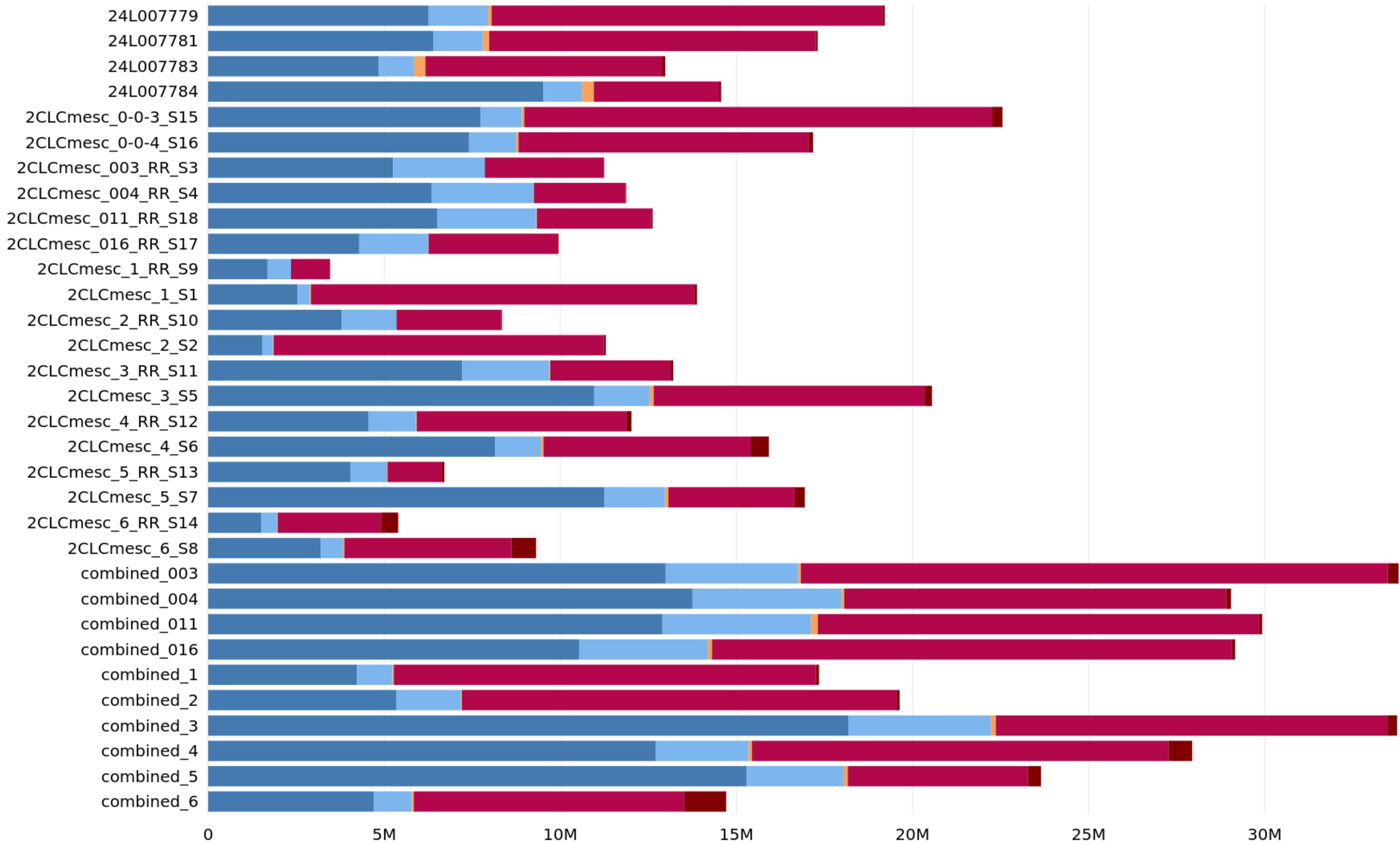


STAR: Alignment Scores

32 samples

- Uniquely mapped
- Mapped to multiple loci
- Mapped to too many loci
- Unmapped: too short
- Unmapped: other



Reads