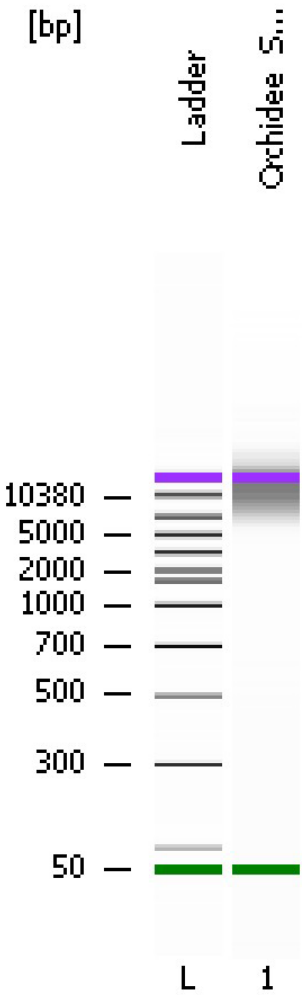
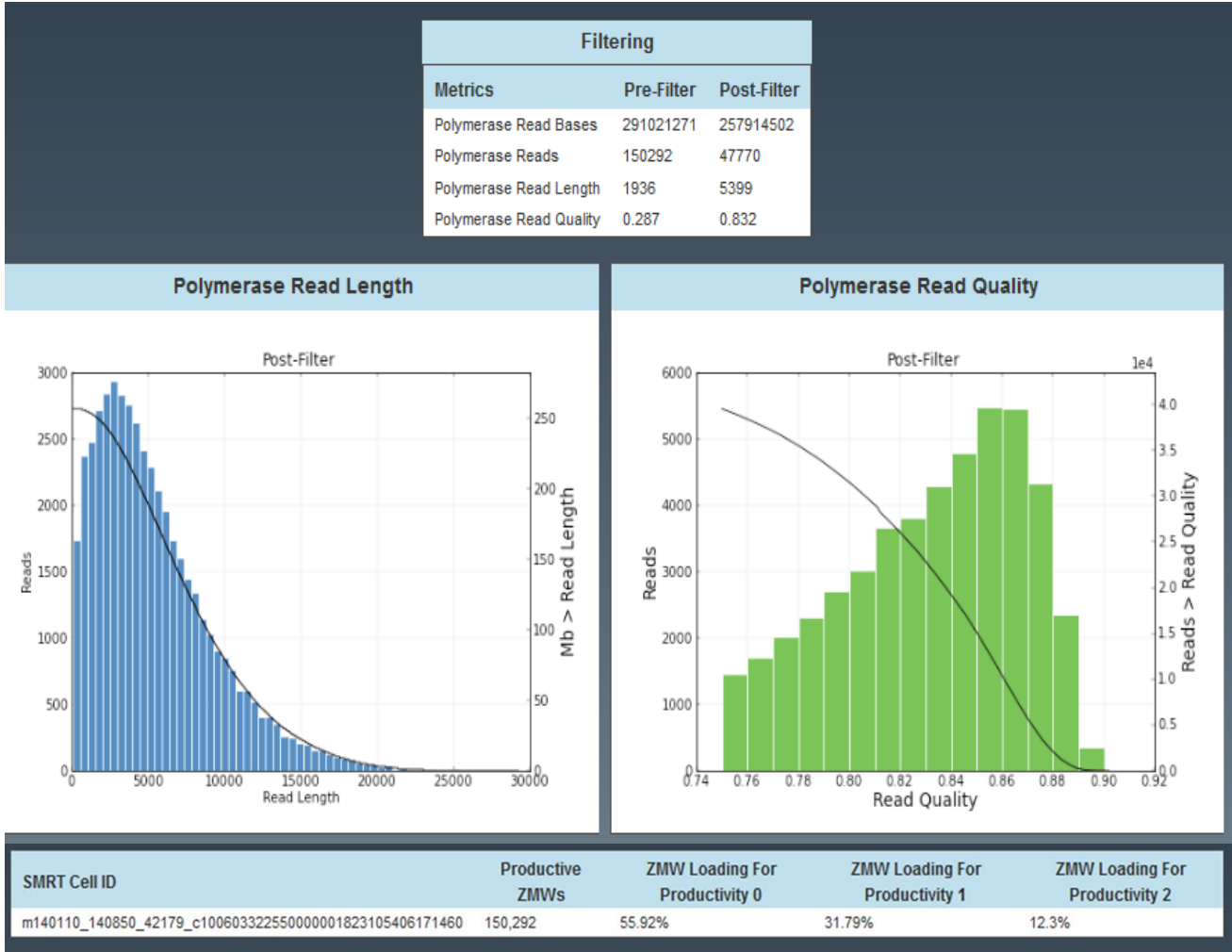
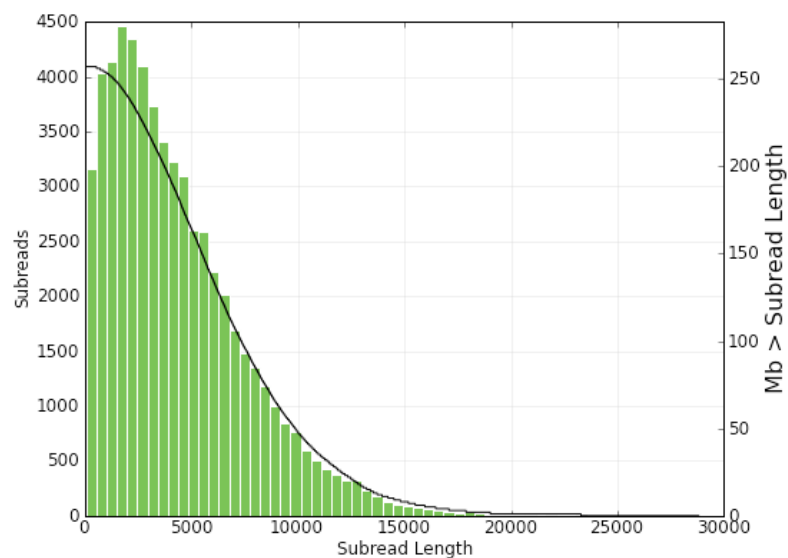


PacBio sequencing *Erycina pusilla*



Mean Subread length	4,351	N50	6,217
Total Number of Bases	257,263,584	Number of Reads	59,123

### Subread Filtering



Adapter Dimers (0-10bp)	0.01%
Short Inserts (11-100bp)	0.0%

### Observed Insert Length Distribution

