a
RH
Pru
ME49
Analyzed
612 cells
Analyzed
2198 cells
Analyzed
1552 cells
Gene count
Mapped reads
b
% mapped
Gene count
(≥ 2 read)
RH
Pru
ME49
Pru
ME49
Uninduced
Induced
c
RH
Pru
ME49
17
17
26
50%
Detection limit
Logreg Fit
Linear fit
$P_{detection}$
r^2: 0.96
r^2: 0.96
r^2: 0.93
CPM ($\log_2$)
Expected abundance ($\log_2$)