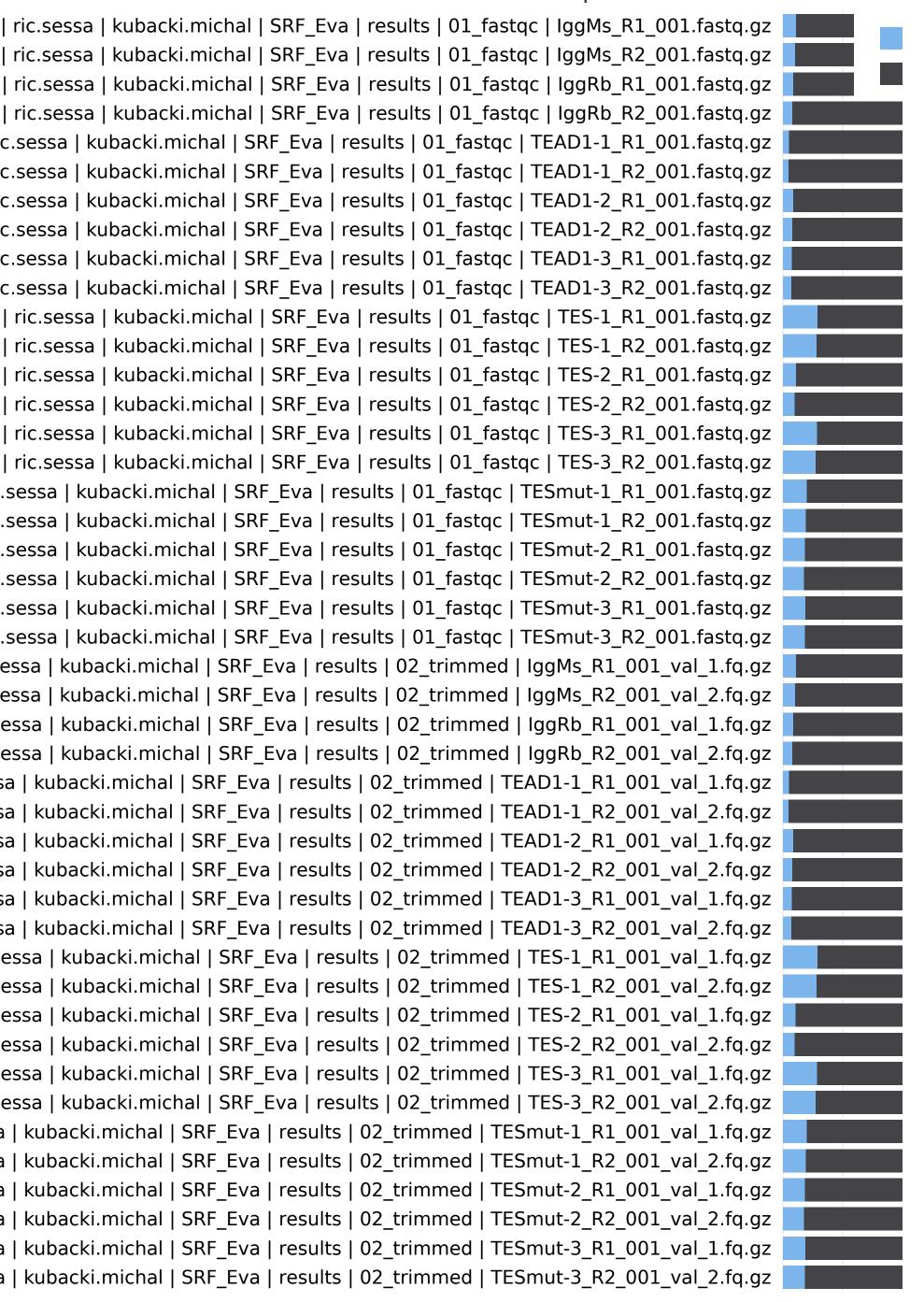
## FastQC: Sequence Counts

44 samples



0% 50% 100%

Unique Reads

**Duplicate Reads**