

A

☒ Preprocess

Smoothing method

Savitzky-Golay (recommended)

Normalization method

None

Cq method

Threshold

☐ Auto Threshold

Threshold RNase P

0.25

B

Color by

Sample

Line Type by

None

Show Targets

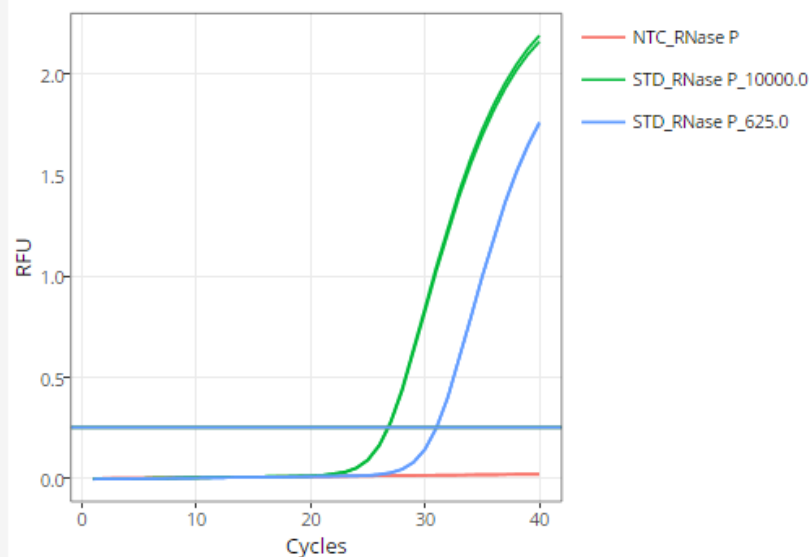
RNase P

Show Cq

None

☐ Log Scale

C



Experiment

Standard Curve

Example

Run

Run001

D

| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|---|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|---|----|----|----|
| A | NTC_RNase P | NTC_RNase P | NTC_RNase P | pop1_RNase P | pop1_RNase P | pop1_RNase P | pop2_RNase P | pop2_RNase P | | | | |
| B | pop2_RNase P | STD_RNase P_100.0 | STD_RNase P_100.0 | STD_RNase P_100.0 | STD_RNase P_500.0 | STD_RNase P_500.0 | STD_RNase P_500.0 | STD_RNase P_250.0 | | | | |
| C | STD_RNase P_250.0 | STD_RNase P_250.0 | STD_RNase P_125.0 | STD_RNase P_125.0 | STD_RNase P_125.0 | STD_RNase P_625.0 | STD_RNase P_625.0 | STD_RNase P_625.0 | | | | |
| D | | | | | | | | | | | | |
| E | | | | | | | | | | | | |
| F | | | | | | | | | | | | |
| G | | | | | | | | | | | | |
| H | | | | | | | | | | | | |

E

Show 25 entries

Search:

| data.name | exp.id | run.id | react.id | position | sample | target | target.dyield | sample.type | cq | quantFluor | cq.mean | |
|-----------------------------|------------------------|--------|----------|----------|-------------|---------|---------------|-------------|-----------|------------|-----------|----|
| A01_NTC_RNase P_ntc_RNase P | Standard Curve Example | Run001 | 1 | A01 | NTC_RNase P | RNase P | FAM | ntc | 995.13163 | 0.25 | 605.35779 | 4. |