

2/22/2019 2:31:30 PM 2/22/2019 2:54:39 PM

Created: Modified:

Assay Class: Eukaryote Total RNA Pico Data Path: C:\... bioanalyzer\2100 expert\data\2019-02-22\2019-02-22_001.xad Electrophoresis File Run Summary (Chip Summary)

Sample Name	Sample Comment	Status	Result Label
HiSeq_1683 -1 RNA with DNaseI	M0D055413_S_I (1:10)	×	RIN: 2.50
Cleanup			
HiSeq_1683 -2 RNA with DNaseI Cleanup	M0D055408_N_I (1:8)	×	RIN: 5.60
HiSeq_1683 -3 RNA with DNaseI Cleanup	M0D055401_G_I (1:10)	×	RIN: 4.80
HiSeq_1683 -4 RNA with DNaseI Cleanup	M0D055402_H_I (1:10)	×	RIN: 4.50
HiSeq_1683 -5 RNA with DNaseI Cleanup	M0D055414_T_I (1:4)	~	RIN: 2.30
HiSeq_1683 -6 RNA with DNaseI Cleanup	M0D055415_U_I (1:10)	~	RIN: 5.10
HiSeq_1683 -7 RNA with DNaseI Cleanup	M0D055302_L_I (1:40)	~	RIN: 3.20
HiSeq_1683 -8 RNA with DNaseI Cleanup	M0D055342_Z_I (1:20_	×	RIN: 2.20
HiSeq_1683 -91 RNA with DNaseI Cleanup	M0D055310_T_I	×	RIN:1
HiSeq_1683 -10 RNA with DNaseI Cleanup	M0D055330_N_I (1:10)	×	RIN: 2.80
HiSeq_1683 -11 RNA with DNaseI Cleanup	M0D055367_Y_I	×	RIN N/A
Ladder		×	All Other Samples

Chip Lot #

Reagent Kit Lot #

Chip Comments :

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Electrophoresis Assay Details

General Analysis Settings

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

Minimum Visible Range [s] : 17Maximum Visible Range [s] : 70Start Analysis Time Range [s] : 18End Analysis Time Range [s] : 69Ladder Concentration $[pg/\mu I] : 1000$ Lower Marker Concentration $[pg/\mu I] : 0$ Upper Marker Concentration $[pg/\mu I] : 0$ 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

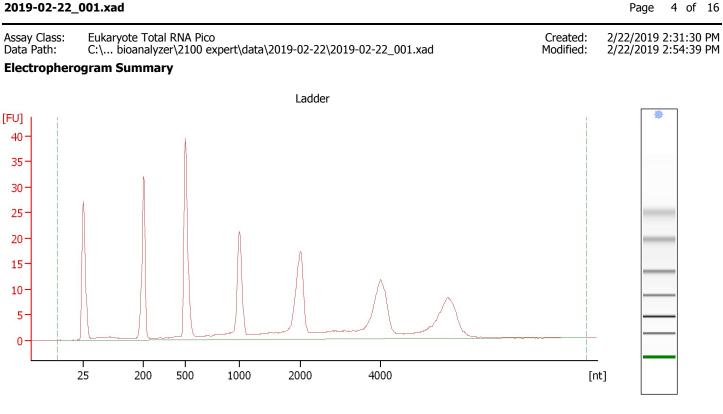
Filter Settings

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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Overall Results for Ladder

RNA Area:	284.8	Result Flagging Color:	
RNA Concentration:	1,000 pg/µl	Result Flagging Label:	All Other Samples

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	ogram Summary Continued	Piodifica.	2/22/2019 2:54:59 111
	HiSeq_1683 -1 RNA with DNaseI Cleanup [M0D055413_S_I (1:10)]		
[FU] 30 -			•
25 -			
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0-			
	25 200 500 1000 2000 4000	[r	t]

Overall Results for sample 1 : HiSeq_1683 -1 RNA with DNaseI Cleanup

RNA Area:	431.7	RNA Integrity Number (RIN):	2.5 (B.02.08)
RNA Concentration:	1,516 pg/µl	Result Flagging Color:	
rRNA Ratio [28s / 18s]:	0.6	Result Flagging Label:	RIN: 2.50

Fragment table for sample 1 : <u>HiSeq 1683 -1 RNA with DNaseI Cleanu</u>				NA with DNaseI Cleanup
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	1,650	1,866	2.1	0.5
28S	2,886	3,293	1.4	0.3

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Assay Class: Data Path:	Eukaryote Total RNA Pico C:\ bioanalyzer\2100 expert\data\201	.9-02-22\2019-02-22_001.xad	Created: 2 Modified: 2	/22/2019 2:31:30 PM /22/2019 2:54:39 PM
Electropher	ogram Summary Continued			
	HiSeq_1683 -2 RNA with DN	aseI Cleanup [M0D055408_N_I (1:8)]		
[FU]			1	
25-				
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	25 200 500 1000	2000 4000	[nt]	
Overall Resu	Ilts for sample 2: <u>HiSeq_1683 -2</u>	RNA with DNaseI Cleanup		
RNA Area:	70.2	RNA Integrity Number (RIN): 5.6	6 (B.02.08)	

RNA Area:	70.2
RNA Concentration:	247 pg/µl
rRNA Ratio [28s / 18s]:	0.5

RNA Integrity Number (RIN):	5.6 (B.02
Result Flagging Color:	
Result Flagging Label:	RIN: 5.60

Fragment table for sample 2 : <u>HiSeq 1683 - 2 RNA with DNaseI Cleanup</u>				
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	1,634	1,881	3.0	4.3
28S	2,827	3,422	1.5	2.1

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Assay Class: Data Path: Electrophere	Eukaryote Total RNA Pico C:\ bioanalyzer\2100 expert\data\2019-02-22\2019-02-22_001.xad ogram Summary Continued	Created: Modified:	2/22/2019 2:31:30 PM 2/22/2019 2:54:39 PM
Liettopher			
[FU] _ 20 - 15 - 10 - 5 -	HiSeq_1683 -3 RNA with DNaseI Cleanup [M0D055401_G_I (1:10)]		
	25 200 500 1000 2000 4000	[n	t]

Overall Results for sample 3 : <u>HiSeq 1683 - 3 RNA with DNaseI Cleanup</u>

RNA Area:	212.6	RNA Integrity Number (RIN):	4.8 (B.02.08)
RNA Concentration:	747 pg/µl	Result Flagging Color:	
rRNA Ratio [28s / 18s]:	0.9	Result Flagging Label:	RIN: 4.80

Fragme	nt table for sample	e3: <u>HiSeq</u> 1	<u> 1683 -3 R</u>	NA with DNaseI Cleanup
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	1,591	1,902	6.1	2.9
28S	2,810	3,398	5.4	2.6

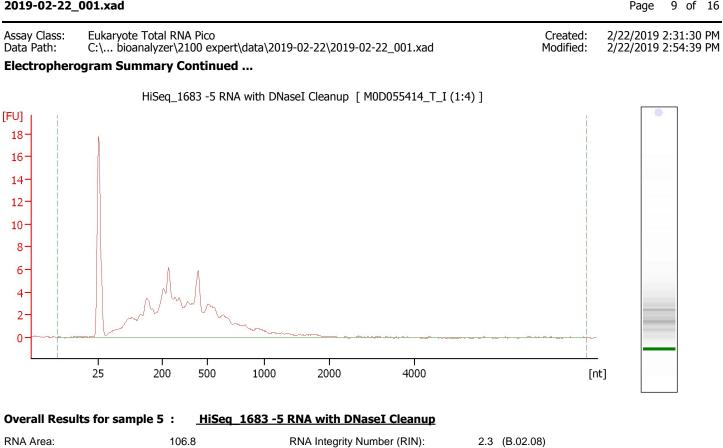
Assay Class: Data Path:	Eukaryote Total RNA Pico C:\ bioanalyzer\2100 expert\data\2019-02-22\2019-02-22_001.xad	Created: Modified:	2/22/2019 2:31:30 PM 2/22/2019 2:54:39 PM
Electropher	ogram Summary Continued		
	HiSeq_1683 -4 RNA with DNaseI Cleanup [M0D055402_H_I (1:10)]		
FU] 20 -			*
15-			
10-			
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0			
	25 200 500 1000 2000 4000	[n	t]

RNA Area:	256.1	RNA Integrity Number (RIN):	4.5 (B.02.08)
RNA Concentration:	899 pg/µl	Result Flagging Color:	
rRNA Ratio [28s / 18s]:	0.3	Result Flagging Label:	RIN: 4.50

Fragme	nt table for sample	e4: <u>HiSeq</u> 1	L683 -4 R	NA with DNaseI Cleanup
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	1,389	1,939	14.3	5.6
28S	2,807	3,390	4.3	1.7

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RNA Concentration: rRNA Ratio [28s / 18s]: 375 pg/µl 0.0

Result Flagging Color: Result Flagging Label: RIN: 2.30

Assay Class: Data Path:	Eukaryote Total RNA Pico C:\ bioanalyzer\2100 expert\data\2019-02-22\2019-02-22_001.xad	Created: Modified:	2/22/2019 2:31:30 PM 2/22/2019 2:54:39 PM
	ogram Summary Continued	r louineu.	
	HiSeq_1683 -6 RNA with DNaseI Cleanup [M0D055415_U_I (1:10)]		
[FU]			
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	25 200 500 1000 2000 4000	[n	t]

Overall Results for sample 6 : <u>HiSeq 1683 - 6 RNA with DNaseI Cleanup</u>
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RNA Area:	221.1	RNA Integrity Number (RIN):	5.1 (B.02.08)
RNA Concentration:	777 pg/µl	Result Flagging Color:	
rRNA Ratio [28s / 18s]:	0.6	Result Flagging Label:	RIN: 5.10

Fragme	nt table for sample	e6: <u>HiSeq</u> 1	L683 -6 R	NA with DNaseI Cleanup
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	1,398	1,984	11.7	5.3
28S	2,790	3,492	6.8	3.1

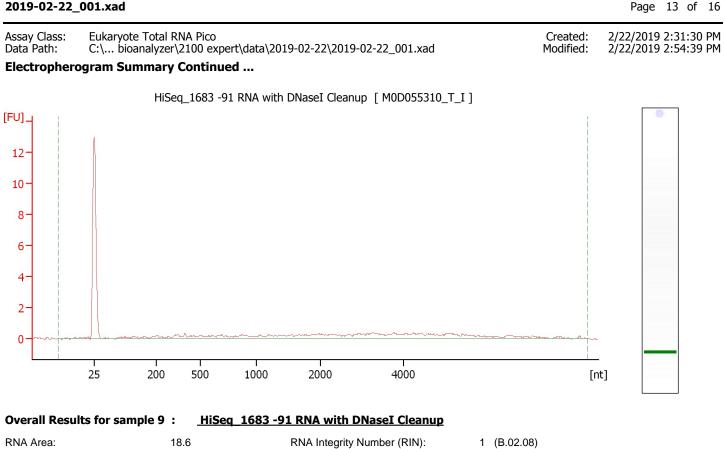
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Assay Class: Data Path:	Eukaryote C:\ bioa	e Total RNA Pico analyzer\2100 expe	rt\data\2019	9-02-22\2019-02	2-22_001.xad	Crea Modi	ited:	2/22/2019 2:31:30 PM 2/22/2019 2:54:39 PM
Electropher	ogram Sun	nmary Continue	ed					
		HiSeq_1683 -7 RN	IA with DNas	seI Cleanup [M	0D055302_L_I (1:40)]		
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0				182	m	<u> </u>		
	25	200 500	1000	2000	4000		[nt]	
Overall Resu	ults for sam	ple 7 : <u>HiSec</u>	<u>1683 -7 </u>	RNA with DNa	seI Cleanup			
RNA Area:		209.2		RNA Integrity Nu	mber (RIN):	3.2 (B.02.08)		
RNA Concentr	ation:	735 pg/µl		Result Flagging (Color:			
rRNA Ratio [28	3s / 18s]:	0.0		Result Flagging I	_abel:	RIN: 3.20		
Fragment ta	ble for sam	ple 7 : <u>HiSe</u>	<u>q 1683 -7</u>	RNA with DNa	seI Cleanup			
Name Sta	art Size [nt]	End Size [nt] Area	% of total	Area			
18S 1,5	587	1,834	4.4	2.1				

Assay Class: Data Path: Electrophere	Eukaryote Total RNA Pico C:\ bioanalyzer\2100 expert\d ogram Summary Continued	ata\2019-02-22\2019-02-22_001.xad •	Created: Modified:	2/22/2019 2:31:30 PM 2/22/2019 2:54:39 PM
	HiSeq_1683 -8 RNA w	ith DNaseI Cleanup [M0D055342_Z_I (1:	20_]	
[FU] 25- 20- 15- 10- 5-				
0-	25 200 500	1000 2000 4000		[nt]
Overall Resu RNA Area: RNA Concentra rRNA Ratio [28	354.3 ation: 1,244 pg/µl	583 -8 RNA with DNaseI Cleanup RNA Integrity Number (RIN): Result Flagging Color: Result Flagging Label:	2.2 (B.02.08)	

Fragme	nt table for sample	e8: <u>HiSeq</u> 1	<u> 1683 -8 R</u>	NA with DNaseI Cleanup
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	1,655	1,906	0.8	0.2

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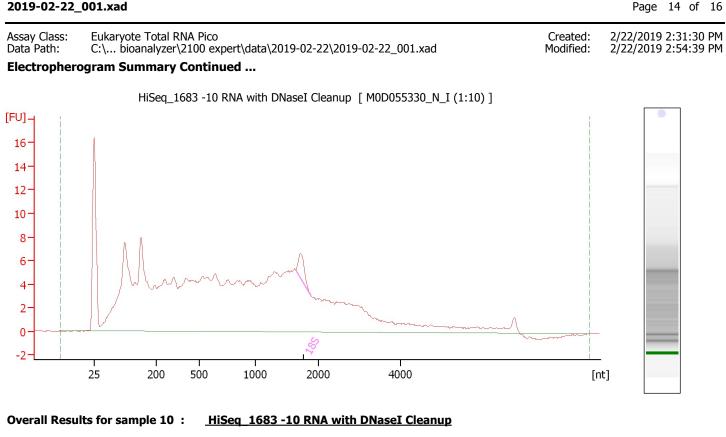


RNA Area: RNA Concentration: rRNA Ratio [28s / 18s]:

65 рg/µl 0.0
 RNA Integrity Number (RIN):
 1
 (B.

 Result Flagging Color:
 Image: Color:
 Image: Color:

 Result Flagging Label:
 RIN:1



 RNA Area:
 300.5

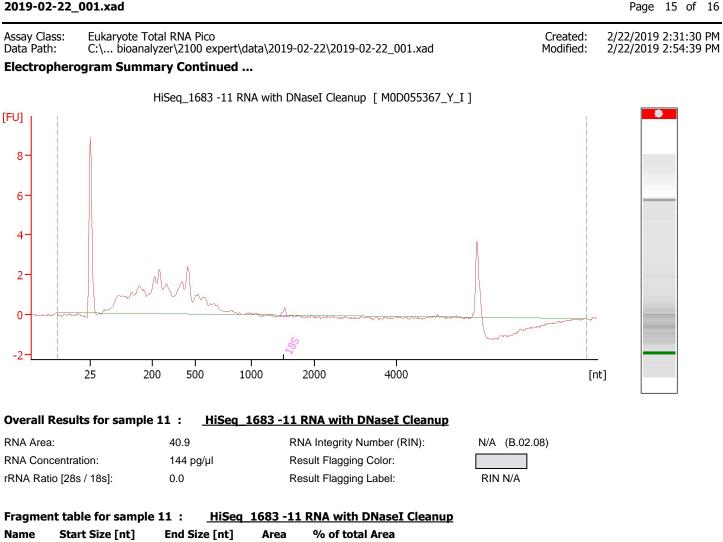
 RNA Concentration:
 1,055 pg/µl

 rRNA Ratio [28s / 18s]:
 0.0

RNA Integrity

RNA Integrity Number (RIN):2.8 (B.02.08)Result Flagging Color:Image: Color:Result Flagging Label:RIN: 2.80

Fragmen	t table for sample	10 : <u>HiSeq</u>	<u> 1683 -10</u>	RNA with DNaseI Cleanup
Name	Start Size [nt]	End Size [nt]	Area	% of total Area
18S	1,643	1,898	3.3	1.1



18S	1,454	1,573	0.4	0.9

