.4 0.0 0.0
1018	NW_066a	0.666 0.666	0.666 0.666	0.666 360	0.666 0.666	63.5	0.0 0.0 0.0	0.666 0.666	66.1 66.1	0.0 0.0 0.0	325.2 2.6	360 1.0 1.0 1.0	95.4 0.0 0.0
1019	NW_073a	0.734 0.734	0.734 0.734	0.734 360	0.734 0.734	70.0	0.0 0.0 0.0	0.734 0.734	72.3 72.3	0.0 0.0 0.0	325.2 2.2	360 1.0 1.0 1.0	95.4 0.0 0.0
1020	NW_080a	0.8 0.8 0.8	0.8 0.8 0.8	0.8 360	0.8 0.8 0.8	76.3	0.0 0.0 0.0	0.8 0.8 0.8	78.1 78.1	0.0 0.0 0.0	325.2 1.8	360 1.0 1.0 1.0	95.4 0.0 0.0
1021	NW_086a	0.866 0.866	0.866 0.866	0.866 360	0.866 0.866	82.6	0.0 0.0 0.0	0.866 0.866	83.9 83.9	0.0 0.0 0.0	325.2 1.3	360 1.0 1.0 1.0	95.4 0.0 0.0
1022	NW_093a	0.933 0.933	0.933 0.933	0.933 360	0.933 0.933	89.0	0.0 0.0 0.0	0.933 0.933	89.7 89.7	0.0 0.0 0.0	325.2 0.6	360 1.0 1.0 1.0	95.4 0.0 0.0
1023	NW_100a	1.0 1.0 1.0	1.0 1.0 1.0	1.0 360	1.0 1.0 1.0	95.4	0.0 0.0 0.0	1.0 1.0 1.0	95.4 95.4	0.0 0.0 0.0	325.2 0.0	360 1.0 1.0 1.0	95.4 0.0 0.0
1024	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	360 1.0 1.0 1.0	95.4 0.0 0.0
1025	NW_006a	0.066 0.066	0.066 0.066	0.066 360	0.066 0.066	6.2	0.0 0.0 0.0	0.066 0.066	6.4 6.4	0.0 0.0 0.0	326.3 1.8	360 1.0 1.0 1.0	95.4 0.0 0.0
1026	NW_013a	0.133 0.133	0.133 0.133	0.133 360	0.133 0.133	12.6	0.0 0.0 0.0	0.133 0.133	12.0 12.0	0.0 0.0 0.0	325.6 0.6	360 1.0 1.0 1.0	95.4 0.0 0.0
1027	NW_020a	0.2 0.2 0.2	0.2 0.2 0.2	0.2 360	0.2 0.2 0.2	19.0	0.0 0.0 0.0	0.2 0.2 0.2	19.7 19.7	0.0 0.0 0.0	325.5 0.6	360 1.0 1.0 1.0	95.4 0.0 0.0
1028	NW_026a	0.266 0.266	0.266 0.266	0.266 360	0.266 0.266	25.3	0.0 0.0 0.0	0.266 0.266	27.0 27.0	0.0 0.0 0.0			

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 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Ma	LabCh*Ma
1053	NW_086a	0.866 0.866 0.866	0.866 0.0	0.866 360	0.866 0.866 0.866	82.6 0.0 0.0	0.866 0.866 0.866	83.9 0.0 0.0	325.2 1.3	360	1.0 1.0 1.0	95.4 0.0 0.0
1054	NW_093a	0.933 0.933 0.933	0.933 0.0	0.933 360	0.933 0.933 0.933	89.0 0.0 0.0	0.933 0.933 0.933	89.7 0.0 0.0	325.2 0.6	360	1.0 1.0 1.0	95.4 0.0 0.0
1055	NW_100a	1.0 1.0 1.0	1.0 0.0	1.0 360	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	325.2 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0
1056	NW_000a	0.0 0.0 0.0	0.0 0.0	0.0 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0
1057	NW_006a	0.066 0.066 0.066	0.066 0.0	0.066 360	0.066 0.066 0.066	6.2 0.0 0.0	0.066 0.066 0.066	4.4 0.0 0.0	326.3 1.8	360	1.0 1.0 1.0	95.4 0.0 0.0
1058	NW_013a	0.133 0.133 0.133	0.133 0.0	0.133 360	0.133 0.133 0.133	12.6 0.0 0.0	0.133 0.133 0.133	12.0 0.0 0.0	325.6 0.6	360	1.0 1.0 1.0	95.4 0.0 0.0
1059	NW_020a	0.2 0.2 0.2	0.2 0.0	0.2 360	0.2 0.2 0.2	19.0 0.0 0.0	0.2 0.2 0.2	19.7 0.0 0.0	325.5 0.6	360	1.0 1.0 1.0	95.4 0.0 0.0
1060	NW_026a	0.266 0.266 0.266	0.266 0.0	0.266 360	0.266 0.266 0.266	25.3 0.0 0.0	0.266 0.266 0.266	27.0 0.0 0.0	325.4 1.6	360	1.0 1.0 1.0	95.4 0.0 0.0
1061	NW_033a	0.333 0.333 0.333	0.333 0.0	0.333 360	0.333 0.333 0.333	31.7 0.0 0.0	0.333 0.333 0.333	34.0 0.0 0.0	325.3 2.2	360	1.0 1.0 1.0	95.4 0.0 0.0
1062	NW_040a	0.4 0.4 0.4	0.4 0.0	0.4 360	0.4 0.4 0.4	38.1 0.0 0.0	0.4 0.4 0.4	40.8 0.0 0.0	325.3 2.6	360	1.0 1.0 1.0	95.4 0.0 0.0
1063	NW_046a	0.466 0.466 0.466	0.466 0.0	0.466 360	0.466 0.466 0.466	44.4 0.0 0.0	0.466 0.466 0.466	47.3 0.0 0.0	325.4 2.8	360	1.0 1.0 1.0	95.4 0.0 0.0
1064	NW_053a	0.533 0.533 0.533	0.533 0.0	0.533 360	0.533 0.533 0.533	50.8 0.0 0.0	0.533 0.533 0.533	53.7 0.0 0.0	325.3 2.9	360	1.0 1.0 1.0	95.4 0.0 0.0
1065	NW_060a	0.6 0.6 0.6	0.6 0.0	0.6 360	0.6 0.6 0.6	57.2 0.0 0.0	0.6 0.6 0.6	60.0 0.0 0.0	325.3 2.8	360	1.0 1.0 1.0	95.4 0.0 0.0
1066	NW_066a	0.666 0.666 0.666	0.666 0.0	0.666 360	0.666 0.666 0.666	63.5 0.0 0.0	0.666 0.666 0.666	66.1 0.0 0.0	325.2 2.6	360	1.0 1.0 1.0	95.4 0.0 0.0
1067	NW_073a	0.734 0.734 0.734	0.734 0.0	0.734 360	0.734 0.734 0.734	70.0 0.0 0.0	0.734 0.734 0.734	72.3 0.0 0.0	325.2 2.2	360	1.0 1.0 1.0	95.4 0.0 0.0
1068	NW_080a	0.8 0.8 0.8	0.8 0.0	0.8 360	0.8 0.8 0.8	76.3 0.0 0.0	0.8 0.8 0.8	78.1 0.0 0.0	325.2 1.8	360	1.0 1.0 1.0	95.4 0.0 0.0
1069	NW_086a	0.866 0.866 0.866	0.866 0.0	0.866 360	0.866 0.866 0.866	82.6 0.0 0.0	0.866 0.866 0.866	83.9 0.0 0.0	325.2 1.3	360	1.0 1.0 1.0	95.4 0.0 0.0
1070	NW_093a	0.933 0.933 0.933	0.933 0.0	0.933 360	0.933 0.933 0.933	89.0 0.0 0.0	0.933 0.933 0.933	89.7 0.0 0.0	325.2 0.6	360	1.0 1.0 1.0	95.4 0.0 0.0
1071	NW_100a	1.0 1.0 1.0	1.0 0.0	1.0 360	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	325.2 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0
1072	NW_000a	0.0 0.0 0.0	0.0 0.0	0.0 360	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0
1073	NW_100a	1.0 1.0 1.0	1.0 0.0	1.0 360	1.0 1.0 1.0	95.4 0.0 0.0	1.0 1.0 1.0	95.4 0.0 0.0	325.2 0.0	360	1.0 1.0 1.0	95.4 0.0 0.0
1074	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0	1.0 0.0 0.0	50.4 76.9 64.5 100.4	39.9 0.0	389	1.0 0.0 0.0	50.4 76.9 64.5 100.4 40.0
1075	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	86.8 -46.1 -13.5 48.1 196.3	0.0 1.0 1.0	86.8 -46.1 -13.5 48.1	196.3 0.0	210	0.0 1.0 1.0	86.8 -46.1 -13.5 48.1 196.3
1076	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	92.6 -20.7 90.7 93.0 102.8	1.0 1.0 0.0	92.6 -20.6 90.7 93.0	102.8 0.0	89	0.0 1.0 0.0	92.6 -20.7 90.7 93.0 102.8
1077	B00R_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2	0.0 0.0 1.0	30.3 76.0 -103.5 128.5	306.2 0.0	270	0.0 0.0 1.0	30.3 76.0 -103.5 128.5 306.2
1078	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	83.6 -82.7 79.8 115.0 136.0	0.0 1.0 0.0	83.6 -82.7 79.8 115.0	136.0 0.0	149	0.0 1.0 0.0	83.6 -82.7 79.8 115.0 136.0
1079	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	57.2 94.3 -58.4 110.9 328.2	1.0 0.0 1.0	57.2 94.3 -58.4 111.0	328.2 0.0	330	1.0 0.0 1.0	57.2 94.3 -58.4 110.9 328.2

delta E* = 1.0

TUB enregistrement: 20130201-PF21/PF21L0NP.PDF /.PS
 application pour la mesure de sortie sur écran, aucune séparation
 TUB matériel: code=rh4ta

