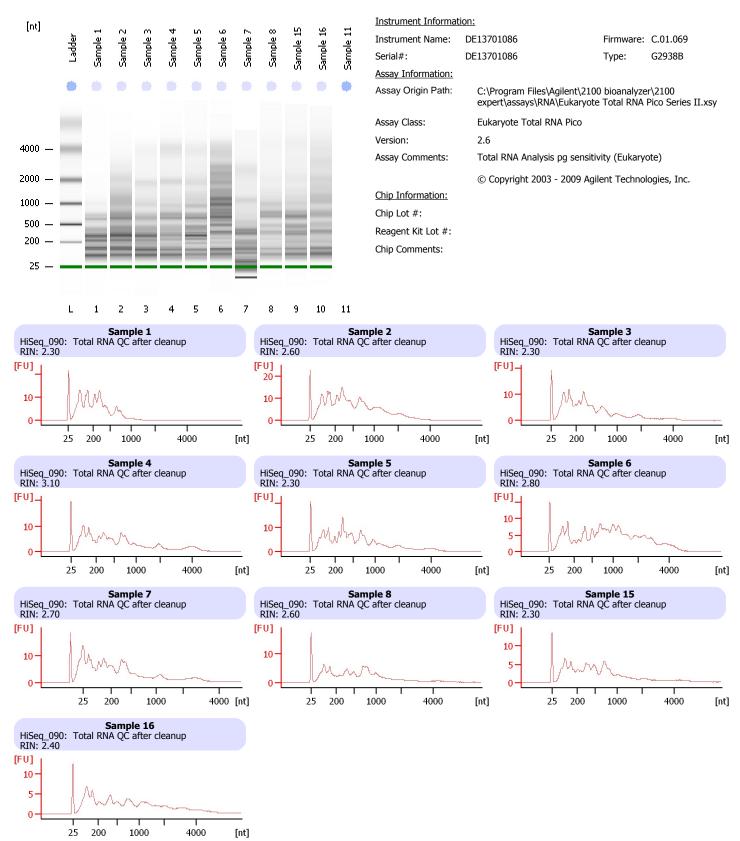
2015-05-14_002.xad Page 1 of 16

Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Cceated: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electrophoresis File Run Summary



5/14/2015 11:41:38 AM

Printed:

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Status	Result Label	Result Color
Sample 1	HiSeq_090: Total RNA QC after cleanup	~	RIN: 2.30	
Sample 2	HiSeq_090: Total RNA QC after cleanup	~	RIN: 2.60	
Sample 3	HiSeq_090: Total RNA QC after cleanup	~	RIN: 2.30	
Sample 4	HiSeq_090: Total RNA QC after cleanup	~	RIN: 3.10	
Sample 5	HiSeq_090: Total RNA QC after cleanup	~	RIN: 2.30	
Sample 6	HiSeq_090: Total RNA QC after cleanup	~	RIN: 2.80	
Sample 7	HiSeq_090: Total RNA QC after cleanup	~	RIN: 2.70	
Sample 8	HiSeq_090: Total RNA QC after cleanup	~	RIN: 2.60	
Sample 15	HiSeq_090: Total RNA QC after cleanup	~	RIN: 2.30	
Sample 16	HiSeq_090: Total RNA QC after cleanup	~	RIN: 2.40	
Sample 11				
Ladder		~	All Other Samples	

Chip Lot # Reagent Kit Lot #

Chip Comments:

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Cc:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-14\2015-05-1

Electrophoresis Assay Details

General Analysis Settings

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

Minimum Visible Range [s]: 17

Maximum Visible Range [s]: 70

Start Analysis Time Range [s]: 18

End Analysis Time Range [s]: 69

Ladder Concentration [pg/µl]: 1000

Lower Marker Concentration [pg/µl]: 0

Upper Marker Concentration [pg/µl]: 0

Used Lower Marker for Quantitation

Standard Curve Fit is Logarithmic

Show Data Aligned to Lower Marker

Integrator Settings

Integration Start Time [s]: 18 Integration End Time [s]: 69 Slope Threshold: 0.6 Height Threshold [FU]: 0.5 Area Threshold: 0.2 Width Threshold [s]: 0.3 Baseline Plateau [s]: 6

Filter Settings

Filter Width [s]: 0.5 Polynomial Order: 4

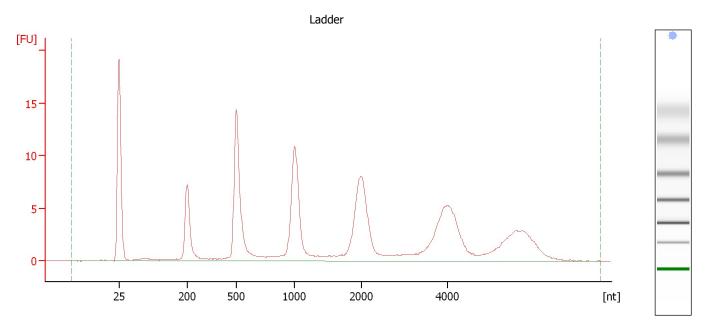
Ladder

Ladder Peak	Size
1	25
2	200
3	500
4	1000
5	2000
6	4000

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electropherogram Summary



Overall Results for Ladder

RNA Area: 150.3 RNA Concentration: 1,000 pg/µl Result Flagging Color: Result Flagging Label:

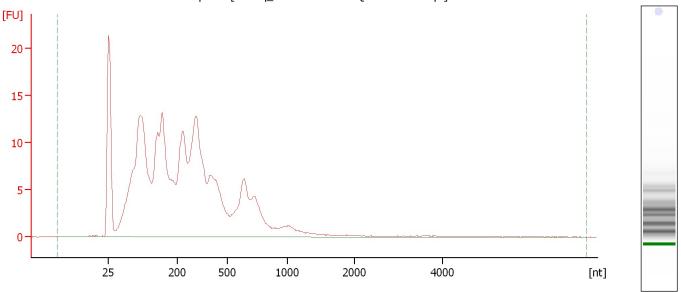
All Other Samples

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electropherogram Summary Continued ...

Sample 1 [HiSeq_090: Total RNA QC after cleanup]



Overall Results for sample 1 : Sample 1

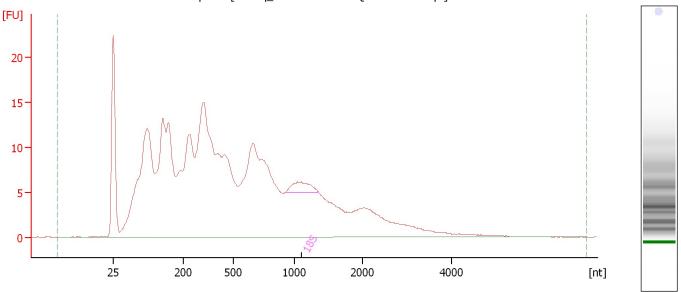
RNA Area: 302.8 RNA Concentration: 2,015 pg/μl rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN): Result Flagging Color: Result Flagging Label: 2.3 (B.02.08) RIN: 2.30

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electropherogram Summary Continued ...

Sample 2 [HiSeq_090: Total RNA QC after cleanup]



Overall Results for sample 2 : <u>Sample 2</u>

Fragment table for sample 2: <u>Sample 2</u>

 Name
 Start Size [nt]
 End Size [nt]
 Area
 % of total Area

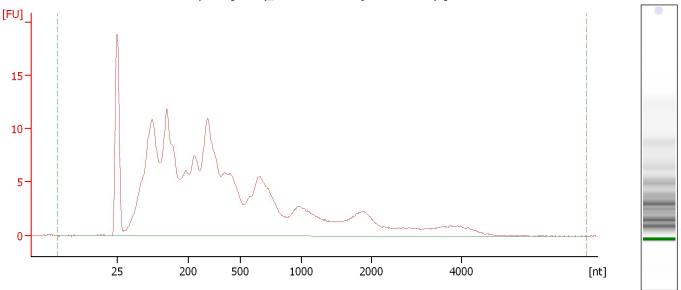
 18S
 923
 1,357
 5.8
 1.2

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electropherogram Summary Continued ...

Sample 3 [HiSeq_090: Total RNA QC after cleanup]



Overall Results for sample 3: Sample 3

RNA Area: 313.7 RNA Concentration: 2,087 pg/μl rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN): Result Flagging Color: Result Flagging Label:

RIN: 2.30

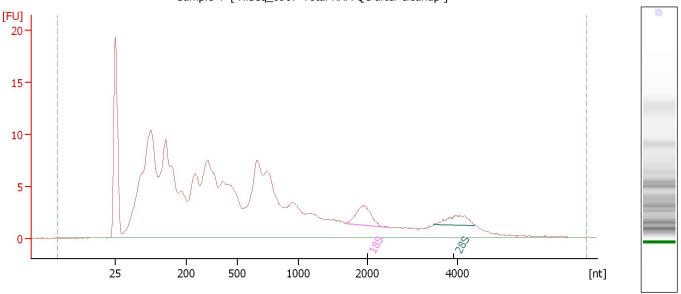
2.3 (B.02.08)

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electropherogram Summary Continued ...

Sample 4 [HiSeq_090: Total RNA QC after cleanup]



Overall Results for sample 4: Sample 4

Fragment table for sample 4: Sample 4

Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	1,675	2,475	6.4	2.0
28S	3,458	4,371	3.7	1.2

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electropherogram Summary Continued ...

Sample 5 [HiSeq_090: Total RNA QC after cleanup]

[FU]
20155-

Overall Results for sample 5 : <u>Sample 5</u>

200

500

1000

RNA Area: 347.2 RNA Concentration: 2,310 pg/μl rRNA Ratio [28s / 18s]: 0.0

25

RNA Integrity Number (RIN): Result Flagging Color: Result Flagging Label:

2000

2.3 (B.02.08) RIN: 2.30 [nt]

Printed: 5/14/2015 11:41:38 AM

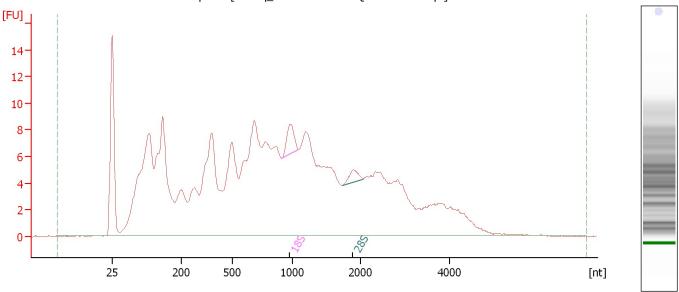
4000

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Assay Class: 5/14/2015 11:08:21 AM 5/14/2015 11:29:50 AM Eukaryote Total RNA Pico Created: C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Data Path: Modified:

Electropherogram Summary Continued ...

Sample 6 [HiSeq_090: Total RNA QC after cleanup]



Overall Results for sample 6: Sample 6

RNA Area: 383.1 RNA Integrity Number (RIN): 2.8 (B.02.08) **RNA Concentration:** 2,550 pg/µl Result Flagging Color: rRNA Ratio [28s / 18s]: 0.4 Result Flagging Label: RIN: 2.80

Fragment table for sample 6: Sample 6

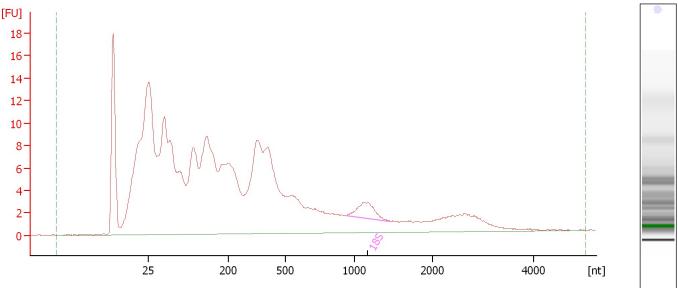
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	911	1,086	4.2	1.1
285	1.730	2.044	1.9	0.5

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electropherogram Summary Continued ...

Sample 7 [HiSeq_090: Total RNA QC after cleanup]



Overall Results for sample 7: Sample 7

Fragment table for sample 7: Sample 7

Name Start Size [nt] End Size [nt] Area % of total Area

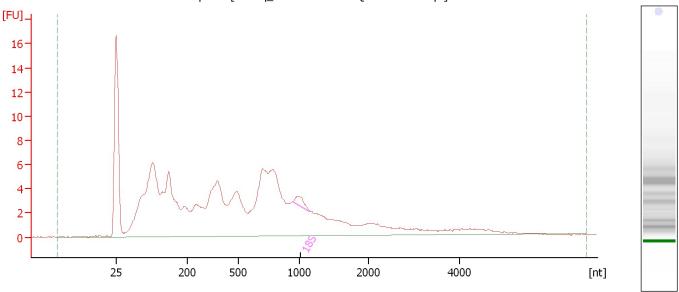
18S 930 1,460 5.8 1.8

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electropherogram Summary Continued ...

Sample 8 [HiSeq_090: Total RNA QC after cleanup]



Overall Results for sample 8 : Sample 8

RNA Area: 192.4 RNA Integrity Number (RIN): 2.6 (B.02.08) RNA Concentration: 1,280 pg/ μ l Result Flagging Color: RNA Ratio [28s / 18s]: 0.0 Result Flagging Label: RIN: 2.60

Fragment table for sample 8: Sample 8

Name Start Size [nt] End Size [nt] Area % of total Area

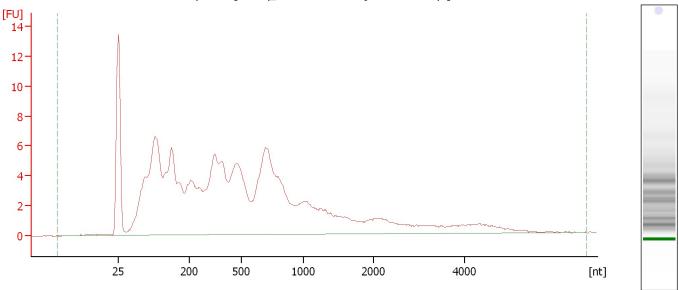
18S 937 1,143 1.7 0.9

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electropherogram Summary Continued ...

Sample 15 [HiSeq_090: Total RNA QC after cleanup]



Overall Results for sample 9: Sample 15

RNA Area: 208.3 RNA Concentration: 1,386 pg/μl rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN): Result Flagging Color: Result Flagging Label:

RIN: 2.30

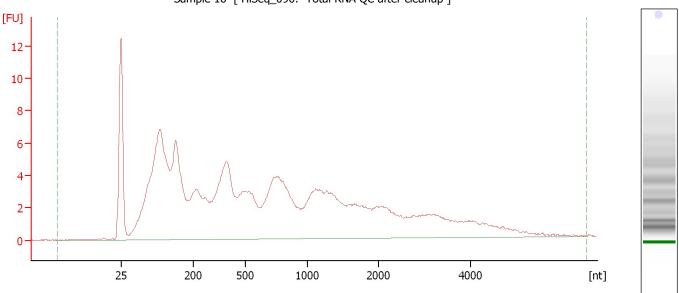
2.3 (B.02.08)

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Assay Class: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: 5/14/2015 11:08:21 AM Modified: 5/14/2015 11:29:50 AM

Electropherogram Summary Continued ...

Sample 16 [HiSeq_090: Total RNA QC after cleanup]



Overall Results for sample 10 : Sample 16

RNA Area: 218.0 RNA Concentration: 1,451 pg/μl rRNA Ratio [28s / 18s]: 0.0 RNA Integrity Number (RIN): Result Flagging Color: Result Flagging Label:

RIN: 2.40

2.4 (B.02.08)

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Assay Class: Data Path: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: Modified: 5/14/2015 11:08:21 AM 5/14/2015 11:29:50 AM **Gel Image** [nt] Sample 16 Sample 11 ٠ ٠ ٠ ۰ ٠ ۰ ٠ ٠ ø • • 4000 — 2000 -1000 — 500 -200 — 25 — 3 5 6 10 11

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Assay Class: Data Path: Eukaryote Total RNA Pico C:\...ents and Settings\Bioanalyzer\2015-05-14\2015-05-14_002.xad Created: Modified: 5/14/2015 11:08:21 AM 5/14/2015 11:29:50 AM

Invalid Samples

Sample 11 has not been run, no results available.