# Stanford University

## Run Information

Run Name:

Project Name:

Flow Cell:

Lane:

Sample:

Operator:

Mapper:

Genome:


## Sequencing Results

Post-Filter Reads:

Mapped PF Reads (Read 1):

Mapped PF Reads (Read 2):

Uniquely-Mapped PF Reads (Read 1):

Uniquely-Mapped PF Reads (Read 2):

Mean Insert Size: