

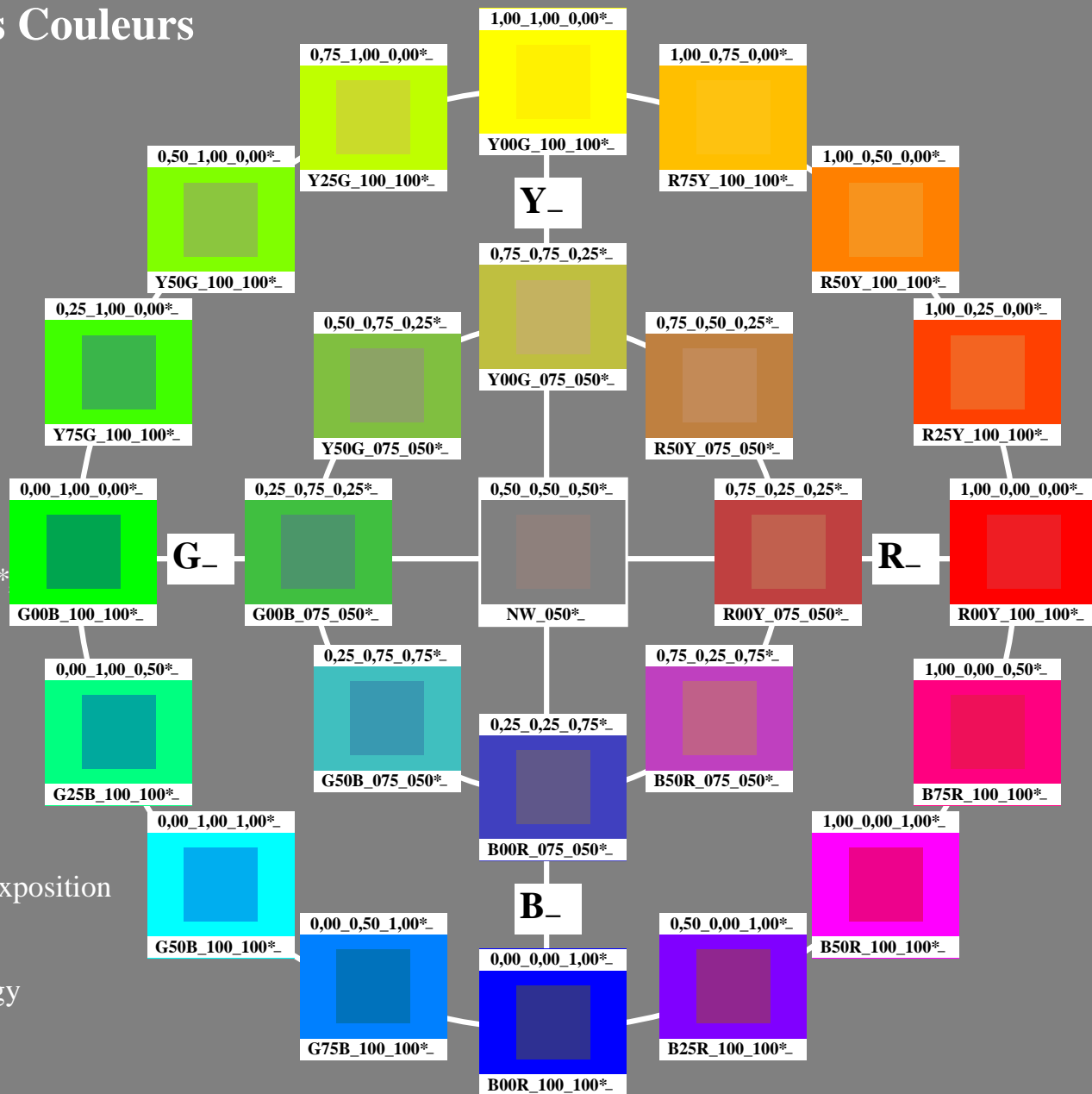
Couleur et la Vision des Couleurs

Couleurs Élémentaires en Informatique

Auteur: Prof. Dr. Klaus Richter

25 couleurs destinée pour D65
cercle chromatique à 16 et 8 étapes
écran standard *sRGB*
rgb data: *rgb**_e (en haut)
couleurs élémentaires *H**, brillance *I**
chromie *C**: *HIC**_e (en bas)

Brochure publiée dans le cadre de l'exposition
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Section Lighting Technology
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voir: <http://www.li.tu-berlin.de>
et <http://130.149.60.45/~farbmetrik>



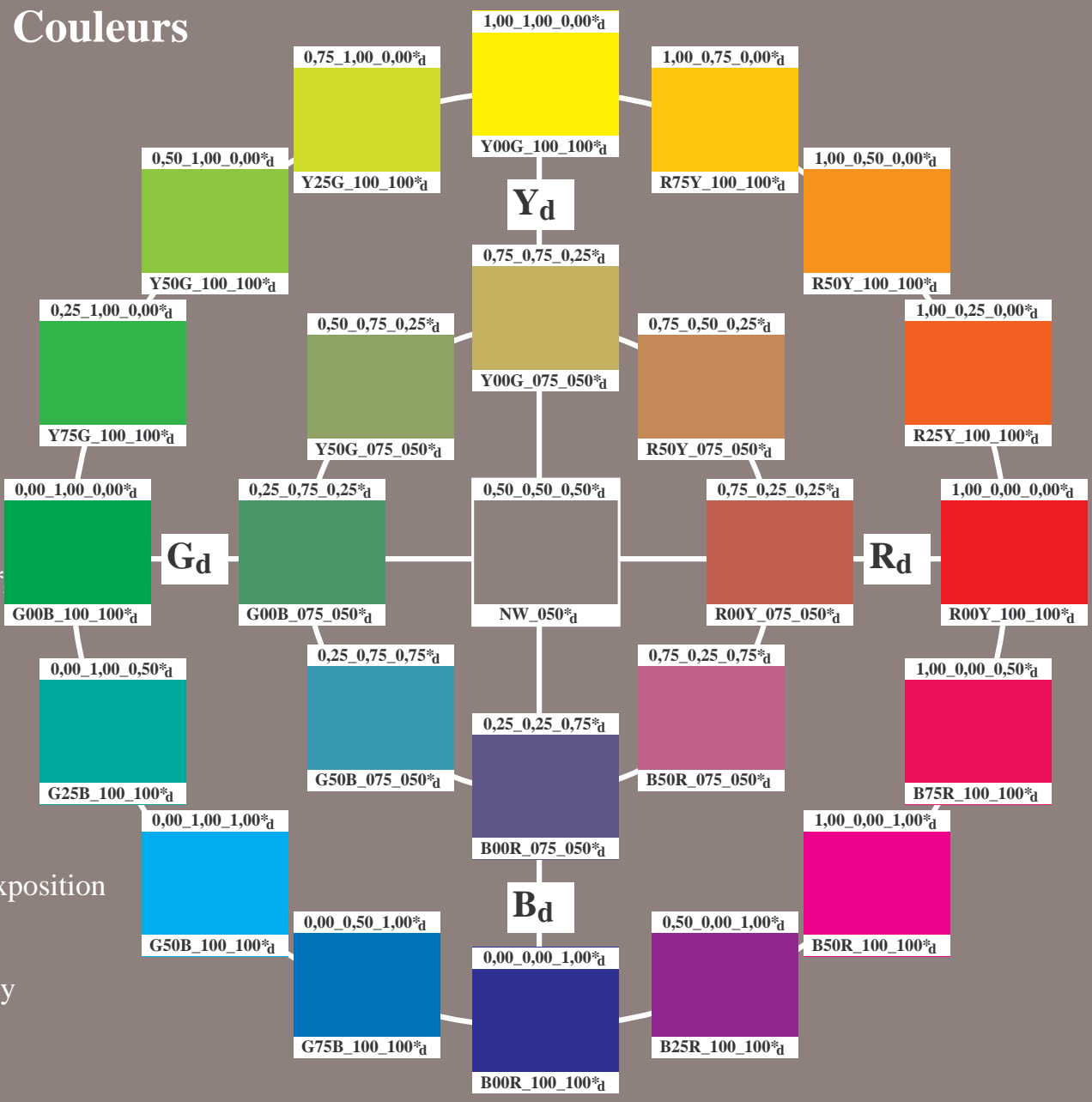
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voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF77/PF77.HTM>
informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

TUB enregistrement: 20130201 -PF77/PF77L0NA.TXT /.PS TUB matériel: code=rh4ta
application pour la mesure des sorties sur offset, séparation cmy0 (CMY0)

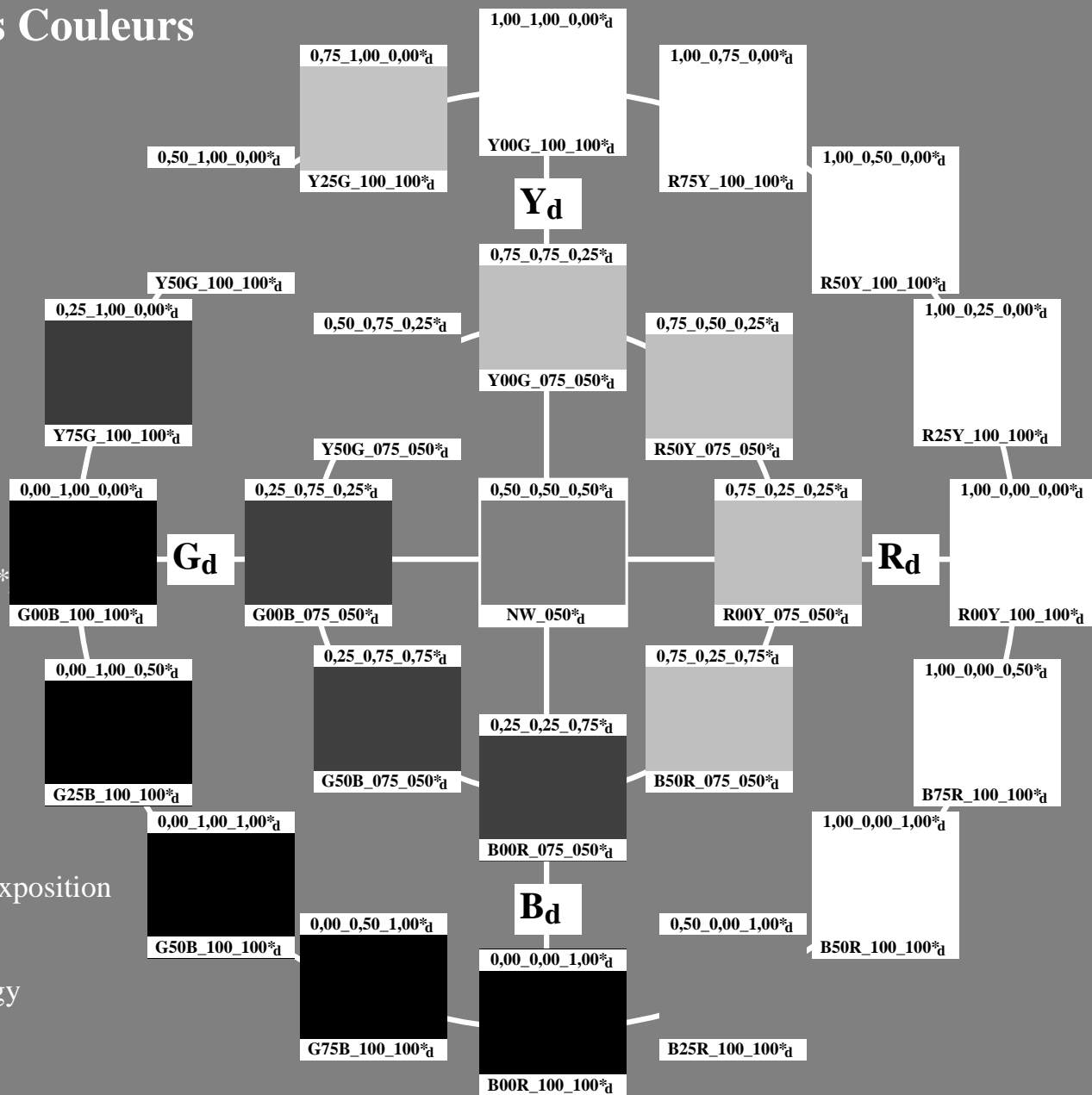
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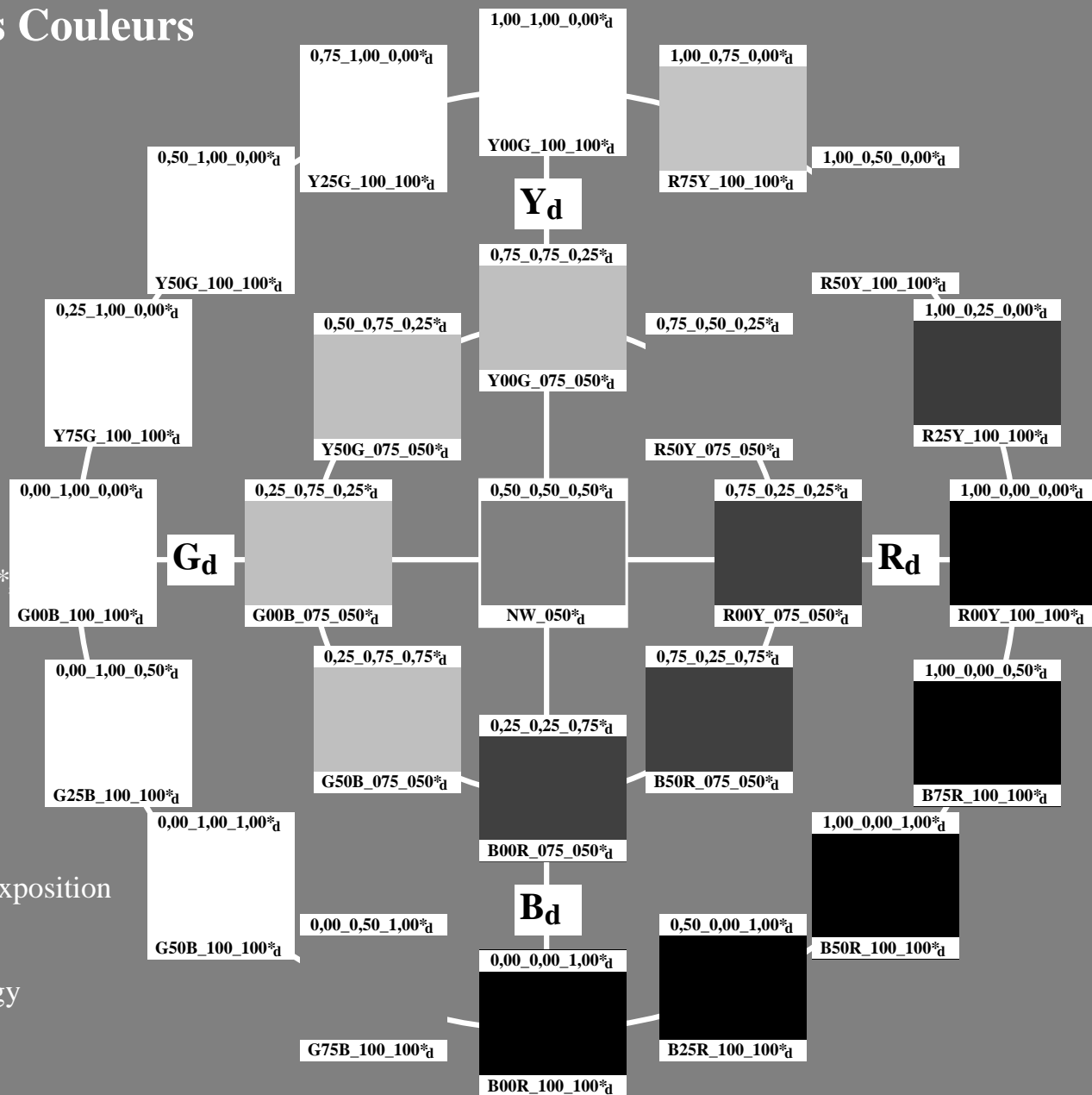
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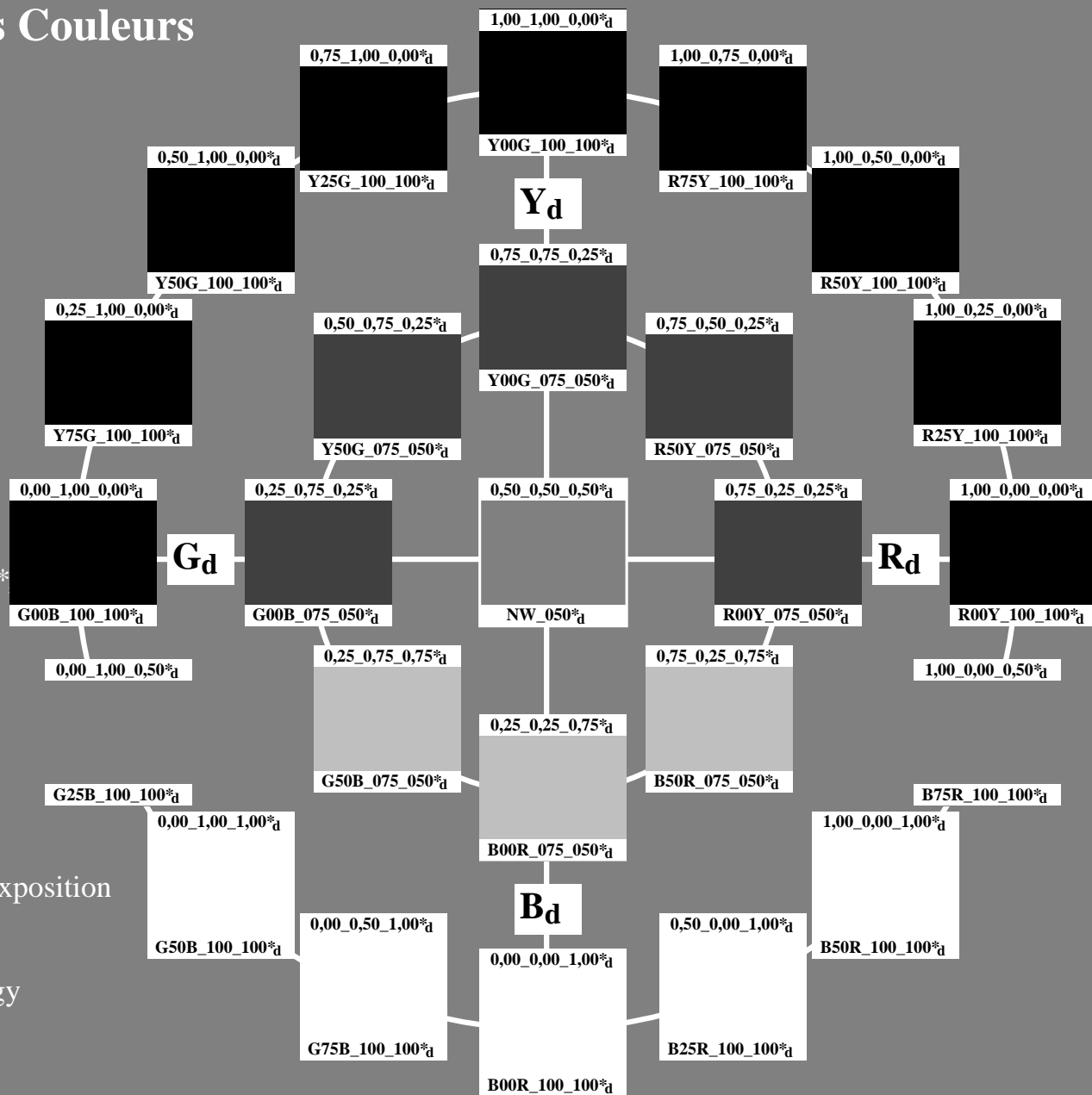
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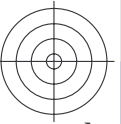
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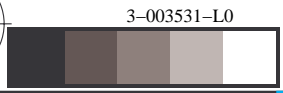
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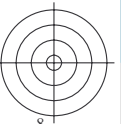
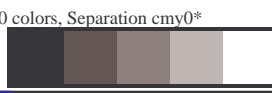
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3-003531-L0 PF770-70
graphique TUB-PF77; cercle de teinte; 16 et 8 étapes
25 couleur de norme pour D65, 3D=0, de=0, cmy0

entrée : *rgb/cmyk* -> *rgb_d*
sortie : transférer à *cmy0_d*



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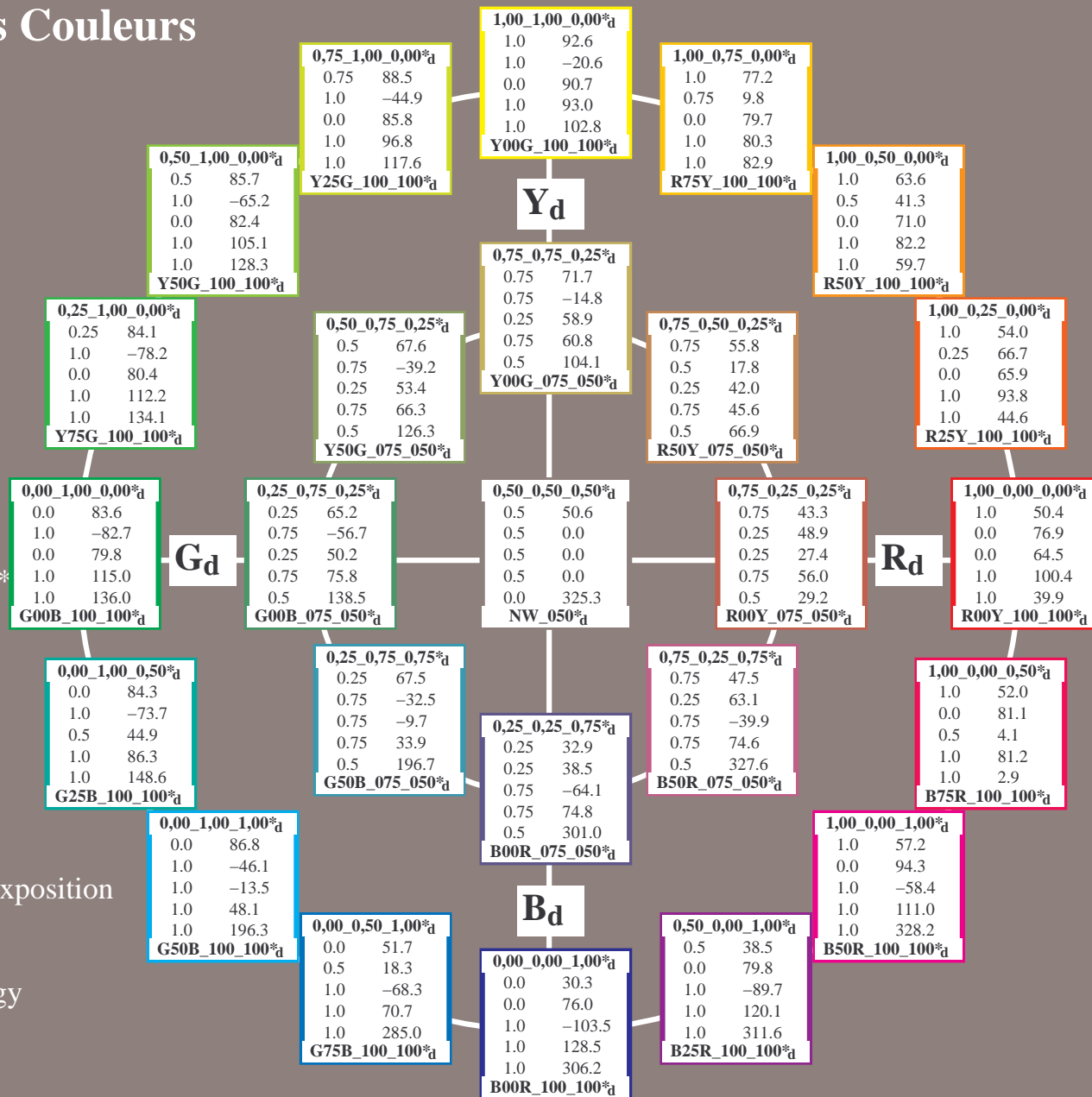
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 chromie *C**: *HIC**_e (en bas)
 code de couleur:
*rgbic*_d; *LabCh**_d

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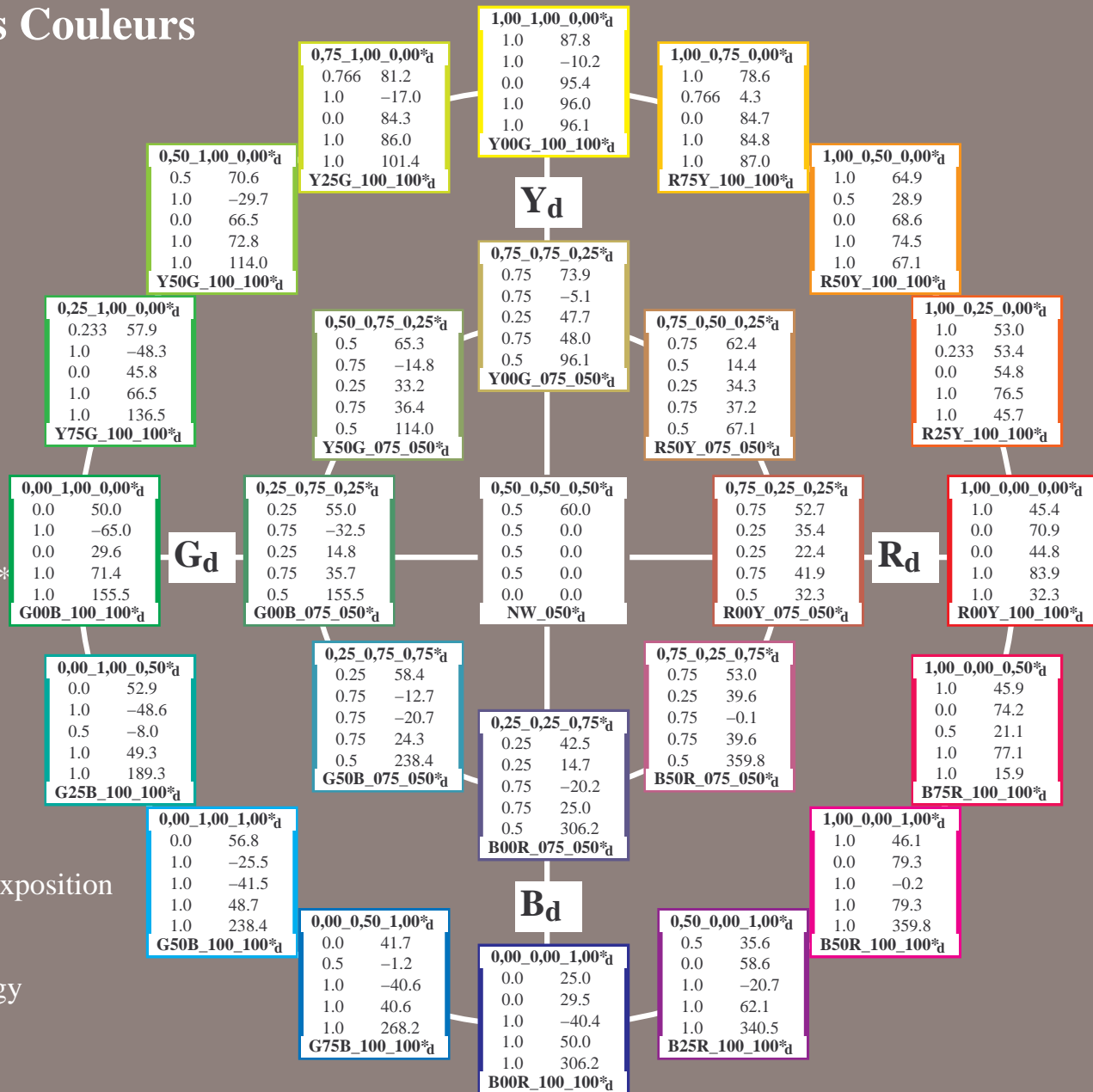
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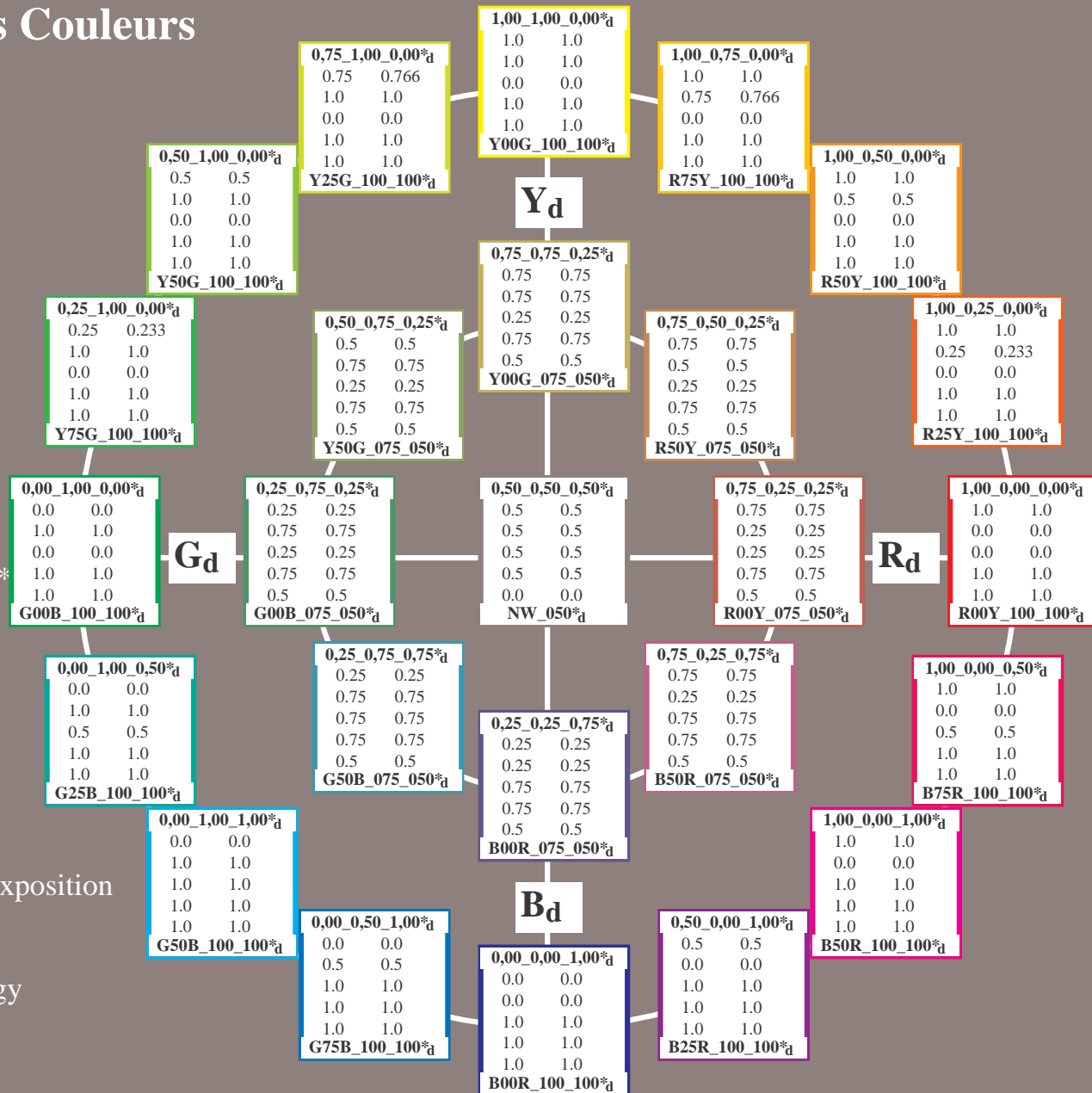
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 code de couleur:
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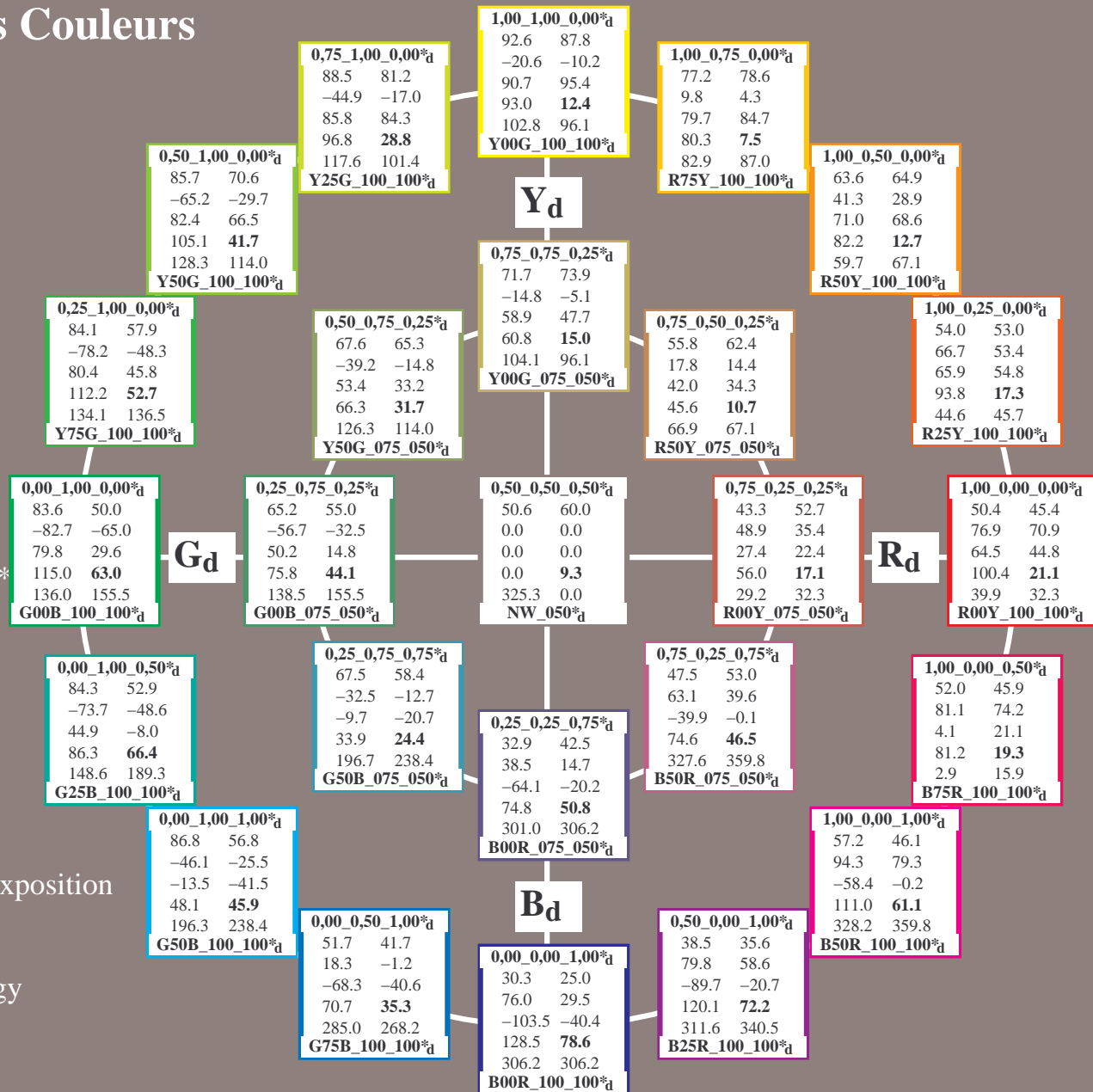
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 couleurs élémentaires H^* , brillance I^*
 chromie C^* : $HIC*_e$ (en bas)
 code de couleur:
 $LabCh*_d$; $Lab*/DE*/h*_d$

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n/fj	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md
0/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	0.0 0.0	389
1/657	R13Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	48.6 63.3 49.1	80.2 37.7	1.0 0.125 0.0	48.9 62.8 49.4	79.9 38.1	0.6 0.6	36
2/666	R25Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7	1.0 0.25 0.0	53.6 51.9 55.5	76.0 46.8	1.7 4.2	51
3/675	R38Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	58.8 41.1 61.7	74.1 56.3	1.0 0.375 0.0	59.1 40.3 62.0	74.0 56.9	0.9 5.1	42
4/684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1	0.0 0.0	59
5/693	R63Y_100_100a	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	72.5 14.8 77.6	79.0 79.1	1.0 0.625 0.0	72.1 15.4 77.1	78.6 78.6	0.8 6.8	68
6/702	R75Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	78.6 4.3 84.7	84.8 87.0	1.0 0.75 0.0	77.9 5.4 83.8	84.0 86.2	1.6 7.7	77
7/711	R88Y_100_100a	1.0 0.875 0.0	1.0 1.0 0.5	83	1.0 0.883 0.0	83.7 -3.8 90.5	90.6 92.4	1.0 0.875 0.0	83.4 -3.4 90.2	90.2 92.1	0.6 8.3	83
8/720	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1	0.0 0.0	89
9/639	Y13G_100_100a	0.875 1.0 0.0	1.0 1.0 0.5	97	0.883 1.0 0.0	84.5 -13.6 89.7	90.7 98.6	0.875 1.0 0.0	84.3 -13.9 89.2	90.3 98.8	0.5 9.6	96
10/558	Y25G_100_100a	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	81.2 -17.0 84.3	86.0 101.4	0.75 1.0 0.0	80.7 -17.5 83.5	83.3 101.8	1.0 1.0	102
11/477	Y38G_100_100a	0.625 1.0 0.0	1.0 1.0 0.5	112	0.633 1.0 0.0	75.6 -23.6 76.2	79.8 107.2	0.625 1.0 0.0	75.3 -24.0 75.7	79.4 107.6	0.7 1.1	111
12/396	Y50G_100_100a	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0	0.0 0.0	119
13/315	Y63G_100_100a	0.375 1.0 0.0	1.0 1.0 0.5	128	0.366 1.0 0.0	65.2 -36.4 57.6	68.2 122.3	0.375 1.0 0.0	65.7 -35.6 58.3	68.3 121.4	1.2 1.2	128
14/234	Y75G_100_100a	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	57.9 -48.3 45.8	66.5 136.5	0.25 1.0 0.0	58.4 -47.3 46.8	66.6 135.3	1.4 1.3	137
15/153	Y88G_100_100a	0.125 1.0 0.0	1.0 1.0 0.5	143	0.116 1.0 0.0	54.4 -54.7 38.0	66.6 145.1	0.125 1.0 0.0	54.7 -53.9 38.5	66.3 144.4	0.9 1.4	143
16/72	G00C_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5	0.0 0.0	149
17/73	G13C_100_100a	0.0 1.0 0.125	1.0 1.0 0.5	157	0.0 1.0 0.116	50.5 -62.9 22.4	66.8 160.4	0.0 1.0 0.125	50.5 -62.8 21.9	66.5 160.7	0.5 1.5	156
18/74	G25C_100_100a	0.0 1.0 0.25	1.0 1.0 0.5	164	0.0 1.0 0.233	51.1 -59.5 13.9	61.1 166.8	0.0 1.0 0.25	51.2 -58.9 12.7	60.3 167.7	1.2 1.6	162
19/75	G38C_100_100a	0.0 1.0 0.375	1.0 1.0 0.5	172	0.0 1.0 0.366	51.9 -54.9 3.7	55.0 176.1	0.0 1.0 0.375	52.0 -54.5 3.1	54.6 176.7	0.6 1.7	171
20/76	G50C_100_100a	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	52.9 -48.6 -8.0	49.3 189.3	0.0 1.0 0.5	52.9 -48.6 -8.0	49.3 189.3	0.0 0.0	180
21/77	G63C_100_100a	0.0 1.0 0.625	1.0 1.0 0.5	188	0.0 1.0 0.633	54.1 -42.0 -18.8	46.0 204.1	0.0 1.0 0.625	54.0 -42.3 -18.1	46.1 203.2	0.7 1.8	188
22/78	G75C_100_100a	0.0 1.0 0.75	1.0 1.0 0.5	196	0.0 1.0 0.766	55.1 -35.4 -28.4	45.4 218.7	0.0 1.0 0.75	55.0 -36.0 -27.4	45.3 217.2	1.1 1.1	197
23/79	G88C_100_100a	0.0 1.0 0.875	1.0 1.0 0.5	203	0.0 1.0 0.883	55.9 -30.4 -35.0	46.3 229.0	0.0 1.0 0.875	55.8 -30.7 -34.5	46.2 228.3	0.5 2.0	203
24/80	C00B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4	0.0 0.0	210
25/71	C13B_100_100a	0.0 0.875 1.0	1.0 1.0 0.5	217	0.0 0.883 1.0	54.3 -21.4 -41.4	46.6 242.6	0.0 0.875 1.0	54.1 -21.1 -41.3	46.4 242.9	0.3 2.1	216
26/62	C25B_100_100a	0.0 0.75 1.0	1.0 1.0 0.5	224	0.0 0.766 1.0	50.9 -16.2 -41.2	44.2 248.4	0.0 0.75 1.0	50.4 -15.5 -41.1	43.9 249.3	0.8 2.2	222
27/53	C38B_100_100a	0.0 0.625 1.0	1.0 1.0 0.5	232	0.0 0.633 1.0	46.8 -9.8 -40.9	42.1 256.4	0.0 0.625 1.0	46.5 -9.4 -40.8	41.9 256.9	0.4 2.3	231
28/44	C50B_100_100a	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	41.7 -1.2 -40.6	40.6 268.2	0.0 0.5 1.0	41.7 -1.2 -40.6	40.6 268.2	0.0 0.0	240
29/35	C63B_100_100a	0.0 0.375 1.0	1.0 1.0 0.5	248	0.0 0.366 1.0	37.0 6.6 -40.2	40.8 279.3	0.0 0.375 1.0	37.3 6.1 -40.2	40.7 278.6	0.6 2.4	248
30/26	C75B_100_100a	0.0 0.25 1.0	1.0 1.0 0.5	256	0.0 0.233 1.0	32.2 15.3 -40.3	43.1 290.8	0.0 0.25 1.0	32.8 14.3 -40.2	42.7 289.6	1.1 2.5	257
31/17	C88B_100_100a	0.0 0.125 1.0	1.0 1.0 0.5	263	0.0 0.116 1.0	28.4 22.8 -40.3	46.3 299.5	0.0 0.125 1.0	28.6 22.4 -40.2	46.1 299.0	0.5 2.6	263
32/8	B00M_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 0.0	270
33/89	B13M_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	27.7 35.6 -36.7	51.1 314.1	0.125 0.0 1.0	27.9 36.0 -36.4	51.2 314.7	0.5 2.7	276
34/170	B25M_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1	0.25 0.0 1.0	28.8 41.9 -32.5	53.1 322.1	0.9 2.8	282
35/251	B38M_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	32.5 51.2 -26.5	57.7 332.6	0.375 0.0 1.0	32.7 51.8 -26.0	58.0 333.3	0.8 2.9	291
36/332	B50M_100_100a	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5	0.0 3.0	300
37/413	B63M_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	38.3 65.8 -13.7	67.2 348.5	0.625 0.0 1.0	38.1 65.4 -14.0	66.9 347.9	0.5 3.0	308
38/494	B75M_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0	0.75 0.0 1.0	41.8 71.0 -9.2	71.6 352.5	0.8 3.1	317
39/575	B88M_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	44.3 75.4 -4.7	75.6 356.3	0.875 0.0 1.0	44.2 75.2 -5.0	75.3 356.1	0.4 3.2	323
40/656	M00R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	0.0 3.3	330
41/655	M13R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	45.9 78.3 3.8	78.4 2.8	1.0 0.0 0.875	45.9 78.2 4.1	78.3 363.0	0.2 3.3	336
42/654	M25R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9	1.0 0.0 0.75	45.9 77.1 8.6	77.6 366.4	0.6 3.4	342
43/653	M38R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	46.0 75.7 14.4	77.1 10.8	1.0 0.0 0.625	46.0 75.6 14.8	77.0 371.1	1.4 3.5	351
44/652	M50R_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9	1.0 0.0 0.5	45.9 74.2 21.1	77.1 375.9	0.0 3.6	360
45/651	M63R_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	45.8 72.9 28.7	78.4 21.5	1.0 0.0 0.375	45.8 72.9 28.3	78.3 381.2	0.4 3.6	368
46/650	M75R_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	45.6 71.1 35.3	80.3 26.1	1.0 0.0 0.25	45.6 72.1 34.6	80.0 385.6	0.7 3.7	377
47/649	M88R_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	45.5 71.4 40.4	82.1 29.5	1.0 0.0 0.125	45.5 71.4 40.1	81.9 389.3	0.3 3.8	383
48/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	1.0 0.0 0.0	45.4 70.9 44.8	83.9 392.3	0.0 3.8	389
49/0	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0	0.0 0.0 0.0	48.9 62.8 49.4	79.9 398.1	83.6 360	360
50/91	NW_013a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	33.2 0.0 0.0	0.0 0.0	0.125 0.125 0.125	24.3 0.0 0.0	78.9 8.9	360 360	360
51/182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	42.1 0.0 0.0	0.0 0.0	0.25 0.25 0.25	29.8 7.2 3.6	8.1 26.3	14.7 360	360
52/273	NW_038a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	51.0 0.0 0.0	0.0 0.0	0.375 0.375 0.375	35.7 7.5 7.1	10.4 43.4	18.5 360	360
53/364	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	60.0 0.0 0.0	0.0 0.0	0.5 0.5 0.5	45.3 10.0 11.0	14.9 47.8	20.9 360	360
54/455	NW_063a	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0	0.625 0.625 0.625	55.1 8.8 9.3	12.8 46.5	18.8 360	360
55/546	NW_075a	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0	0.75 0.75 0.75	64.6 6.7 9.1	11.3 53.7	17.4 360	360
56/637	NW_088a	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0	0.875 0.875 0.875	76.3 4.7 5.9	7.6 51.3	12.8 360	360
57/728	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0	1.0 1.0 1.0	86.7 1.6 2.9	3.3 60.9	9.5 360	360

delta E* = 4.0

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF77/PF77L0NA.TXT> / .PS
 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

TUB enregistrement: 20130201-PF77/PF77L0NA.TXT / .PS
 application pour la mesure des sorties sur offset, séparation cmy0 (CMY0)
 TUB matériel: code=rh4ta

n/j	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md		
0/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	0.0 0.0	45.4 70.9 44.8	83.9 32.3	
1/666	R25Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7	1.0 0.25 0.0	53.6 51.9 55.5	76.0 46.8 1.7	42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7
2/684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1	0.0 59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
3/702	R75Y_100_100a	1.0 0.75 0.0	1.0 1.0 0.5	76	1.0 0.766 0.0	78.6 4.3 84.7	84.8 87.0	1.0 0.75 0.0	77.9 5.4 83.8	84.0 86.2 1.6	77	1.0 0.766 0.0	78.6 4.3 84.7	84.8 87.0
4/720	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1	0.0 89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
5/558	Y25G_100_100a	0.75 1.0 0.0	1.0 1.0 0.5	104	0.766 1.0 0.0	81.2 -17.0 84.3	86.0 101.4	0.75 1.0 0.0	80.7 -17.5 83.5	85.3 101.8 1.0	102	0.766 1.0 0.0	81.2 -17.0 84.3	86.0 101.4
6/396	Y50G_100_100a	0.5 1.0 0.0	1.0 1.0 0.5	120	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0	0.0 119	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0
7/234	Y75G_100_100a	0.25 1.0 0.0	1.0 1.0 0.5	136	0.233 1.0 0.0	57.9 -48.3 45.8	66.5 136.5	0.25 1.0 0.0	58.4 -47.3 46.8	66.6 135.3 1.4	137	0.233 1.0 0.0	57.9 -48.3 45.8	66.5 136.5
8/72	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5	0.0 149	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5
9/72	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5	0.0 149	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5
10/76	G25B_100_100a	0.0 1.0 0.5	1.0 1.0 0.5	180	0.0 1.0 0.5	52.9 -48.6 -8.0	49.3 189.3	0.0 1.0 0.5	52.9 -48.6 -8.0	49.3 189.3	0.0 180	0.0 1.0 0.5	52.9 -48.6 -8.0	49.3 189.3
11/80	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4	0.0 210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4
12/44	G75B_100_100a	0.0 0.5 1.0	1.0 1.0 0.5	240	0.0 0.5 1.0	41.7 -1.2 -40.6	40.6 268.2	0.0 0.5 1.0	41.7 -1.2 -40.6	40.6 268.2	0.0 240	0.0 0.5 1.0	41.7 -1.2 -40.6	40.6 268.2
13/8	B00R_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
14/332	B25R_100_100a	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5	0.0 300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
15/656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	0.0 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
16/652	B75R_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9	0.0 360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
17/648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	0.0 389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
18/688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	70.5 35.4 22.4	41.9 32.3	1.0 0.5 0.5	68.0 29.9 28.7	41.5 32.8 8.7	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
19/706	R50Y_100_050a	1.0 0.75 0.5	1.0 0.5 0.75	60	1.0 0.75 0.5	80.2 14.4 34.3	37.2 67.1	1.0 0.75 0.5	80.4 9.0 35.3	36.5 75.5 5.4	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
20/724	Y00G_100_050a	1.0 1.0 0.5	1.0 0.5 0.75	90	1.0 1.0 0.5	91.7 -5.1 47.7	48.0 96.1	1.0 1.0 0.5	91.4 -7.7 42.5	43.2 100.3 5.8	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
21/562	Y50G_100_050a	0.75 1.0 0.5	1.0 0.5 0.75	120	0.75 1.0 0.5	83.1 -14.8 33.2	36.4 114.0	0.75 1.0 0.5	84.2 -14.1 31.5	34.5 114.0 2.1	119	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0
22/400	G00B_100_050a	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	72.8 -32.5 14.8	35.7 155.5	0.5 1.0 0.5	73.9 -23.7 19.9	31.0 140.0 10.1	149	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5
23/404	G50B_100_050a	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	76.2 -12.7 -20.7	24.3 238.4	0.5 1.0 1.0	78.7 -11.6 -18.3	21.7 237.6 3.6	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4
24/368	B00R_100_050a	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	60.3 14.7 -20.2	25.0 306.2	0.5 0.5 1.0	57.9 18.3 -20.7	27.7 311.4 4.3	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
25/692	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	70.8 39.6 -0.1	39.6 359.8	1.0 0.5 1.0	70.7 35.2 -3.7	35.4 353.9 5.7	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
26/688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	70.5 35.4 22.4	41.9 32.3	1.0 0.5 0.5	68.0 29.9 28.7	41.5 32.8 8.7	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
27/506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	52.7 35.4 22.4	41.9 32.3	0.75 0.25 0.25	50.4 39.4 31.9	50.7 38.9 10.5	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
28/524	R50Y_075_050a	0.75 0.5 0.25	0.75 0.5 0.5	60	0.75 0.5 0.25	62.4 14.4 34.3	37.2 67.1	0.75 0.5 0.25	61.2 18.1 39.5	43.4 65.3 6.4	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
29/542	Y00G_075_050a	0.75 0.75 0.25	0.75 0.5 0.5	90	0.75 0.75 0.25	73.9 -5.1 47.7	48.0 96.1	0.75 0.75 0.25	72.4 -1.4 48.0	48.0 91.7 3.9	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
30/380	Y50G_075_050a	0.5 0.75 0.25	0.75 0.5 0.5	120	0.5 0.75 0.25	65.3 -14.8 33.2	36.4 114.0	0.5 0.75 0.25	63.2 -12.6 35.5	37.7 109.6 3.7	119	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0
31/218	G00B_075_050a	0.25 0.75 0.25	0.75 0.5 0.5	150	0.25 0.75 0.25	55.0 -32.5 14.8	35.7 155.5	0.25 0.75 0.25	53.0 -20.9 21.7	35.3 142.0 8.5	149	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5
32/222	G50B_075_050a	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.75	48.4 -12.7 -20.7	24.3 238.4	0.25 0.75 0.75	55.9 -14.3 -16.3	21.7 228.6 5.3	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4
33/186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	52.5 14.7 -20.2	25.0 306.2	0.25 0.25 0.75	37.5 18.9 -20.4	27.9 312.8 6.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
34/510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	53.0 39.6 -0.1	39.6 359.8	0.75 0.25 0.75	52.4 44.4 0.5	44.4 0.6 4.8	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
35/506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	52.7 35.4 22.4	41.9 32.3	0.75 0.25 0.25	50.4 39.4 31.9	50.7 38.9 10.5	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
36/324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.0	34.9 35.4 22.4	41.9 32.3	0.5 0.0 0.0	34.8 44.7 22.4	50.0 26.6 9.2	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
37/342	R50Y_050_050a	0.5 0.25 0.0	0.5 0.5 0.25	60	0.5 0.25 0.0	44.6 14.4 34.3	37.2 67.1	0.5 0.25 0.0	43.4 24.2 33.3	41.2 53.9 9.9	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
38/360	Y00G_050_050a	0.5 0.5 0.0	0.5 0.5 0.25	90	0.5 0.5 0.0	56.1 -5.1 47.7	48.0 96.1	0.5 0.5 0.0	52.6 3.9 44.2	44.3 84.8 10.3	89	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1
39/198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.5 0.25	120	0.25 0.5 0.0	47.4 -14.8 33.2	36.4 114.0	0.25 0.5 0.0	43.1 -14.1 28.4	31.7 116.4 6.5	119	0.5 1.0 0.0	70.6 -29.7 66.5	72.8 114.0
40/36	G00B_050_050a	0.0 0.5 0.0	0.5 0.5 0.25	150	0.0 0.5 0.0	37.2 -32.5 14.8	35.7 155.5	0.0 0.5 0.0	37.3 -36.4 15.2	39.5 157.2 3.9	149	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5
41/40	G50B_050_050a	0.0 0.5 0.5	0.5 0.5 0.25	210	0.0 0.5 0.5	40.5 -12.7 -20.7	24.3 238.4	0.0 0.5 0.5	39.1 -11.5 -13.3	25.3 211.8 11.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4
42/4	B00R_050_050a	0.0 0.0 0.5	0.5 0.5 0.25	270	0.0 0.0 0.5	24.7 14.7 -20.2	25.0 306.2	0.0 0.0 0.5	24.3 11.6 -18.9	22.1 301.5 3.4	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
43/328	B50R_050_050a	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	35.2 39.6 -0.1	39.6 359.8	0.5 0.0 0.5	35.0 49.8 0.6	49.8 0.7 10.2	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
44/324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.0	34.9 35.4 22.4	41.9 32.3	0.5 0.0 0.0	34.8 44.7 22.4	50.0 26.6 9.2	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
45/0	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	360	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0 0.0	360	1.0 1.0 1.0	95.6 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	33.2 0.0 0.0	0.0 0.0	0.125 0.125 0.125	29.8 7.2 3.6	8.1 26.3 8.7	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
47/182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	42.1 0.0 0.0	0.0 0.0	0.25 0.25 0.25	35.7 7.5 7.1	10.4 43.4 12.2	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
48/273	NW													

n=j	HIC*Fd	rgb_Fd	iet_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md
0	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 360	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1	B00R_012_012a	0.0 0.0 0.125	0.125 0.125 0.125	0.062 270	0.0 0.0 0.125	24.4 3.6 -5.0	6.2 306.2	0.0 0.0 0.125	23.8 2.3 -3.5	4.2 303.1 2.1	27.0 29.5 -40.4	50.0 50.0 306.2
2	B00R_025_025a	0.0 0.0 0.25	0.25 0.25 0.125	0.187 270	0.0 0.0 0.25	24.5 7.3 -10.1	12.5 306.2	0.0 0.0 0.25	23.9 4.8 -8.0	9.4 300.8 3.3	27.0 29.5 -40.4	50.0 306.2
3	B00R_037_037a	0.0 0.0 0.375	0.375 0.375 0.187	0.270 270	0.0 0.0 0.375	24.6 11.0 -15.1	18.7 306.2	0.0 0.0 0.375	24.1 6.9 -12.1	13.9 299.8 5.1	27.0 29.5 -40.4	50.0 306.2
4	B00R_050_050a	0.0 0.0 0.5	0.5 0.5 0.25	0.270 270	0.0 0.0 0.5	24.7 14.7 -20.2	25.0 306.2	0.0 0.0 0.5	24.3 11.6 -18.9	22.1 301.5 3.4	27.0 29.5 -40.4	50.0 306.2
5	B00R_062_062a	0.0 0.0 0.625	0.625 0.625 0.312	0.270 270	0.0 0.0 0.625	24.8 18.4 -25.2	31.3 306.2	0.0 0.0 0.625	24.6 15.8 -24.6	29.2 307.7 2.7	27.0 29.5 -40.4	50.0 306.2
6	B00R_075_075a	0.0 0.0 0.75	0.75 0.75 0.375	0.270 270	0.0 0.0 0.75	24.9 22.1 -30.3	37.5 306.2	0.0 0.0 0.75	24.7 20.7 -30.7	37.0 303.9 1.5	27.0 29.5 -40.4	50.0 306.2
7	B00R_087_087a	0.0 0.0 0.875	0.875 0.875 0.437	0.270 270	0.0 0.0 0.875	24.9 25.8 -35.3	43.8 306.2	0.0 0.0 0.875	24.8 25.5 -35.9	44.0 303.5 0.7	27.0 29.5 -40.4	50.0 306.2
8	B00R_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	0.270 270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2 0.0	27.0 29.5 -40.4	50.0 306.2
9	G00B_012_012a	0.0 0.125 0.0	0.125 0.125 0.062	0.150 210	0.0 0.125 0.0	27.5 -8.1 3.7	8.9 155.5	0.0 0.125 0.0	27.1 -8.2 2.9	8.7 160.0 0.8	149 50.0 -65.0	29.6 71.4 155.5
10	G50B_012_012a	0.0 0.125 0.125	0.125 0.125 0.062	0.210 210	0.0 0.125 0.125	28.4 -3.1 -5.1	6.0 238.4	0.0 0.125 0.125	26.7 -5.9 -1.1	6.0 190.5 5.2	210 56.8 -25.5	-41.5 48.7 238.4
11	G75B_025_025a	0.0 0.125 0.25	0.25 0.25 0.125	0.240 210	0.0 0.125 0.25	28.7 -0.3 -10.1	10.1 268.2	0.0 0.125 0.25	27.1 -3.6 -5.7	6.8 237.4 5.7	240 41.7 -12.2	-40.6 40.6 268.2
12	G84B_037_037a	0.0 0.125 0.375	0.375 0.375 0.187	0.251 210	0.0 0.118 0.375	28.4 3.7 -15.1	15.6 283.7	0.0 0.125 0.375	27.1 -0.2 -10.8	10.8 268.5 5.9	251 35.2 9.9	-40.4 41.6 283.7
13	G88B_050_050a	0.0 0.125 0.5	0.5 0.5 0.25	0.256 210	0.0 0.116 0.5	28.3 7.6 -20.1	21.5 290.8	0.0 0.125 0.5	27.3 4.4 -17.8	18.3 284.1 4.1	257 32.2 15.3	-40.3 43.1 290.8
14	G90B_062_062a	0.0 0.125 0.625	0.625 0.625 0.312	0.259 210	0.0 0.114 0.625	28.2 11.6 -25.2	27.8 294.6	0.0 0.125 0.625	27.8 8.7 -24.2	25.7 289.8 3.0	260 30.6 18.5	-40.4 44.5 294.6
15	G92B_075_075a	0.0 0.125 0.75	0.75 0.75 0.375	0.261 210	0.0 0.112 0.75	28.2 15.5 -30.3	34.0 291.1	0.0 0.125 0.75	28.1 13.4 -30.2	33.0 293.9 2.1	262 29.5 20.7	-40.4 45.4 297.1
16	G93B_087_087a	0.0 0.125 0.875	0.875 0.875 0.437	0.262 210	0.0 0.116 0.875	28.3 19.1 -35.2	40.1 298.4	0.0 0.125 0.875	28.3 18.0 -35.9	39.9 296.8 1.1	262 28.9 21.8	-40.3 45.8 298.4
17	G94B_100_100a	0.0 0.125 1.0	1.0 1.0 0.5	0.263 210	0.0 0.116 1.0	28.4 22.8 -40.3	46.3 299.5	0.0 0.125 1.0	28.6 22.4 -40.2	46.1 299.0 0.5	263 28.4 22.8	-40.3 46.3 299.5
18	G00B_025_025a	0.0 0.25 0.0	0.25 0.25 0.125	0.180 210	0.0 0.25 0.0	30.7 -16.2 7.4	17.8 155.5	0.0 0.25 0.0	30.5 -18.5 7.5	20.0 157.7 2.2	149 50.0 -65.0	29.6 71.4 155.5
19	G25B_025_025a	0.0 0.25 0.125	0.25 0.25 0.125	0.180 210	0.0 0.25 0.125	31.5 -12.1 -2.0	12.3 189.3	0.0 0.25 0.125	30.7 -16.4 2.9	16.6 169.8 6.5	180 56.8 -25.5	-41.5 48.7 238.4
20	G50B_025_025a	0.0 0.25 0.25	0.25 0.25 0.125	0.210 210	0.0 0.25 0.25	32.4 -6.3 -10.3	12.1 238.4	0.0 0.25 0.25	31.1 -13.5 -2.5	13.7 190.8 10.6	210 56.8 -25.5	-41.5 48.7 238.4
21	G65B_037_037a	0.0 0.25 0.375	0.375 0.375 0.187	0.229 210	0.0 0.256 0.375	33.3 -4.6 -15.4	16.0 253.3	0.0 0.25 0.375	31.7 -11.0 -8.3	13.7 217.0 9.7	228 48.3 -12.2	-41.1 42.9 253.3
22	G75B_050_050a	0.0 0.25 0.5	0.5 0.5 0.25	0.240 210	0.0 0.25 0.5	33.0 0.6 -20.3	20.3 268.2	0.0 0.25 0.5	31.8 -5.6 -15.7	16.7 250.1 6.9	240 41.7 -1.2	-40.6 40.6 268.2
23	G80B_062_062a	0.0 0.25 0.625	0.625 0.625 0.312	0.247 210	0.0 0.239 0.625	32.6 3.5 -25.1	25.4 277.9	0.0 0.25 0.625	32.1 -0.6 -22.5	22.5 268.3 4.9	247 37.6 5.6	-40.3 40.7 277.9
24	G84B_075_075a	0.0 0.25 0.75	0.75 0.75 0.375	0.251 210	0.0 0.237 0.75	32.5 7.4 -30.3	31.2 283.7	0.0 0.25 0.75	32.2 4.8 -29.1	29.5 279.4 2.8	251 35.2 9.9	-40.4 41.6 283.7
25	G88B_087_087a	0.0 0.25 0.875	0.875 0.875 0.437	0.254 210	0.0 0.233 0.875	32.3 11.5 -35.2	37.1 288.1	0.0 0.25 0.875	32.3 9.9 -34.9	36.3 285.8 1.6	255 33.2 13.2	-40.3 42.4 288.1
26	G88B_100_100a	0.0 0.25 1.0	1.0 1.0 0.5	0.256 210	0.0 0.233 1.0	32.2 15.3 -40.3	43.1 290.8	0.0 0.25 1.0	32.8 14.3 -40.2	42.7 289.6 1.1	257 32.2 15.3	-40.3 43.1 290.8
27	G00B_037_037a	0.0 0.375 0.0	0.375 0.375 0.187	0.150 210	0.0 0.375 0.0	34.0 -24.3 11.1	26.7 155.5	0.0 0.375 0.0	33.9 -27.6 11.4	29.8 157.4 3.2	149 50.0 -65.0	29.6 71.4 155.5
28	G15B_037_037a	0.0 0.375 0.125	0.375 0.375 0.187	0.169 210	0.0 0.375 0.118	34.6 -21.3 2.7	21.4 172.5	0.0 0.375 0.125	34.2 -25.2 6.6	26.4 165.4 5.7	168 51.6 -56.8	7.4 57.3 172.5
29	G34B_037_037a	0.0 0.375 0.25	0.375 0.375 0.187	0.191 210	0.0 0.375 0.256	35.6 -14.8 8.5	17.1 209.7	0.0 0.375 0.25	34.7 -22.1 -0.5	22.1 181.3 10.8	191 54.5 -39.7	-22.7 45.7 209.7
30	G50B_037_037a	0.0 0.375 0.375	0.375 0.375 0.187	0.210 210	0.0 0.375 0.375	36.5 -9.5 -15.5	18.2 238.4	0.0 0.375 0.375	34.9 -18.4 -6.6	19.6 199.8 12.6	210 56.8 -25.5	-41.5 48.7 238.4
31	G61B_050_050a	0.0 0.375 0.5	0.5 0.5 0.25	0.224 210	0.0 0.383 0.5	37.6 -8.1 -20.6	22.1 248.4	0.0 0.375 0.5	35.7 -14.1 -14.6	23.3 226.1 8.6	222 50.9 -16.2	-41.2 44.2 248.4
32	G69B_062_062a	0.0 0.375 0.625	0.625 0.625 0.312	0.233 210	0.0 0.385 0.625	38.0 -5.5 -25.5	26.1 257.7	0.0 0.375 0.625	36.6 -10.0 -21.5	23.7 245.0 6.1	232 46.2 -8.9	-40.9 41.8 257.7
33	G75B_075_075a	0.0 0.375 0.75	0.75 0.75 0.375	0.240 210	0.0 0.375 0.75	37.3 0.9 -30.4	30.5 268.2	0.0 0.375 0.75	36.5 -4.0 -28.4	28.6 261.8 3.8	240 41.7 -1.2	-40.6 40.6 268.2
34	G79B_087_087a	0.0 0.375 0.875	0.875 0.875 0.437	0.245 210	0.0 0.364 0.875	37.0 3.2 -35.4	35.6 275.1	0.0 0.375 0.875	36.5 2.0 -34.7	34.8 273.3 1.4	245 38.8 3.6	-40.5 40.6 275.1
35	G81B_100_100a	0.0 0.375 1.0	1.0 1.0 0.5	0.248 210	0.0 0.366 1.0	37.0 6.6 -40.2	40.8 279.3	0.0 0.375 1.0	37.3 6.1 -40.2	40.7 278.6 0.6	248 37.0 6.6	-40.2 40.8 279.3
36	G00B_050_050a	0.0 0.5 0.0	0.5 0.5 0.25	0.150 210	0.0 0.5 0.0	37.2 -32.5 14.8	35.7 155.5	0.0 0.5 0.0	37.3 -36.4 15.2	39.5 157.2 3.9	149 50.0 -65.0	29.6 71.4 155.5
37	G11B_050_050a	0.0 0.5 0.125	0.5 0.5 0.25	0.164 210	0.0 0.5 0.116	37.7 -29.7 6.9	30.5 166.8	0.0 0.5 0.125	37.6 -34.1 9.9	35.5 163.8 5.2	162 51.1 -59.5	13.9 61.1 166.8
38	G25B_050_050a	0.0 0.5 0.25	0.5 0.5 0.25	0.180 210	0.0 0.5 0.25	38.6 -24.3 4.0	24.6 189.3	0.0 0.5 0.25	38.1 -30.3 2.2	30.4 175.7 8.7	180 51.6 -56.8	-8.0 49.3 189.3
39	G38B_050_050a	0.0 0.5 0.375	0.5 0.5 0.25	0.196 210	0.0 0.5 0.383	39.7 -17.7 -14.2	22.7 218.7	0.0 0.5 0.375	38.7 -26.0 -5.6	26.6 192.2 12.0	197 55.1 -35.4	-28.4 45.4 218.7
40	G50B_050_050a	0.0 0.5 0.5	0.5 0.5 0.25	0.210 210	0.0 0.5 0.5	40.5 -12.7 -20.7	24.3 238.4	0.0 0.5 0.5	39.1 -21.5 -13.3	25.3 211.8 11.5	210 56.8 -25.5	-41.5 48.7 238.4
41	G59B_062_062a	0.0 0.625 0.625	0.625 0.625 0.312	0.221 210	0.0 0.51 0.625	41.9 -11.5 -25.8	28.3 245.8	0.0 0.625 0.403	40.7 -17.0 -21.0	27.1 231.0 7.4	219 52.4 -18.5	-41.3 45.3 245.8
42	G65B_075_075a	0.0 0.625 0.75	0.75 0.75 0.375	0.229 210	0.0 0.512 0.75	42.3 -9.2 -30.8	32.1 253.3	0.0 0.625 0.75	41.1 -12.1 -28.0	30.5 246.4 4.2	228 48.3 -12.2	-41.1 42.9 253.3
43	G70B_087_087a	0.0 0.625 0.875	0.875 0.875 0.437	0.235 210	0.0 0.51 0.875	42.3 -5.8 -35.8	36.3 260.7	0.0 0.625 0.875	41.6 -6.8 -34.8	35.4 258.8 1.6	234 44.9 -6.6	-41.0 41.5 260.7
44	G75B_100_100a	0.0 0.625 1.0	1.0 1.0 0.5	0.240 210	0.0 0.51 1.0	41.7 -1.2 -40.6	40.6 268.2	0.0 0.625 1.0	41.7 -1.2 -40.6	40.6 268.2 0.0	240 41.7 -1.2	-40.6 40.6 268.2
45	G00B_062_062a	0.0 0.625 0.0	0.625 0.625 0.312	0.150 210	0.0 0.625 0.0	40.4 -40.6 18.5	44.6 155.5	0.0 0.625 0.0	41.4 -45.8 19.8	49.9 156.6 5.5	149 50.0 -65.0	29.6 71.4 155.5
46	G09B_062_062a	0.0 0.625 0.125	0.625 0.625 0.312	0.161 210	0.0 0.625 0.114	40.9 -38.2 10.9	39.7 164.0	0.0 0.625 0.125	41.6 -43.4 13.8	45.5 162.2 5.9	159 50.8 -61.1	17.4 63.6 164.0
47	G19B_062_062a	0.0 0.625 0.25	0.625 0.625 0.312	0.173 210	0.0 0.625 0.239	41.6 -33.9 1.4	33.9 177.5	0.0 0.625 0.25	42.1 -39.7 5.6	40.1 171.9 7.1	172 52.0 0.0	0.0 0.0 177.5
48	G30B_062_062a	0.0 0.625 0.375	0.625 0.625 0.312	0.187 210	0.0 0.625 0.385	42.8 -26.7 -10.9	28.9 202.2	0.0 0.625 0.375	42.9 -34.7 -3.7	34.9 186.1 10.7	187 54.3 -33.3	-31.3 54.3 202.2
49	G40B_062_062a	0.0 0.625 0.5	0.625 0.625 0.312	0.199 210	0.0 0.625 0.51	43.8 -20.8 -15.9	28.6 233.1	0.0 0.625 0.5	43.4 -29.7 -12.3	32.1 202.4 11.4	200 55.4 -33.3	-31.3 45.7 233.1
50	G50B_062_062a	0.0 0.625 0.625	0.625 0.625 0.312	0.210 210	0.0 0.625 0.625	44.6 -15.9 -25.9	30.4 238.4	0.0 0.625 0.625	44.2 -25.0 -19.8	31.9 218.3 10.9	210 56.8 -25.5	-41.5 48.7 238.4
51	G57B_075_075a	0.0 0.625 0.75	0.75 0.75 0.375	0.219 210	0.0 0.637 0.75	46.1 -15.0 -31.0	34.4 244.1	0.0 0.625 0.75	45.4 -19.9 -27.5	34.0 234.0 6.1	217 53.4 -20.0	-41.3 45.9 244.1
52	G63B_087_087a	0.0 0.625 0.875	0.875 0.875 0.437	0.226 210	0.0 0.641 0.875	46.7 -12.8 -36.0	38.2 250.3	0.0 0.625 0.875	46.0 -14.7 -34.6	37.6 246.9 2.3	224 53.4 -20.0	-41.1 43.6 250.3
53	G68B_100_100a	0.0 0.625 1.0	1.0 1.0 0.5	0.232 210	0.0 0.633 1.0	46.8 -9.8 -40.9	42.1 256.4	0.0				

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF77/PF77.HTM>
 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

TUB enregistrement: 20130201 -PF77/PF77L0NA.TXT /.PS TUB matériel: code=rh4ta
 application pour la mesure des sorties sur offset, séparation cmy0 (CMY0)

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md		
81	R00Y_012_012a	0.125 0.0 0.0	0.125 0.125 0.062	390	0.125 0.0 0.0	27.0 8.8 5.6	10.4 32.3	0.125 0.0 0.0	26.6 14.6 4.2	15.2 16.1 5.9	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
82	B50R_012_012a	0.125 0.0 0.125	0.125 0.125 0.062	330	0.125 0.0 0.125	27.0 9.9 0.0	9.9 359.8	0.125 0.0 0.125	26.7 15.8 0.3	15.8 11.1 5.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
83	B25R_025_025a	0.125 0.0 0.25	0.25 0.25 0.125	300	0.125 0.0 0.25	27.1 14.6 -5.1	15.5 340.5	0.125 0.0 0.25	26.9 17.8 -4.5	18.4 345.8 3.2	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
84	B15R_037_037a	0.125 0.0 0.375	0.375 0.375 0.187	289	0.118 0.0 0.375	26.8 17.7 -11.0	20.9 328.1	0.125 0.0 0.375	26.6 19.3 -9.3	21.5 334.2 2.3	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
85	B11R_050_050a	0.125 0.0 0.5	0.5 0.5 0.25	284	0.116 0.0 0.5	26.5 20.6 -16.5	26.4 321.1	0.125 0.0 0.5	27.0 21.7 -15.4	26.6 324.6 1.7	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
86	B09R_062_062a	0.125 0.0 0.625	0.625 0.625 0.312	281	0.114 0.0 0.625	26.8 24.2 -21.7	32.5 318.2	0.125 0.0 0.625	27.1 25.2 -21.3	33.1 319.7 1.0	279	0.183 0.0 1.0	28.3 38.8 -34.7	52.1 318.2
87	B07R_075_075a	0.125 0.0 0.75	0.75 0.75 0.375	279	0.112 0.0 0.75	27.1 27.9 -26.8	38.7 316.2	0.125 0.0 0.75	27.4 29.1 -26.9	39.7 317.2 1.2	278	0.15 0.0 1.0	28.1 37.2 -35.7	51.6 316.2
88	B06R_087_087a	0.125 0.0 0.875	0.875 0.875 0.437	278	0.116 0.0 0.875	27.5 31.9 -31.6	44.9 315.2	0.125 0.0 0.875	27.4 33.0 -32.0	46.0 315.8 1.1	277	0.133 0.0 1.0	27.9 36.4 -36.2	51.3 315.2
89	B05R_100_100a	0.125 0.0 1.0	1.0 1.0 0.5	277	0.116 0.0 1.0	27.7 35.6 -36.7	51.1 314.1	0.125 0.0 1.0	27.9 36.0 -36.4	51.2 314.7 0.5	276	0.116 0.0 1.0	27.7 35.6 -36.7	51.1 314.1
90	Y00G_012_012a	0.125 0.125 0.0	0.125 0.125 0.062	90	0.125 0.125 0.0	32.3 -1.2	11.9 120 96.1	0.125 0.125 0.0	29.6 5.9 7.7	9.7 52.8 8.6	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
91	NW_012a	0.125 0.125 0.125	0.125 0.0 0.125	360	0.125 0.125 0.125	33.2 0.0 0.0	0.0 0.0 0.0	0.125 0.125 0.125	29.8 7.2 3.6	8.1 26.3 8.7	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
92	BO0R_025_012a	0.125 0.125 0.25	0.25 0.125 0.187	270	0.124 0.124 0.25	33.3 3.6 -5.0	6.2 306.2	0.125 0.125 0.25	30.0 8.9 -1.7	9.1 34.9 7.0	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
93	BO0R_037_025a	0.125 0.125 0.375	0.375 0.25 0.25	270	0.124 0.124 0.375	33.4 7.3 -10.1	12.5 306.2	0.125 0.125 0.375	30.4 11.8 -7.5	14.0 327.5 5.9	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
94	BO0R_050_037a	0.125 0.125 0.5	0.5 0.375 0.312	270	0.124 0.124 0.5	33.5 11.0 -15.1	18.7 306.2	0.125 0.125 0.5	30.5 14.5 -14.1	20.3 315.8 4.7	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
95	BO0R_062_050a	0.125 0.125 0.625	0.625 0.5 0.375	270	0.125 0.125 0.625	33.6 14.7 -20.2	25.0 306.2	0.125 0.125 0.625	30.9 17.9 -20.2	27.0 311.4 4.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
96	BO0R_075_062a	0.125 0.125 0.75	0.75 0.625 0.437	270	0.125 0.125 0.75	33.7 18.4 -25.2	31.3 306.2	0.125 0.125 0.75	31.5 21.1 -26.2	33.7 308.7 3.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
97	BO0R_087_075a	0.125 0.125 0.875	0.875 0.75 0.5	270	0.125 0.125 0.875	33.8 22.1 -30.3	37.5 306.2	0.125 0.125 0.875	31.5 25.0 -31.5	40.2 308.4 3.8	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
98	BO0R_100_087a	0.125 0.125 1.0	1.0 0.875 0.562	270	0.125 0.125 1.0	33.9 25.8 -35.3	43.8 306.2	0.125 0.125 1.0	32.0 28.2 -36.5	46.0 307.8 3.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
99	Y50G_025_025a	0.125 0.25 0.0	0.25 0.25 0.125	150	0.125 0.25 0.0	35.9 -7.4	16.6 182 114.0	0.125 0.25 0.0	33.7 -4.5	12.9 13.6 109.2	5.2 119	0.5 1.0 0.0	70.6 -69.7	66.5 72.8 114.0
100	GO0B_025_012a	0.125 0.25 0.125	0.25 0.125 0.187	210	0.124 0.25 0.124	36.4 -8.1	3.7 8.9 155.5	0.125 0.25 0.125	33.9 -3.6	8.3 9.1 113.6	6.9 149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
101	G50B_025_012a	0.125 0.25 0.25	0.25 0.125 0.187	210	0.124 0.25 0.25	37.3 -3.1	-5.1 6.0 238.4	0.125 0.25 0.25	34.4 -1.1	1.6 2.0 124.6	7.7 210	0.0 0.0 1.0	56.8 -25.5	-41.5 48.7 238.4
102	G75B_037_025a	0.125 0.25 0.375	0.375 0.25 0.25	240	0.124 0.25 0.375	37.6 -0.3	-10.1 10.1 268.2	0.125 0.25 0.375	34.7 1.3 -4.5	4.7 286.1 6.5	240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6 268.2
103	G84B_050_037a	0.125 0.25 0.5	0.5 0.375 0.312	251	0.124 0.243 0.5	37.3 3.7 -15.1	15.6 283.7	0.125 0.25 0.5	35.0 4.5 -11.8	12.7 291.0 4.1	251	0.0 0.316 1.0	35.2 9.9 -40.4	41.6 283.7
104	G88B_062_050a	0.125 0.25 0.625	0.625 0.5 0.375	256	0.125 0.241 0.625	37.2 7.6 -20.1	21.5 290.8	0.125 0.25 0.625	35.2 8.5 -18.0	20.0 295.3 2.9	257	0.0 0.233 1.0	32.2 15.3 -40.3	43.1 290.8
105	G90B_075_062a	0.125 0.25 0.75	0.75 0.625 0.437	259	0.125 0.239 0.75	37.1 11.6 -25.2	27.8 294.6	0.125 0.25 0.75	35.7 12.5 -24.8	27.8 296.7 1.7	260	0.0 0.183 1.0	30.6 18.5 -40.4	44.5 294.6
106	G92B_087_075a	0.125 0.25 0.875	0.875 0.75 0.5	261	0.125 0.237 0.875	37.1 15.5 -30.3	34.0 297.1	0.125 0.25 0.875	36.1 16.4 -30.6	34.8 298.2 1.3	262	0.0 0.15 1.0	29.5 20.7 -40.4	45.4 297.1
107	G93B_100_087a	0.125 0.25 1.0	1.0 0.875 0.562	262	0.125 0.241 1.0	37.2 19.1 -35.2	40.1 294.4	0.125 0.25 1.0	36.4 19.7 -35.8	40.8 298.8 1.1	262	0.0 0.133 1.0	28.9 21.8 -40.3	45.8 298.4
108	Y68G_037_037a	0.125 0.375 0.0	0.375 0.375 0.187	131	0.118 0.375 0.0	38.6 -15.5	19.9 25.3 127.8	0.125 0.375 0.0	37.4 -15.0	17.0 22.7 131.3	3.1 131	0.316 1.0 0.0	62.3 -41.4	53.2 67.5 127.8
109	GO0B_037_025a	0.125 0.375 0.125	0.375 0.25 0.25	150	0.124 0.375 0.124	39.7 -16.2	7.4 17.8 155.5	0.125 0.375 0.125	37.6 -12.8	11.7 17.3 137.3	5.9 149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
110	G25B_037_025a	0.125 0.375 0.25	0.375 0.25 0.25	180	0.124 0.375 0.25	40.4 -12.1	-2.0 12.3 189.3	0.125 0.375 0.25	38.4 -10.8	5.2 12.0 154.3	7.6 180	0.0 1.0 0.5	52.9 -48.6	-8.0 49.3 189.3
111	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	41.3 -6.3	-10.3 12.1 238.4	0.125 0.375 0.375	38.8 -7.8	-2.3 8.2 196.2	8.6 210	0.0 1.0 1.0	56.8 -25.5	-41.5 48.7 238.4
112	G65B_050_037a	0.125 0.375 0.5	0.5 0.375 0.312	229	0.124 0.381 0.5	42.2 -4.6	-15.4 16.0 253.3	0.125 0.375 0.5	39.7 -5.2	-9.5 10.8 241.1	6.4 228	0.0 0.683 1.0	48.3 -12.2	-41.1 42.9 253.3
113	G75B_062_050a	0.125 0.375 0.625	0.625 0.5 0.375	240	0.125 0.375 0.625	41.9 -0.6	-20.3 20.3 268.2	0.125 0.375 0.625	39.7 -0.9	-16.6 16.6 266.8	4.2 240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6 268.2
114	G80B_075_062a	0.125 0.375 0.75	0.75 0.625 0.437	247	0.125 0.364 0.75	41.5 3.5 -20.1	25.4 277.9	0.125 0.375 0.75	39.8 4.0 -24.0	24.4 279.5 2.0	247	0.0 0.383 1.0	37.6 5.6 -40.3	40.7 277.9
115	G84B_087_075a	0.125 0.375 0.875	0.875 0.75 0.5	251	0.125 0.362 0.875	41.4 7.4 -30.3	31.2 283.7	0.125 0.375 0.875	40.3 8.1 -30.2	31.3 285.1 1.3	251	0.0 0.316 1.0	35.2 9.9 -40.4	41.6 283.7
116	G86B_100_087a	0.125 0.375 1.0	1.0 0.875 0.562	254	0.125 0.358 1.0	41.2 11.5 -35.2	37.1 288.1	0.125 0.375 1.0	40.4 12.6 -35.8	37.9 289.4 1.4	255	0.0 0.266 1.0	33.4 13.2 -40.3	42.4 288.1
117	Y76G_050_050a	0.125 0.5 0.0	0.5 0.5 0.25	136	0.116 0.5 0.0	41.1 -24.1	22.9 33.2 136.5	0.125 0.5 0.0	41.0 -23.7	21.5 32.0 137.7	1.4 137	0.233 1.0 0.0	57.9 -48.3	45.8 66.5 136.5
118	GO0B_050_037a	0.125 0.5 0.125	0.5 0.375 0.312	150	0.124 0.5 0.124	42.9 -24.3	11.1 26.7 155.5	0.125 0.5 0.125	41.5 -21.6	15.4 26.6 144.4	5.3 149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
119	G15B_050_037a	0.125 0.5 0.25	0.5 0.375 0.312	169	0.124 0.5 0.243	43.5 -21.3	27.1 21.4 172.5	0.125 0.5 0.25	42.1 -19.2	8.0 20.8 157.3	5.8 169	0.0 1.0 0.316	51.6 -56.8	7.4 57.3 172.5
120	G34B_050_037a	0.125 0.5 0.375	0.5 0.375 0.312	191	0.124 0.5 0.381	44.5 -14.8	-8.5 17.1 209.7	0.125 0.5 0.375	42.7 -15.8	-0.3 15.8 181.2	8.4 191	0.0 1.0 0.683	54.5 -39.7	-22.7 45.7 209.7
121	G50B_050_037a	0.125 0.5 0.5	0.5 0.375 0.312	210	0.124 0.5 0.5	45.4 -9.5	-15.5 18.2 238.4	0.125 0.5 0.5	43.0 -12.4	-8.0 14.7 212.9	8.3 210	0.0 1.0 1.0	56.8 -25.5	-41.5 48.7 238.4
122	G61B_062_050a	0.125 0.5 0.625	0.625 0.5 0.375	224	0.125 0.508 0.625	46.5 -8.1	-20.6 22.1 248.4	0.125 0.5 0.625	44.2 -9.4	-15.4 18.1 238.4	5.7 222	0.0 0.766 1.0	50.9 -16.2	-41.2 44.2 248.4
123	G69B_075_062a	0.125 0.5 0.75	0.75 0.625 0.437	233	0.125 0.51 0.75	46.9 -5.5	-25.5 26.1 257.7	0.125 0.5 0.75	45.0 -5.2	-22.9 23.4 257.0	3.3 232	0.0 0.616 1.0	46.2 -8.9	-40.9 41.8 257.7
124	G75B_087_075a	0.125 0.5 0.875	0.875 0.75 0.5	240	0.125 0.5 0.875	46.2 -0.9	-30.4 30.5 268.2	0.125 0.5 0.875	45.2 -0.4	-29.7 29.7 269.1	1.4 240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6 268.2
125	G79B_100_087a	0.125 0.5 1.0	1.0 0.875 0.562	245	0.125 0.489 1.0	45.9 3.2 -35.4	35.6 275.1	0.125 0.5 1.0	45.4 4.0 -35.8	36.1 276.5 1.0	245	0.0 0.416 1.0	38.8 3.6 -40.5	40.6 275.1
126	Y81G_062_062a	0.125 0.625 0.0	0.625 0.625 0.312	139	0.114 0.625 0.0	44.4 -31.9	26.6 41.5 140.1	0.125 0.625 0.0	45.0 -33.3	26.4 42.5 141.5	1.5 140	0.183 1.0 0.0	56.4 -51.0	42.5 66.4 140.1
127	GO0B_062_050a	0.125 0.625 0.125	0.625 0.5 0.375	150	0.125 0.625 0.125	46.1 -33.2	14.8 35.7 155.5	0.125 0.625 0.125	45.9 -30.8	20.2 36.9 146.8	5.6 149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
128	G11B_062_050a	0.125 0.625 0.25	0.625 0.5 0.375	164	0.125 0.625 0.241	46.6 -29.7	6.9 30.5 166.8	0.125 0.625 0.25	46.5 -28.6	12.2 31.1 156.7	5.4 162	0.0 1.		

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF77/PF77.HTM>
 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	icf_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md		
162	R00Y_025_025a	0.25 0.0 0.0	0.25 0.25 0.125	390	0.25 0.0 0.0	29.6 17.7 11.2	20.9 32.3	0.25 0.0 0.0	28.1 24.0 7.8	25.2 18.0 7.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
163	R00Y_025_025a	0.25 0.0 0.125	0.25 0.25 0.125	360	0.25 0.0 0.125	29.7 18.5 5.2	19.2 15.0	0.25 0.0 0.125	28.1 25.5 4.4	25.9 17.7 7.2	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
164	B50R_025_025a	0.25 0.0 0.25	0.25 0.25 0.125	330	0.25 0.0 0.25	29.8 19.8 0.0	19.8 359.8	0.25 0.0 0.25	28.3 27.3 -0.1	27.3 359.7 7.6	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
165	B34R_037_037a	0.25 0.0 0.375	0.375 0.375 0.187	310	0.256 0.0 0.375	30.1 25.5 -4.4	25.9 350.0	0.25 0.0 0.375	28.5 29.3 -4.3	29.6 351.6 4.1	310	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
166	B25R_050_050a	0.25 0.0 0.5	0.5 0.5 0.25	300	0.25 0.0 0.5	29.9 29.3 -10.3	31.0 340.5	0.25 0.0 0.5	28.5 30.6 -10.4	32.3 341.1 1.9	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
167	B19R_062_062a	0.25 0.0 0.625	0.625 0.625 0.312	293	0.239 0.0 0.625	29.7 32.7 -16.0	36.4 333.8	0.25 0.0 0.625	28.5 32.6 -17.0	36.8 332.3 1.5	292	0.383 0.0 1.0	32.9 52.3 -25.7	58.3 333.8
168	B15R_075_075a	0.25 0.0 0.75	0.75 0.75 0.375	289	0.237 0.0 0.75	29.3 35.5 -22.0	41.8 328.1	0.25 0.0 0.75	28.7 36.0 -23.1	42.8 327.3 1.3	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
169	B13R_087_087a	0.25 0.0 0.875	0.875 0.875 0.437	286	0.233 0.0 0.875	28.7 37.9 -27.8	47.0 323.6	0.25 0.0 0.875	28.6 39.2 -28.1	48.2 324.3 1.3	284	0.266 0.0 1.0	29.4 43.3 -31.8	53.8 323.6
170	B11R_100_100a	0.25 0.0 1.0	1.0 1.0 0.5	284	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1	0.25 0.0 1.0	28.8 41.9 -32.5	53.1 322.1 0.9	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
171	R50Y_025_025a	0.25 0.125 0.0	0.25 0.25 0.125	60	0.25 0.125 0.0	34.5 7.2 17.1	18.6 67.1	0.25 0.125 0.0	31.0 16.0 11.8	19.9 36.4 10.8	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
172	R00Y_025_012a	0.25 0.125 0.125	0.25 0.125 0.187	390	0.25 0.124 0.124	35.9 8.8 5.6	10.4 32.3	0.25 0.125 0.125	31.0 16.8 8.0	18.6 25.3 9.7	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
173	B50R_025_012a	0.25 0.125 0.25	0.25 0.125 0.187	330	0.25 0.124 0.25	36.0 9.9 0.0	9.9 359.8	0.25 0.125 0.25	31.5 18.6 3.2	18.9 9.7 10.3	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
174	B25R_037_025a	0.25 0.125 0.375	0.375 0.25 0.25	300	0.25 0.124 0.375	36.0 14.6 -5.1	15.5 340.5	0.25 0.125 0.375	31.7 20.5 -2.2	20.6 353.7 7.9	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
175	B15R_050_037a	0.25 0.125 0.5	0.5 0.375 0.312	289	0.243 0.124 0.5	35.7 17.7 -11.0	20.9 328.1	0.25 0.125 0.5	31.9 22.3 -9.9	24.4 335.9 6.0	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
176	B11R_062_050a	0.25 0.125 0.625	0.625 0.5 0.375	284	0.241 0.125 0.625	35.4 20.6 -16.5	26.4 321.1	0.25 0.125 0.625	32.1 24.7 -16.1	29.5 326.9 5.3	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
177	B09R_075_062a	0.25 0.125 0.75	0.75 0.625 0.437	281	0.239 0.125 0.75	35.7 24.2 -21.7	32.5 318.2	0.25 0.125 0.75	32.4 28.2 -22.2	35.9 321.8 5.1	279	0.183 0.0 1.0	28.3 38.8 -34.7	52.1 318.2
178	B07R_087_057a	0.25 0.125 0.875	0.875 0.75 0.5	279	0.237 0.125 0.875	36.0 27.9 -26.8	38.7 316.2	0.25 0.125 0.875	32.8 30.9 -27.8	41.6 318.0 4.5	278	0.15 0.0 1.0	28.1 37.2 -35.7	51.6 316.2
179	B06R_100_087a	0.25 0.125 1.0	1.0 0.875 0.562	278	0.241 0.125 1.0	36.4 31.9 -31.6	44.9 315.2	0.25 0.125 1.0	33.1 33.9 -32.5	47.0 316.2 3.9	277	0.133 0.0 1.0	27.9 36.4 -36.2	51.3 315.2
180	Y00G_025_012a	0.25 0.25 0.0	0.25 0.25 0.125	90	0.25 0.25 0.0	40.2 -2.5 23.8	24.0 96.1	0.25 0.25 0.0	35.1 5.0 17.5	18.2 74.0 11.1	89	1.0 1.0 1.0	87.8 -10.2	95.4 96.0 96.1
181	Y00G_025_012a	0.25 0.25 0.125	0.25 0.125 0.187	90	0.25 0.25 0.124	41.2 -1.2 11.9	12.0 96.1	0.25 0.25 0.125	35.4 5.7 13.1	14.6 66.3 9.1	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
182	NW_025a	0.25 0.25 0.25	0.25 0.0 0.25	360	0.25 0.25 0.25	42.1 0.0 0.0	0.0 0.0	0.0 0.25 0.25	35.7 7.5 7.1	10.4 43.4 12.2	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
183	B00R_037_012a	0.25 0.25 0.375	0.375 0.125 0.312	270	0.249 0.249 0.375	42.2 3.6 -5.0	6.2 306.2	0.25 0.25 0.375	36.3 9.3 0.7	9.4 4.5 10.0	270	1.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
184	B00R_050_025a	0.25 0.25 0.5	0.5 0.25 0.375	270	0.249 0.249 0.5	42.3 7.3 -10.1	12.5 306.2	0.25 0.25 0.5	36.4 12.3 -6.8	14.1 330.9 8.3	270	1.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
185	B00R_062_037a	0.25 0.25 0.625	0.625 0.375 0.437	270	0.245 0.25 0.625	42.4 11.0 -15.1	18.7 306.2	0.25 0.25 0.625	37.0 15.3 -14.0	20.9 318.0 7.1	270	1.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
186	B00R_075_050a	0.25 0.25 0.75	0.75 0.5 0.5	270	0.25 0.25 0.75	42.5 14.7 -20.2	25.0 306.2	0.25 0.25 0.75	37.5 18.9 -20.4	27.9 312.8 6.5	270	1.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
187	B00R_087_062a	0.25 0.25 0.875	0.875 0.625 0.562	270	0.25 0.25 0.875	42.6 18.4 -25.2	31.3 306.2	0.25 0.25 0.875	38.0 22.0 -26.5	34.4 309.7 5.9	270	1.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
188	B00R_100_075a	0.25 0.25 1.0	1.0 0.75 0.625	270	0.25 0.25 1.0	42.7 -22.1 -30.3	37.5 306.2	0.25 0.25 1.0	38.2 25.3 -31.6	40.5 308.7 5.6	270	1.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
189	Y31G_037_037a	0.25 0.375 0.0	0.375 0.375 0.187	109	0.256 0.375 0.0	44.4 -7.9 29.8	30.8 104.9	0.25 0.375 0.0	39.4 -4.2 23.2	23.6 100.4 9.0	108	0.683 1.0 0.0	77.8 -21.1	79.4 82.2 104.9
190	Y50G_037_025a	0.25 0.375 0.125	0.375 0.25 0.25	120	0.25 0.375 0.124	44.8 -7.4 16.6	18.2 114.0	0.25 0.375 0.125	39.4 -3.5 17.7	18.1 101.1 6.7	119	0.5 1.0 0.0	70.6 -29.7	66.5 72.8 114.0
191	G00B_037_012a	0.25 0.375 0.25	0.375 0.125 0.312	150	0.249 0.375 0.249	45.4 -8.1 3.7	8.9 155.5	0.25 0.375 0.25	40.0 -1.6 10.9	11.0 98.5 11.0	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
192	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	46.2 -3.1 -5.1	6.0 238.4	0.25 0.375 0.375	40.5 0.2 4.0	4.0 87.0 11.3	210	0.0 1.0 1.0	56.8 -25.5	-41.5 48.7 238.4
193	G75B_050_025a	0.25 0.375 0.5	0.5 0.25 0.375	240	0.249 0.375 0.5	46.5 -0.3 -10.1	10.1 268.2	0.25 0.375 0.5	40.8 3.5 -4.5	5.8 307.8 8.8	240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6 268.2
194	G84B_062_037a	0.25 0.375 0.625	0.625 0.375 0.437	251	0.25 0.368 0.625	46.2 3.7 -10.1	15.6 283.7	0.25 0.375 0.625	40.7 7.2 -11.9	13.9 301.0 7.2	251	0.0 0.316 1.0	35.2 9.9	-40.4 41.6 283.7
195	G88B_075_050a	0.25 0.375 0.75	0.75 0.5 0.5	256	0.25 0.366 0.75	46.1 7.6 -20.1	21.5 290.8	0.25 0.375 0.75	41.7 10.7 -19.4	22.2 299.0 5.4	257	0.0 0.233 1.0	32.2 15.3	-40.3 43.1 290.8
196	G90B_087_062a	0.25 0.375 0.875	0.875 0.625 0.562	259	0.25 0.364 0.875	46.0 11.6 -25.2	27.8 294.6	0.25 0.375 0.875	41.7 14.8 -26.1	30.1 299.6 5.5	260	0.0 0.183 1.0	30.6 18.5	-40.4 44.5 294.6
197	G92B_100_075a	0.25 0.375 1.0	1.0 0.75 0.625	261	0.25 0.362 1.0	46.0 15.5 -30.3	34.0 297.1	0.25 0.375 1.0	42.1 18.3 -31.8	36.7 299.9 5.0	262	0.0 0.15 1.0	29.5 20.7	-40.4 45.4 297.1
198	Y50G_050_050a	0.25 0.5 0.0	0.5 0.25 0.125	120	0.25 0.5 0.0	47.4 -14.8 33.2	36.4 114.0	0.25 0.5 0.0	43.1 -14.1	28.4 31.7	119	0.5 1.0 0.0	70.6 -29.7	66.5 72.8 114.0
199	Y68G_050_037a	0.25 0.5 0.125	0.5 0.375 0.312	131	0.243 0.5 0.124	47.5 -15.5 19.9	25.3 127.8	0.25 0.5 0.125	43.2 -13.5 21.7	25.6 121.9 4.9	141	0.316 1.0 0.0	62.3 -41.4	53.2 67.5 127.8
200	G00B_050_025a	0.25 0.5 0.25	0.5 0.25 0.375	150	0.249 0.5 0.249	48.6 -16.2 7.4	17.8 155.5	0.25 0.5 0.25	44.2 -10.9 14.5	18.2 126.7 9.9	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
201	G25B_050_025a	0.25 0.5 0.375	0.5 0.25 0.375	180	0.249 0.5 0.375	49.3 -12.1 -2.0	12.3 189.3	0.25 0.5 0.375	44.9 -8.5 6.8	10.9 141.0 10.5	180	0.0 1.0 0.5	52.9 -48.6	-8.0 49.3 189.3
202	G50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	50.2 -6.3 -10.3	12.1 238.4	0.25 0.5 0.5	45.1 -5.3 -2.5	5.9 205.3 9.3	210	0.0 1.0 1.0	56.8 -25.5	-41.5 48.7 238.4
203	G63B_062_037a	0.25 0.5 0.625	0.625 0.375 0.437	229	0.25 0.506 0.625	51.1 -4.6 -15.4	16.0 253.3	0.25 0.5 0.625	46.3 -2.7 -10.5	10.8 255.5 7.1	228	0.0 0.683 1.0	48.3 -12.2	-41.1 42.9 253.3
204	G75B_075_050a	0.25 0.5 0.75	0.75 0.5 0.5	240	0.25 0.5 0.75	50.8 -0.6 -20.3	20.3 268.2	0.25 0.5 0.75	46.6 1.2 -18.6	18.6 273.6 4.8	240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6 268.2
205	G80B_087_062a	0.25 0.5 0.875	0.875 0.625 0.562	247	0.25 0.489 0.875	50.4 3.5 -20.1	25.4 277.9	0.25 0.5 0.875	46.6 6.3 -25.4	26.2 283.9 4.7	247	0.0 0.383 1.0	37.6 5.6	-40.3 40.7 277.9
206	G84B_100_075a	0.25 0.5 1.0	1.0 0.75 0.625	251	0.25 0.487 1.0	50.3 7.4 -35.3	31.2 283.7	0.25 0.5 1.0	47.4 9.7 -31.7	33.1 287.1 3.9	251	0.0 0.316 1.0	35.2 9.9	-40.4 41.6 283.7
207	Y61G_062_062a	0.25 0.625 0.0	0.625 0.625 0.312	127	0.239 0.625 0.0	50.4 -22.0 36.7	42.8 120.9	0.25 0.625 0.0	47.5 -24.9 33.3	41.6 126.7 5.2	127	0.383 1.0 0.0	66.0 -35.2	58.8 68.6 120.9
208	Y76G_062_050a	0.25 0.625 0.125	0.625 0.5 0.375	136	0.241 0.625 0.125	50.0 -24.1 22.9	33.2 136.5	0.25 0.625 0.125	47.9 -23.7 26.3	35.4 132.0 4.0	137	0.233 1.0 0.0	57.9 -48.3	45.8 66.5 136.5
209	G00B_062_037a	0.25 0.625 0.25	0.625 0.375 0.437	150	0.25 0.625 0.25	51.8 -24.3 11.1	16.7 155.5	0.25 0.625 0.25	48.7 -21.1 18.4	28.0 138.9 8.5	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
210	G15B_062_037a	0.25 0.625 0.375	0.625 0.375 0.437	169	0.25 0.625 0.368	51.4 -21.3 2.7	21.4 172.5	0.25 0.625 0.375	49.6 -18.5 9.9					

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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		
243	R00Y_037_037a	0.375 0.0 0.0	0.375 0.375 0.187	390	0.375 0.0 0.0	32.2 26.6 16.8	31.4 32.3	0.375 0.0 0.0	31.7 36.2 17.7	40.3 26.1 9.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
244	R18Y_037_037a	0.375 0.0 0.125	0.375 0.375 0.187	371	0.375 0.0 0.118	32.3 27.2 11.7	29.6 23.2	0.375 0.0 0.125	31.6 36.7 13.2	39.0 19.8 9.6	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
245	B65R_037_037a	0.375 0.0 0.25	0.375 0.375 0.187	349	0.375 0.0 0.256	32.4 28.6 4.4	29.0 8.9	0.375 0.0 0.25	31.7 38.5 8.1	39.3 11.9 10.5	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
246	B50R_037_037a	0.375 0.0 0.375	0.375 0.375 0.187	330	0.375 0.0 0.375	32.5 29.7 0.0	29.7 35.9	0.375 0.0 0.375	31.7 39.8 3.0	39.9 4.3 10.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
247	B38R_050_050a	0.375 0.0 0.5	0.5 0.5 0.25	316	0.383 0.0 0.5	33.2 35.8 -4.3	36.0 353.0	0.375 0.0 0.5	32.2 42.9 -3.3	43.0 355.5 7.3	317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0
248	B30R_062_062a	0.375 0.0 0.625	0.625 0.625 0.312	307	0.385 0.0 0.625	32.8 40.6 -9.0	41.6 347.4	0.375 0.0 0.625	32.4 45.1 -9.5	46.1 348.0 4.5	307	0.616 0.0 1.0	37.9 65.0 -14.5	66.6 347.4
249	B25R_075_075a	0.375 0.0 0.75	0.75 0.75 0.375	300	0.375 0.0 0.75	32.7 43.9 -15.5	46.6 340.5	0.375 0.0 0.75	32.5 47.1 -15.8	49.6 344.4 3.1	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
250	B20R_087_087a	0.375 0.0 0.875	0.875 0.875 0.437	295	0.364 0.0 0.875	32.5 47.4 -21.5	51.9 335.7	0.375 0.0 0.875	32.6 49.3 -21.4	53.8 336.5 1.9	294	0.416 0.0 1.0	33.7 54.1 -24.4	59.4 335.7
251	B18R_100_100a	0.375 0.0 1.0	1.0 1.0 0.5	292	0.366 0.0 1.0	32.5 51.2 -26.5	57.7 332.6	0.375 0.0 1.0	32.7 51.8 -26.0	58.0 333.3 0.8	291	0.366 0.0 1.0	32.5 51.2 -26.5	57.7 332.6
252	R31Y_037_037a	0.375 0.125 0.0	0.375 0.375 0.187	49	0.375 0.118 0.0	36.4 17.1 22.2	28.1 52.2	0.375 0.125 0.0	34.8 28.0 21.3	35.2 37.3 10.9	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
253	R00Y_037_025a	0.375 0.125 0.125	0.375 0.25 0.25	390	0.375 0.124 0.124	38.5 17.7 11.2	20.9 32.3	0.375 0.125 0.125	35.3 28.3 16.7	32.9 30.6 12.4	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
254	R00Y_037_025a	0.375 0.125 0.25	0.375 0.25 0.25	360	0.375 0.124 0.25	38.6 18.5 5.2	19.2 15.9	0.375 0.125 0.25	35.3 29.6 10.7	31.5 19.8 12.7	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
255	B50R_037_025a	0.375 0.125 0.375	0.375 0.25 0.25	330	0.375 0.124 0.375	38.7 19.8 0.0	19.8 359.8	0.375 0.125 0.375	35.5 31.2 5.0	31.6 9.2 12.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
256	B34R_050_037a	0.375 0.125 0.5	0.5 0.375 0.312	311	0.381 0.124 0.5	39.0 25.5 -4.4	25.9 350.0	0.375 0.125 0.5	36.2 33.7 -2.3	33.7 355.9 8.9	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
257	B25R_062_050a	0.375 0.125 0.625	0.625 0.5 0.375	300	0.375 0.125 0.625	38.8 29.3 -10.3	31.0 340.5	0.375 0.125 0.625	36.2 35.2 -9.0	36.3 345.6 6.6	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
258	B19R_075_062a	0.375 0.125 0.75	0.75 0.625 0.437	293	0.364 0.125 0.75	38.6 32.7 -16.0	36.4 333.8	0.375 0.125 0.75	36.6 37.1 -15.7	40.3 337.0 4.8	292	0.383 0.0 1.0	32.9 52.3 -25.7	58.3 333.8
259	B15R_087_075a	0.375 0.125 0.875	0.875 0.75 0.5	289	0.362 0.125 0.875	38.2 35.5 -22.0	41.8 328.1	0.375 0.125 0.875	36.9 39.8 -21.4	45.2 331.6 4.5	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
260	B13R_100_087a	0.375 0.125 1.0	1.0 0.875 0.562	286	0.358 0.125 1.0	37.6 37.9 -27.8	47.0 323.6	0.375 0.125 1.0	36.8 42.2 -26.6	49.9 327.7 4.5	284	0.266 0.0 1.0	29.4 43.3 -31.8	53.8 323.6
261	R68Y_037_037a	0.375 0.25 0.0	0.375 0.375 0.187	71	0.375 0.256 0.0	43.2 4.1 30.1	30.4 82.1	0.375 0.25 0.0	39.9 16.0 27.6	31.9 59.7 12.6	71	1.0 0.683 0.0	74.8 11.0 80.4	81.1 82.1
262	R50Y_037_025a	0.375 0.25 0.125	0.375 0.25 0.25	60	0.375 0.25 0.124	43.4 7.2 17.1	18.6 67.1	0.375 0.25 0.125	39.9 17.1 21.7	27.7 51.6 11.5	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
263	R00Y_037_012a	0.375 0.25 0.25	0.375 0.125 0.312	390	0.375 0.249 0.249	44.8 8.8 5.6	10.4 32.3	0.375 0.25 0.25	40.0 18.4 15.1	23.9 39.3 14.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
264	B50R_037_012a	0.375 0.25 0.375	0.375 0.125 0.312	330	0.375 0.249 0.375	44.9 9.9 0.0	9.9 359.8	0.375 0.25 0.375	40.7 19.7 8.1	21.3 22.2 13.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
265	B25R_050_025a	0.375 0.25 0.5	0.5 0.25 0.375	300	0.375 0.249 0.5	44.9 14.6 -5.1	15.5 340.5	0.375 0.25 0.5	41.2 22.1 -0.1	22.1 359.9 9.7	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
266	B15R_062_037a	0.375 0.25 0.625	0.625 0.375 0.437	289	0.368 0.25 0.625	44.6 17.7 -11.0	20.9 328.1	0.375 0.25 0.625	41.6 23.9 -7.1	25.0 343.2 7.9	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
267	B11R_075_050a	0.375 0.25 0.75	0.75 0.5 0.5	284	0.366 0.25 0.75	44.3 20.6 -16.5	26.4 321.1	0.375 0.25 0.75	42.1 26.2 -14.0	29.7 331.7 6.5	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
268	B09R_087_062a	0.375 0.25 0.875	0.875 0.625 0.562	281	0.364 0.25 0.875	44.6 24.2 -21.7	32.5 318.2	0.375 0.25 0.875	42.9 28.9 -20.3	35.3 324.8 5.1	279	0.183 0.0 1.0	27.9 38.8 -34.7	52.1 318.2
269	B07R_100_075a	0.375 0.25 1.0	1.0 0.75 0.625	279	0.362 0.25 1.0	44.9 27.9 -24.8	38.7 312.2	0.375 0.25 1.0	43.1 31.3 -26.0	40.7 320.3 3.9	278	0.15 0.0 1.0	28.1 37.2 -35.7	51.6 316.2
270	Y00G_037_037a	0.375 0.375 0.0	0.375 0.375 0.187	90	0.375 0.375 0.0	48.1 -3.8 35.8	36.0 96.1	0.375 0.375 0.0	44.1 6.7 33.2	33.8 78.5 11.5	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
271	Y00G_037_025a	0.375 0.375 0.125	0.375 0.25 0.25	90	0.375 0.375 0.124	49.1 -2.5 38.8	24.0 96.1	0.375 0.375 0.125	44.5 7.0 26.3	27.2 75.0 10.9	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
272	Y00G_037_012a	0.375 0.375 0.25	0.375 0.125 0.312	90	0.375 0.375 0.249	50.1 -1.2 11.9	12.0 96.1	0.375 0.375 0.25	44.7 8.5 18.5	20.4 65.3 12.9	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
273	NW_037a	0.375 0.375 0.375	0.375 0.0 0.375	360	0.375 0.375 0.375	51.0 0.0 0.0	0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
274	B00R_050_012a	0.375 0.375 0.5	0.5 0.125 0.437	270	0.375 0.375 0.5	51.1 3.6 -5.0	6.2 306.2	0.375 0.375 0.5	46.1 12.2 2.1	12.3 10.0 12.2	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
275	B00R_062_025a	0.375 0.375 0.625	0.625 0.25 0.5	270	0.375 0.375 0.625	51.2 7.3 -10.1	12.5 306.2	0.375 0.375 0.625	46.7 14.8 -5.3	15.7 340.2 9.9	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
276	B00R_075_037a	0.375 0.375 0.75	0.75 0.375 0.562	270	0.375 0.375 0.75	51.3 11.0 -15.1	18.7 306.2	0.375 0.375 0.75	47.4 17.2 -12.5	21.3 323.8 7.7	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
277	B00R_087_050a	0.375 0.375 0.875	0.875 0.5 0.625	270	0.375 0.375 0.875	51.4 14.7 -20.2	25.0 306.2	0.375 0.375 0.875	48.1 19.9 -19.3	27.7 315.9 6.1	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
278	B00R_100_062a	0.375 0.375 1.0	1.0 0.625 0.687	270	0.375 0.375 1.0	51.5 18.4 -25.2	31.3 306.2	0.375 0.375 1.0	48.4 23.0 -25.3	34.2 312.3 5.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
279	Y23G_050_050a	0.375 0.5 0.0	0.5 0.5 0.25	104	0.383 0.5 0.0	52.8 8.5 42.1	43.0 101.4	0.375 0.5 0.0	49.1 -2.0 38.9	38.9 92.9 8.1	102	0.766 1.0 0.0	81.2 -17.0	84.3 86.0 101.4
280	Y31G_050_037a	0.375 0.5 0.125	0.5 0.375 0.312	109	0.381 0.5 0.124	53.3 -7.9 29.8	30.8 104.9	0.375 0.5 0.125	49.5 -1.7 31.0	31.0 93.2 7.3	108	0.683 1.0 0.0	77.8 -21.1	79.4 82.2 104.9
281	Y50G_050_025a	0.375 0.5 0.25	0.5 0.25 0.375	120	0.375 0.5 0.249	53.7 -7.4 16.6	18.2 114.0	0.375 0.5 0.25	49.7 -0.9 22.3	22.3 92.5 9.4	119	0.5 1.0 0.0	70.6 -29.7	66.5 72.8 114.0
282	G00B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.5 0.375	54.3 -8.1 3.7	8.9 155.5	0.375 0.5 0.375	50.4 0.8 13.6	13.6 86.3 13.9	149	0.0 1.0 1.0	50.0 -65.0	29.6 71.4 155.5
283	G50B_050_012a	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.5	55.1 -3.1 -5.1	6.0 238.4	0.375 0.5 0.5	51.1 2.9 4.1	5.0 54.4 11.8	210	0.0 1.0 1.0	56.8 -25.5	-41.5 48.7 238.4
284	G75B_062_025a	0.375 0.5 0.625	0.625 0.25 0.5	240	0.375 0.5 0.625	55.4 -3.3 -10.1	10.1 268.2	0.375 0.5 0.625	51.7 5.8 -4.1	7.1 324.8 9.3	240	0.0 0.5 1.0	41.7 -1.2	-40.6 40.6 268.2
285	G84B_075_037a	0.375 0.5 0.75	0.75 0.375 0.562	251	0.375 0.493 0.75	55.1 3.7 -15.1	15.6 283.7	0.375 0.5 0.75	52.4 8.7 -11.9	14.7 306.3 6.5	251	0.0 0.316 1.0	35.2 39.9	-40.4 41.6 283.7
286	G88B_087_050a	0.375 0.5 0.875	0.875 0.5 0.625	256	0.375 0.491 0.875	55.0 7.6 -20.1	21.5 290.8	0.375 0.5 0.875	52.9 12.1 -18.6	22.2 303.1 5.1	257	0.0 0.233 1.0	32.2 15.3	-40.3 43.1 290.8
287	G90B_100_062a	0.375 0.5 1.0	1.0 0.625 0.687	259	0.375 0.489 1.0	54.9 11.6 -25.2	27.8 294.6	0.375 0.5 1.0	53.6 15.1 -25.2	29.4 301.0 3.8	260	0.0 0.183 1.0	30.6 18.5	-40.4 44.5 294.6
288	Y38G_062_062a	0.375 0.625 0.0	0.625 0.625 0.312	113	0.385 0.625 0.0	56.0 -15.3 46.9	49.4 108.0	0.375 0.625 0.0	54.2 -12.9 44.7	46.5 106.1 3.7	112	0.616 1.0 0.0	75.0 -24.4	75.1 79.0 108.0
289	Y50G_062_050a	0.375 0.625 0.125	0.625 0.5 0.375	120	0.375 0.625 0.125	56.4 -14.8 33.2	36.4 114.0	0.375 0.625 0.125	54.5 -12.7 36.0	38.2 109.5 3.9	119	0.5 1.0 0.0	70.6 -29.7	66.5 72.8 114.0
290	Y68G_062_037a	0.375 0.625 0.25	0.625 0.375 0.437	131	0.368 0.625 0.25	56.4 -15.5 19.9	25.3 127.8	0.375 0.625 0.25	54.9 -11.6 26.1	28.6 114.0 7.3	131	0.316 1.0 0.0	62.3 -41.4	53.2 67.5 127.8
291	G00B_062_025a	0.375 0.625 0.												

n	HIC*Fd	rgb_Fd	icf_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md		
324	R00Y_050_050a	0.5 0.0 0.0	0.5 0.5 0.25	390	0.5 0.0 0.0	34.9 35.4 22.4	41.9 32.3	0.5 0.0 0.0	34.8 44.7 22.4	50.0 26.6 9.2	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
325	R26Y_050_050a	0.5 0.0 0.125	0.5 0.5 0.25	376	0.5 0.0 0.116	35.0 36.0 17.6	40.1 26.1	0.5 0.0 0.125	34.7 45.7 18.0	49.1 21.5 9.6	377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1
326	R00Y_050_050a	0.5 0.0 0.25	0.5 0.5 0.25	360	0.5 0.0 0.25	35.1 37.1 10.5	38.5 15.9	0.5 0.0 0.25	34.8 46.7 12.4	48.3 14.9 9.7	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
327	B61R_050_050a	0.5 0.0 0.375	0.5 0.5 0.25	344	0.5 0.0 0.383	35.1 38.6 4.0	38.8 5.9	0.5 0.0 0.375	34.8 48.4 6.7	48.9 7.8 10.1	342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
328	B50R_050_050a	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	35.2 39.6 -0.1	39.6 35.9	0.5 0.0 0.5	35.0 49.8 0.6	49.8 0.7 10.2	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 35.9
329	B40R_062_062a	0.5 0.0 0.625	0.625 0.625 0.312	319	0.51 0.0 0.625	36.0 45.8 -4.4	46.0 35.4	0.5 0.0 0.625	35.3 52.5 -4.7	52.7 35.4 6.7	320	0.816 0.0 1.0	43.1 73.2 -7.0	73.6 35.4
330	B34R_075_075a	0.5 0.0 0.75	0.75 0.75 0.375	311	0.512 0.0 0.75	35.9 51.0 -8.9	51.8 35.0	0.5 0.0 0.75	35.7 54.4 -10.3	55.4 34.2 3.6	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 35.0
331	B29R_087_087a	0.5 0.0 0.875	0.875 0.875 0.437	305	0.51 0.0 0.875	35.6 55.3 -14.3	57.1 345.4	0.5 0.0 0.875	35.8 56.7 -15.7	58.8 34.4 1.9	305	0.583 0.0 1.0	37.2 63.2 -16.9	65.3 345.4
332	B25R_100_100a	0.5 0.0 1.0	1.0 1.0 0.5	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5 0.0	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
333	R23Y_050_050a	0.5 0.125 0.0	0.5 0.5 0.25	44	0.5 0.116 0.0	38.7 26.7 27.4	38.2 45.7	0.5 0.125 0.0	38.2 36.5 26.8	45.3 36.2 9.9	42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7
334	R00Y_050_037a	0.5 0.125 0.125	0.5 0.375 0.312	390	0.5 0.124 0.124	41.1 26.6 16.8	31.4 32.3	0.5 0.125 0.125	38.6 36.6 21.7	42.6 30.7 11.4	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
335	R18Y_050_037a	0.5 0.125 0.25	0.5 0.375 0.312	371	0.5 0.124 0.243	41.2 27.2 11.7	29.6 23.2	0.5 0.125 0.25	38.5 37.3 15.9	40.6 23.1 11.3	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
336	B63R_050_037a	0.5 0.125 0.375	0.5 0.375 0.312	349	0.5 0.124 0.381	41.3 28.6 4.4	29.0 8.9	0.5 0.125 0.375	38.8 39.2 8.8	40.2 12.6 11.7	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
337	B50R_050_037a	0.5 0.125 0.5	0.5 0.375 0.312	330	0.5 0.124 0.5	41.4 29.7 0.0	29.7 359.8	0.5 0.125 0.5	39.3 40.7 1.9	40.8 2.7 11.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
338	B38R_062_050a	0.5 0.125 0.625	0.625 0.5 0.375	316	0.508 0.125 0.625	42.1 35.8 -4.3	36.0 353.0	0.5 0.125 0.625	39.5 42.6 -4.1	42.8 35.4 7.3	317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0
339	B30R_075_062a	0.5 0.125 0.75	0.75 0.625 0.437	307	0.51 0.125 0.75	41.7 40.6 -9.0	41.6 347.4	0.5 0.125 0.75	40.4 44.7 -10.1	45.8 34.1 4.4	307	0.616 0.0 1.0	37.9 65.0 -14.5	66.6 347.4
340	B25R_087_075a	0.5 0.125 0.875	0.875 0.75 0.5	300	0.5 0.125 0.875	41.7 43.9 -15.5	46.6 340.5	0.5 0.125 0.875	40.2 46.8 -16.1	49.5 34.0 3.3	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
341	B20R_100_087a	0.5 0.125 1.0	1.0 0.875 0.562	295	0.489 0.125 1.0	41.4 47.4 -21.3	51.9 335.7	0.5 0.125 1.0	40.3 48.4 -21.7	53.0 335.8 1.5	294	0.416 0.0 1.0	33.7 54.1 -24.4	59.4 335.7
342	R50Y_050_050a	0.5 0.25 0.0	0.5 0.5 0.25	60	0.5 0.25 0.0	44.6 14.4 34.3	37.2 67.1	0.5 0.25 0.0	43.4 24.2 33.3	41.2 53.9 9.9	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
343	R31Y_050_037a	0.5 0.25 0.125	0.5 0.375 0.312	49	0.5 0.243 0.124	45.3 17.1 22.2	28.1 52.2	0.5 0.25 0.125	43.4 25.3 26.7	36.8 46.5 9.5	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
344	R00Y_050_025a	0.5 0.25 0.25	0.5 0.25 0.375	390	0.5 0.249 0.249	47.4 17.7 11.2	20.9 32.3	0.5 0.25 0.25	44.0 25.7 19.7	32.4 37.4 12.1	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
345	R00Y_050_025a	0.5 0.25 0.375	0.5 0.25 0.375	360	0.5 0.249 0.375	47.5 18.5 5.2	19.2 15.9	0.5 0.25 0.375	44.3 27.0 12.6	29.8 25.1 11.6	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
346	B50R_050_025a	0.5 0.25 0.5	0.5 0.25 0.375	330	0.5 0.249 0.5	47.6 19.8 0.0	19.8 359.8	0.5 0.25 0.5	44.8 28.7 4.6	29.0 9.2 10.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
347	B34R_062_037a	0.5 0.25 0.625	0.625 0.375 0.437	311	0.506 0.25 0.625	47.9 25.5 -4.4	25.9 350.0	0.5 0.25 0.625	45.5 30.6 -2.0	30.7 35.6 6.1	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
348	B25R_075_050a	0.5 0.25 0.75	0.75 0.5 0.375	300	0.5 0.25 0.75	47.8 29.3 -10.3	31.0 340.5	0.5 0.25 0.75	45.9 32.2 -9.6	33.6 34.3 3.5	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
349	B19R_087_062a	0.5 0.25 0.875	0.875 0.625 0.293	293	0.489 0.25 0.875	47.5 32.7 -16.0	36.4 333.8	0.5 0.25 0.875	46.1 34.4 -15.8	37.9 33.2 2.2	292	0.383 0.0 1.0	32.9 52.3 -25.7	58.3 333.8
350	B15R_100_075a	0.5 0.25 1.0	1.0 0.75 0.625	289	0.487 0.25 1.0	47.1 35.3 -22.0	41.8 328.1	0.5 0.25 1.0	46.6 36.7 -21.3	42.4 29.8 1.5	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
351	R76Y_050_050a	0.5 0.375 0.0	0.5 0.5 0.25	76	0.5 0.383 0.0	51.5 21.4 42.3	42.4 87.0	0.5 0.375 0.0	48.2 12.8 39.3	41.4 71.8 11.5	77	1.0 0.766 0.0	78.6 4.3 84.7	84.8 87.0
352	R68Y_050_037a	0.5 0.375 0.125	0.5 0.375 0.312	71	0.5 0.381 0.124	52.2 4.1 30.1	30.4 82.1	0.5 0.375 0.125	48.7 13.5 32.0	34.7 67.1 10.1	71	1.0 0.683 0.0	78.8 11.0 80.4	81.1 82.1
353	R50Y_050_025a	0.5 0.375 0.25	0.5 0.25 0.375	60	0.5 0.375 0.249	52.3 7.2 17.1	18.6 67.1	0.5 0.375 0.25	48.7 15.3 23.6	28.1 56.9 10.9	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
354	R00Y_050_012a	0.5 0.375 0.375	0.5 0.125 0.437	390	0.5 0.375 0.375	53.7 8.8 5.6	10.4 32.3	0.5 0.375 0.375	49.3 16.6 15.4	22.7 42.7 13.2	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
355	B50R_050_012a	0.5 0.375 0.5	0.5 0.125 0.437	330	0.5 0.375 0.5	53.8 9.9 0.0	9.9 359.8	0.5 0.375 0.5	50.0 18.1 6.9	19.4 21.0 11.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
356	B25R_062_025a	0.5 0.375 0.625	0.625 0.25 0.5	300	0.5 0.375 0.625	53.9 14.6 -5.1	15.5 340.5	0.5 0.375 0.625	50.6 20.3 -0.7	20.3 35.7 7.9	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
357	B15R_075_037a	0.5 0.375 0.75	0.75 0.375 0.562	289	0.493 0.375 0.75	53.5 17.7 -11.0	20.9 328.1	0.5 0.375 0.75	51.3 22.1 -8.5	23.7 33.8 5.5	288	0.316 0.0 1.0	30.9 47.3 -29.4	55.7 328.1
358	B11R_087_050a	0.5 0.375 0.875	0.875 0.5 0.625	284	0.491 0.375 0.875	53.2 20.6 -16.5	26.4 321.1	0.5 0.375 0.875	51.7 24.3 -15.1	28.6 32.8 4.2	282	0.233 0.0 1.0	28.7 41.2 -33.1	52.9 321.1
359	B09R_100_062a	0.5 0.375 1.0	1.0 0.625 0.687	281	0.489 0.375 1.0	53.5 24.2 -21.7	32.5 318.2	0.5 0.375 1.0	52.1 26.7 -21.3	24.2 32.1 2.8	279	0.183 0.0 1.0	28.3 38.8 -34.7	52.1 318.2
360	Y00G_050_050a	0.5 0.5 0.0	0.5 0.5 0.25	90	0.5 0.5 0.0	56.1 -5.1 47.7	48.0 96.1	0.5 0.5 0.0	52.6 3.9 44.2	44.3 84.8 10.3	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
361	Y00G_050_037a	0.5 0.5 0.125	0.5 0.375 0.312	90	0.5 0.5 0.124	57.0 -3.8 35.8	36.0 96.1	0.5 0.5 0.125	53.0 4.5 36.2	36.5 82.8 9.3	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
362	Y00G_050_025a	0.5 0.5 0.25	0.5 0.25 0.375	90	0.5 0.5 0.249	58.0 -2.5 23.8	24.0 96.1	0.5 0.5 0.25	53.6 5.7 27.6	28.2 78.1 10.1	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
363	Y00G_050_012a	0.5 0.5 0.375	0.5 0.125 0.437	90	0.5 0.5 0.375	59.0 -1.2 11.9	12.0 96.1	0.5 0.5 0.375	54.5 6.9 19.0	20.2 69.9 11.7	89	1.0 1.0 0.0	87.8 -10.2	95.4 96.0 96.1
364	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	60.0 0.0 0.0	0.0 0.0	0.5 0.5 0.5	55.1 8.8 9.3	12.8 46.5 13.7	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
365	B00R_062_012a	0.5 0.5 0.625	0.625 0.125 0.562	270	0.5 0.5 0.625	60.0 3.6 -5.0	6.2 306.2	0.5 0.5 0.625	55.7 11.2 0.8	11.2 4.5 10.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
366	B00R_075_025a	0.5 0.5 0.75	0.75 0.25 0.625	270	0.5 0.5 0.75	60.1 7.3 -10.1	12.5 306.2	0.5 0.5 0.75	56.5 13.3 -7.1	15.1 33.1 7.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
367	B00R_087_037a	0.5 0.5 0.875	0.875 0.375 0.687	270	0.5 0.5 0.875	60.2 11.0 -15.1	18.7 306.2	0.5 0.5 0.875	57.2 15.8 -14.2	21.3 31.1 5.7	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
368	B00R_100_050a	0.5 0.5 1.0	1.0 0.5 0.75	270	0.5 0.5 1.0	60.3 14.7 -20.2	25.0 306.2	0.5 0.5 1.0	57.9 18.3 -20.7	27.7 31.4 4.3	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2
369	Y18G_062_062a	0.5 0.625 0.0	0.625 0.625 0.312	101	0.51 0.625 0.0	60.8 -9.7 54.1	55.0 100.2	0.5 0.625 0.0	58.2 -6.1 51.8	52.1 96.8 5.0	99	0.816 1.0 0.0	82.6 -15.6	86.6 88.0 100.2
370	Y23G_062_050a	0.5 0.625 0.125	0.625 0.5 0.375	104	0.508 0.625 0.125	61.7 -8.5 42.1	43.0 101.4	0.5 0.625 0.125	58.8 -5.8 42.5	42.9 97.8 3.9	102	0.766 1.0 0.0	81.2 -17.0	84.3 86.0 101.4
371	Y31G_062_037a	0.5 0.625 0.25	0.625 0.375 0.437	109	0.506 0.625 0.25	62.2 -7.9 29.8	30.8 104.9	0.5 0.625 0.25	59.3 -4.8 32.3	32.7 98.5 4.9	108	0.683 1.0 0.0	77.8 -21.1	79.4 82.2 104.9
372	Y50G_062_025a	0.5 0.625 0.375	0.625 0.25 0.5	120	0.5 0.625 0.375	62.6 -7.4 16.6	18.2 114.0	0.5 0.625 0.375	59.7 -3.4 22.2	22.4 98.7 7.4	119	0.5 1.0 0.0	70.6 -29.7	66.5 72.8 114.0
373	G00B_062_012a	0.5 0.625 0.5	0.625 0.125 0.562	150	0.5 0.625 0.5	63.2 -8.1 3.7	8.9 155.5	0.5 0.625 0.5	60.6 -1.5 12.5	12.6 96.8 11.3	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
374	G50B_062_012a	0.5 0.625 0.625	0.625 0.125 0.562	210	0.5 0.625 0.625									

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF77/PF77.HTM>
 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	37.5 44.3 28.0	52.4 32.3	0.625 0.0 0.0	37.2 53.3 28.6	60.5 28.2 9.0	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3				
406	R31Y_062_062a	0.625 0.0 0.125	0.625 0.625 0.312	379	0.625 0.0 0.114	37.6 44.9 23.4	50.6 27.5	0.625 0.0 0.125	37.4 54.0 24.4	59.3 28.3 9.2	380	1.0 0.0 0.183	45.5 71.8 37.5	81.0 27.5				
407	R11Y_062_062a	0.625 0.0 0.25	0.625 0.625 0.312	367	0.625 0.0 0.239	37.7 45.6 17.4	48.8 20.8	0.625 0.0 0.25	37.3 54.8 19.5	58.2 19.6 9.4	367	1.0 0.0 0.383	45.8 73.0 27.8	78.2 20.8				
408	B69R_062_062a	0.625 0.0 0.375	0.625 0.625 0.312	353	0.625 0.0 0.385	37.8 47.2 9.5	48.1 11.4	0.625 0.0 0.375	37.4 56.1 13.0	57.6 13.0 9.5	352	1.0 0.0 0.616	46.0 75.5 15.2	77.1 11.4				
409	B59R_062_062a	0.625 0.0 0.5	0.625 0.625 0.312	341	0.625 0.0 0.51	37.8 48.6 3.9	48.7 4.6	0.625 0.0 0.5	37.4 57.9 6.5	58.2 6.4 9.6	339	1.0 0.0 0.816	45.9 77.7 6.2	78.0 4.6				
410	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	37.9 49.5	-0.1 49.5	0.625 0.0 0.625	37.4 59.3 1.1	59.3 1.0 9.8	330	1.0 0.0 1.0	46.1 79.3	-0.2 79.3	359.8			
411	B42R_075_075a	0.625 0.0 0.75	0.75 0.75 0.375	321	0.637 0.0 0.75	38.9 55.7	-4.4 55.9	0.625 0.0 0.75	37.9 61.6	-4.2 61.8	356.0 5.9	322	0.85 0.0 1.0	43.7 74.3	-5.9 74.6	355.4		
412	B36R_087_087a	0.625 0.0 0.875	0.875 0.875 0.437	314	0.641 0.0 0.875	39.2 61.5	-8.7 62.1	0.625 0.0 0.875	38.3 64.0	-9.1 64.6	351.8 2.6	315	0.733 0.0 1.0	41.3 70.3	-9.9 71.0	351.9		
413	B31R_100_100a	0.625 0.0 1.0	1.0 1.0 0.5	308	0.633 0.0 1.0	38.3 65.8	-13.7 67.2	0.625 0.0 1.0	38.1 65.4	-14.0 66.9	347.9 0.5	308	0.633 0.0 1.0	38.3 65.8	-13.7 67.2	348.2		
414	R18Y_062_062a	0.625 0.125 0.0	0.625 0.625 0.312	41	0.625 0.114 0.0	41.1 36.1	32.8 48.8	0.625 0.125 0.0	40.5 45.1	32.7 55.7	35.9 9.0	39	1.0 0.183 0.0	51.1 57.8	52.5 78.1	42.1		
415	R00Y_062_050a	0.625 0.125 0.125	0.625 0.5 0.375	390	0.625 0.125 0.125	43.8 35.4	22.4 41.9	0.625 0.125 0.125	41.0 44.9	28.0 53.0	31.9 11.3	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3		
416	R26Y_062_050a	0.625 0.125 0.25	0.625 0.5 0.375	376	0.625 0.125 0.241	43.9 36.0	17.6 40.1	0.625 0.125 0.25	41.0 45.8	22.3 51.0	25.9 11.2	377	1.0 0.0 0.233	45.6 72.1	35.3 80.3	26.1		
417	R00Y_062_050a	0.625 0.125 0.375	0.625 0.5 0.375	360	0.625 0.125 0.375	44.0 37.1	10.5 38.5	0.625 0.125 0.375	41.1 47.2	15.5 49.7	18.2 11.6	360	1.0 0.0 0.5	45.9 74.2	21.1 77.1	15.9		
418	B61R_062_050a	0.625 0.125 0.5	0.625 0.5 0.375	344	0.625 0.125 0.508	44.0 38.6	4.0 38.8	0.625 0.125 0.5	41.4 48.6	7.7 49.3	9.0 11.0	342	1.0 0.0 0.766	45.9 77.3	8.0 77.7	5.9		
419	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	44.1 39.6	-0.1 39.6	0.625 0.125 0.625	41.7 50.4	1.6 50.4	11.1 11.1	330	1.0 0.0 1.0	46.1 79.3	-0.2 79.3	359.8		
420	B40R_075_062a	0.625 0.125 0.75	0.75 0.625 0.437	319	0.635 0.125 0.75	44.9 45.8	-4.4 46.0	0.625 0.125 0.75	42.7 52.1	-4.3 52.3	355.2 6.7	320	0.816 0.0 1.0	43.1 73.2	-7.0 73.6	354.4		
421	B34R_087_075a	0.625 0.125 0.875	0.875 0.75 0.5	311	0.637 0.125 0.875	44.5 51.0	-8.9 51.8	0.625 0.125 0.875	42.7 54.6	-10.3 55.5	349.2 4.3	311	0.683 0.0 1.0	39.8 68.1	-11.9 69.1	350.0		
422	B29R_100_087a	0.625 0.125 1.0	1.0 0.875 0.562	305	0.635 0.125 1.0	44.5 55.3	-14.3 57.1	0.625 0.125 1.0	43.0 56.2	-15.1 58.2	344.9 1.9	305	0.583 0.0 1.0	37.2 63.2	-16.4 65.3	345.4		
423	R38Y_062_062a	0.625 0.25 0.0	0.625 0.625 0.312	53	0.625 0.239 0.0	46.3 24.7	39.1 46.2	0.625 0.25 0.0	45.1 34.1	38.7 51.6	48.5 9.5	52	1.0 0.383 0.0	59.5 39.5	62.5 74.0	57.6		
424	R23Y_062_050a	0.625 0.25 0.125	0.625 0.5 0.375	44	0.625 0.241 0.125	47.6 26.7	27.4 38.2	0.625 0.25 0.125	45.7 34.0	33.2 47.6	44.3 9.6	42	1.0 0.233 0.0	53.0 53.4	54.8 76.5	47.7		
425	R00Y_062_037a	0.625 0.25 0.25	0.625 0.375 0.437	390	0.625 0.25 0.25	50.1 26.6	16.8 31.4	0.625 0.25 0.25	46.1 34.0	26.2 43.0	37.6 12.6	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3		
426	R18Y_062_037a	0.625 0.25 0.375	0.625 0.375 0.437	371	0.625 0.25 0.368	50.2 27.2	11.7 29.6	0.625 0.25 0.375	46.5 35.2	19.1 40.1	28.4 11.4	371	1.0 0.0 0.316	45.7 72.6	31.2 79.1	22.2		
427	B65R_062_037a	0.625 0.25 0.5	0.625 0.375 0.437	349	0.625 0.25 0.506	50.2 28.6	4.4 29.0	0.625 0.25 0.5	46.9 37.0	10.1 38.4	15.3 10.6	348	1.0 0.0 0.683	45.9 76.4	11.9 77.3	8.9		
428	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	50.3 29.7	0.0 29.7	0.625 0.25 0.625	47.5 38.1 3.1	38.3 4.7	9.4 3.4	330	1.0 0.0 1.0	46.1 79.3	-0.2 79.3	359.8		
429	B38R_075_050a	0.625 0.25 0.75	0.75 0.5 0.5	316	0.633 0.25 0.75	51.0 35.8	-4.3 36.0	0.625 0.25 0.75	48.6 39.4	-3.6 39.6	354.4 4.4	317	0.766 0.0 1.0	42.1 71.6	-8.7 72.1	353.0		
430	B30R_087_062a	0.625 0.25 0.875	0.875 0.625 0.307	307	0.635 0.25 0.875	50.6 40.6	-9.0 41.6	0.625 0.25 0.875	49.0 42.1	-9.7 43.2	345.9 2.2	307	0.616 0.0 1.0	37.9 65.0	-14.5 66.6	347.4		
431	B25R_100_075a	0.625 0.25 1.0	1.0 0.75 0.625	300	0.625 0.25 1.0	50.6 43.9	-15.5 46.6	0.625 0.25 1.0	49.1 43.7	-15.5 46.4	340.3 1.5	300	0.5 0.0 1.0	35.6 58.6	-20.7 62.1	340.5		
432	R61Y_062_062a	0.625 0.375 0.0	0.625 0.625 0.312	67	0.625 0.385 0.0	53.9 10.2	47.9 49.0	0.625 0.375 0.0	50.8 21.2	46.0 50.6	65.2 11.5	67	1.0 0.616 0.0	71.6 16.4	76.6 78.4	77.8		
433	R50Y_062_050a	0.625 0.375 0.125	0.625 0.5 0.375	60	0.625 0.375 0.125	53.5 14.4	34.3 37.2	0.61 0.1	0.625 0.375 0.125	50.7 22.7	38.2 44.5	59.2 9.5	59	1.0 0.5 0.0	64.9 28.9	68.6 74.5	61.1	
434	R31Y_062_037a	0.625 0.375 0.25	0.625 0.375 0.437	49	0.625 0.368 0.25	54.2 17.1	22.2 28.1	0.625 0.375 0.25	50.9 23.8	30.0 38.3	51.5 10.8	48	1.0 0.316 0.0	56.6 45.8	59.2 74.9	52.2		
435	R00Y_062_025a	0.625 0.375 0.375	0.625 0.25 0.5	390	0.625 0.375 0.375	56.2 17.7	11.2 20.9	0.32	0.625 0.375 0.375	51.6 24.4	22.1 33.0	42.1 13.7	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3	
436	R00Y_062_025a	0.625 0.375 0.5	0.625 0.25 0.5	360	0.625 0.375 0.5	56.4 18.5	5.2 19.2	0.59	0.625 0.375 0.5	52.0 26.1	13.2 29.2	26.9 11.8	360	1.0 0.0 0.5	45.9 74.2	21.1 77.1	15.9	
437	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	56.5 19.8	0.0 19.8	0.59	0.625 0.375 0.625	52.6 27.8	4.7 28.2	9.6 10.1	330	1.0 0.0 1.0	46.1 79.3	-0.2 79.3	359.8	
438	B34R_075_037a	0.625 0.375 0.75	0.75 0.375 0.562	311	0.631 0.375 0.75	56.8 25.5	-4.4 25.9	0.625 0.375 0.75	53.8 29.6	-2.9 29.8	354.2 5.3	311	0.683 0.0 1.0	39.8 68.1	-11.9 69.1	350.0		
439	B25R_087_050a	0.625 0.375 0.875	0.875 0.5 0.625	300	0.625 0.375 0.875	56.7 29.3	-10.3 31.0	0.625 0.375 0.875	54.2 31.4	-9.8 32.9	342.6 3.3	300	0.5 0.0 1.0	35.6 58.6	-20.7 62.1	340.5		
440	B19R_100_062a	0.625 0.375 1.0	1.0 0.625 0.687	293	0.614 0.375 1.0	56.4 32.7	-16.0 36.4	0.625 0.375 1.0	54.3 32.9	-16.3 36.8	333.5 2.1	292	0.383 0.0 1.0	32.9 52.3	-25.7 58.3	333.8		
441	R81Y_062_062a	0.625 0.5 0.0	0.625 0.625 0.312	79	0.625 0.51 0.0	59.7 0.5	54.6 54.6	0.625 0.5 0.0	55.7 11.1	52.4 53.6	77.9 11.5	80	1.0 0.816 0.0	80.8 0.8	87.3 87.3	89.4		
442	R76Y_062_050a	0.625 0.5 0.125	0.625 0.5 0.375	76	0.625 0.508 0.125	60.4 2.1	42.3 42.4	0.625 0.5 0.125	56.2 11.5	43.8 45.3	75.3 10.3	77	1.0 0.766 0.0	78.6 4.3	84.7 84.8	87.0		
443	R68Y_062_037a	0.625 0.5 0.25	0.625 0.375 0.437	71	0.625 0.506 0.25	61.1 4.1	30.1 30.4	0.625 0.5 0.25	56.7 12.5	34.7 36.9	70.0 10.5	71	1.0 0.683 0.0	74.8 11.0	80.4 81.1	82.1		
444	R50Y_062_025a	0.625 0.5 0.375	0.625 0.25 0.5	60	0.625 0.5 0.375	61.2 7.2	17.1 18.6	0.61 0.1	0.625 0.5 0.375	57.0 14.3	25.0 28.8	60.2 11.4	59	1.0 0.5 0.0	64.9 28.9	68.6 74.5	61.1	
445	R00Y_062_012a	0.625 0.5 0.5	0.625 0.125 0.562	390	0.625 0.5 0.5	62.6 8.8	5.6 10.4	0.32	0.625 0.5 0.5	57.5 16.1	15.5 22.3	44.0 13.3	389	1.0 0.0 0.0	45.4 70.9	44.8 83.9	32.3	
446	B50R_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	62.7 9.9	0.0 9.9	0.59	0.625 0.5 0.625	58.3 18.1	6.5 19.3	19.8 11.4	330	1.0 0.0 1.0	46.1 79.3	-0.2 79.3	359.8	
447	B25R_075_025a	0.625 0.5 0.75	0.75 0.25 0.625	300	0.625 0.5 0.75	62.8 14.6	-5.1 15.5	0.34	0.625 0.5 0.75	58.9 19.9	-1.9 19.9	354.3 7.2	300	0.5 0.0 1.0	35.6 58.6	-20.7 62.1	340.5	
448	B15R_087_037a	0.625 0.5 0.875	0.875 0.375 0.687	289	0.618 0.5 0.875	62.4 17.7	-11.0 20.9	0.32	0.625 0.5 0.875	59.3 21.8	-9.6 23.8	336.0 5.3	288	0.316 0.0 1.0	30.9 47.3	-29.4 55.7	328.1	
449	B11R_100_050a	0.625 0.5 1.0	1.0 0.5 0.75	284	0.616 0.5 1.0	62.1 20.6	-16.5 26.4	0.31	0.625 0.5 1.0	59.7 24.4	-16.2 29.3	326.2 4.5	282	0.233 0.0 1.0	28.7 41.2	-33.1 52.9	321.1	
450	Y00G_062_062a	0.625 0.625 0.0	0.625 0.625 0.312	90	0.625 0.625 0.0	64.0	-6.3 59.6	0.60	0.61 0.0	0.625 0.625 0.0	61.0 0.3	58.3 58.3	89.6 7.4	89	1.0 1.0 0.0	87.8	-10.2 95.4	96.0 96.1
451	Y00G_062_050a	0.625 0.625 0.125	0.625 0.5 0.375	90	0.625 0.625 0.125	65.0	-5.1 47.7	48.0	0.61 0.0	0.625 0.625 0.125	61.5 0.9	49.3 49.3	88.9 7.1	89	1.0 1.0 0.0	87.8	-10.2 95.4	96.0 96.1
452	Y00G_062_037a	0.625 0.625 0.25	0.625 0.375															

n	HIC*Fa	rgb_Fa	icf_Fa	hsi_Fa	rgb*Fa	LabCh*Fa	rgb*Fa	LabCh*Fa	DE*Fa	hsiMd	rgb*Md	LabCh*Md		
486	R00Y_075_075a	0.75 0.0 0.0	0.75 0.75 0.375	390	0.75 0.0 0.0	40.2 53.2 33.6	62.9 32.3	0.75 0.0 0.0	40.7 59.2 36.3	69.4 31.5 6.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
487	R35Y_075_075a	0.75 0.0 0.125	0.75 0.75 0.375	381	0.75 0.0 0.112	40.2 53.7 29.2	61.1 28.5	0.75 0.0 0.125	40.6 60.2 31.6	68.0 27.7 6.9	382	1.0 0.0 0.15	45.5 71.6 39.0	81.5 28.5
488	R18Y_075_075a	0.75 0.0 0.25	0.75 0.75 0.375	371	0.75 0.0 0.237	40.4 54.5 23.4	59.3 23.2	0.75 0.0 0.25	40.9 61.1 25.5	66.2 22.6 6.9	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
489	R00Y_075_075a	0.75 0.0 0.375	0.75 0.75 0.375	360	0.75 0.0 0.375	40.5 55.6 18.8	57.8 15.9	0.75 0.0 0.375	41.0 62.2 19.2	65.1 17.1 7.4	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
490	B65R_075_075a	0.75 0.0 0.5	0.75 0.75 0.375	349	0.75 0.0 0.512	40.5 57.3 8.9	58.0 8.9	0.75 0.0 0.5	40.9 64.0 11.4	65.1 10.1 7.2	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
491	B57R_075_075a	0.75 0.0 0.625	0.75 0.75 0.375	339	0.75 0.0 0.637	40.5 58.5 3.7	58.6 3.7	0.75 0.0 0.625	41.1 65.4 5.1	65.6 4.4 7.0	337	1.0 0.0 0.85	45.9 78.0 5.0	78.2 3.7
492	B50R_075_075a	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	40.6 59.4 -0.1	59.4 359.8	0.75 0.0 0.75	41.1 66.9 0.0	66.9 0.0 7.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
493	B43R_087_087a	0.75 0.0 0.875	0.875 0.875 0.437	322	0.758 0.0 0.875	41.6 65.5 -4.6	65.7 355.9	0.75 0.0 0.875	41.4 69.0 -4.7	69.2 356.0 3.4	322	0.866 0.0 1.0	44.0 74.9 -5.3	75.1 355.9
494	B38R_100_100a	0.75 0.0 1.0	1.0 1.0 0.5	316	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0	0.75 0.0 1.0	41.8 71.0 -9.2	71.6 352.5 0.8	317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0
495	R15Y_075_075a	0.75 0.125 0.0	0.75 0.75 0.375	39	0.75 0.112 0.0	43.4 45.5 38.0	59.3 39.9	0.75 0.125 0.0	43.9 51.3 40.0	65.1 37.8 6.1	37	1.0 0.15 0.0	49.8 60.7 50.7	79.1 39.9
496	R00Y_075_062a	0.75 0.125 0.125	0.75 0.625 0.437	390	0.75 0.125 0.125	46.4 44.3 28.0	52.4 32.3	0.75 0.125 0.125	44.5 50.6 34.5	61.3 34.3 9.2	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
497	R31Y_075_062a	0.75 0.125 0.25	0.75 0.625 0.437	379	0.75 0.125 0.239	46.5 44.9 23.4	50.6 27.5	0.75 0.125 0.25	44.8 51.4 28.4	58.8 28.9 8.3	380	1.0 0.0 0.183	45.5 71.8 37.5	81.0 27.5
498	R11Y_075_062a	0.75 0.125 0.375	0.75 0.625 0.437	367	0.75 0.125 0.364	46.6 45.6 17.4	48.8 20.8	0.75 0.125 0.375	45.0 52.4 21.2	56.5 22.0 7.9	367	1.0 0.0 0.383	45.8 73.0 27.8	78.2 20.8
499	B69R_087_062a	0.75 0.125 0.5	0.75 0.625 0.437	353	0.75 0.125 0.51	46.8 47.2 9.5	48.1 11.4	0.75 0.125 0.5	45.4 54.0 12.4	55.4 12.9 7.5	352	1.0 0.0 0.616	46.0 75.5 15.2	77.1 11.4
500	B59R_075_062a	0.75 0.125 0.625	0.75 0.625 0.437	341	0.75 0.125 0.635	46.7 48.6 3.9	48.7 4.6	0.75 0.125 0.625	45.8 55.0 5.5	55.3 5.7 6.6	339	1.0 0.0 0.816	45.9 77.7 6.2	78.0 4.6
501	B50R_075_062a	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	46.8 49.5 -0.1	49.5 359.8	0.75 0.125 0.75	45.9 56.5 -0.2	56.5 359.7 7.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
502	B42R_087_075a	0.75 0.125 0.875	0.875 0.75 0.5	321	0.762 0.125 0.875	47.8 55.7 -4.4	55.9 355.4	0.75 0.125 0.875	46.6 58.6 -5.6	58.9 354.5 3.3	322	0.85 0.0 1.0	43.7 74.3 -5.9	74.6 355.4
503	B36R_100_087a	0.75 0.125 1.0	1.0 0.875 0.562	314	0.766 0.125 1.0	48.1 61.5 -8.7	61.2 351.9	0.75 0.125 1.0	47.0 60.4 -10.4	61.3 350.2 2.2	315	0.733 0.0 1.0	41.3 70.3 -9.9	71.0 351.9
504	R31Y_075_075a	0.75 0.25 0.0	0.75 0.75 0.375	49	0.75 0.237 0.0	48.5 34.3 44.4	56.2 52.2	0.75 0.25 0.0	48.9 39.7 46.7	61.3 49.6 5.8	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
505	R18Y_075_062a	0.75 0.25 0.125	0.75 0.625 0.437	41	0.75 0.239 0.125	50.0 36.1 32.8	48.8 42.2	0.75 0.25 0.125	49.3 39.8 39.4	56.1 44.7 7.6	39	1.0 0.183 0.0	51.1 57.8 52.5	78.1 42.2
506	R00Y_075_050a	0.75 0.25 0.25	0.75 0.5 0.5	390	0.75 0.25 0.25	52.7 35.4 22.4	41.9 32.3	0.75 0.25 0.25	50.4 39.4 31.9	50.7 38.9 10.5	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
507	R26Y_075_050a	0.75 0.25 0.375	0.75 0.5 0.5	376	0.75 0.25 0.366	52.8 36.0 17.6	40.1 26.1	0.75 0.25 0.375	51.0 39.9 24.4	46.8 31.4 8.0	377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1
508	R00Y_075_050a	0.75 0.25 0.5	0.75 0.5 0.5	360	0.75 0.25 0.5	52.9 37.1 10.5	38.5 15.9	0.75 0.25 0.5	51.3 41.4 15.2	44.1 20.2 6.5	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
509	B61R_075_050a	0.75 0.25 0.625	0.75 0.5 0.5	344	0.75 0.25 0.633	52.9 38.6 4.0	38.8 5.9	0.75 0.25 0.625	52.0 42.7 7.1	43.3 9.4 5.1	342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
510	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	53.0 39.6 -0.1	39.6 359.8	0.75 0.25 0.75	52.4 44.4 0.5	44.4 0.6 4.8	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
511	B40R_087_062a	0.75 0.25 0.875	0.875 0.625 0.562	319	0.76 0.25 0.875	53.9 45.8 -4.4	46.0 354.4	0.75 0.25 0.875	53.4 46.0 -5.4	46.3 353.2 1.1	320	0.816 0.0 1.0	43.1 73.2 -7.0	73.6 354.4
512	B34R_100_075a	0.75 0.25 1.0	1.0 0.75 0.625	311	0.762 0.25 1.0	53.7 51.0 -8.9	51.8 350.0	0.75 0.25 1.0	53.7 47.7 -10.9	48.9 347.1 3.9	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
513	R50Y_075_075a	0.75 0.375 0.0	0.75 0.75 0.375	60	0.75 0.375 0.0	54.7 21.6 51.5	55.9 67.1	0.75 0.375 0.0	54.3 28.1 53.1	60.1 62.1 6.6	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
514	R38Y_075_062a	0.75 0.375 0.125	0.75 0.625 0.437	53	0.75 0.364 0.125	55.2 24.7 39.1	46.2 57.6	0.75 0.375 0.125	54.7 28.8 44.2	52.8 56.8 6.6	52	1.0 0.383 0.0	59.5 39.5 62.5	74.0 57.6
515	R23Y_075_050a	0.75 0.375 0.25	0.75 0.5 0.5	44	0.75 0.366 0.25	56.5 26.7 27.4	38.2 45.7	0.75 0.375 0.25	55.2 29.4 35.2	45.9 50.0 8.3	42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7
516	R00Y_075_037a	0.75 0.375 0.375	0.75 0.375 0.562	390	0.75 0.375 0.375	59.0 26.6 16.8	31.4 32.3	0.75 0.375 0.375	56.5 29.0 26.5	39.3 42.3 10.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
517	R18Y_075_037a	0.75 0.375 0.5	0.75 0.375 0.562	371	0.75 0.375 0.493	59.1 27.2 11.7	29.6 23.2	0.75 0.375 0.5	56.9 30.5 18.0	35.4 30.6 7.4	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
518	B65R_075_037a	0.75 0.375 0.625	0.75 0.375 0.562	349	0.75 0.375 0.631	59.1 28.6 4.4	29.0 8.9	0.75 0.375 0.625	57.9 31.7 8.4	32.8 14.8 5.1	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
519	B50R_075_037a	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	59.2 29.7 0.0	29.7 359.8	0.75 0.375 0.75	58.3 33.3 1.5	33.4 2.6 4.1	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
520	B38R_087_050a	0.75 0.375 0.875	0.875 0.5 0.625	316	0.758 0.375 0.875	59.9 35.8 -4.3	36.0 353.0	0.75 0.375 0.875	59.1 35.6 -4.8	35.9 352.3 0.9	317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0
521	B30R_100_062a	0.75 0.375 1.0	1.0 0.625 0.687	307	0.76 0.375 1.0	59.5 40.6 -9.0	41.6 347.4	0.75 0.375 1.0	59.9 36.8 -10.8	38.4 343.5 4.2	307	0.616 0.0 1.0	37.9 65.0 -14.5	66.6 347.4
522	R68Y_075_075a	0.75 0.5 0.0	0.75 0.75 0.375	71	0.75 0.512 0.0	62.2 8.2 60.3	60.8 82.1	0.75 0.5 0.0	60.6 15.9 60.3	62.4 75.2 7.8	71	1.0 0.683 0.0	74.8 11.0 80.4	81.1 82.1
523	R61Y_075_062a	0.75 0.5 0.125	0.75 0.625 0.437	67	0.75 0.51 0.125	62.8 10.2 47.9	49.0 77.8	0.75 0.5 0.125	61.1 16.4 50.3	52.9 71.9 6.7	67	1.0 0.616 0.0	71.6 16.4 76.6	78.4 77.8
524	R50Y_075_050a	0.75 0.5 0.25	0.75 0.5 0.5	60	0.75 0.5 0.25	62.4 14.4 34.3	37.2 67.1	0.75 0.5 0.25	61.2 18.1 39.5	43.4 65.3 6.4	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
525	R31Y_075_037a	0.75 0.5 0.375	0.75 0.375 0.562	49	0.75 0.493 0.375	63.1 17.1 22.2	28.1 52.2	0.75 0.5 0.375	61.9 19.2 29.9	35.5 57.3 8.0	48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
526	R00Y_075_025a	0.75 0.5 0.5	0.75 0.25 0.625	390	0.75 0.5 0.5	65.2 17.7 11.2	20.9 32.3	0.75 0.5 0.5	62.8 20.1 19.9	28.3 44.7 9.3	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
527	R00Y_075_025a	0.75 0.5 0.625	0.75 0.25 0.625	360	0.75 0.5 0.625	65.3 18.5 5.2	19.2 15.9	0.75 0.5 0.625	63.6 21.9 10.7	24.4 26.2 6.6	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
528	B50R_075_025a	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	65.4 19.8 0.0	19.8 359.8	0.75 0.5 0.75	64.0 23.8 2.5	24.0 6.1 5.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
529	B34R_087_037a	0.75 0.5 0.875	0.875 0.375 0.687	311	0.756 0.5 0.875	65.7 25.5 -4.4	25.9 350.0	0.75 0.5 0.875	65.2 25.4 -4.4	25.8 350.1 0.5	311	0.683 0.0 1.0	39.8 68.1 -11.9	69.1 350.0
530	B25R_100_050a	0.75 0.5 1.0	1.0 0.5 0.75	300	0.75 0.5 1.0	65.6 29.3 -10.3	31.0 340.5	0.75 0.5 1.0	65.7 26.9 -11.2	29.2 337.4 2.4	300	0.5 0.0 1.0	35.6 58.6 -20.7	62.1 340.5
531	R85Y_075_075a	0.75 0.625 0.0	0.75 0.75 0.375	81	0.75 0.637 0.0	67.8 -1.1 66.7	66.7 91.0	0.75 0.625 0.0	66.7 4.4 67.2	67.4 86.2 5.7	81	1.0 0.85 0.0	82.3 -1.5 89.0	89.0 91.0
532	R81Y_075_062a	0.75 0.625 0.125	0.75 0.625 0.437	79	0.75 0.635 0.125	68.6 0.5 54.6	54.6 67.9	0.75 0.625 0.125	67.6 4.8 56.3	56.5 85.0 4.7	80	1.0 0.816 0.0	80.8 0.8 87.3	87.3 89.4
533	R76Y_075_050a	0.75 0.625 0.25	0.75 0.5 0.5	76	0.75 0.633 0.25	69.3 2.1 42.3	42.4 87.0	0.75 0.625 0.25	68.3 5.5 44.9	45.3 82.9 4.3	77	1.0 0.766 0.0	78.6 4.3 84.7	84.8 87.0
534	R68Y_075_037a	0.75 0.625 0.375	0.75 0.375 0.562	71	0.75 0.631 0.375	70.0 4.1 30.1	30.4 82.1	0.75 0.625 0.375	68.8 7.0 34.2	34.9 78.3 5.1	71	1.0 0.683 0.0	74.8 11.0 80.4	81.1 82.1
535	R50Y_075_025a	0.75 0.625 0.5	0.75 0.25 0.625	60	0.75 0.625 0.5	70.1 7.2 17.1	18.6 67.1	0.75 0.625 0.5	69.5 8.8 22.7	24.4 68.7 5.8	59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
536	R00Y_075_012a	0.75 0.625 0.625	0											

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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		
567	R00Y_087_087a	0.875 0.0 0.0	0.875 0.875 0.437	390	0.875 0.0 0.0	42.8 62.0 39.2	73.4 32.3	0.875 0.0 0.0	43.2 65.4 40.5	76.9 31.8 3.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
568	R36Y_087_087a	0.875 0.0 0.125	0.875 0.875 0.437	382	0.875 0.0 0.116	42.9 62.0 34.7	71.6 29.0	0.875 0.0 0.125	43.3 66.0 35.3	74.9 28.1 3.5	382	1.0 0.0 0.133	45.5 71.5 39.7	81.8 29.0
569	R23Y_087_087a	0.875 0.0 0.25	0.875 0.875 0.437	374	0.875 0.0 0.233	43.0 63.2 29.5	69.8 25.0	0.875 0.0 0.25	43.6 66.5 29.6	72.8 23.9 3.3	375	1.0 0.0 0.266	45.6 72.3 33.8	79.8 25.0
570	R08Y_087_087a	0.875 0.0 0.375	0.875 0.875 0.437	365	0.875 0.0 0.364	43.1 64.2 22.7	68.1 19.4	0.875 0.0 0.375	43.6 67.7 23.3	71.6 19.0 3.5	365	1.0 0.0 0.416	45.8 73.4 25.9	77.9 19.4
571	B70R_087_087a	0.875 0.0 0.5	0.875 0.875 0.437	355	0.875 0.0 0.51	43.2 65.8 14.8	67.4 12.7	0.875 0.0 0.5	43.7 69.3 16.0	71.2 13.0 3.7	354	1.0 0.0 0.583	45.9 75.2 16.9	77.1 12.7
572	B63R_087_087a	0.875 0.0 0.625	0.875 0.875 0.437	346	0.875 0.0 0.641	43.2 67.3 8.3	67.8 7.0	0.875 0.0 0.625	43.8 70.8 9.3	71.4 7.5 3.6	344	1.0 0.0 0.733	45.9 77.0 9.4	77.5 7.0
573	B56R_087_087a	0.875 0.0 0.75	0.875 0.875 0.437	338	0.875 0.0 0.758	43.2 68.4 3.8	68.5 3.2	0.875 0.0 0.75	43.8 72.3 4.2	72.5 3.3 4.0	337	1.0 0.0 0.866	45.9 78.1 4.4	78.3 3.2
574	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	43.4 69.4 -0.1	69.4 359.8	0.875 0.0 0.875	44.0 73.5 -0.8	73.5 359.3 4.2	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
575	B44R_100_100a	0.875 0.0 1.0	1.0 1.0 0.5	323	0.883 0.0 1.0	44.3 75.4 -4.7	75.6 356.3	0.875 0.0 1.0	44.2 75.2 -5.0	75.3 356.1 0.4	323	0.883 0.0 1.0	44.3 75.4 -4.7	75.6 356.3
576	R13Y_087_087a	0.875 0.125 0.0	0.875 0.875 0.437	38	0.875 0.116 0.0	46.1 54.3 43.6	69.7 38.7	0.875 0.125 0.0	47.3 56.4 44.0	71.5 38.0 2.4 37	1.0 0.133 0.0	49.2 61.2 49.8	79.6 38.7	
577	R00Y_087_075a	0.875 0.125 0.125	0.875 0.75 0.5	390	0.875 0.125 0.125	49.1 53.2 33.6	62.9 32.3	0.875 0.125 0.125	47.4 56.0 38.5	67.9 34.5 5.8	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
578	R35Y_087_075a	0.875 0.125 0.25	0.875 0.75 0.5	381	0.875 0.125 0.237	49.1 53.7 29.2	61.1 28.5	0.875 0.125 0.25	47.9 56.7 32.6	65.4 29.8 4.6	382	1.0 0.0 0.15	45.5 71.6 39.0	81.5 28.5
579	R18Y_087_075a	0.875 0.125 0.375	0.875 0.75 0.5	371	0.875 0.125 0.362	49.3 54.5 23.4	59.3 23.2	0.875 0.125 0.375	48.2 57.5 25.3	62.8 23.7 3.7	371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
580	R00Y_087_075a	0.875 0.125 0.5	0.875 0.75 0.5	360	0.875 0.125 0.5	49.4 55.6 15.8	57.8 15.9	0.875 0.125 0.5	48.4 59.1 16.9	61.5 15.9 3.7	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
581	B65R_087_075a	0.875 0.125 0.625	0.875 0.75 0.5	349	0.875 0.125 0.637	49.4 57.3 8.9	58.0 8.9	0.875 0.125 0.625	48.8 60.3 9.3	61.0 8.8 3.1	348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
582	B57R_087_075a	0.875 0.125 0.75	0.875 0.75 0.5	339	0.875 0.125 0.762	49.4 58.5 3.7	58.6 3.7	0.875 0.125 0.75	48.9 62.0 2.9	62.0 2.7 3.6	337	1.0 0.0 0.85	45.9 78.0 5.0	78.2 3.7
583	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	49.5 59.4 -0.1	59.4 359.8	0.875 0.125 0.875	49.3 62.9 -2.0	62.9 358.1 3.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
584	B43R_100_087a	0.875 0.125 1.0	1.0 0.875 0.562	322	0.883 0.125 1.0	50.5 65.5 -4.6	65.7 355.9	0.875 0.125 1.0	49.6 64.5 -6.6	64.9 354.1 2.3	322	0.866 0.0 1.0	46.0 74.9 -5.3	75.1 355.9
585	R26Y_087_087a	0.875 0.25 0.0	0.875 0.875 0.437	46	0.875 0.233 0.0	50.6 44.1 49.4	66.2 48.2	0.875 0.25 0.0	51.7 45.6 50.7	68.2 48.0 2.3 44	1.0 0.266 0.0	54.4 50.4 56.5	75.7 48.2	
586	R15Y_087_075a	0.875 0.25 0.125	0.875 0.75 0.5	39	0.875 0.237 0.125	52.4 45.5 38.0	59.3 39.9	0.875 0.25 0.125	52.6 45.0 43.6	62.7 44.1 5.6 37	1.0 0.15 0.0	49.8 60.7 50.7	79.1 39.9	
587	R00Y_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	55.3 44.3 28.0	52.4 32.3	0.875 0.25 0.25	53.7 44.1 35.9	56.8 39.1 8.0	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
588	R31Y_087_062a	0.875 0.25 0.375	0.875 0.625 0.562	379	0.875 0.25 0.364	55.4 44.9 23.4	50.6 27.5	0.875 0.25 0.375	54.3 44.5 28.2	52.7 32.3 4.8	380	1.0 0.0 0.183	45.5 71.8 37.5	81.0 27.5
589	R11Y_087_062a	0.875 0.25 0.5	0.875 0.625 0.562	367	0.875 0.25 0.489	55.6 45.6 17.4	48.8 20.8	0.875 0.25 0.5	54.5 45.9 19.9	50.0 23.4 2.7 367	1.0 0.0 0.383	45.8 73.0 27.8	78.2 20.8	
590	B69R_087_062a	0.875 0.25 0.625	0.875 0.625 0.562	353	0.875 0.25 0.635	55.7 47.2 9.5	48.1 11.4	0.875 0.25 0.625	55.1 47.5 10.8	48.7 12.8 1.4	352	1.0 0.0 0.616	46.0 75.5 15.2	77.1 11.4
591	B59R_087_062a	0.875 0.25 0.75	0.875 0.625 0.562	341	0.875 0.25 0.76	55.6 48.6 3.9	48.7 4.6	0.875 0.25 0.75	55.4 48.8 4.0	49.0 4.6 0.3	339	1.0 0.0 0.816	45.9 77.7 6.2	78.0 4.6
592	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	55.7 49.5 -0.1	49.5 359.8	0.875 0.25 0.875	56.0 49.9 -1.8	49.9 357.9 1.7	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
593	B42R_100_075a	0.875 0.25 1.0	1.0 0.75 0.625	321	0.887 0.25 1.0	56.7 55.7 -4.4	55.9 355.4	0.875 0.25 1.0	56.7 51.9 -6.8	52.3 352.4 4.5	322	0.85 0.0 1.0	43.7 74.3 -5.9	74.6 355.4
594	R41Y_087_087a	0.875 0.375 0.0	0.875 0.875 0.437	55	0.875 0.364 0.0	56.5 32.0 56.4	64.9 60.3	0.875 0.375 0.0	57.5 33.5 67.7	66.8 59.8 2.2 54	1.0 0.416 0.0	61.0 36.6 64.5	74.1 60.3	
595	R31Y_087_075a	0.875 0.375 0.125	0.875 0.75 0.5	49	0.875 0.362 0.125	57.4 34.3 44.4	56.2 52.2	0.875 0.375 0.125	57.9 33.6 48.9	59.4 55.5 4.6 48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2	
596	R18Y_087_062a	0.875 0.375 0.25	0.875 0.625 0.562	41	0.875 0.364 0.25	58.9 36.1 32.8	48.8 42.2	0.875 0.375 0.25	58.6 34.1 39.3	52.1 49.0 6.8 39	1.0 0.183 0.0	51.1 57.8 52.5	78.1 42.2	
597	R00Y_087_050a	0.875 0.375 0.375	0.875 0.5 0.625	390	0.875 0.375 0.375	61.6 35.4 22.4	41.9 32.3	0.875 0.375 0.375	59.7 33.8 30.7	45.6 42.2 8.6	389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
598	R26Y_087_050a	0.875 0.375 0.5	0.875 0.5 0.625	376	0.875 0.375 0.491	61.7 36.0 17.6	40.1 26.1	0.875 0.375 0.5	60.3 34.8 21.9	41.1 32.1 4.6 377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1	
599	R00Y_087_050a	0.875 0.375 0.625	0.875 0.5 0.625	360	0.875 0.375 0.625	61.8 37.1 10.5	38.5 15.9	0.875 0.375 0.625	61.1 36.1 12.9	38.3 19.7 2.6 360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9	
600	B61R_087_050a	0.875 0.375 0.75	0.875 0.5 0.625	344	0.875 0.375 0.758	61.8 38.6 4.0	38.8 5.9	0.875 0.375 0.75	61.4 37.8 4.6	38.1 7.0 1.0	342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
601	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	61.9 39.6 -0.1	39.6 359.8	0.875 0.375 0.875	62.3 38.7 -1.4	38.7 357.8 1.6	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
602	B40R_100_062a	0.875 0.375 1.0	1.0 0.625 0.687	319	0.885 0.375 1.0	62.8 45.8 -4.4	46.0 354.4	0.875 0.375 1.0	63.0 40.3 -7.2	40.9 349.7 6.1	320	0.816 0.0 1.0	43.1 73.2 -7.0	73.6 354.4
603	R58Y_087_087a	0.875 0.5 0.0	0.875 0.875 0.437	65	0.875 0.51 0.0	64.0 57.7 65.2	67.6 74.8	0.875 0.5 0.0	63.7 21.0 64.7	68.1 72.0 3.3 65	1.0 0.583 0.0	69.7 20.2 74.6	77.3 74.8	
604	R50Y_087_075a	0.875 0.5 0.125	0.875 0.75 0.5	60	0.875 0.5 0.125	63.6 21.6 51.5	55.9 67.1	0.875 0.5 0.125	63.9 22.1 53.8	58.2 67.6 2.4 59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1	
605	R38Y_087_062a	0.875 0.5 0.25	0.875 0.625 0.562	53	0.875 0.489 0.25	64.1 24.7 39.1	46.2 57.6	0.875 0.5 0.25	64.0 23.7 43.4	49.4 61.3 4.4 52	1.0 0.383 0.0	59.5 39.5 62.5	74.0 57.6	
606	R23Y_087_050a	0.875 0.5 0.375	0.875 0.5 0.625	44	0.875 0.491 0.375	65.4 26.7 27.4	38.2 45.7	0.875 0.5 0.375	64.9 24.1 33.4	41.4 54.1 6.5 42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7	
607	R00Y_087_037a	0.875 0.5 0.5	0.875 0.375 0.687	390	0.875 0.5 0.5	67.9 26.6 16.8	31.4 32.3	0.875 0.5 0.5	65.9 24.7 24.0	44.2 44.2 7.7 389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	
608	R18Y_087_037a	0.875 0.5 0.625	0.875 0.375 0.687	371	0.875 0.5 0.618	68.0 27.2 11.7	29.6 23.2	0.875 0.5 0.625	66.7 26.0 14.9	29.9 29.8 3.6 371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2	
609	B65R_087_037a	0.875 0.5 0.75	0.875 0.375 0.687	349	0.875 0.5 0.756	68.1 28.6 4.4	29.0 8.9	0.875 0.5 0.75	67.4 27.8 5.7	28.4 11.6 1.6 348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9	
610	B50R_087_037a	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	68.1 29.7 0.0	29.7 359.8	0.875 0.5 0.875	68.2 29.1 -0.9	29.1 358.1 1.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
611	B38R_100_050a	0.875 0.5 1.0	1.0 0.5 0.75	316	0.883 0.5 1.0	68.8 35.8 -4.3	36.0 353.0	0.875 0.5 1.0	69.1 30.9 -7.1	31.7 346.9 5.6 317	0.766 0.0 1.0	42.1 71.6 -8.7	72.1 353.0	
612	R73Y_087_087a	0.875 0.625 0.0	0.875 0.875 0.437	74	0.875 0.641 0.0	70.5 6.0 72.6	72.9 85.2	0.875 0.625 0.0	70.1 9.2 72.5	73.1 82.7 3.2 75	1.0 0.733 0.0	77.1 6.9 83.0	83.3 85.2	
613	R68Y_087_075a	0.875 0.625 0.125	0.875 0.75 0.5	71	0.875 0.637 0.125	71.1 8.2 60.3	60.8 82.1	0.875 0.625 0.125	70.5 9.9 60.9	61.7 80.7 1.8 71	1.0 0.683 0.0	74.8 11.0 80.4	81.1 82.1	
614	R61Y_087_062a	0.875 0.625 0.25	0.875 0.625 0.562	67	0.875 0.635 0.25	71.7 10.2 47.9	49.0 77.8	0.875 0.625 0.25	71.4 10.4 49.1	50.2 78.0 1.2 67	1.0 0.616 0.0	71.6 16.4 76.6	78.4 77.8	
615	R50Y_087_050a	0.875 0.625 0.375	0.875 0.5 0.625	60	0.875 0.625 0.375	71.3 14.4 34.3	37.2 67.1	0.875 0.625 0.375	71.7 11.9 38.5					

n	HIC*Fd	rgb_Fd	icf_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md		
648	R00Y_100_100a	1.0 0.0 0.0	1.0 1.0 0.5	390	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3	0.0 389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
649	R38Y_100_100a	1.0 0.0 0.125	1.0 1.0 0.5	383	1.0 0.0 0.116	45.5 71.4 40.4	82.1 29.5	1.0 0.0 0.125	45.5 71.4 40.1	81.9 29.3	0.3 383	1.0 0.0 0.116	45.5 71.4 40.4	82.1 29.5
650	R26Y_100_100a	1.0 0.0 0.25	1.0 1.0 0.5	376	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1	1.0 0.0 0.25	45.6 72.1 34.6	80.0 25.6	0.7 377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1
651	R13Y_100_100a	1.0 0.0 0.375	1.0 1.0 0.5	368	1.0 0.0 0.366	45.8 72.9 28.7	78.4 21.5	1.0 0.0 0.375	45.8 72.9 28.3	78.3 21.2	0.4 368	1.0 0.0 0.366	45.8 72.9 28.7	78.4 21.5
652	R00Y_100_100a	1.0 0.0 0.5	1.0 1.0 0.5	360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9	0.0 360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
653	B68R_100_100a	1.0 0.0 0.625	1.0 1.0 0.5	352	1.0 0.0 0.633	46.0 75.7 14.4	77.1 10.8	1.0 0.0 0.625	46.0 75.6 14.8	77.0 11.1	0.4 351	1.0 0.0 0.633	46.0 75.7 14.4	77.1 10.8
654	B61R_100_100a	1.0 0.0 0.75	1.0 1.0 0.5	344	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9	1.0 0.0 0.75	45.9 77.1 8.6	77.6 6.4	0.6 342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
655	B55R_100_100a	1.0 0.0 0.875	1.0 1.0 0.5	337	1.0 0.0 0.883	45.9 78.3 3.8	78.4 2.8	1.0 0.0 0.875	45.9 78.2 4.1	78.3 3.0	0.2 336	1.0 0.0 0.883	45.9 78.3 3.8	78.4 2.8
656	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	0.0 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
657	R11Y_100_100a	1.0 0.125 0.0	1.0 1.0 0.5	37	1.0 0.116 0.0	48.6 63.3 49.1	80.7 37.7	1.0 0.125 0.0	48.9 62.8 49.4	79.9 38.1	0.6 36	1.0 0.116 0.0	48.6 63.3 49.1	80.7 37.7
658	R00Y_100_087a	1.0 0.125 0.125	1.0 0.875 0.562	390	1.0 0.125 0.125	51.7 62.0 39.2	73.4 32.3	1.0 0.125 0.125	49.6 62.3 43.6	76.1 34.9	4.8 389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
659	R36Y_100_087a	1.0 0.125 0.25	1.0 0.875 0.562	382	1.0 0.125 0.241	51.8 62.5 34.7	71.6 29.0	1.0 0.125 0.25	49.6 61.3 36.9	73.1 30.3	3.0 382	1.0 0.0 0.133	45.5 71.5 39.7	81.8 29.0
660	R23Y_100_087a	1.0 0.125 0.375	1.0 0.875 0.562	374	1.0 0.125 0.358	51.9 63.2 29.5	69.8 25.0	1.0 0.125 0.375	50.0 63.5 30.1	70.3 25.3	2.0 375	1.0 0.0 0.266	45.6 72.3 33.8	79.8 25.0
661	R08Y_100_087a	1.0 0.125 0.5	1.0 0.875 0.562	365	1.0 0.125 0.489	52.0 64.2 22.7	68.1 19.4	1.0 0.125 0.5	50.2 64.7 22.4	68.5 19.1	1.9 365	1.0 0.0 0.416	45.8 73.4 25.9	77.9 19.4
662	B70R_100_087a	1.0 0.125 0.625	1.0 0.875 0.562	355	1.0 0.125 0.635	52.1 65.8 14.8	67.4 12.7	1.0 0.125 0.625	50.6 65.8 14.3	67.3 12.2	1.6 354	1.0 0.0 0.583	45.9 75.2 16.9	77.1 12.7
663	B63R_100_087a	1.0 0.125 0.75	1.0 0.875 0.562	346	1.0 0.125 0.766	52.1 67.3 8.3	67.8 7.0	1.0 0.125 0.75	50.9 66.9 7.4	67.3 6.3	1.5 344	1.0 0.0 0.733	45.9 77.0 9.4	77.5 7.0
664	B56R_100_087a	1.0 0.125 0.875	1.0 0.875 0.562	338	1.0 0.125 0.883	52.1 68.4 3.8	68.5 3.2	1.0 0.125 0.875	51.0 68.3 2.4	68.3 2.0	1.9 337	1.0 0.0 0.866	45.9 78.1 4.4	78.3 3.2
665	B50R_100_087a	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	52.3 69.4 -0.1	69.4 359.8	1.0 0.125 1.0	51.3 69.1 -2.3	69.2 358.0	2.4 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
666	R23Y_100_100a	1.0 0.25 0.0	1.0 1.0 0.5	44	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7	1.0 0.25 0.0	53.6 51.9 55.5	76.0 46.8	1.7 42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7
667	R13Y_100_087a	1.0 0.25 0.125	1.0 0.875 0.562	38	1.0 0.241 0.125	55.0 54.3 43.6	69.7 38.7	1.0 0.25 0.125	54.4 51.3 48.5	70.6 43.3	5.7 37	1.0 0.133 0.0	49.2 61.1 49.8	79.6 38.7
668	R00Y_100_075a	1.0 0.25 0.25	1.0 0.75 0.625	390	1.0 0.25 0.25	58.0 53.2 33.6	62.9 32.3	1.0 0.25 0.25	55.3 50.6 40.6	64.9 38.7	7.8 389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
669	R35Y_100_075a	1.0 0.25 0.375	1.0 0.75 0.625	381	1.0 0.25 0.362	58.0 53.7 29.2	61.1 28.5	1.0 0.25 0.375	55.8 50.9 33.0	60.7 32.9	5.2 382	1.0 0.0 0.15	45.5 71.6 39.0	81.5 28.5
670	R18Y_100_075a	1.0 0.25 0.5	1.0 0.75 0.625	371	1.0 0.25 0.487	58.2 54.5 23.4	59.3 23.2	1.0 0.25 0.5	56.4 51.4 24.6	57.0 25.5	3.6 371	1.0 0.0 0.316	45.7 72.6 31.2	79.1 23.2
671	R00Y_100_075a	1.0 0.25 0.625	1.0 0.75 0.625	360	1.0 0.25 0.625	58.3 55.6 15.8	57.8 15.9	1.0 0.25 0.625	56.8 52.8 15.9	57.2 16.7	3.1 360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
672	B65R_100_075a	1.0 0.25 0.75	1.0 0.75 0.625	349	1.0 0.25 0.762	58.3 57.3 8.9	58.0 8.9	1.0 0.25 0.75	57.1 54.5 7.8	55.1 8.1	3.2 348	1.0 0.0 0.683	45.9 76.4 11.9	77.3 8.9
673	B57R_100_075a	1.0 0.25 0.875	1.0 0.75 0.625	339	1.0 0.25 0.887	58.3 58.5 3.7	58.6 3.7	1.0 0.25 0.875	57.6 55.4 1.7	55.5 1.7	3.7 337	1.0 0.0 0.85	45.9 78.0 5.0	78.2 3.7
674	B50R_100_075a	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	58.4 59.4 -0.1	59.4 359.8	1.0 0.25 1.0	57.0 58.0 3.2	56.3 3.2	3.6 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
675	R36Y_100_100a	1.0 0.375 0.0	1.0 1.0 0.5	52	1.0 0.366 0.0	58.8 41.1 61.7	74.1 56.3	1.0 0.375 0.0	59.1 40.3 62.0	74.0 56.9	0.9 51	1.0 0.366 0.0	58.8 41.1 61.7	74.1 56.3
676	R26Y_100_087a	1.0 0.375 0.125	1.0 0.875 0.562	46	1.0 0.358 0.125	59.5 44.1 49.4	66.2 48.2	1.0 0.375 0.125	59.2 41.2 53.0	67.1 52.1	4.6 44	1.0 0.266 0.0	54.4 50.4 56.5	75.7 48.2
677	R15Y_100_075a	1.0 0.375 0.25	1.0 0.75 0.625	39	1.0 0.362 0.25	61.3 45.5 38.0	59.3 39.9	1.0 0.375 0.25	59.8 41.2 44.0	60.3 46.8	7.4 37	1.0 0.15 0.0	49.8 60.7 50.7	79.1 39.9
678	R00Y_100_062a	1.0 0.375 0.375	1.0 0.625 0.687	390	1.0 0.375 0.375	64.2 44.3 28.0	52.4 32.3	1.0 0.375 0.375	61.2 40.1 35.6	53.7 41.6	9.2 389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
679	R31Y_100_062a	1.0 0.375 0.5	1.0 0.625 0.687	379	1.0 0.375 0.489	64.3 44.9 23.4	50.6 27.5	1.0 0.375 0.5	61.7 40.7 27.1	48.9 33.6	6.1 380	1.0 0.0 0.183	45.5 71.8 37.5	81.0 27.5
680	R11Y_100_062a	1.0 0.375 0.625	1.0 0.625 0.687	367	1.0 0.375 0.614	64.5 45.6 17.4	48.8 20.8	1.0 0.375 0.625	62.6 41.7 17.7	45.3 23.0	4.4 367	1.0 0.0 0.383	45.8 73.0 27.8	78.2 20.8
681	B69R_100_062a	1.0 0.375 0.75	1.0 0.625 0.687	353	1.0 0.375 0.76	64.6 47.2 9.5	48.1 11.4	1.0 0.375 0.75	63.0 43.5 8.8	44.4 11.4	4.1 352	1.0 0.0 0.616	46.0 75.5 15.2	77.1 11.4
682	B59R_100_062a	1.0 0.375 0.875	1.0 0.625 0.687	341	1.0 0.375 0.885	64.5 48.6 3.9	48.7 4.6	1.0 0.375 0.875	63.9 44.3 1.6	44.3 2.1	4.8 339	1.0 0.0 0.816	45.9 77.7 6.2	78.0 4.6
683	B50R_100_062a	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	64.6 49.5 -0.1	49.5 359.8	1.0 0.375 1.0	64.6 45.0 -3.7	45.2 355.2	5.7 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
684	R50Y_100_100a	1.0 0.5 0.0	1.0 1.0 0.5	60	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1	0.0 59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
685	R41Y_100_087a	1.0 0.5 0.125	1.0 0.875 0.562	55	1.0 0.489 0.125	65.4 32.0 56.4	64.9 60.3	1.0 0.5 0.125	64.9 29.9 58.6	65.9 62.9	3.1 54	1.0 0.416 0.0	61.0 36.6 64.5	74.1 60.3
686	R31Y_100_075a	1.0 0.5 0.25	1.0 0.75 0.625	49	1.0 0.487 0.25	66.3 34.3 44.4	56.2 52.2	1.0 0.5 0.25	65.7 30.0 48.4	57.0 58.2	5.9 48	1.0 0.316 0.0	56.6 45.8 59.2	74.9 52.2
687	R18Y_100_062a	1.0 0.5 0.375	1.0 0.625 0.687	41	1.0 0.489 0.375	67.8 36.1 32.8	48.8 42.2	1.0 0.5 0.375	66.5 30.2 39.0	49.3 52.2	8.6 39	1.0 0.183 0.0	51.1 57.8 52.5	78.1 42.2
688	R00Y_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	70.5 35.4 22.4	41.9 32.3	1.0 0.5 0.5	68.0 29.9 28.7	41.5 43.8	8.7 389	1.0 0.0 0.0	45.4 70.9 44.8	83.9 32.3
689	R26Y_100_050a	1.0 0.5 0.625	1.0 0.5 0.75	376	1.0 0.5 0.616	70.6 36.0 17.6	40.1 26.1	1.0 0.5 0.625	68.6 31.2 19.2	36.6 31.5	5.4 377	1.0 0.0 0.233	45.6 72.1 35.3	80.3 26.1
690	R00Y_100_050a	1.0 0.5 0.75	1.0 0.5 0.75	360	1.0 0.5 0.75	70.7 37.1 10.5	38.5 15.9	1.0 0.5 0.75	69.1 32.9 10.3	34.5 17.4	4.4 360	1.0 0.0 0.5	45.9 74.2 21.1	77.1 15.9
691	B61R_100_050a	1.0 0.5 0.875	1.0 0.5 0.75	344	1.0 0.5 0.883	70.7 38.6 4.0	38.8 5.9	1.0 0.5 0.875	70.2 34.0 2.5	34.1 4.2	4.9 342	1.0 0.0 0.766	45.9 77.3 8.0	77.7 5.9
692	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	70.8 39.6 -0.1	39.6 359.8	1.0 0.5 1.0	70.7 35.2 -3.7	35.4 353.9	5.7 330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
693	R63Y_100_100a	1.0 0.625 0.0	1.0 1.0 0.5	68	1.0 0.633 0.0	72.5 14.8 77.6	79.0 79.1	1.0 0.625 0.0	72.1 15.4 77.1	78.6 78.6	0.8 68	1.0 0.633 0.0	72.5 14.8 77.6	79.0 79.1
694	R58Y_100_087a	1.0 0.625 0.125	1.0 0.875 0.562	65	1.0 0.635 0.125	72.9 17.7 65.2	67.6 74.8	1.0 0.625 0.125	73.0 15.1 66.5	68.2 77.1	2.8 65	1.0 0.583 0.0	69.7 20.2 74.6	77.3 74.8
695	R50Y_100_075a	1.0 0.625 0.25	1.0 0.75 0.625	60	1.0 0.625 0.25	72.5 21.6 51.5	55.9 67.1	1.0 0.625 0.25	73.3 16.2 54.7	57.1 73.4	6.3 59	1.0 0.5 0.0	64.9 28.9 68.6	74.5 67.1
696	R38Y_100_062a	1.0 0.625 0.375	1.0 0.625 0.687	53	1.0 0.614 0.375	73.0 24.7 39.1	46.2 57.6	1.0 0.625 0.375	73.7 17.5 43.5	46.9 68.0	8.4 52	1.0 0.383 0.0	59.5 39.5 62.5	74.0 57.6
697	R23Y_100_050a	1.0 0.625 0.5	1.0 0.5 0.75	44	1.0 0.616 0.5	74.3 26.7 27.4	38.2 45.7	1.0 0.625 0.5	74.7 18.3 32.2	37.0 60.3	9.6 42	1.0 0.233 0.0	53.0 53.4 54.8	76.5 45.7
698	R00Y_100_037a	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	76.8 26.6 16.8	31.4 32.3	1.0 0.625 0.625	76.0 19.0 21.7	28.9 48.7	9.0 389	1.0 0.0 0.		

n	HIC*Fd	rgb_Fd	icf_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md
729	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.5 0.0 0.0	0.0 0.0 0.0	112.0 0.1 360
730	G50B_100_012a	0.875 1.0 1.0	1.0 0.125 0.937	210	0.875 1.0 1.0	90.7 -3.1 -5.1	6.0 6.0 238.4	0.875 1.0 1.0	91.9 -2.9 -4.1	5.1 5.0 234.3	1.6 210 0.0	1.0 1.0 1.0
731	G50B_100_025a	0.75 1.0 1.0	1.0 0.25 0.875	210	0.75 1.0 1.0	85.9 -6.3 -10.3	12.1 238.4	0.75 1.0 1.0	87.8 -5.7 -8.6	10.3 236.4 2.7	210 0.0 1.0	1.0 1.0 1.0
732	G50B_100_037a	0.625 1.0 1.0	1.0 0.375 0.812	210	0.625 1.0 1.0	81.0 -9.5 -15.5	18.2 238.4	0.625 1.0 1.0	83.2 -8.6 -13.4	15.9 237.2 3.2	210 0.0 1.0	1.0 1.0 1.0
733	G50B_100_050a	0.5 1.0 1.0	1.0 0.5 0.75	210	0.5 1.0 1.0	76.2 -12.7 -20.7	24.3 238.4	0.5 1.0 1.0	77.6 -12.2 -19.4	22.9 237.6 2.0	210 0.0 1.0	1.0 1.0 1.0
734	G50B_100_062a	0.375 1.0 1.0	1.0 0.625 0.687	210	0.375 1.0 1.0	71.3 -15.9 -25.9	30.4 238.4	0.375 1.0 1.0	72.3 -15.5 -24.9	29.4 238.1 1.4	210 0.0 1.0	1.0 1.0 1.0
735	G50B_100_075a	0.25 1.0 1.0	1.0 0.75 0.625	210	0.25 1.0 1.0	66.5 -19.1 -31.1	36.5 238.4	0.25 1.0 1.0	66.5 -19.1 -31.2	36.6 238.4 0.0	210 0.0 1.0	1.0 1.0 1.0
736	G50B_100_087a	0.125 1.0 1.0	1.0 0.875 0.562	210	0.125 1.0 1.0	61.6 -22.3 -36.3	42.6 238.4	0.125 1.0 1.0	61.2 -21.8 -36.5	42.5 239.0 0.6	210 0.0 1.0	1.0 1.0 1.0
737	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4	0.0 1.0 1.0	55.3 -24.7 -42.3	49.0 239.6 1.7	210 0.0 1.0	1.0 1.0 1.0
738	ROOY_100_012a	1.0 0.875 0.875	1.0 0.125 0.937	390	1.0 0.875 0.875	89.3 8.8 5.6	10.4 32.3	1.0 0.875 0.875	89.7 4.4 7.8	9.0 60.1 4.9	389 1.0 0.0	0.0 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	86.7 0.0 0.0	0.0 0.0	0.875 0.875 0.875	86.1 1.2 3.6	3.8 70.9 3.8	360 1.0 1.0	1.0 1.0 1.0
740	G50B_087_012a	0.75 0.875 0.875	0.875 0.125 0.812	210	0.75 0.875 0.875	81.8 -3.1 -5.1	6.0 6.0 238.4	0.75 0.875 0.875	82.2 -1.9 -0.8	2.1 204.3 4.4	210 0.0 1.0	1.0 1.0 1.0
741	G50B_087_025a	0.625 0.875 0.875	0.875 0.25 0.75	210	0.625 0.875 0.875	77.0 -6.3 -10.3	12.1 238.4	0.625 0.875 0.875	77.9 -5.4 -5.5	7.8 225.6 4.9	210 0.0 1.0	1.0 1.0 1.0
742	G50B_087_037a	0.5 0.875 0.875	0.875 0.375 0.687	210	0.5 0.875 0.875	72.1 -9.5 -15.5	18.2 238.4	0.5 0.875 0.875	72.8 -9.5 -11.3	14.8 229.9 4.2	210 0.0 1.0	1.0 1.0 1.0
743	G50B_087_050a	0.375 0.875 0.875	0.875 0.5 0.625	210	0.375 0.875 0.875	67.3 -12.7 -20.7	24.3 238.4	0.375 0.875 0.875	67.6 -13.7 -16.9	21.8 230.9 3.9	210 0.0 1.0	1.0 1.0 1.0
744	G50B_087_062a	0.25 0.875 0.875	0.875 0.625 0.562	210	0.25 0.875 0.875	62.4 -15.9 -25.9	30.4 238.4	0.25 0.875 0.875	62.2 -18.3 -23.4	29.8 231.9 3.4	210 0.0 1.0	1.0 1.0 1.0
745	G50B_087_075a	0.125 0.875 0.875	0.875 0.75 0.5	210	0.125 0.875 0.875	57.6 -19.1 -31.1	36.5 238.4	0.125 0.875 0.875	57.2 -22.1 -28.6	36.1 232.2 3.9	210 0.0 1.0	1.0 1.0 1.0
746	G50B_087_087a	0.0 0.875 0.875	0.875 0.875 0.437	210	0.0 0.875 0.875	52.7 -22.3 -36.3	42.6 238.4	0.0 0.875 0.875	51.9 -26.3 -34.9	43.7 232.9 4.3	210 0.0 1.0	1.0 1.0 1.0
747	ROOY_100_025a	1.0 0.75 0.75	1.0 0.25 0.875	390	1.0 0.75 0.75	83.0 17.7 11.2	20.9 32.3	1.0 0.75 0.75	82.3 11.7 15.1	19.1 52.1 7.1	389 1.0 0.0	0.0 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	80.4 8.8 5.6	10.4 32.3	0.875 0.75 0.75	79.1 8.0 10.9	13.6 53.6 5.5	389 1.0 0.0	0.0 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	77.8 0.0 0.0	0.0 0.0	0.75 0.75 0.75	75.6 4.4 6.7	8.0 56.1 8.3	360 1.0 1.0	1.0 1.0 1.0
750	G50B_075_012a	0.625 0.75 0.75	0.75 0.125 0.687	210	0.625 0.75 0.75	72.9 -3.1 -5.1	6.0 6.0 238.4	0.625 0.75 0.75	71.2 0.3 1.9	2.0 79.0 8.2	210 0.0 1.0	1.0 1.0 1.0
751	G50B_075_025a	0.5 0.75 0.75	0.75 0.25 0.625	210	0.5 0.75 0.75	68.1 -6.3 -10.3	12.1 238.4	0.5 0.75 0.75	66.4 -4.7 -3.8	6.1 219.4 6.9	210 0.0 1.0	1.0 1.0 1.0
752	G50B_075_037a	0.375 0.75 0.75	0.75 0.375 0.562	210	0.375 0.75 0.75	63.2 -9.5 -15.5	18.2 238.4	0.375 0.75 0.75	61.8 -9.3 -9.6	13.4 225.8 6.0	210 0.0 1.0	1.0 1.0 1.0
753	G50B_075_050a	0.25 0.75 0.75	0.75 0.5 0.5	210	0.25 0.75 0.75	58.4 -12.7 -20.7	24.3 238.4	0.25 0.75 0.75	56.5 -15.2 -16.0	22.1 226.3 5.6	210 0.0 1.0	1.0 1.0 1.0
754	G50B_075_062a	0.125 0.75 0.75	0.75 0.625 0.437	210	0.125 0.75 0.75	53.5 -15.9 -25.9	30.4 238.4	0.125 0.75 0.75	52.2 -19.8 -21.1	28.9 226.8 6.3	210 0.0 1.0	1.0 1.0 1.0
755	G50B_075_075a	0.0 0.75 0.75	0.75 0.75 0.375	210	0.0 0.75 0.75	48.7 -19.1 -31.1	36.5 238.4	0.0 0.75 0.75	47.3 -25.7 -27.2	37.5 226.6 7.8	210 0.0 1.0	1.0 1.0 1.0
756	ROOY_100_037a	1.0 0.625 0.625	1.0 0.375 0.812	390	1.0 0.625 0.625	76.8 26.6 16.8	31.4 32.3	1.0 0.625 0.625	76.1 18.3 22.9	29.3 51.3 10.2	389 1.0 0.0	0.0 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	74.1 17.7 11.2	20.9 32.3	0.875 0.625 0.625	73.0 14.4 18.5	23.5 52.0 8.0	389 1.0 0.0	0.0 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	71.5 8.8 5.6	10.4 32.3	0.75 0.625 0.625	69.8 10.1 14.0	17.3 54.0 8.6	389 1.0 0.0	0.0 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	68.9 0.0 0.0	0.0 0.0	0.625 0.625 0.625	65.4 5.8 9.1	10.9 57.3 11.4	360 1.0 1.0	1.0 1.0 1.0
760	G50B_062_012a	0.5 0.625 0.625	0.625 0.125 0.562	210	0.5 0.625 0.625	64.0 -3.1 -5.1	6.0 6.0 238.4	0.5 0.625 0.625	61.0 0.4 3.7	3.7 83.2 10.1	210 0.0 1.0	1.0 1.0 1.0
761	G50B_062_025a	0.375 0.625 0.625	0.625 0.25 0.5	210	0.375 0.625 0.625	59.2 -6.3 -10.3	12.1 238.4	0.375 0.625 0.625	56.7 -5.3 -2.1	5.7 201.6 8.6	210 0.0 1.0	1.0 1.0 1.0
762	G50B_062_037a	0.25 0.625 0.625	0.625 0.375 0.437	210	0.25 0.625 0.625	54.3 -9.5 -15.5	18.2 238.4	0.25 0.625 0.625	51.9 -12.3 -8.5	14.9 214.7 7.9	210 0.0 1.0	1.0 1.0 1.0
763	G50B_062_050a	0.125 0.625 0.625	0.625 0.5 0.375	210	0.125 0.625 0.625	49.4 -12.7 -20.7	24.3 238.4	0.125 0.625 0.625	48.0 -18.0 -13.9	22.8 217.6 8.7	210 0.0 1.0	1.0 1.0 1.0
764	G50B_062_062a	0.0 0.625 0.625	0.625 0.625 0.312	210	0.0 0.625 0.625	44.6 -15.9 -25.9	30.4 238.4	0.0 0.625 0.625	43.3 -25.1 -20.1	32.1 218.6 10.9	210 0.0 1.0	1.0 1.0 1.0
765	ROOY_100_050a	1.0 0.5 0.5	1.0 0.5 0.75	390	1.0 0.5 0.5	70.5 35.4 22.4	41.9 32.3	1.0 0.5 0.5	68.2 29.0 29.0	41.1 45.0 9.5	389 1.0 0.0	0.0 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	67.9 26.6 16.8	31.4 32.3	0.875 0.5 0.5	65.3 24.5 25.2	35.1 45.7 9.0	389 1.0 0.0	0.0 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	65.2 17.7 11.2	20.9 32.3	0.75 0.5 0.5	62.2 20.1 20.1	28.5 45.0 9.7	389 1.0 0.0	0.0 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	62.6 8.8 5.6	10.4 32.3	0.625 0.5 0.5	58.7 14.9 15.6	21.6 46.3 12.3	389 1.0 0.0	0.0 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	60.0 0.0 0.0	0.0 0.0	0.5 0.5 0.5	54.3 8.9 10.1	13.5 48.5 14.6	360 1.0 1.0	1.0 1.0 1.0
770	G50B_050_012a	0.375 0.5 0.5	0.5 0.125 0.437	210	0.375 0.5 0.5	55.1 -3.1 -5.1	6.0 6.0 238.4	0.375 0.5 0.5	50.6 1.9 4.3	4.7 65.2 11.7	210 0.0 1.0	1.0 1.0 1.0
771	G50B_050_025a	0.25 0.5 0.5	0.5 0.25 0.375	210	0.249 0.5 0.5	50.2 -6.3 -10.3	12.1 238.4	0.25 0.5 0.5	46.0 -5.6 -2.0	6.0 199.5 9.3	210 0.0 1.0	1.0 1.0 1.0
772	G50B_050_037a	0.125 0.5 0.5	0.5 0.375 0.312	210	0.124 0.5 0.5	45.4 -9.5 -15.5	18.2 238.4	0.125 0.5 0.5	42.3 -12.7 -7.7	14.9 211.3 8.9	210 0.0 1.0	1.0 1.0 1.0
773	G50B_050_050a	0.0 0.5 0.5	0.5 0.5 0.25	210	0.0 0.5 0.5	40.5 -12.7 -20.7	24.3 238.4	0.0 0.5 0.5	38.5 -21.4 -13.9	25.5 213.0 11.2	210 0.0 1.0	1.0 1.0 1.0
774	ROOY_100_062a	1.0 0.375 0.375	1.0 0.625 0.687	390	1.0 0.375 0.375	64.2 44.3 28.0	52.4 32.3	1.0 0.375 0.375	61.4 39.0 35.7	52.9 42.4 9.7	389 1.0 0.0	0.0 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	61.6 35.4 22.4	41.9 32.3	0.875 0.375 0.375	58.9 33.9 31.5	46.3 42.8 9.6	389 1.0 0.0	0.0 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	59.0 26.6 16.8	31.4 32.3	0.75 0.375 0.375	55.9 29.2 26.8	39.7 42.5 10.8	389 1.0 0.0	0.0 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	56.3 17.7 11.2	20.9 32.3	0.625 0.375 0.375	52.5 23.8 21.9	32.3 42.6 12.9	389 1.0 0.0	0.0 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	53.7 8.8 5.6	10.4 32.3	0.5 0.375 0.375	48.7 16.8 16.1	23.3 43.7 14.1	389 1.0 0.0	0.0 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	51.0 0.0 0.0	0.0 0.0	0.375 0.375 0.375	45.0 9.7 10.1	14.0 46.0 15.3	360 1.0 1.0	1.0 1.0 1.0
780	G50B_037_012a	0.25 0.375 0.375	0.375 0.125 0.312	210	0.249 0.375 0.375	46.2 -3.1 -5.1	6.0 6.0 238.4	0.25 0.375 0.375	40.9 0.8 3.7	3.8 77.4 11.1	210 0.0 1.0	1.0 1.0 1.0
781	G50B_037_025a	0.125 0.375 0.375	0.375 0.25 0.25	210	0.124 0.375 0.375	41.3 -6.3 -10.3	12.1 238.4	0.125 0.375 0.375	37.7 -7.6 -1.5	7.8 191.5 9.6	210 0.0 1.0	1.0 1.0 1.0
782	G50B_037_037a	0.0 0.375 0.375	0.375 0.375 0.187	210	0.0 0.375 0.375	36.5 -9.5 -15.5	18.2 238.4	0.0 0.375 0.375	34.4 -17.9 -8.0	19.6 204.0 11.5	210 0.0 1.0	1.0 1.0 1.0
783	ROOY_100_075a	1.0 0.25 0.25	1.0 0.75 0.625	390	1.0 0.25 0.25	58.0 53.2 33.6	62.9 32.3	1.0 0.25 0.25	55.3 49.7 40.6	64.1 39.2 8.2	389 1.0 0.0	0.0 0.0 0.0
784	ROOY_087_062a	0.875 0.25 0.25	0.875 0.625 0.562	390	0.875 0.25 0.25	55.3 44.3 28.0	52.4 32.3	0.875 0.25 0.25	52.8 44			

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF77/PF77.HTM>
 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

TUB enregistrement: 20130201 -PF77/PF77L0NA.TXT /.PS
 application pour la mesure des sorties sur offset, séparation cmy0 (CMY0)
 TUB matériel: code=rh4ta

n	HIC*Fd	rgb_Fd	icf_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md
810	NW_100d	1.0 1.0 1.0	1.0 0.0 1.0	0.0 1.0 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.6 0.0 0.1	116.7 0.1 360	1.0 1.0 1.0	95.6 0.0 0.0
811	BOOR_100_012d	0.875 0.875 1.0	1.0 0.125 0.937	270 360	0.875 0.875 1.0	86.8 3.6 -5.0	6.2 306.2	0.875 0.875 1.0	87.2 3.8 -5.3	6.6 305.3	0.5 270 0.0	25.0 29.5 -40.4
812	BOOR_100_025d	0.75 0.75 1.0	1.0 0.25 0.875	270 360	0.75 0.75 1.0	77.9 7.3 -10.1	12.5 306.2	0.75 0.75 1.0	76.6 9.6 -10.6	14.3 312.1	2.6 270 0.0	25.0 29.5 -40.4
813	BOOR_100_037d	0.625 0.625 1.0	1.0 0.375 0.812	270 360	0.625 0.625 1.0	69.1 11.0 -15.1	18.7 306.2	0.625 0.625 1.0	67.2 13.6 -15.6	20.8 310.1	3.2 270 0.0	25.0 29.5 -40.4
814	BOOR_100_050d	0.5 0.5 1.0	1.0 0.5 0.75	270 360	0.5 0.5 1.0	60.3 14.7 -20.2	25.0 306.2	0.5 0.5 1.0	55.8 19.6 -21.4	29.1 312.4	6.7 270 0.0	25.0 29.5 -40.4
815	BOOR_100_062d	0.375 0.375 1.0	1.0 0.625 0.687	270 360	0.375 0.375 1.0	51.5 18.4 -25.2	31.3 306.2	0.375 0.375 1.0	45.8 24.1 -26.3	35.7 312.5	8.1 270 0.0	25.0 29.5 -40.4
816	BOOR_100_075d	0.25 0.25 1.0	1.0 0.75 0.625	270 360	0.25 0.25 1.0	42.7 22.1 -30.3	37.5 306.2	0.25 0.25 1.0	37.4 26.6 -31.6	41.3 310.1	6.9 270 0.0	25.0 29.5 -40.4
817	BOOR_100_087d	0.125 0.125 1.0	1.0 0.875 0.562	270 360	0.125 0.125 1.0	33.9 25.8 -35.3	43.8 306.2	0.125 0.125 1.0	28.7 31.4 -36.1	47.8 311.0	7.6 270 0.0	25.0 29.5 -40.4
818	BOOR_100_100d	0.0 0.0 1.0	1.0 1.0 0.5	270 360	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 0.0 1.0	23.4 30.0 -39.6	50.1 307.6	2.0 270 0.0	25.0 29.5 -40.4
819	Y00G_100_012d	1.0 1.0 0.875	1.0 0.125 0.937	90 90	1.0 1.0 0.875	94.6 -1.2 11.9	12.0 96.1	1.0 1.0 0.875	94.6 -2.5 9.9	10.2 104.1	2.3 89 1.0	1.0 1.0 0.878
820	NW_087d	0.875 0.875 0.875	0.875 0.0 0.875	360 360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	86.3 1.2 3.7	3.9 71.1	3.9 360 1.0	1.0 1.0 0.956
821	BOOR_087_012d	0.75 0.75 0.875	0.875 0.125 0.812	270 360	0.75 0.75 0.875	77.9 3.6 -5.0	6.2 306.2	0.75 0.75 0.875	76.0 6.9 -2.3	7.3 341.0	4.5 270 0.0	25.0 29.5 -40.4
822	BOOR_087_025d	0.625 0.625 0.875	0.875 0.25 0.75	270 360	0.625 0.625 0.875	69.0 7.3 -10.1	12.5 306.2	0.625 0.625 0.875	66.7 11.0 -8.0	13.6 323.8	4.7 270 0.0	25.0 29.5 -40.4
823	BOOR_087_037d	0.5 0.5 0.875	0.875 0.375 0.687	270 360	0.5 0.5 0.875	60.2 11.0 -15.1	18.7 306.2	0.5 0.5 0.875	55.5 16.6 -14.6	22.1 318.6	7.2 270 0.0	25.0 29.5 -40.4
824	BOOR_087_050d	0.375 0.375 0.875	0.875 0.5 0.625	270 360	0.375 0.375 0.875	51.4 14.7 -20.2	25.0 306.2	0.375 0.375 0.875	45.6 21.0 -20.4	29.2 315.8	8.5 270 0.0	25.0 29.5 -40.4
825	BOOR_087_062d	0.25 0.25 0.875	0.875 0.625 0.562	270 360	0.25 0.25 0.875	42.6 18.4 -25.2	31.3 306.2	0.25 0.25 0.875	37.1 23.2 -26.2	35.0 311.5	7.3 270 0.0	25.0 29.5 -40.4
826	BOOR_087_075d	0.125 0.125 0.875	0.875 0.75 0.5	270 360	0.125 0.125 0.875	33.8 22.1 -30.3	37.5 306.2	0.125 0.125 0.875	29.0 26.9 -31.2	41.2 310.8	6.8 270 0.0	25.0 29.5 -40.4
827	BOOR_087_087d	0.0 0.0 0.875	0.875 0.875 0.437	270 360	0.0 0.0 0.875	24.9 25.8 -35.3	43.8 306.2	0.0 0.0 0.875	23.4 26.1 -35.1	43.8 306.6	1.6 270 0.0	25.0 29.5 -40.4
828	Y00G_100_025d	1.0 1.0 0.75	1.0 0.25 0.875	90 90	1.0 1.0 0.75	93.6 -2.5 23.8	24.0 96.1	1.0 1.0 0.75	93.5 -4.4 20.0	20.4 102.4	4.2 89 1.0	1.0 1.0 0.878
829	Y00G_087_012d	0.875 0.875 0.75	0.875 0.125 0.812	90 90	0.875 0.875 0.75	85.7 -1.2 11.9	12.0 96.1	0.875 0.875 0.75	85.2 -0.7 13.0	13.1 93.4	1.3 89 1.0	1.0 1.0 0.878
830	NW_075d	0.75 0.75 0.75	0.75 0.0 0.75	360 360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	75.1 4.6 6.6	8.1 54.7	8.5 360 1.0	1.0 1.0 0.956
831	BOOR_075_012d	0.625 0.625 0.75	0.75 0.125 0.687	270 360	0.625 0.625 0.75	68.9 3.6 -5.0	6.2 306.2	0.625 0.625 0.75	66.1 8.4 0.2	8.4 17.7	7.7 270 0.0	25.0 29.5 -40.4
832	BOOR_075_025d	0.5 0.5 0.75	0.75 0.25 0.625	270 360	0.5 0.5 0.75	60.1 7.3 -10.1	12.5 306.2	0.5 0.5 0.75	54.8 13.8 -6.8	15.4 333.6	8.9 270 0.0	25.0 29.5 -40.4
833	BOOR_075_037d	0.375 0.375 0.75	0.75 0.375 0.562	270 360	0.375 0.375 0.75	51.3 11.0 -15.1	18.7 306.2	0.375 0.375 0.75	45.6 17.2 -13.3	21.7 322.1	8.5 270 0.0	25.0 29.5 -40.4
834	BOOR_075_050d	0.25 0.25 0.75	0.75 0.5 0.5	270 360	0.25 0.25 0.75	42.5 14.7 -20.2	25.0 306.2	0.25 0.25 0.75	37.2 19.3 -19.7	27.6 314.5	7.0 270 0.0	25.0 29.5 -40.4
835	BOOR_075_062d	0.125 0.125 0.75	0.75 0.625 0.437	270 360	0.125 0.125 0.75	33.7 18.4 -25.2	31.3 306.2	0.125 0.125 0.75	29.3 22.6 -25.7	34.2 311.4	6.1 270 0.0	25.0 29.5 -40.4
836	BOOR_075_075d	0.0 0.0 0.75	0.75 0.75 0.375	270 360	0.0 0.0 0.75	24.9 22.1 -30.3	37.5 306.2	0.0 0.0 0.75	23.6 21.0 -30.2	36.9 304.8	1.6 270 0.0	25.0 29.5 -40.4
837	Y00G_100_037d	1.0 1.0 0.625	1.0 0.375 0.812	90 90	1.0 1.0 0.625	92.6 -3.8 35.8	36.0 96.1	1.0 1.0 0.625	92.4 -6.1 30.9	31.6 101.2	5.3 89 1.0	1.0 1.0 0.878
838	Y00G_087_025d	0.875 0.875 0.625	0.875 0.25 0.75	90 90	0.875 0.875 0.625	84.7 -2.5 23.8	24.0 96.1	0.875 0.875 0.625	84.2 -2.8 23.6	23.8 96.7	0.5 89 1.0	1.0 1.0 0.878
839	Y00G_075_012d	0.75 0.75 0.625	0.75 0.125 0.687	90 90	0.75 0.75 0.625	76.8 -1.2 11.9	12.0 96.1	0.75 0.75 0.625	74.4 2.4 16.3	16.5 81.4	6.2 89 1.0	1.0 1.0 0.878
840	NW_062d	0.625 0.625 0.625	0.625 0.0 0.625	360 360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	65.5 5.9 9.4	11.1 57.6	11.6 360 1.0	1.0 1.0 0.956
841	BOOR_062_012d	0.5 0.5 0.625	0.625 0.125 0.562	270 360	0.5 0.5 0.625	60.0 3.6 -5.0	6.2 306.2	0.5 0.5 0.625	54.5 11.4 1.1	11.4 5.8	11.3 270 0.0	25.0 29.5 -40.4
842	BOOR_062_025d	0.375 0.375 0.625	0.625 0.25 0.5	270 360	0.375 0.375 0.625	51.2 7.3 -10.1	12.5 306.2	0.375 0.375 0.625	45.2 14.8 -6.0	16.0 337.7	10.3 270 0.0	25.0 29.5 -40.4
843	BOOR_062_037d	0.25 0.25 0.625	0.625 0.375 0.437	270 360	0.25 0.25 0.625	42.4 11.0 -15.1	18.7 306.2	0.25 0.25 0.625	36.9 16.3 -13.2	21.0 320.9	7.8 270 0.0	25.0 29.5 -40.4
844	BOOR_062_050d	0.125 0.125 0.625	0.625 0.5 0.375	270 360	0.125 0.125 0.625	33.6 14.7 -20.2	25.0 306.2	0.125 0.125 0.625	29.1 19.3 -19.9	27.7 314.1	6.3 270 0.0	25.0 29.5 -40.4
845	BOOR_062_062d	0.0 0.0 0.625	0.625 0.625 0.312	270 360	0.0 0.0 0.625	24.8 18.4 -25.2	31.3 306.2	0.0 0.0 0.625	23.5 16.8 -24.9	30.0 304.0	2.1 270 0.0	25.0 29.5 -40.4
846	Y00G_100_050d	1.0 1.0 0.5	1.0 0.5 0.75	90 90	1.0 1.0 0.5	91.7 -5.1 47.7	48.0 96.1	1.0 1.0 0.5	91.2 -7.6 43.4	44.1 100.0	5.0 89 1.0	1.0 1.0 0.878
847	Y00G_087_037d	0.875 0.875 0.5	0.875 0.375 0.687	90 90	0.875 0.875 0.5	83.7 -3.8 35.8	36.0 96.1	0.875 0.875 0.5	83.1 -4.5 35.6	35.8 97.2	0.9 89 1.0	1.0 1.0 0.878
848	Y00G_075_025d	0.75 0.75 0.5	0.75 0.25 0.625	90 90	0.75 0.75 0.5	75.8 -2.5 23.8	24.0 96.1	0.75 0.75 0.5	73.6 0.4 27.0	27.0 88.9	4.9 89 1.0	1.0 1.0 0.878
849	Y00G_062_012d	0.625 0.625 0.5	0.625 0.125 0.562	90 90	0.625 0.625 0.5	67.9 -1.2 11.9	12.0 96.1	0.625 0.625 0.5	64.7 3.9 19.0	19.4 78.1	9.3 89 1.0	1.0 1.0 0.878
850	NW_050d	0.5 0.5 0.5	0.5 0.0 0.5	360 360	0.5 0.5 0.5	60.0 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	54.3 9.1 9.8	13.4 47.1	14.5 360 1.0	1.0 1.0 0.956
851	BOOR_050_012d	0.375 0.375 0.5	0.5 0.125 0.437	270 360	0.375 0.375 0.5	51.1 3.6 -5.0	6.2 306.2	0.375 0.375 0.5	45.1 12.0 1.6	12.1 7.7	12.2 270 0.0	25.0 29.5 -40.4
852	BOOR_050_025d	0.25 0.25 0.5	0.5 0.25 0.375	270 360	0.25 0.25 0.5	42.3 7.3 -10.1	12.5 306.2	0.25 0.25 0.5	36.8 13.1 -6.7	14.7 332.9	8.6 270 0.0	25.0 29.5 -40.4
853	BOOR_050_037d	0.125 0.125 0.5	0.5 0.375 0.312	270 360	0.125 0.125 0.5	33.5 11.0 -15.1	18.7 306.2	0.125 0.125 0.5	29.0 15.8 -14.1	21.2 318.3	6.6 270 0.0	25.0 29.5 -40.4
854	BOOR_050_050d	0.0 0.0 0.5	0.5 0.5 0.25	270 360	0.0 0.0 0.5	24.7 14.7 -20.2	25.0 306.2	0.0 0.0 0.5	23.6 12.6 -19.4	23.2 302.9	2.5 270 0.0	25.0 29.5 -40.4
855	Y00G_100_062d	1.0 1.0 0.375	1.0 0.625 0.687	90 90	1.0 1.0 0.375	90.7 -6.3 59.6	60.0 96.1	1.0 1.0 0.375	89.9 -8.6 55.9	56.5 96.8	4.4 89 1.0	1.0 1.0 0.878
856	Y00G_087_050d	0.875 0.875 0.375	0.875 0.5 0.625	90 90	0.875 0.875 0.375	82.8 -5.1 47.7	48.0 96.1	0.875 0.875 0.375	81.9 -5.6 47.6	47.9 96.7	1.0 89 1.0	1.0 1.0 0.878
857	Y00G_075_037d	0.75 0.75 0.375	0.75 0.375 0.562	90 90	0.75 0.75 0.375	74.8 -3.8 35.8	36.0 96.1	0.75 0.75 0.375	72.6 -0.8 38.3	38.3 91.1	4.5 89 1.0	1.0 1.0 0.878
858	Y00G_062_025d	0.625 0.625 0.375	0.625 0.25 0.5	90 90	0.625 0.625 0.375	66.9 -2.5 23.8	24.0 96.1	0.625 0.625 0.375	64.1 2.1 29.3	29.4 85.7	7.7 89 1.0	1.0 1.0 0.878
859	Y00G_050_012d	0.5 0.5 0.375	0.5 0.125 0.437	90 90	0.5 0.5 0.375	59.0 -1.2 11.9	12.0 96.1	0.5 0.5 0.375	53.6 6.9 18.8	20.1 69.7	12.0 89 1.0	1.0 1.0 0.878
860	NW_037d	0.375 0.375 0.375	0.375 0.0 0.375	360 360	0.375 0.375 0.375	51.0 0.0 0.0	0.0 0.0 0.0	0.375 0.375 0.375	44.7 10.1 9.6	14.0 43.4	15.3 360 1.0	1.0 1.0 0.956
861	BOOR_037_012d	0.25 0.25 0.375	0.375 0.125 0.312	270 360	0.25 0.25 0.375	42.2 3.6 -5.0	6.2 306.2	0.25 0.25 0.375	36.9 10.5 0.5	10.5 3.1	10.3 270 0.0	25.0 29.5 -40.4
862	BOOR_037_025d	0.125 0.125 0.375	0.375 0.25 0.25	270 360	0.125 0.125 0.375	33.4 7.3 -10.1	12.5 306.2	0.125 0.125 0.375	28.8 12.5 -7.8	14.8 328.0	7.2 270 0.0	25.0 29.5 -40.4
863	BOOR_037_037d	0.0 0.0 0.375	0.375 0.375 0.187	270 360	0.0 0.0 0.375	24.6 11.0 -15.1	18.7 306.2	0.0 0.0 0.375	23.3 8.6 -14.0	16.5 301.4	2.9 270 0.0	25.0 29.5 -40.4
864												

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF77/PF77.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	hsi_Md	rgb*Md	LabCh*Md		
891	NW_100a	1.0 1.0 1.0	1.0 0.0 1.0	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.6 0.0 0.1	0.1 111.4 0.1	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
892	B50R_100_012a	1.0 0.875 1.0	1.0 0.125 0.937	330	1.0 0.875 1.0	89.4 9.9 0.0	9.9 359.8	1.0 0.875 1.0	90.7 6.8 -1.4	6.9 348.2 3.6	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
893	B50R_100_025a	1.0 0.75 1.0	1.0 0.25 0.875	330	1.0 0.75 1.0	83.2 19.8 0.0	19.8 359.8	1.0 0.75 1.0	84.2 15.6 -2.4	15.8 351.1 4.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
894	B50R_100_037a	1.0 0.625 1.0	1.0 0.375 0.812	330	1.0 0.625 1.0	77.0 29.7 0.0	29.7 359.8	1.0 0.625 1.0	78.5 23.6 -3.2	23.8 352.2 7.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
895	B50R_100_050a	1.0 0.5 1.0	1.0 0.5 0.75	330	1.0 0.5 1.0	70.8 39.6 -0.1	39.6 359.8	1.0 0.5 1.0	70.6 35.6 -3.8	35.8 353.8 5.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
896	B50R_100_062a	1.0 0.375 1.0	1.0 0.625 0.687	330	1.0 0.375 1.0	64.6 49.5 -0.1	49.5 359.8	1.0 0.375 1.0	63.5 46.7 -3.8	46.9 353.7 4.7	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
897	B50R_100_075a	1.0 0.25 1.0	1.0 0.75 0.625	330	1.0 0.25 1.0	58.4 59.4 -0.1	59.4 359.8	1.0 0.25 1.0	57.0 58.1 -2.9	58.1 357.1 3.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
898	B50R_100_087a	1.0 0.125 1.0	1.0 0.875 0.562	330	1.0 0.125 1.0	52.3 69.4 -0.1	69.4 359.8	1.0 0.125 1.0	50.3 70.4 -1.6	70.4 358.6 2.6	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
899	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	1.0 0.0 1.0	45.4 79.5 1.0	79.5 0.7 1.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
900	GO0B_100_012a	0.875 1.0 0.875	1.0 0.125 0.937	150	0.875 1.0 0.875	89.9 -8.1 3.7	8.9 155.5	0.875 1.0 0.875	90.9 -5.6 5.6	7.9 135.3 3.2	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
901	NW_087a	0.875 0.875 0.875	0.875 0.0 0.875	360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	86.2 1.2 3.6	3.8 71.0 3.8	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
902	B50R_087_012a	0.875 0.75 0.875	0.875 0.125 0.812	330	0.875 0.75 0.875	80.5 9.9 0.0	9.9 359.8	0.875 0.75 0.875	80.1 10.0 2.1	10.2 11.8 2.1	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
903	B50R_087_025a	0.875 0.625 0.875	0.875 0.25 0.75	330	0.875 0.625 0.875	74.3 19.8 0.0	19.8 359.8	0.875 0.625 0.875	74.6 18.0 0.9	18.1 2.9 2.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
904	B50R_087_037a	0.875 0.5 0.875	0.875 0.375 0.687	330	0.875 0.5 0.875	68.1 29.7 0.0	29.7 359.8	0.875 0.5 0.875	66.7 30.6 -0.6	30.6 358.7 1.7	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
905	B50R_087_050a	0.875 0.375 0.875	0.875 0.5 0.625	330	0.875 0.375 0.875	61.9 39.6 -0.1	39.6 359.8	0.875 0.375 0.875	60.5 40.8 -1.0	40.8 358.5 2.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
906	B50R_087_062a	0.875 0.25 0.875	0.875 0.625 0.562	330	0.875 0.25 0.875	55.7 49.5 -0.1	49.5 359.8	0.875 0.25 0.875	54.0 52.3 -1.0	52.3 358.7 3.3	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
907	B50R_087_075a	0.875 0.125 0.875	0.875 0.75 0.5	330	0.875 0.125 0.875	49.5 59.4 -0.1	59.4 359.8	0.875 0.125 0.875	47.7 64.4 -0.5	64.4 359.4 5.3	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
908	B50R_087_087a	0.875 0.0 0.875	0.875 0.875 0.437	330	0.875 0.0 0.875	43.4 69.4 -0.1	69.4 359.8	0.875 0.0 0.875	42.9 73.7 1.1	73.7 0.8 4.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
909	GO0B_100_025a	0.75 1.0 0.75	1.0 0.25 0.875	150	0.75 1.0 0.75	84.2 -16.2 7.4	17.8 155.5	0.75 1.0 0.75	85.6 -11.0 10.4	15.2 136.5 6.2	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
910	GO0B_087_012a	0.75 0.875 0.75	0.875 0.125 0.812	150	0.75 0.875 0.75	81.0 -8.1 3.7	8.9 155.5	0.75 0.875 0.75	81.1 -4.3 8.3	9.4 117.5 5.9	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
911	NW_075a	0.75 0.75 0.75	0.75 0.0 0.75	360	0.75 0.75 0.75	77.8 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	75.6 4.3 6.4	7.8 56.1 8.1	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
912	B50R_075_012a	0.75 0.625 0.75	0.75 0.125 0.687	330	0.75 0.625 0.75	71.6 9.9 0.0	9.9 359.8	0.75 0.625 0.75	70.5 12.2 4.7	13.1 21.4 5.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
913	B50R_075_025a	0.75 0.5 0.75	0.75 0.25 0.625	330	0.75 0.5 0.75	65.4 19.8 0.0	19.8 359.8	0.75 0.5 0.75	63.2 23.9 2.7	24.1 6.6 5.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
914	B50R_075_037a	0.75 0.375 0.75	0.75 0.375 0.562	330	0.75 0.375 0.75	59.2 29.7 0.0	29.7 359.8	0.75 0.375 0.75	57.3 34.4 1.7	34.4 2.9 5.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
915	B50R_075_050a	0.75 0.25 0.75	0.75 0.5 0.5	330	0.75 0.25 0.75	53.0 39.6 -0.1	39.6 359.8	0.75 0.25 0.75	50.7 45.7 0.7	45.8 0.9 6.6	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
916	B50R_075_062a	0.75 0.125 0.75	0.75 0.625 0.437	330	0.75 0.125 0.75	46.8 49.5 -0.1	49.5 359.8	0.75 0.125 0.75	44.9 57.7 0.1	57.7 0.1 8.4	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
917	B50R_075_075a	0.75 0.0 0.75	0.75 0.75 0.375	330	0.75 0.0 0.75	40.6 59.4 -0.1	59.4 359.8	0.75 0.0 0.75	40.3 67.0 1.0	67.0 0.8 7.6	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
918	GO0B_100_037a	0.625 1.0 0.625	1.0 0.375 0.812	150	0.625 1.0 0.625	78.5 -24.3 11.1	26.7 155.5	0.625 1.0 0.625	79.8 -17.2 15.5	23.2 137.8 8.5	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
919	GO0B_087_025a	0.625 0.875 0.625	0.875 0.25 0.75	150	0.625 0.875 0.625	75.3 -16.2 7.4	17.8 155.5	0.625 0.875 0.625	76.0 -10.5 12.9	16.7 129.1 8.0	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
920	GO0B_075_012a	0.625 0.75 0.625	0.75 0.125 0.687	150	0.625 0.75 0.625	72.1 -8.1 3.7	8.9 155.5	0.625 0.75 0.625	70.7 -2.0 10.9	11.1 100.3 9.5	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
921	NW_062a	0.625 0.625 0.625	0.625 0.0 0.625	360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	66.0 5.6 8.9	10.5 57.5 10.9	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
922	B50R_062_012a	0.625 0.5 0.625	0.625 0.125 0.562	330	0.625 0.5 0.625	62.7 9.9 0.0	9.9 359.8	0.625 0.5 0.625	59.5 17.0 6.1	18.1 19.9 9.9	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
923	B50R_062_025a	0.625 0.375 0.625	0.625 0.25 0.5	330	0.625 0.375 0.625	56.5 19.8 0.0	19.8 359.8	0.625 0.375 0.625	53.7 26.9 4.3	27.3 9.1 8.8	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
924	B50R_062_037a	0.625 0.25 0.625	0.625 0.375 0.437	330	0.625 0.25 0.625	50.3 29.7 0.0	29.7 359.8	0.625 0.25 0.625	47.9 38.2 2.9	38.3 4.3 9.3	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
925	B50R_062_050a	0.625 0.125 0.625	0.625 0.5 0.375	330	0.625 0.125 0.625	44.1 39.6 -0.1	39.6 359.8	0.625 0.125 0.625	42.0 50.1 1.3	50.1 1.5 10.7	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
926	B50R_062_062a	0.625 0.0 0.625	0.625 0.625 0.312	330	0.625 0.0 0.625	37.9 49.5 -0.1	49.5 359.8	0.625 0.0 0.625	37.5 59.5 0.8	59.5 0.7 10.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
927	GO0B_100_050a	0.5 1.0 0.5	1.0 0.5 0.75	150	0.5 1.0 0.5	72.8 -32.5 14.8	35.7 155.5	0.5 1.0 0.5	73.8 -24.0 19.6	31.0 140.7 9.7	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
928	GO0B_087_037a	0.5 0.875 0.5	0.875 0.375 0.687	150	0.5 0.875 0.5	69.6 -24.3 11.1	26.7 155.5	0.5 0.875 0.5	70.0 -18.0 17.2	24.9 136.3 8.8	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
929	GO0B_075_025a	0.5 0.75 0.5	0.75 0.25 0.625	150	0.5 0.75 0.5	66.4 -16.2 7.4	17.8 155.5	0.5 0.75 0.5	65.3 -9.6 14.9	17.7 122.9 10.0	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
930	GO0B_062_012a	0.5 0.625 0.5	0.625 0.125 0.562	150	0.5 0.625 0.5	63.2 -8.1 3.7	8.9 155.5	0.5 0.625 0.5	61.0 -2.3 12.4	12.6 100.7 10.6	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
931	NW_050a	0.5 0.5 0.5	0.5 0.0 0.5	360	0.5 0.5 0.5	60.0 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	54.8 8.7 9.3	12.7 47.0 13.7	360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
932	B50R_050_012a	0.5 0.375 0.5	0.5 0.125 0.437	330	0.5 0.375 0.5	53.8 9.9 0.0	9.9 359.8	0.5 0.375 0.5	49.6 18.6 6.7	19.8 19.7 11.8	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
933	B50R_050_025a	0.5 0.25 0.5	0.5 0.25 0.375	330	0.5 0.25 0.5	47.6 19.8 0.0	19.8 359.8	0.5 0.25 0.5	44.1 29.4 4.1	29.7 7.9 11.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
934	B50R_050_037a	0.5 0.125 0.5	0.5 0.375 0.312	330	0.5 0.125 0.5	41.4 29.7 0.0	29.7 359.8	0.5 0.125 0.5	38.7 41.2 1.8	41.3 2.5 12.0	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
935	B50R_050_050a	0.5 0.0 0.5	0.5 0.5 0.25	330	0.5 0.0 0.5	35.2 39.6 -0.1	39.6 359.8	0.5 0.0 0.5	34.5 50.1 0.7	50.1 0.8 10.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8
936	GO0B_100_062a	0.375 1.0 0.375	1.0 0.625 0.887	150	0.375 1.0 0.375	67.1 -40.6 18.5	44.6 155.5	0.375 1.0 0.375	67.5 -31.6 23.8	39.6 143.0 10.4	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
937	GO0B_087_050a	0.375 0.875 0.375	0.875 0.5 0.625	150	0.375 0.875 0.375	63.9 -32.5 14.8	35.7 155.5	0.375 0.875 0.375	64.2 -26.0 21.1	33.5 140.9 9.0	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
938	GO0B_075_037a	0.375 0.75 0.375	0.75 0.375 0.562	150	0.375 0.75 0.375	60.7 -24.3 11.1	26.7 155.5	0.375 0.75 0.375	60.0 -17.7 18.6	25.7 133.4 10.1	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
939	GO0B_062_025a	0.375 0.625 0.375	0.625 0.25 0.5	150	0.375 0.625 0.375	57.5 -16.2 7.4	17.8 155.5	0.375 0.625 0.375	56.3 -10.4 16.0	19.1 123.1 10.4	149	0.0 1.0 0.0	50.0 -65.0	29.6 71.4 155.5
940	GO0B_050_012a	0.375 0.5 0.375	0.5 0.125 0.437	150	0.375 0.									

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF77/PF77.HTM>
informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	icf_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md			
972	NW_000a	0.0 0.0 0.0	0.0 0.0 0.0	0.0 360	0.0 0.0 0.0	24.3 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	23.1 1.0	-1.6 1.9	302.0 2.2 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
973	NW_012a	0.125 0.125 0.125	0.125 0.125 0.125	0.125 360	0.125 0.125 0.125	33.2 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.125 0.125 0.125	28.5 8.0	4.0 8.9	26.4 10.1 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	42.1 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.25 0.25 0.25	36.5 9.3	8.5 12.6	42.5 13.9 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
975	NW_037a	0.375 0.375 0.375	0.375 0.375 0.375	0.375 360	0.375 0.375 0.375	51.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.375 0.375 0.375	45.3 10.1	10.9 14.8	47.1 15.9 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	60.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	55.2 8.8	10.0 13.3	48.4 14.2 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
977	NW_062a	0.625 0.625 0.625	0.625 0.625 0.625	0.625 360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	66.4 5.6	9.0 10.6	58.3 10.9 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	77.8 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	76.2 3.9	6.3 7.5	57.9 7.6 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
979	NW_087a	0.875 0.875 0.875	0.875 0.875 0.875	0.875 360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	86.5 1.1	3.3 3.6	70.5 3.6 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	95.6 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.6 0.0	0.0 0.1	126.7 0.1 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	24.3 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	22.9 1.2	-0.6 1.4	332.7 2.0 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
982	NW_012a	0.125 0.125 0.125	0.125 0.125 0.125	0.125 360	0.125 0.125 0.125	33.2 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.125 0.125 0.125	28.4 8.3	4.3 9.4	27.2 10.5 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	42.1 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.25 0.25 0.25	35.9 9.7	9.1 13.3	43.2 14.7 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
984	NW_037a	0.375 0.375 0.375	0.375 0.375 0.375	0.375 360	0.375 0.375 0.375	51.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.375 0.375 0.375	45.6 9.9	11.0 14.9	47.9 15.8 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	60.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	55.1 8.6	9.9 13.1	49.1 14.0 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
986	NW_062a	0.625 0.625 0.625	0.625 0.625 0.625	0.625 360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	66.2 5.6	9.1 10.7	58.2 11.1 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	77.8 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	76.0 4.1	6.1 7.4	56.0 7.6 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
988	NW_087a	0.875 0.875 0.875	0.875 0.875 0.875	0.875 360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	86.6 1.2	3.4 3.6	70.8 3.6 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	95.6 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.6 0.0	0.0 0.0	133.9 0.1 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	24.3 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	23.0 0.5	-0.7 0.9	307.9 1.6 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
991	NW_012a	0.125 0.125 0.125	0.125 0.125 0.125	0.125 360	0.125 0.125 0.125	33.2 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.125 0.125 0.125	28.1 7.9	4.7 9.2	30.9 10.6 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	42.1 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.25 0.25 0.25	36.3 9.2	9.2 13.0	45.2 14.3 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
993	NW_037a	0.375 0.375 0.375	0.375 0.375 0.375	0.375 360	0.375 0.375 0.375	51.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.375 0.375 0.375	44.9 10.0	11.2 15.1	48.2 16.3 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	60.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	54.7 8.9	9.9 13.3	48.3 14.3 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
995	NW_062a	0.625 0.625 0.625	0.625 0.625 0.625	0.625 360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	66.3 5.6	9.3 10.9	59.0 11.2 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	77.8 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	75.8 4.1	6.3 7.5	56.9 7.8 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
997	NW_087a	0.875 0.875 0.875	0.875 0.875 0.875	0.875 360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	86.3 1.1	3.4 3.6	71.6 3.6 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	95.6 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.7 0.0	0.1 0.1	120.9 0.2 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	24.3 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	22.8 0.5	-0.5 0.8	317.5 1.7 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1000	NW_012a	0.125 0.125 0.125	0.125 0.125 0.125	0.125 360	0.125 0.125 0.125	33.2 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.125 0.125 0.125	27.9 8.0	4.4 9.1	28.8 10.5 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	42.1 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.25 0.25 0.25	35.8 9.1	9.3 13.0	45.5 14.5 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1002	NW_037a	0.375 0.375 0.375	0.375 0.375 0.375	0.375 360	0.375 0.375 0.375	51.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.375 0.375 0.375	44.9 10.0	11.4 15.2	48.7 16.4 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	60.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.5 0.5 0.5	54.7 9.1	10.4 13.8	48.7 14.8 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1004	NW_062a	0.625 0.625 0.625	0.625 0.625 0.625	0.625 360	0.625 0.625 0.625	68.9 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.625 0.625 0.625	66.0 5.6	9.5 11.1	59.3 11.4 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	77.8 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.75 0.75 0.75	75.7 4.1	6.4 7.6	57.3 7.9 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1006	NW_087a	0.875 0.875 0.875	0.875 0.875 0.875	0.875 360	0.875 0.875 0.875	86.7 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.875 0.875 0.875	86.3 1.1	3.5 3.7	71.9 3.8 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	95.6 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 0.0	113.6 0.1 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	24.3 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	23.1 1.4	-1.9 2.4	306.9 2.7 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1009	NW_006a	0.066 0.066 0.066	0.066 0.066 0.066	0.066 360	0.066 0.066 0.066	29.0 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.066 0.066 0.066	26.0 5.8	0.2 5.8	2.4 6.6 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1010	NW_013a	0.133 0.133 0.133	0.133 0.133 0.133	0.133 360	0.133 0.133 0.133	33.8 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.133 0.133 0.133	28.8 8.4	3.0 9.0	19.7 10.3 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	38.6 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.2 0.2 0.2	32.3 9.7	5.8 11.4	30.8 13.0 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1012	NW_026a	0.266 0.266 0.266	0.266 0.266 0.266	0.266 360	0.266 0.266 0.266	43.3 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.266 0.266 0.266	37.0 9.1	8.3 12.3	42.4 13.8 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1013	NW_033a	0.333 0.333 0.333	0.333 0.333 0.333	0.333 360	0.333 0.333 0.333	48.1 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.333 0.333 0.333	41.4 10.4	9.4 14.0	42.0 15.5 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	52.8 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.4 0.4 0.4	47.5 8.9	9.8 13.3	47.7 14.3 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1015	NW_046a	0.466 0.466 0.466	0.466 0.466 0.466	0.466 360	0.466 0.466 0.466	57.5 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.466 0.466 0.466	52.0 8.9	10.0 13.4	48.0 14.5 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1016	NW_053a	0.533 0.533 0.533	0.533 0.533 0.533	0.533 360	0.533 0.533 0.533	62.3 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.533 0.533 0.533	57.0 7.2	10.0 12.3	53.9 13.4 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 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	67.1 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.6 0.6 0.6	64.2 5.6	8.6 10.3	57.1 10.7 360	1.0 1.0 1.0	95.6 0.0 0.0	0.0 0.0 0.0
1018	NW_066a	0.666 0.666 0.666	0.666 0.666 0.666	0.666 360	0.666 0.666 0.666	71.8 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	0.666 0.666 0.666	69.7 5.2	8.2 9.7				

voir fichiers similaires: <http://130.149.60.45/~farbmetrik/PF77/PF77.HTM>
 informations techniques: <http://www.ps.bam.de> ou <http://130.149.60.45/~farbmetrik>

n	HIC*Fd	rgb_Fd	icf_Fd	hsi_Fd	rgb*Fd	LabCh*Fd	rgb*Fd	LabCh*Fd	DE*Fd	hsiMd	rgb*Md	LabCh*Md
1053	NW_086a	0.866 0.866 0.866	0.866 0.0 0.866	360	0.866 0.866 0.866	86.0 0.0 0.0	0.0 0.0 0.0	0.866 0.866 0.866	86.1 1.2 3.4	3.7 69.9 3.7	360	95.6 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	90.8 0.0 0.0	0.0 0.0 0.0	0.933 0.933 0.933	90.8 0.4 1.4	1.5 71.6 1.5	360	95.6 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	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.6 0.0 0.1	0.1 114.3 0.1	360	95.6 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	24.3 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	23.0 0.7 -0.9	1.1 308.5 1.7	360	95.6 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	29.0 0.0 0.0	0.0 0.0 0.0	0.066 0.066 0.066	25.6 5.5 0.6	5.5 6.7 6.5	360	95.6 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	33.8 0.0 0.0	0.0 0.0 0.0	0.133 0.133 0.133	28.2 8.3 3.4	9.0 22.4 10.6	360	95.6 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	38.6 0.0 0.0	0.0 0.0 0.0	0.2 0.2 0.2	32.0 10.0 5.8	11.6 30.4 13.3	360	95.6 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	43.3 0.0 0.0	0.0 0.0 0.0	0.266 0.266 0.266	36.7 8.8 8.7	12.4 44.7 14.0	360	95.6 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	48.1 0.0 0.0	0.0 0.0 0.0	0.333 0.333 0.333	40.7 10.4 8.9	13.4 40.4 15.5	360	95.6 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	52.8 0.0 0.0	0.0 0.0 0.0	0.4 0.4 0.4	46.8 8.7 10.2	13.4 49.7 14.7	360	95.6 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	57.5 0.0 0.0	0.0 0.0 0.0	0.466 0.466 0.466	51.8 8.8 9.9	13.3 48.4 14.5	360	95.6 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	62.3 0.0 0.0	0.0 0.0 0.0	0.533 0.533 0.533	57.5 7.3 9.2	11.8 51.6 12.7	360	95.6 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	67.1 0.0 0.0	0.0 0.0 0.0	0.6 0.6 0.6	63.6 6.0 9.2	11.0 56.7 11.5	360	95.6 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	71.8 0.0 0.0	0.0 0.0 0.0	0.666 0.666 0.666	69.3 5.2 8.3	9.8 57.5 10.1	360	95.6 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	76.6 0.0 0.0	0.0 0.0 0.0	0.734 0.734 0.734	74.5 4.8 6.5	8.1 53.5 8.3	360	95.6 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	81.3 0.0 0.0	0.0 0.0 0.0	0.8 0.8 0.8	80.5 2.7 5.2	5.9 62.0 5.9	360	95.6 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	86.0 0.0 0.0	0.0 0.0 0.0	0.866 0.866 0.866	86.1 1.2 3.4	3.6 69.4 3.6	360	95.6 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	90.8 0.0 0.0	0.0 0.0 0.0	0.933 0.933 0.933	90.7 0.4 1.4	1.5 71.7 1.5	360	95.6 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	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.7 0.0 0.0	0.1 118.4 0.1	360	95.6 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	24.3 0.0 0.0	0.0 0.0 0.0	0.0 0.0 0.0	23.3 1.3 -2.4	2.8 299.2 2.9	360	95.6 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	95.6 0.0 0.0	0.0 0.0 0.0	1.0 1.0 1.0	95.7 0.0 0.0	0.0 138.7 0.0	360	95.6 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	45.4 70.9 44.8	83.9 32.3	1.0 0.0 0.0	45.4 70.5 45.5	83.9 32.8 0.7	389	1.0 0.0 0.0
1075	G50B_100_100a	0.0 1.0 1.0	1.0 1.0 0.5	210	0.0 1.0 1.0	56.8 -25.5 -41.5	48.7 238.4	0.0 1.0 1.0	56.4 -25.2 -41.8	48.8 238.9 0.5	210	0.0 1.0 1.0
1076	Y00G_100_100a	1.0 1.0 0.0	1.0 1.0 0.5	90	1.0 1.0 0.0	87.8 -10.2 95.4	96.0 96.1	1.0 1.0 0.0	87.5 -10.0 95.1	95.7 96.0 0.4	89	1.0 1.0 0.0
1077	B00R_100_100a	0.0 0.0 1.0	1.0 1.0 0.5	270	0.0 0.0 1.0	25.0 29.5 -40.4	50.0 306.2	0.0 0.0 1.0	24.7 29.8 -40.1	49.9 306.6 0.5	270	0.0 0.0 1.0
1078	G00B_100_100a	0.0 1.0 0.0	1.0 1.0 0.5	150	0.0 1.0 0.0	50.0 -65.0 29.6	71.4 155.5	0.0 1.0 0.0	49.2 -65.4 28.0	71.2 156.7 1.8	149	0.0 1.0 0.0
1079	B50R_100_100a	1.0 0.0 1.0	1.0 1.0 0.5	330	1.0 0.0 1.0	46.1 79.3 -0.2	79.3 359.8	1.0 0.0 1.0	45.8 79.2 -0.2	79.2 359.8 0.2	330	1.0 0.0 1.0

delta E* = 5.8

3-0032531-F0

PF770-7N, 26/26-F

PE4600L_120830.TXT, 1080 colors, Separation cmy0*

graphique TUB-PF77; cercle de teinte; 16 et 8 étapes
 couleurs et différences, ΔE^* , 3D=0, de=0, cmy0

entrée : $rgb/cmyk \rightarrow rgb_d$
 sortie : transférer à $cmy0_d$

3-0032531-F0