Appendix C. Detailed segmentation statistics.

The tables is this appendix show distribution statistics, by finger position, for the segmentation algorithms tested as compared to the hand marked ground truth for 3-inch slap images. The differences between the segmentation algorithm and ground truth are sorted into bins based on the tolerances allowed for correct segmentation. Specifically, the left/right edges must be within -32/+64 pixels of the ground truth, top edge -64/+64 and bottom edge -64/+128. For each finger position there is a column for each of the four segmentation box edges (L, R, T and B).

The first row (“No Finger Found”) shows the counts for when a finger was not detected by the segmentation algorithm. The next four rows show statistics for segmentation edges that are within the specified minimum (MN) and maximum (MX) pixel tolerances compared to the ground truth, so these are considered good segmentations. Rows 1 (MN <= d < 0) and 3 (0 <= d <= MX) show the average value for all differences in that range and rows 3 and 5 show the total count occurring in that range.

Rows 6-9 also show average difference values and bin counts but for ranges MN-32 <= d < MN and MX < d <= MX+32, which are just outside the accepted tolerance ranges. Rows 10-13 tally everything greater than 32 pixels away from the accepted tolerance range, d < MN-32 and d > MX+32.

The last three rows show the total count for each bin, the overall average difference value and the standard deviation of all the difference values.

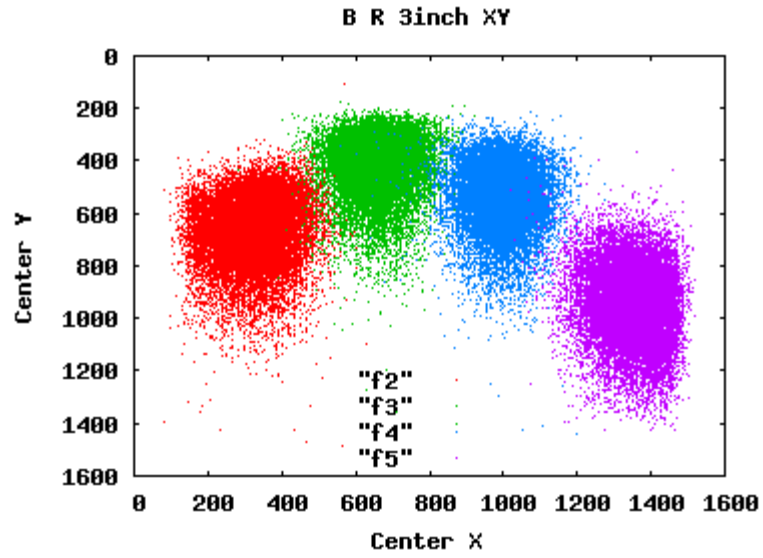
B

	R. Thumb				R. Index				R Middle				R. Ring				R. Little			
No Finger Found	**3**				**3**				**22**				**7**				**35**			
	L	**R**	**T**	**B**	**L**	**R**	**T**	**B**	**L**	**R**	**T**	**B**	**L**	**R**	**T**	**B**	**L**	**R**	**T**	**B**
MN <= d < 0	-6.58	-7.54	-5.27	-17.34	-5.52	-6.42	-6.58	-12.57	-5.19	-6.36	-6.30	-18.40	-5.18	-7.80	-6.52	-19.45	-5.68	-7.27	-7.83	-15.24
#	73	300	3983	5260	1448	1280	12371	1102	1375	1595	11723	1534	629	2732	11201	1827	1912	5426	11084	2614
0 <= d <= MX	30.30	24.49	12.82	38.66	15.05	16.57	8.34	39.07	16.22	16.30	9.40	45.03	18.11	14.76	9.13	45.90	13.44	11.21	8.87	33.01
#	24273	24042	20384	18769	23489	23609	12557	23739	23554	23280	13211	23198	24292	22146	13722	22785	22964	19398	13825	22159
MN-32 <= d < MN	-55.00	-42.91	-81.36	-76.79	-40.14	-38.17	-80.83	-77.56	-34.00	-38.22	-91.00	-77.35	-37.00	-36.70	-85.83	-76.72	-42.50	-38.43	-74.00	-77.78
#	3	11	11	205	7	6	6	16	1	9	3	67	1	46	6	145	8	90	8	72
MX < d <= MX+32	70.09	77.09	76.25	139.80	#DIV/0!	71.27	78.50	138.84	75.25	71.02	80.50	137.46	76.50	80.20	88.50	139.60	76.04	69.50	66.67	139.61
#	32	29	4	60	0	48	6	60	4	48	9	98	8	5	2	122	35	4	3	36
d < MN-32	-382.40	-839.57	-504.04	-204.12	-255.37	-216.25	-680.78	-196.92	-334.00	-811.44	-504.14	-383.98	-430.57	-662.00	-465.71	-167.78	-236.50	-922.91	-389.00	-561.03
#	26	15	34	81	19	2	18	24	15	18	7	41	14	20	17	51	6	44	2	66
d > MX+32	468.43	661.46	893.92	601.22	137.20	374.24	297.60	710.61	515.95	396.33	298.73	311.17	446.13	368.47	192.80	367.53	836.47	154.83	625.93	219.62
#	15	25	6	47	5	23	10	27	19	18	15	30	24	19	20	38	43	6	46	21
Total #	24422	24422	24422	24422	24968	24968	24968	24968	24968	24968	24968	24968	24968	24968	24968	24968	24968	24968	24968	24968
Average	30.06	24.25	9.34	26.15	13.66	15.79	0.56	37.45	15.22	14.61	2.07	40.78	17.70	11.94	1.91	40.92	13.40	5.41	2.54	26.38
Std Dev	23.01	39.69	29.10	53.06	12.47	16.60	24.75	38.37	19.57	27.49	17.14	39.39	20.79	32.49	18.77	38.77	41.08	52.97	31.13	47.21

	L. Thumb				L. Index				L. Middle				L. Ring				L. Little			
No Finger Found	**26**				**0**				**9**				**14**				**19**			
	L	**R**	**T**	**B**	**L**	**R**	**T**	**B**	**L**	**R**	**T**	**B**	**L**	**R**	**T**	**B**	**L**	**R**	**T**	**B**
MN <= d < 0	-8.12	-5.40	-5.09	-16.44	-4.97	-8.57	-6.67	-12.66	-4.78	-7.12	-6.56	-18.11	-4.87	-7.25	-6.32	-19.02	-7.67	-7.30	-8.24	-14.70
#	171	129	5548	5718	1165	2424	12213	1124	1494	1831	11640	1685	1908	1176	10772	2458	4396	2468	10624	3077
0 <= d <= MX	27.60	26.08	11.76	37.55	16.41	15.28	8.34	40.22	16.23	16.83	9.50	45.50	15.16	18.39	10.24	42.80	12.30	14.34	9.85	30.95
#	24121	24226	18818	18335	23721	22439	12708	23726	23410	23069	13285	23048	23006	23709	14141	22203	20505	22351	14277	21719
MN-32 <= d < MN	-48.67	-46.13	-69.25	-77.79	-49.50	-37.15	-79.94	-77.09	-36.75	-36.25	-74.14	-77.53	-43.00	-40.07	-74.00	-76.67	-40.50	-38.65	-85.20	-73.98
#	12	8	4	147	2	65	8	35	4	16	7	83	2	14	3	158	15	46	5	41
MX < d <= MX+32	73.71	74.75	#DIV/0!	143.34	72.41	66.00	81.50	140.82	73.75	70.30	78.00	141.13	72.20	69.56	72.75	138.83	82.36	74.37	#DIV/0!	140.43
#	77	4	0	65	44	2	2	28	16	10	5	63	5	25	4	48	11	49	0	35
d < MN-32	-444.04	-410.14	-391.29	-410.10	-890.50	-270.05	-550.20	-196.09	-665.50	-507.33	-340.93	-249.59	-416.58	-586.86	-536.54	-264.03	-247.04	-268.45	-501.31	-625.92
#	13	43	26	96	13	20	15	23	7	30	22	29	13	29	23	64	13	30	31	39
d > MX+32	306.38	650.79	842.35	312.57	312.05	1080.54	238.33	574.41	428.82	924.81	204.60	381.76	386.37	620.77	301.19	484.80	182.65	341.58	684.04	462.80
#	28	12	26	61	19	14	18	28	33	8	5	56	30	11	21	33	24	20	27	53
Total #	24422	24422	24422	24422	24964	24964	24964	24964	24964	24964	24964	24964	24964	24964	24964	24964	24964	24964	24964	24964
Average	27.53	25.44	8.37	23.43	15.26	13.20	0.81	38.17	15.36	14.72	1.73	41.45	13.86	16.77	2.84	35.94	8.81	12.14	2.23	25.19
Std Dev	19.75	29.46	34.67	57.72	26.00	31.14	20.21	34.89	23.16	29.40	17.22	39.52	20.73	29.88	24.73	44.42	14.27	20.01	33.64	47.75

Appendix D. Plots of 3-inch segmentation box centers.

The plots in this appendix show the distribution of the segmentation box centers (x,y) for the 3-inch data. There is a combined plot for each slap image and then a smaller plot for each finger position. The individual finger plots are better for seeing the full “spread” of x,y positions detected. The plot for the ground truth (GT) is included as a baseline for comparison. The blank lines that appear in some of the plots are most likely caused by the segmentation algorithm doing some level of sampling of the input image. The reason the lines are not evenly distributed in some plots is an artifact of the sampling when scaling the images for displaying in the report.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Center X
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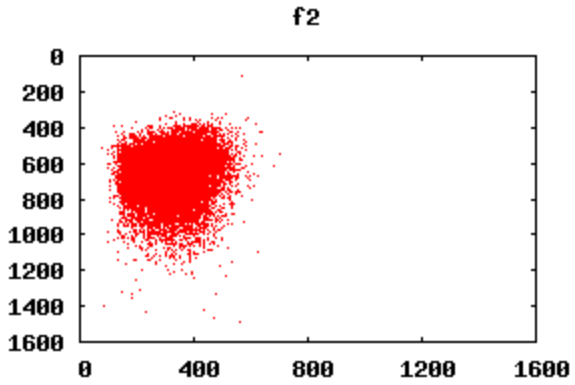
"f3"
"f4"
"f5"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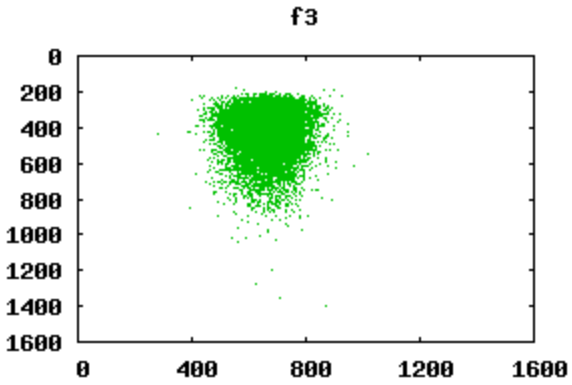
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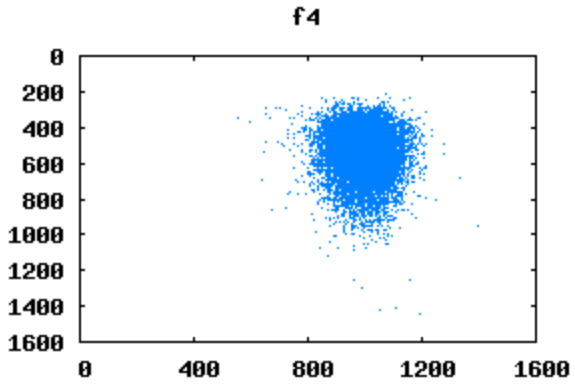
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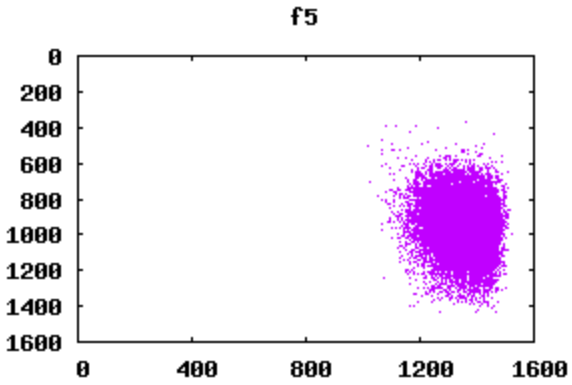
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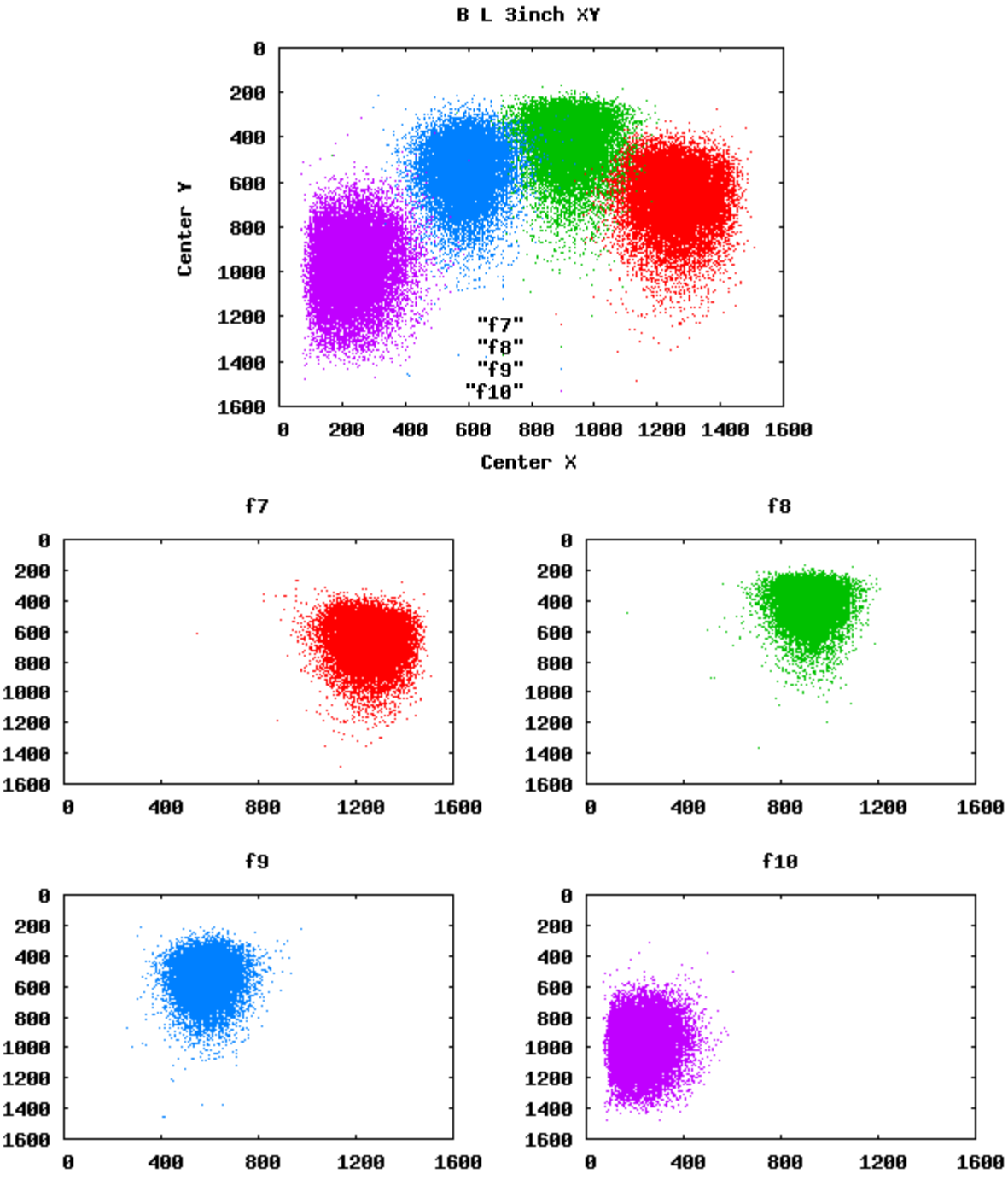
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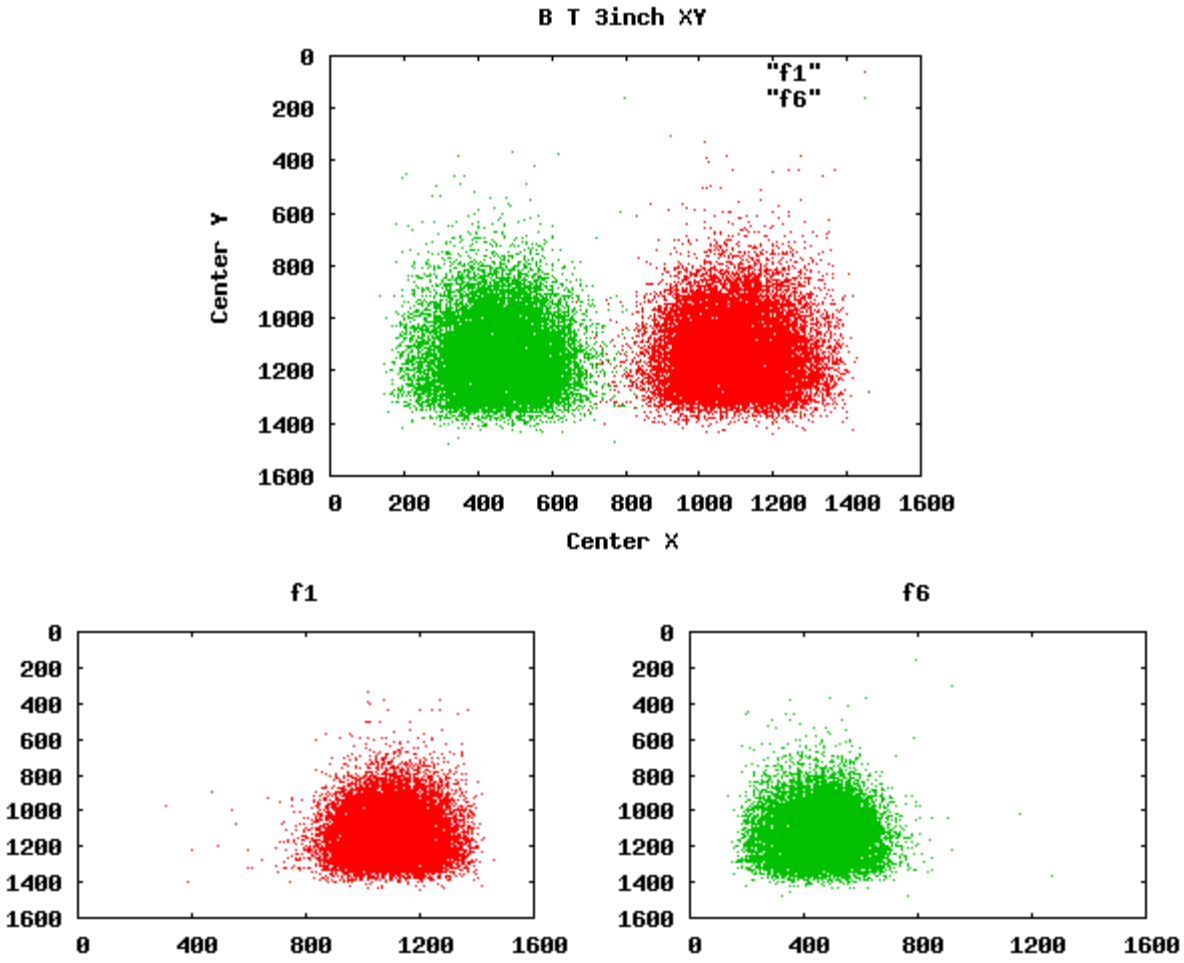
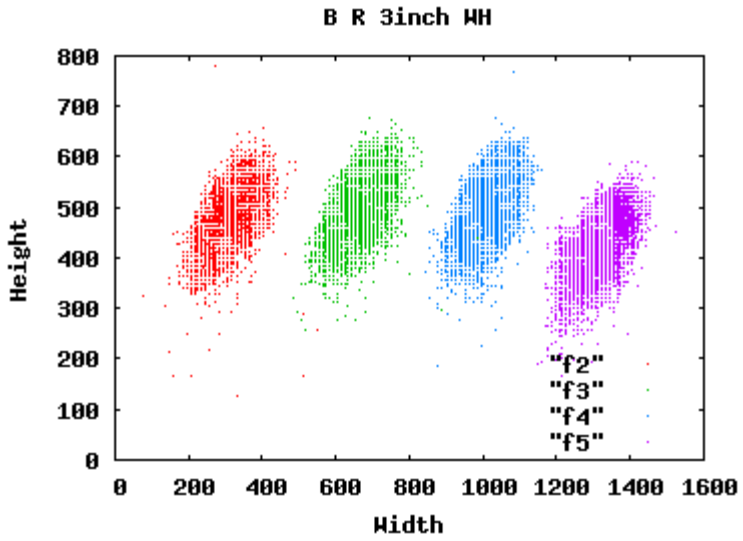
Center Y
Center X
"f7"
"f8"
"f9"
"f10"
f7
f8
f9
f10


B T 3inch XY
"f1"
"f6"
Center Y
Center X
0
200
400
600
800
1000
1200
1400
1600
f1
f6

Appendix E. Plots of 3-inch segmentation box widths and heights.

The plots in this appendix show the distribution of the segmentation box widths and heights for the 3-inch data. There is a combined plot for each slap image and then a smaller plot for each finger position. The individual finger plots are better for seeing the full "spread" of widths and heights detected. The widths are "spread out" on the plot by adding 350, 750 and 1050 to the 2nd, 3rd, and 4th widths plotted. The plot for the ground truth (GT) is included as a baseline for comparison. The blank lines that appear in some of the plots are most likely caused by the segmentation algorithm doing some level of sampling of the input image. The reason the lines are not evenly distributed in some plots is an artifact of the sampling when scaling the images for displaying in the report.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Height
Width
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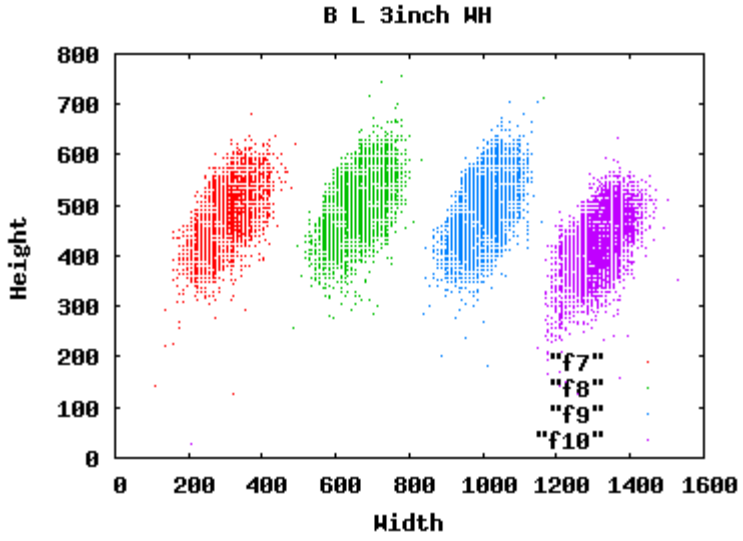
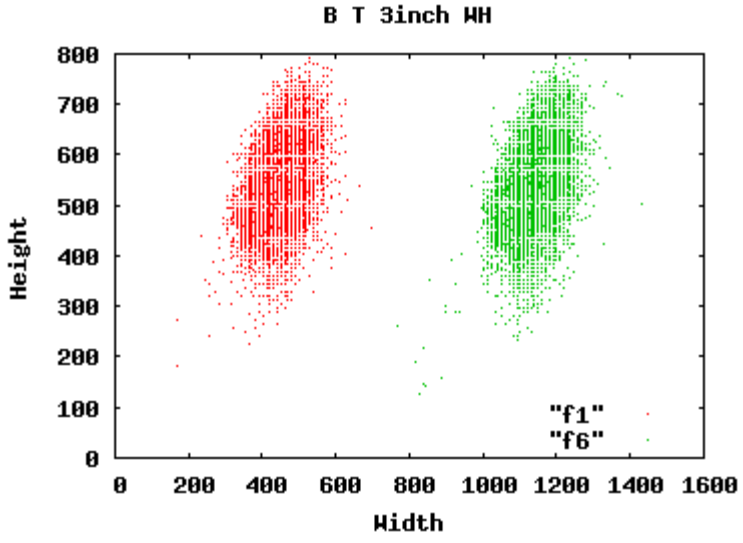
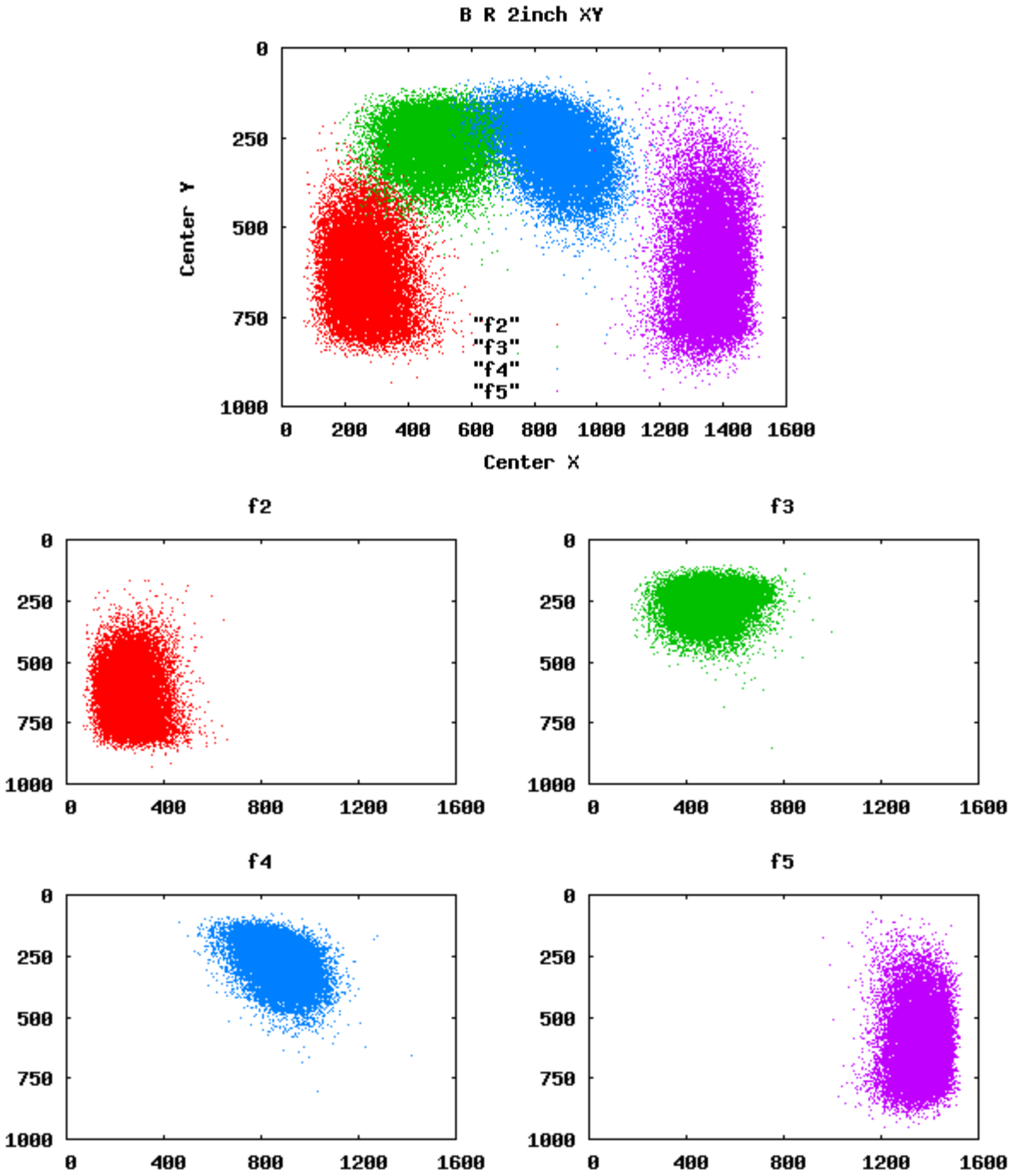
"f3"
"f4"
"f5"


B L 3inch WH
Height
Width
"f7"
"f8"
"f9"
"f10"


B T 3inch WH
Height
Width
"f1"
"f6"

Appendix F. Plots of 2-inch segmentation box centers.

The plots in this appendix show the distribution of the segmentation box centers (x,y) for the 2-inch data. There is a combined plot for each slap image and then a smaller plot for each finger position. The individual finger plots are better for seeing the full “spread” of x,y positions detected. The plot for the ground truth (GT) is included as a baseline for comparison. The blank lines that appear in some of the plots are most likely caused by the segmentation algorithm doing some level of sampling of the input image. The reason the lines are not evenly distributed in some plots is an artifact of the sampling when scaling the images for displaying in the report.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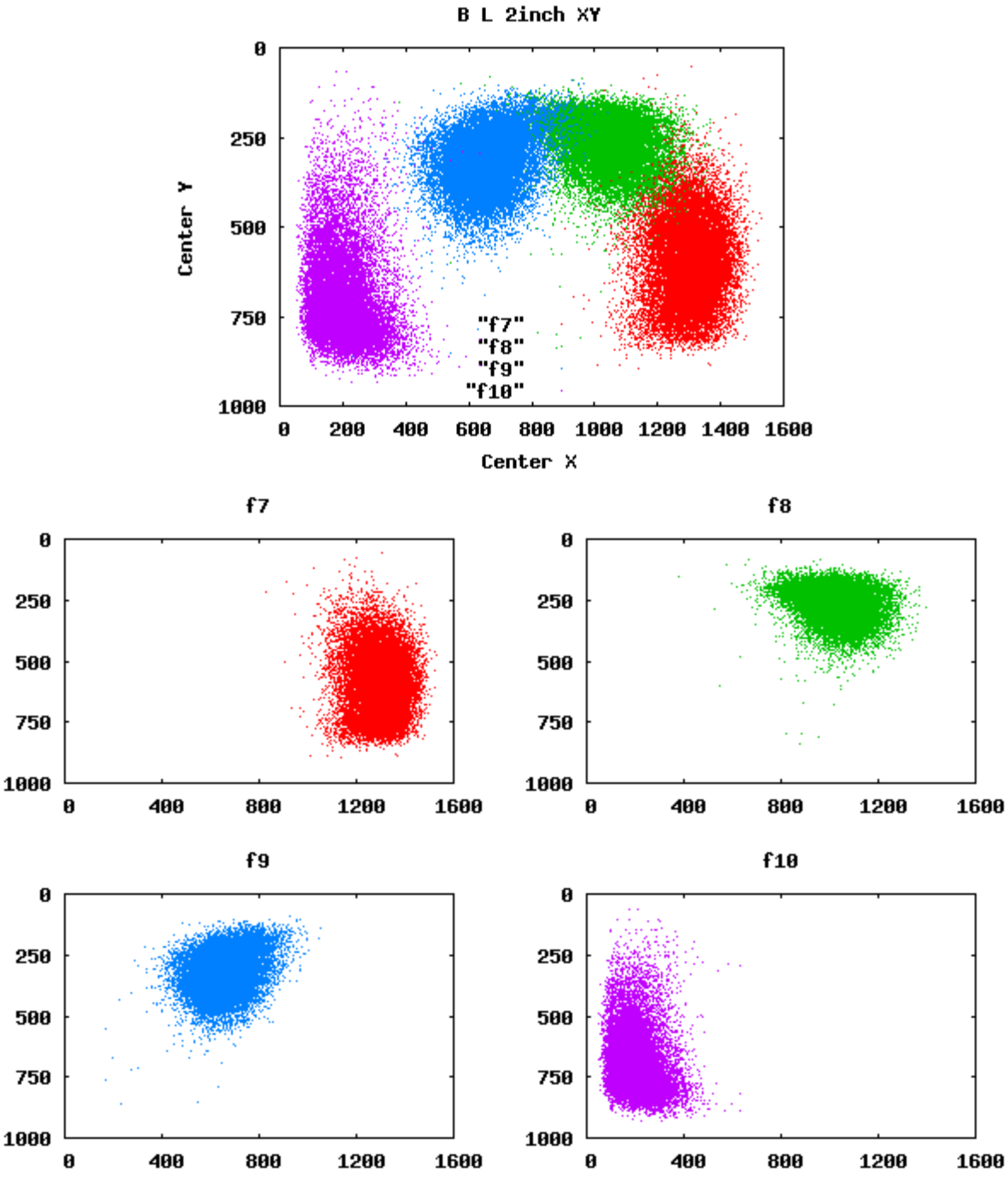
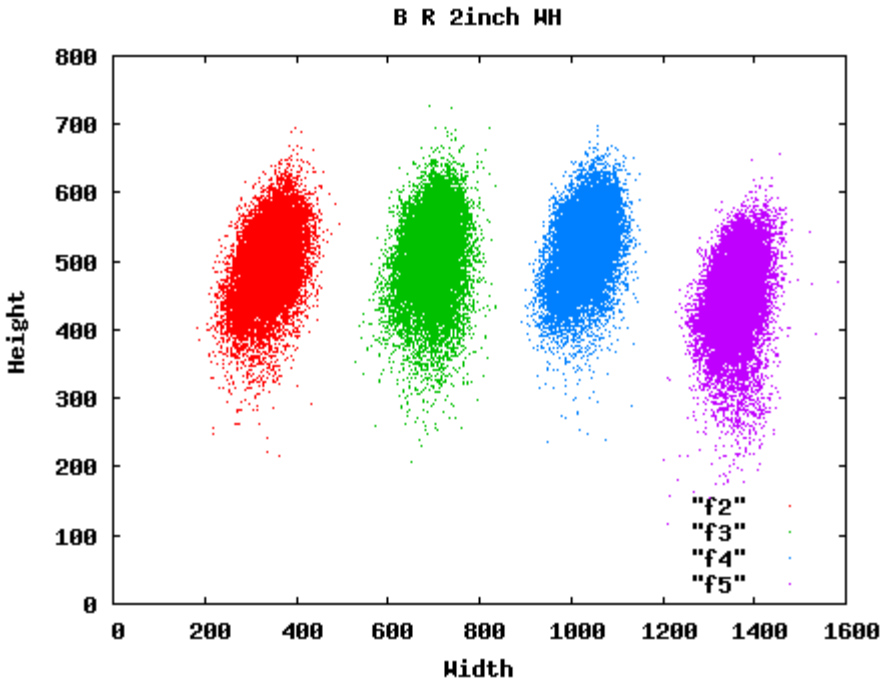
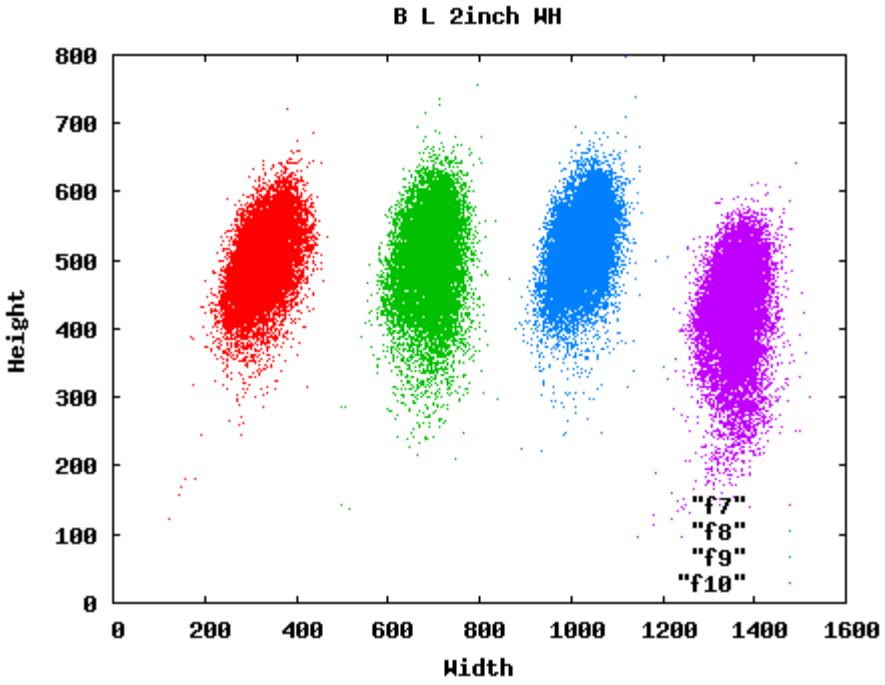
Center Y
Center X
0
250
500
750
1000
0
200
400
600
800
1000
1200
1400
1600
"f2"
"f3"
"f4"
"f5"
f2
f3
f4
f5
0
400
800
1200
1600


B L 2inch XY
Center Y
Center X
0
250
500
750
1000
0
200
400
600
800
1000
1200
1400
1600
"f7"
"f8"
"f9"
"f10"
f7
f8
f9
f10

Appendix G. Plots of 2-inch segmentation box widths and heights.

The plots in this appendix show the distribution of the segmentation box widths and heights for the 2-inch data. There is a combined plot for each slap image and then a smaller plot for each finger position. The individual finger plots are better for seeing the full "spread" of widths and heights detected. The widths are "spread out" on the plot by adding 350, 750 and 1050 to the 2nd, 3rd, and 4th widths plotted. The plot for the ground truth (GT) is included as a baseline for comparison. The blank lines that appear in some of the plots are most likely caused by the segmentation algorithm doing some level of sampling of the input image. The reason the lines are not evenly distributed in some plots is an artifact of the sampling when scaling the images for displaying in the report.


B R 2inch WH
Height
Width
0
100
200
300
400
500
600
700
800
0
200
400
600
800
1000
1200
1400
1600
"f2"
"f3"
"f4"
"f5"


B L 2inch WH
Height
Width
0
100
200
300
400
500
600
700
800
0
200
400
600
800
1000
1200
1400
1600
"f7"
"f8"
"f9"
"f10"