## **QC Report - Agilent Technologies : 2 Color Gene Expression**

Image Protocol

Thursday, May 24, 2018 - 11:35 BG Method SG15074465\_257480913948\_S001 [2\_2] Background Detrend GE2\_1200\_Jun14\_PEG200\_JRmod Multiplicative Detrend

(Editable)

NA

NA

No Background  $On (Feat NCR ange, \, LoP ass)$ 

User Name Grid FE Version Sample(red/green) DyeNorm List

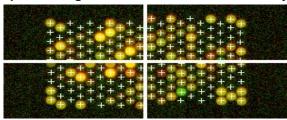
No of Probes in DyeNorm List

ExternalFullGEML\_074809\_D\_F\_20171030 Linear DyeNorm Factor

Admin Dye Norm 12.0.1.1 Additive Error Saturation Value

Linear 11.7(Red) 21.8(Green) 17(Red)38(Green) 776684 (r), 777880 (g)

#### Spot Finding of the Four Corners of the Array

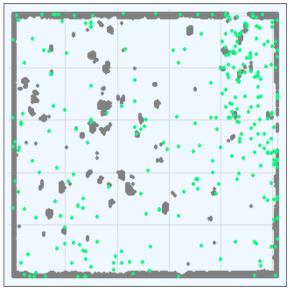


#### **Grid Normal**

	Feature	Feature		Local Background	
	Red	Green	Red	Green	
Non Uniform	1	0	1543	0	
Population	322	285	3658	3561	

## Spatial Distribution of All Outliers on the Array

384 rows x 164 columns



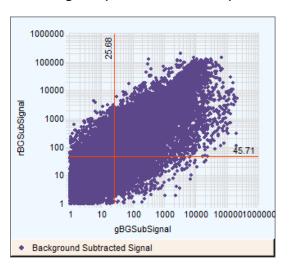
#### **Net Signal Statistics**

	Red	Green
# Saturated Features	0	0
99% of Sig. Distrib.	57383	12963
50% of Sig. Distrib.	57	46
1% of Sig. Distrib.	9	17

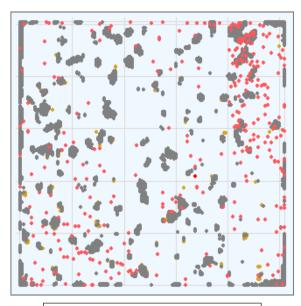
#### **Negative Control Stats**

	Red	Green
Average Net Cincele	0.00	40.54
Average Net Signals	9.66	18.51
StdDev Net Signals	1.16	1.70
Average BG Sub Signal	-1.56	-2.29
StdDev BG Sub Signal	1.15	1.62

#### **Red and Green Background Corrected** Signals (Non-Control Inliers)



# Features (NonCtrl) with BGSubSignals < 0: 3799 (Red); 5388 (Green)



# FeatureNonUnif (Red or Green) = 1(0.00%) # GeneNonUnif (Red or Green) = 1 (0.002 %)

BG NonUniform

**BGSubSignal** ProcessedSignal

- •Red FeaturePopulation Green Feature Population

- BG PopulationRed Feature NonUniformGreen Feature NonUniform

#### Local Bkg (inliers)

	Red	Green
Number	59180	59415
Number	59160	59415
Avg	23.31	24.95
SD	1.97	0.91

#### **Foreground Surface Fit**

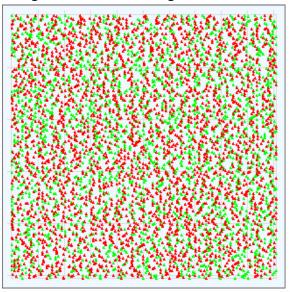
	Red	Green
RMS_Fit	0.23	0.58
RMS_Resid	1.49	1.75
Avg_Fit	21.23	30.81

### Reproducibility: %CV for Replicated Probes

Median %CV Signal (inliers)

	ian 700 v Olginai (ilillolo)
	Non-Control probes
Red	Green
5.05	5.76
5.05	5.70
5.05	3.57

#### Spatial Distribution of Significantly Up-Regulated and Down-Regulated Features

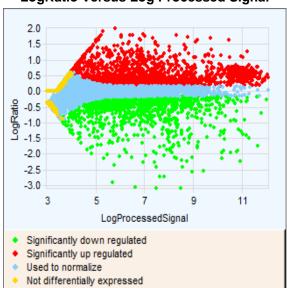


#Up-Regulated:15199 (Red); #Down-Regulated:12797 (Green)

Up-Regulated Down-Regulated

Up Reg: 25.63% of NonCtrl Features : Random (Value 0.81) Down Reg: 21.58% of NonCtrl Features: Random (Value 1.08)

#### **LogRatio Versus Log Processed Signal**



## **Array Uniformity: LogRatios**

Non-Control

AbsAvgLogRatio	0.55
AverageS/N	29.80

# Evaluation Metrics for GE2\_QCMT\_Jun14 Good (8); Evaluate (1)

Metric Name	Value	Excellent	Good	Evaluate
IsGoodGrid	1.00		>1	<1
AnyColorPrcntFeatNonU	0.00		<1	>1
gNegCtrlAveBGSubSig	-2.29		-20 to 10	<-20 or >10
gNegCtrlSDevBGSubSig	1.62		<15	>15
rNegCtrlAveBGSubSig	-1.56		-20 to 4	<-20 or >4
rNegCtrlSDevBGSubSig	1.15		<6	>6
gNonCntrlMedCVBkSubSi.	5.76		0 to 18	<0 or >18
rNonCntrlMedCVBkSubSi	5.05		0 to 18	<0 or >18
gE1aMedCVBkSubSignal	-		0 to 18	<0 or >18
rE1aMedCVBkSubSignal	-		0 to 18	<0 or >18
absE1aObsVsExpCorr	-		>0.86	<0.86
absE1aObsVsExpSlope	-		>0.85	<0.85
LogRatioImbalance	-16.33	-0.26 to 0.26	(-0.75 to -0.2	<-0.75 or >0.75

Excellent
Good
Evaluate