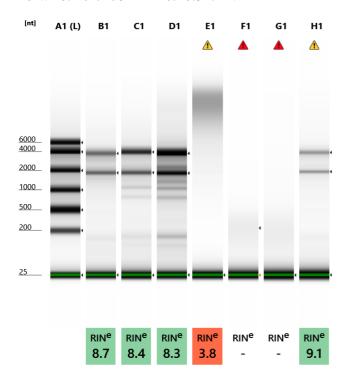
Filename: 2019-08-12 - 16.16.31.RNA

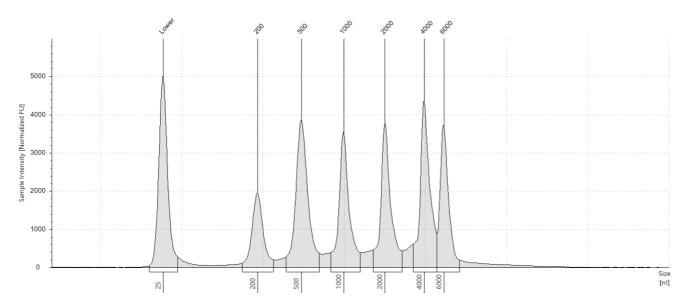


Default image (Contrast 100%)

Sample Info

| Well | RINe | 28S/18S (Area) | Conc. [ng/µl] | Sample Description | Alert | Observations |
|------|------|----------------|---------------|-----------------------------|----------|---|
| A1 | - | - | 119 | Ladder | | Ladder |
| B1 | 8.7 | 1.3 | 26.1 | 2875 Mcap Soft (301) | | |
| C1 | 8.4 | 1.4 | 30.6 | 1090 Pacuta Soft (313) | | |
| D1 | 8.3 | 1.7 | 65.5 | 1047 Pacuta Soft (325) | | |
| El | 3.8 | - | 23.2 | 1250 Mcap Soft (337) DNA | A | The lower ribosomal fragment is missing. Sample concentration outside functional range for RINe and the assay |
| F1 | - | - | 4.41 | 221 cap BS | A | Sample concentration outside functional range for RINe and the assay |
| Gl | - | - | 5.06 | 1431 cap bS | A | Sample concentration outside functional range for RINe and the assay |
| H1 | 9.1 | 1.4 | 11.3 | 1250 Mcap Soft (337) RNA | <u>^</u> | Sample concentration outside functional range for RINe and the assay |

A1: Ladder

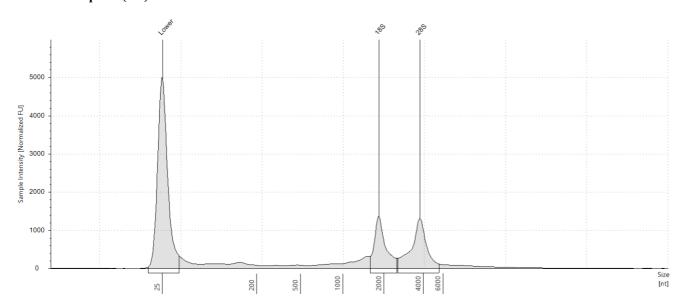


Sample Table

| Well | RINe | 28S/18S (Area) | Conc. [ng/µl] | Sample Description | Alert | Observations |
|------|------|----------------|---------------|--------------------|-------|--------------|
| A1 | - | - | 119 | Ladder | | Ladder |

| Size [nt] | Calibrated Conc. [ng/µl] | Assigned Conc. [ng/µl] | Peak Molarity [nmol/l] | % Integrated Area | Peak Comment | Observations |
|-----------|-----------------------------|------------------------|------------------------|-------------------|--------------|--------------|
| 25 | 40.0 | 40.0 | 4710 | = | | Lower Marker |
| 200 | 10.9 | - | 160 | 10.18 | | |
| 500 | 23.2 | - | 136 | 21.62 | | |
| 1000 | 17.8 | - | 52.2 | 16.56 | | |
| 2000 | 18.4 | - | 27.0 | 17.15 | | |
| 4000 | 20.9 | - | 15.4 | 19.52 | | |
| 6000 | 16.0 | - | 7.87 | 14.97 | | |

B1: 2875 Mcap Soft (301)

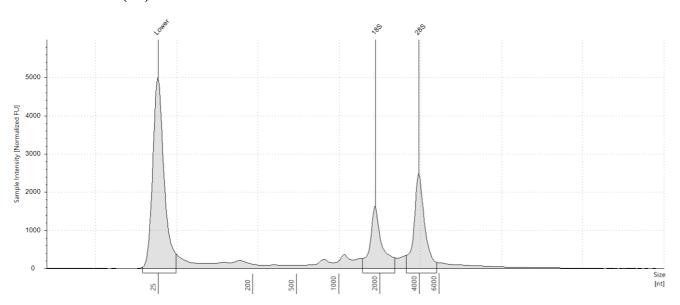


Sample Table

| ĺ | Well | RINe | 28S/18S (Area) | Conc. [ng/µl] | Sample Description | Alert | Observations |
|---|------|------|----------------|---------------|----------------------|-------|--------------|
| | B1 | 8.7 | 1.3 | 26.1 | 2875 Mcap Soft (301) | | |

| | Size [nt] | Calibrated Conc. [ng/μl] | Assigned Conc. [ng/μl] | Peak Molarity [nmol/l] | % Integrated Area | Peak Comment | Observations |
|---|-----------|-----------------------------|------------------------|------------------------|-------------------|--------------|--------------|
| Ī | 25 | 40.0 | 40.0 | 4710 | - | | Lower Marker |
| Ī | 1836 | 6.66 | - | 10.7 | 43.67 | | 18S |
| Ī | 3766 | 8.59 | - | 6.71 | 56.33 | | 28S |

C1: 1090 Pacuta Soft (313)

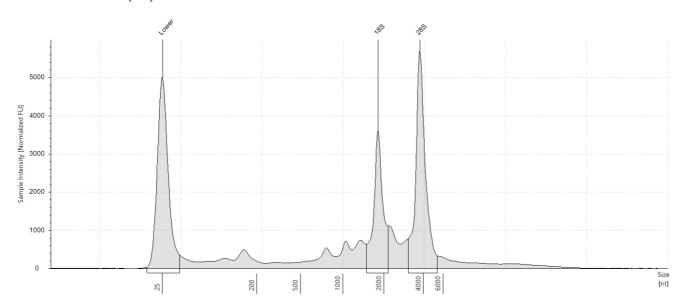


Sample Table

| Well | RINe | 28S/18S (Area) | Conc. [ng/µl] | Sample Description | Alert | Observations |
|------|------|----------------|---------------|------------------------|-------|--------------|
| C1 | 8.4 | 1.4 | 30.6 | 1090 Pacuta Soft (313) | | |

| Ì | Size [nt] | Calibrated Conc. [ng/μl] | Assigned Conc. [ng/µl] | Peak Molarity [nmol/l] | % Integrated Area | Peak Comment | Observations |
|---|-----------|-----------------------------|------------------------|------------------------|-------------------|--------------|--------------|
| Ī | 25 | 40.0 | 40.0 | 4710 | - | | Lower Marker |
| Ī | 1861 | 7.27 | - | 11.5 | 41.26 | | 18S |
| Ī | 3981 | 10.3 | - | 7.64 | 58.74 | | 28S |

D1: 1047 Pacuta Soft (325)

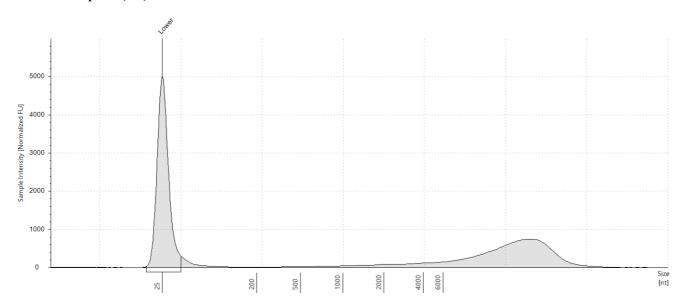


Sample Table

| Well | RINe | 28S/18S (Area) | Conc. [ng/µl] | Sample Description | Alert | Observations |
|------|------|----------------|---------------|------------------------|-------|--------------|
| D1 | 8.3 | 1.7 | 65.5 | 1047 Pacuta Soft (325) | | |

| Size [nt] | Calibrated Conc. [ng/μl] | Assigned Conc. [ng/μl] | Peak Molarity [nmol/l] | % Integrated Area | Peak Comment | Observations |
|-----------|-----------------------------|------------------------|------------------------|-------------------|--------------|--------------|
| 25 | 40.0 | 40.0 | 4710 | - | | Lower Marker |
| 1806 | 13.2 | - | 21.6 | 37.26 | | 18S |
| 3757 | 22.3 | - | 17.4 | 62.74 | | 28S |

E1: 1250 Mcap Soft (337) DNA

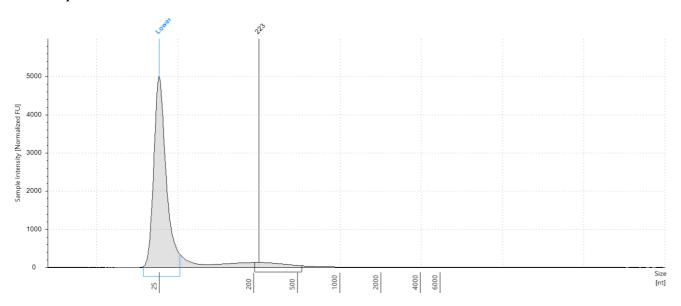


Sample Table

| Well | RINe | 28S/18S (Area) | Conc. [ng/µl] | Sample Description | Alert | Observations |
|------|------|----------------|---------------|-----------------------------|----------|---|
| E1 | 3.8 | - | 23.2 | 1250 Mcap Soft (337) DNA | <u> </u> | The lower ribosomal fragment is missing; Sample concentration outside functional range for RINe and the assay |

| Size [nt] | Calibrated Conc. [ng/µl] | Assigned Conc. [ng/µl] | Peak Molarity [nmol/l] | % Integrated Area | Peak Comment | Observations |
|-----------|-----------------------------|------------------------|------------------------|-------------------|--------------|--------------|
| 25 | 40.0 | 40.0 | 4710 | - | | Lower Marker |

F1: 221 cap BS

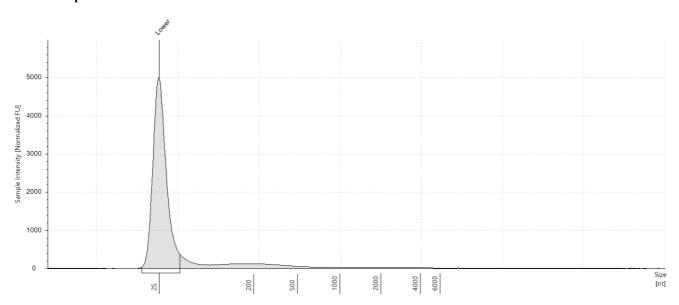


Sample Table

| Well | RINe | 28S/18S (Area) | Conc. [ng/µl] | Sample Description | Alert | Observations |
|------|------|----------------|---------------|--------------------|----------|--|
| F1 | - | - | 4.41 | 221 cap BS | <u> </u> | Sample concentration outside functional range for RINe and the assay |

| Size [nt] | Calibrated Conc. [ng/μl] | Assigned Conc. [ng/µl] | Peak Molarity [nmol/l] | % Integrated Area | Peak Comment | Observations |
|-----------|-----------------------------|------------------------|------------------------|-------------------|--------------|--------------|
| 25 | 40.0 | 40.0 | 4710 | - | | Lower Marker |
| 223 | 1.38 | - | 18.3 | 100.00 | | |

G1: 1431 cap bS

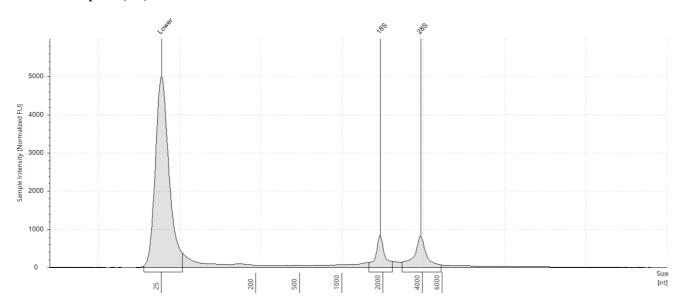


Sample Table

| Well | RINe | 28S/18S (Area) | Conc. [ng/µl] | Sample Description | Alert | Observations |
|------|------|----------------|---------------|--------------------|----------|--|
| Gl | - | - | 5.06 | 1431 cap bS | <u> </u> | Sample concentration outside functional range for RINe and the assay |

| Size [nt] | Calibrated Conc. [ng/μl] | Assigned Conc. [ng/µl] | Peak Molarity [nmol/l] | % Integrated Area | Peak Comment | Observations |
|-----------|-----------------------------|------------------------|------------------------|-------------------|--------------|--------------|
| 25 | 40.0 | 40.0 | 4710 | | | Lower Marker |

H1: 1250 Mcap Soft (337) RNA



Sample Table

| Well | RINe | 28S/18S (Area) | Conc. [ng/µl] | Sample Description | Alert | Observations |
|------|------|----------------|---------------|-----------------------------|----------|--|
| Hl | 9.1 | 1.4 | 11.3 | 1250 Mcap Soft (337) RNA | <u> </u> | Sample concentration outside functional range for RINe and the assay |

| Ì | Size [nt] | Calibrated Conc. [ng/μl] | Assigned Conc. [ng/μl] | Peak Molarity [nmol/l] | % Integrated Area | Peak Comment | Observations |
|---|-----------|-----------------------------|------------------------|------------------------|-------------------|--------------|--------------|
| Ī | 25 | 40.0 | 40.0 | 4710 | - | | Lower Marker |
| Ī | 1911 | 2.30 | - | 3.55 | 41.38 | | 18S |
| Ī | 3901 | 3.26 | - | 2.46 | 58.62 | | 28S |