



C

Image Level results: [edit](#)

Cellular Analysis(Dig.Ph.Cont_70)

Well	Value
Well ID	G2
Name	
Cell Count	115
Object Sum Area	3.51E+05
Cell Count(Subpopulation 1)	84
Object Sum Area(Subpopulation 1)	2.77E+05

Object: [edit](#) [Select All](#)

Subpopulation 1

#	Size	Area	Mean	Circula	StdDev	Ph
52	57.8	2.62E+03	20188	0.934	5147	3
100	40.7	1.27E+03	21492	0.546	5669	4
94	45.0	1.5E+03	23931	0.373	5695	3
93	98.5	7.59E+03	23221	0.794	5816	4
8	72.2	4.09E+03	20430	0.898	5824	3
22	58.9	2.72E+03	24689	0.838	5839	3
50	45.2	1.54E+03	24192	0.433	6077	3
63	54.6	2.1E+03	22847	0.250	6090	3
95	40.4	1.08E+03	23324	0.88	6156	3
87	64.9	3.3E+03	25442	0.823	6171	4
88	53.2	2.2E+03	23341	0.657	6205	3
86	41.4	1.26E+03	24932	0.254	6391	3
67	43.5	1.42E+03	20025	0.44	6471	3

Highlight pixels: Saturated In the well In ROI Customized mask...

NAVIGATION

Time: ROI:

Read: (0:00:00)

Slide show: Image delay: sec Loop

Highlight objects: In the Plug Threshold outliers

Channel: Dig.Ph.Cont_70

Mask: Primary Secondary

Spots: Center Threshold outliers

Show index Fill objects

1000 μm