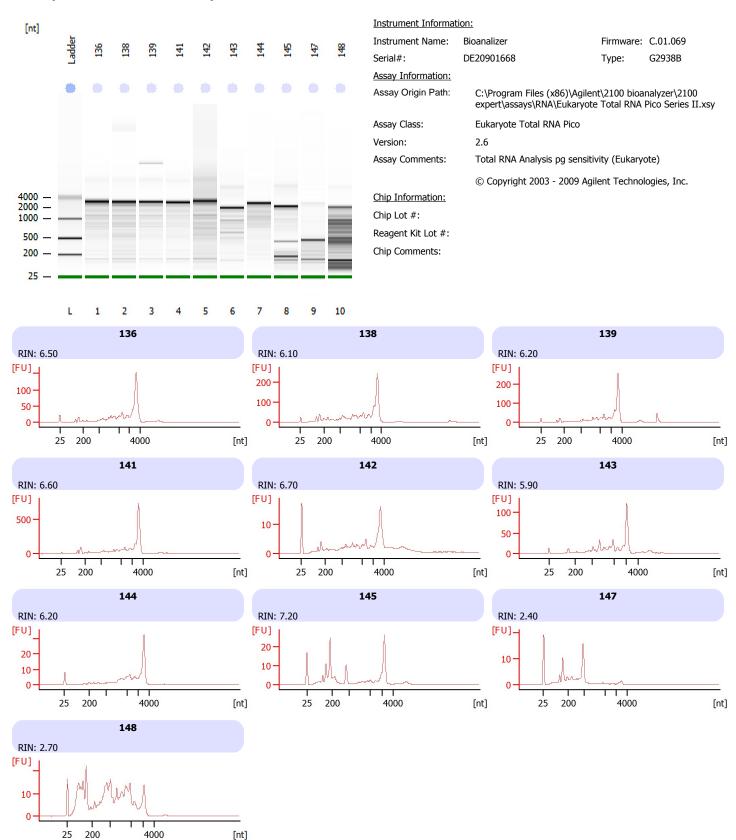
Assay Class: Eukaryote Total RNA Pico C:\...han\_1-29-2016\_Selected Samples\_Eukaryote Total RNA Pico.xad Created: 1/29/2016 10:28:19 AM Modified: 1/29/2016 2:46:43 PM

# **Electrophoresis File Run Summary**



Assay Class: Eukaryote Total RNA Pico C:\...han\_1-29-2016\_Selected Samples\_Eukaryote Total RNA Pico.xad Created: 1/29/2016 10:28:19 AM Modified: 1/29/2016 2:46:43 PM

# **Electrophoresis File Run Summary (Chip Summary)**

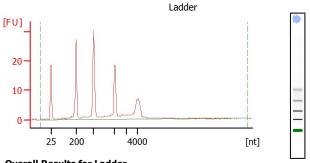
Sample Name	Sample Comment		Result Label	Result Color
136		~	RIN: 6.50	
138		~	RIN: 6.10	
139		<b>V</b>	RIN: 6.20	
141		~	RIN: 6.60	
142		~	RIN: 6.70	
143		~	RIN: 5.90	
144		~	RIN: 6.20	
145		~	RIN: 7.20	
147		~	RIN: 2.40	
148		~	RIN: 2.70	
Ladder		~	All Other	
			Samples	

Chip Lot # Reagent Kit Lot #

**Chip Comments:** 

Assay Class: Eukaryote Total RNA Pico C:\...han\_1-29-2016\_Selected Samples\_Eukaryote Total RNA Pico.xad Cc:\...han\_1-29-2016\_Selected Samples\_Eukaryote Total RNA Pico.xad Cc:\...han\_1-29-2016\_Selected Samples\_Eukaryote Total RNA Pico.xad Modified: 1/29/2016 2:46:43 PM

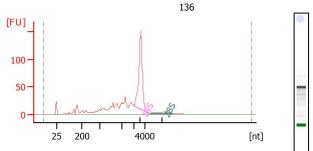
# **Electropherogram Summary**



#### **Overall Results for Ladder**

RNA Area: RNA Concentration: Result Flagging Color: Result Flagging Label:





# Overall Results for sample 1 : 136

RNA Area: 944.8 RNA Concentration: 5,096 pg/µl rRNA Ratio [28s / 18s]: 0.1

RNA Integrity Number (RIN): 6.5 (B.02.08, Anomaly Threshold(s) manually adapted)

<u>136</u>

Result Flagging Color:

Result Flagging Label:

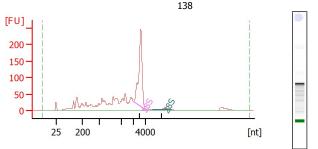
RIN: 6.50

Fragment table for sample 1:

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

 18S
 1,997
 4,203
 316.9
 33.5

 28S
 4,891
 8,730
 17.5
 1.9



#### Overall Results for sample 2: 138

RNA Area: 1,821.2 RNA Concentration: 9,824 pg/µl rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 6.1 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color:

Result Flagging Label:

RIN: 6.10

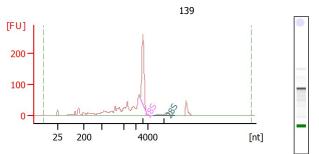
Fragment table for sample 2:

138

### lame Start Size [nt] End Size [nt] Area % of total Area

 18S
 1,975
 4,066
 495.2
 27.2

 28S
 5,251
 8,482
 18.7
 1.0



#### Overall Results for sample 3: 139

RNA Area: 1,274.7
RNA Concentration: 6,876 pg/µl
rRNA Ratio [28s / 18s]: 0.1

RNA Integrity Number (RIN): 6.2 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color:
Result Flagging Label:
Rin: 6.20

Fragment table for sample 3: 139

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

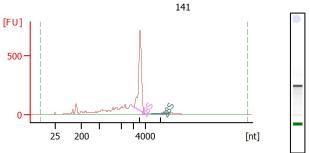
 18S
 2,712
 3,784
 305.2
 23.9

 28S
 4,821
 8,716
 34.9
 2.7

5

Assay Class: Eukaryote Total RNA Pico Created: 1/29/2016 10:28:19 AM C:\...han\_1-29-2016\_Selected Samples\_Eukaryote Total RNA Pico.xad Data Path: Modified: 1/29/2016 2:46:43 PM

# **Electropherogram Summary Continued ...**



Overall Results for sample 4: <u>141</u>

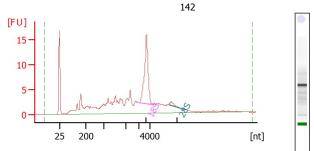
RNA Area: 3,435.1 18,530 pg/µl RNA Concentration: rRNA Ratio [28s / 18s]:

RNA Integrity Number (RIN): 6.6 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: Result Flagging Label: RIN: 6.60

Fragment table for sample 4: <u> 141</u> Start Size [nt] End Size [nt] Area % of total Area Name

18S 1,939 3,970 1,245.1 36.2 28S 5,035 8,053 48.5 1.4



Overall Results for sample 5: <u> 142</u>

RNA Area: 164.6 888 pg/µl RNA Concentration: rRNA Ratio [28s / 18s]:

RNA Integrity Number (RIN): 6.7 (B.02.08, Anomaly Threshold(s) manually adapted)

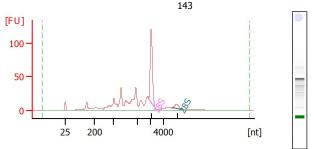
<u>142</u>

Result Flagging Color: Result Flagging Label: RIN: 6.70

Fragment table for sample 5:

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

18S 2,130 4,503 31.4 19.1 285 7.125 9.853 1.7 1.0



Overall Results for sample 6: <u>143</u>

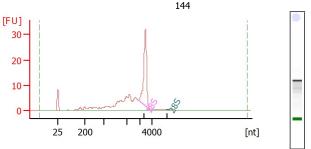
669.3 RNA Area: RNA Concentration: 3,610 pg/µl rRNA Ratio [28s / 18s]: 0.1

RNA Integrity Number (RIN): 5.9 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: RIN: 5.90 Result Flagging Label: Fragment table for sample 6: 143

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

18S 1,839 2,746 139.2 20.8 28S 5,326 6,927 12.1 1.8



Overall Results for sample 7: <u> 144</u>

RNA Area: 161.8 RNA Concentration: 873 pg/µl rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 6.2 (B.02.08, Anomaly Threshold(s) manually adapted)

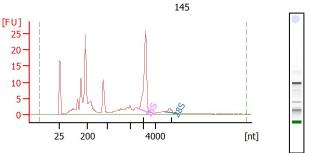
Result Flagging Color: Result Flagging Label: RIN: 6.20 Fragment table for sample 7: 144

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

18S 1,911 3,630 53.4 33.0 28S 6,369 6,780 0.1 0.1

Assay Class: Eukaryote Total RNA Pico Created: 1/29/2016 10:28:19 AM C:\...han\_1-29-2016\_Selected Samples\_Eukaryote Total RNA Pico.xad Data Path: Modified: 1/29/2016 2:46:43 PM

# **Electropherogram Summary Continued ...**



Overall Results for sample 8: <u>145</u>

RNA Area: 203.4 1,097 pg/µl RNA Concentration: rRNA Ratio [28s / 18s]: 0.1

RNA Integrity Number (RIN): 7.2 (B.02.08, Anomaly Threshold(s) manually adapted)

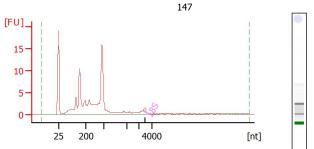
<u>145</u>

Result Flagging Color: Result Flagging Label: RIN: 7.20

Fragment table for sample 8:

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

18S 1,606 3,025 43.0 21.2 28S 5,334 7,610 2.8 1.4



Overall Results for sample 9: <u> 147</u>

RNA Area: 109.3 RNA Concentration: 590 pg/μl rRNA Ratio [28s / 18s]: 0.0

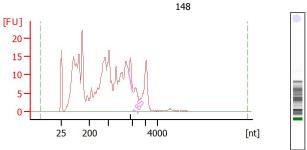
2.4 (B.02.08, Anomaly Threshold(s) manually adapted) RNA Integrity Number (RIN):

<u> 147</u>

Result Flagging Color: Result Flagging Label: RIN: 2.40 Fragment table for sample 9:

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

18S 3,335 2,572 1.3 1.2



Overall Results for sample 10: 148

RNA Area: 496.9 RNA Concentration: 2,681 pg/µl rRNA Ratio [28s / 18s]: 0.0

RNA Integrity Number (RIN): 2.7 (B.02.08, Anomaly Threshold(s) manually adapted)

Result Flagging Color: RIN: 2.70 Result Flagging Label: Fragment table for sample 10: 148

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

18S 929 1,051 6.8 1.4