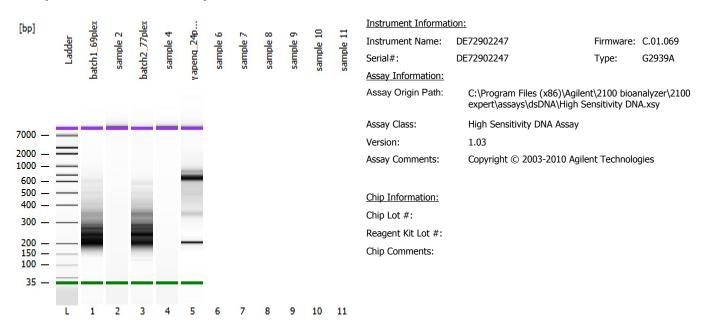
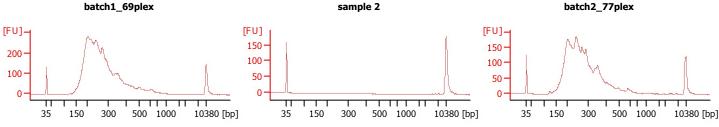
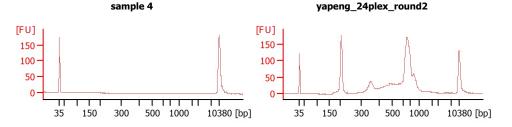
Assay Class: High Sensitivity DNA Assay Created: 4/20/2020 12:02:44 PM Data Path: T:\...gh Sensitivity DNA Assay_DE72902247_2020-04-20_12-02-44.xad Created: 4/20/2020 12:39:01 PM

Electrophoresis File Run Summary







Printed:

4/20/2020 12:39:32 PM

Printed: 4/20/2020 12:39:32 PM

High Sensitivity DNA Assay T:\...gh Sensitivity DNA Assay_DE72902247_2020-04-20_12-02-44.xad Assay Class: Created: 4/20/2020 12:02:44 PM Data Path: Modified: 4/20/2020 12:39:01 PM **Electrophoresis File Run Summary (Chip Summary)** Sample Rest. **Stat Observation** Result **Result Color Sample Name** Comment Digest us Label batch1_69plex sample 2 batch2_77plex sample 4 yapeng_24plex_round sample 6

Chip Lot # Reagent Kit Lot #

Chip Comments:

sample 7 sample 8 sample 9 sample 10 sample 11 Ladder

Printed: 4/20/2020 12:39:32 PM

Assay Class: High Sensitivity DNA Assay Created: 4/20/2020 12:02:44 PM Data Path: T:\...gh Sensitivity DNA Assay_DE72902247_2020-04-20_12-02-44.xad Created: 4/20/2020 12:39:01 PM

Electrophoresis Assay Details

General Analysis Settings

Number of Available Sample and Ladder Wells (Max.): 12

Minimum Visible Range [s]: 32
Maximum Visible Range [s]: 138
Start Analysis Time Range [s]: 33
End Analysis Time Range [s]: 137.5
Ladder Concentration [pg/µl]: 1950
Uses Standard Area for Ladder Fragments
Lower Marker Concentration [pg/µl]: 125
Upper Marker Concentration [pg/µl]: 75
Used Upper Marker for Quantitation
Standard Curve Fit is Point to Point
Show Data Aligned to Lower and Upper Marker

Integrator Settings

Integration Start Time [s]: 33.05
Integration End Time [s]: 137
Slope Threshold: 0.8
Height Threshold [FU]: 5
Area Threshold: 0.1
Width Threshold [s]: 0.6
Baseline Plateau [s]: 0.5

Filter Settings

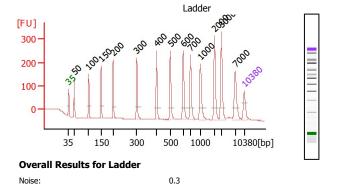
Filter Width [s]: 0.5 Polynomial Order: 4

Ladder

Ladder Peak	Size	Area
1	35	160
2	50	210
3	100	208
4	150	221
5	200	242
6	300	270
7	400	305
8	500	306
9	600	336
10	700	321
11	1000	366
12	2000	413
13	3000	411
14	7000	400
15	10380	214

Assay Class: High Sensitivity DNA Assay Created: 4/20/2020 12:02:44 PM Data Path: T:\...gh Sensitivity DNA Assay_DE72902247_2020-04-20_12-02-44.xad Created: 4/20/2020 12:02:49 PM

Electropherogram Summary



Peak table for Ladder						
Peak		Size [bp]	Conc. [pg/µl]	Molarity [pmol/l]	Observation s	
1	•	35	125.00	5,411.3	Lower Marker	
2	L	50	150.00	4,545.5	Ladder Peak	
3	L	100	150.00	2,272.7	Ladder Peak	
4	L	150	150.00	1,515.2	Ladder Peak	
5	L	200	150.00	1,136.4	Ladder Peak	
6	L	300	150.00	757.6	Ladder Peak	
7	L	400	150.00	568.2	Ladder Peak	
8	L	500	150.00	454.5	Ladder Peak	
9	L	600	150.00	378.8	Ladder Peak	
10	L	700	150.00	324.7	Ladder Peak	
11	L	1,000	150.00	227.3	Ladder Peak	
12	L	2,000	150.00	113.6	Ladder Peak	
13	L	3,000	150.00	75.8	Ladder Peak	

Peak		Size [bp] Conc.		Molarity	Observation
14	L	7,000	[pg/μl] 150.00	[pmol/l] 32.5	s Ladder Peak
15		10,380	75.00	10.9	Upper Marker

4/20/2020 12:02:44 PM

4/20/2020 12:39:01 PM

Created:

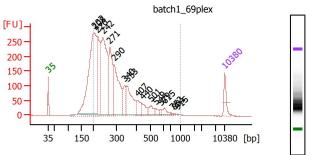
Modified:

batch1_69plex

High Sensitivity DNA Assay Assay Class:

T:\...gh Sensitivity DNA Assay_DE72902247_2020-04-20_12-02-44.xad Data Path:

Electropherogram Summary Continued ...



Overall Results for sample 1: batch1_69plex

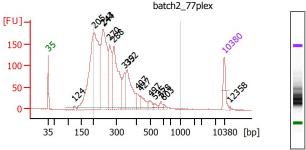
Number of peaks found: 17 0.4 Noise: Corr. Area 1: 5,221.0

Region table for sample 1: batch1_69plex

From To Corr. % of Average Size Conc. Molarit Co [bp] [bp] Area Total Size distribution [pg/µl y [pmol/l r [bp] in CV [%] 1,000 5,221.0 81 301 34.9 4,348.33 24,333.4

Peak	Size [bp	o] Conc. [pg/µl]	Molarity [pmol/l]	Observation s
1	35	125.00	5,411.3	Lower Marker
2	207	6,389.13	46,766.0	
3	208	1,745.85	12,695.5	
4	220	570.39	3,925.5	
5	242	1,072.53	6,704.3	
6	271	569.74	3,191.0	
7	290	569.03	2,977.3	
8	340	155.13	691.7	
9	353	329.00	1,410.4	
10	407	99.76	371.7	
11	440	73.76	253.9	
12	501	67.05	202.7	
13	539	31.16	87.6	
14	579	16.37	42.8	
15	615	47.51	117.1	
16	752	4.41	8.9	
17	801	3.88	7.3	
18	915	2.04	3.4	
19	10,380	75.00	10.9	Upper Marker

Peak table for sample 1:



Overall Results for sample 3: batch2_77plex

Number of peaks found: 15 Noise: 0.3 3,535.2 Corr. Area 1:

Region table for sample 3: batch2_77plex

From To Corr. % of Average Size Conc. Molarit Co distribution [pg/µl y [bp] [bp] Area Total Size lo in CV [%]] [pmol/l r [bp] 31.1 3,140.93 17,416.6 1,000 3,535.2 83 299 200

Peak table for sample 3:				batch2_77plex		
Peak		Size [bp] Conc. [pg/µl]	Molarity [pmol/l]	Observation s	
1	•	35	125.00	5,411.3	Lower Marker	
2		243	3,996.45	24,933.3		
3		124	17.72	216.6		
4		205	1,413.70	10,473.3		
5		244	836.17	5,187.2		
6		270	348.50	1,958.2		
7		288	522.17	2,750.3		
8		339	139.67	624.2		
9		352	307.54	1,324.7		
10		407	55.98	208.3		
11		427	73.19	259.7		
12		497	36.08	110.0		
13		535	17.74	50.3		
14		578	20.70	54.2		
15		603	11.24	28.2		
16		10,380	75.00	10.9	Upper Marker	
17		12,358	0.00	0.0		

