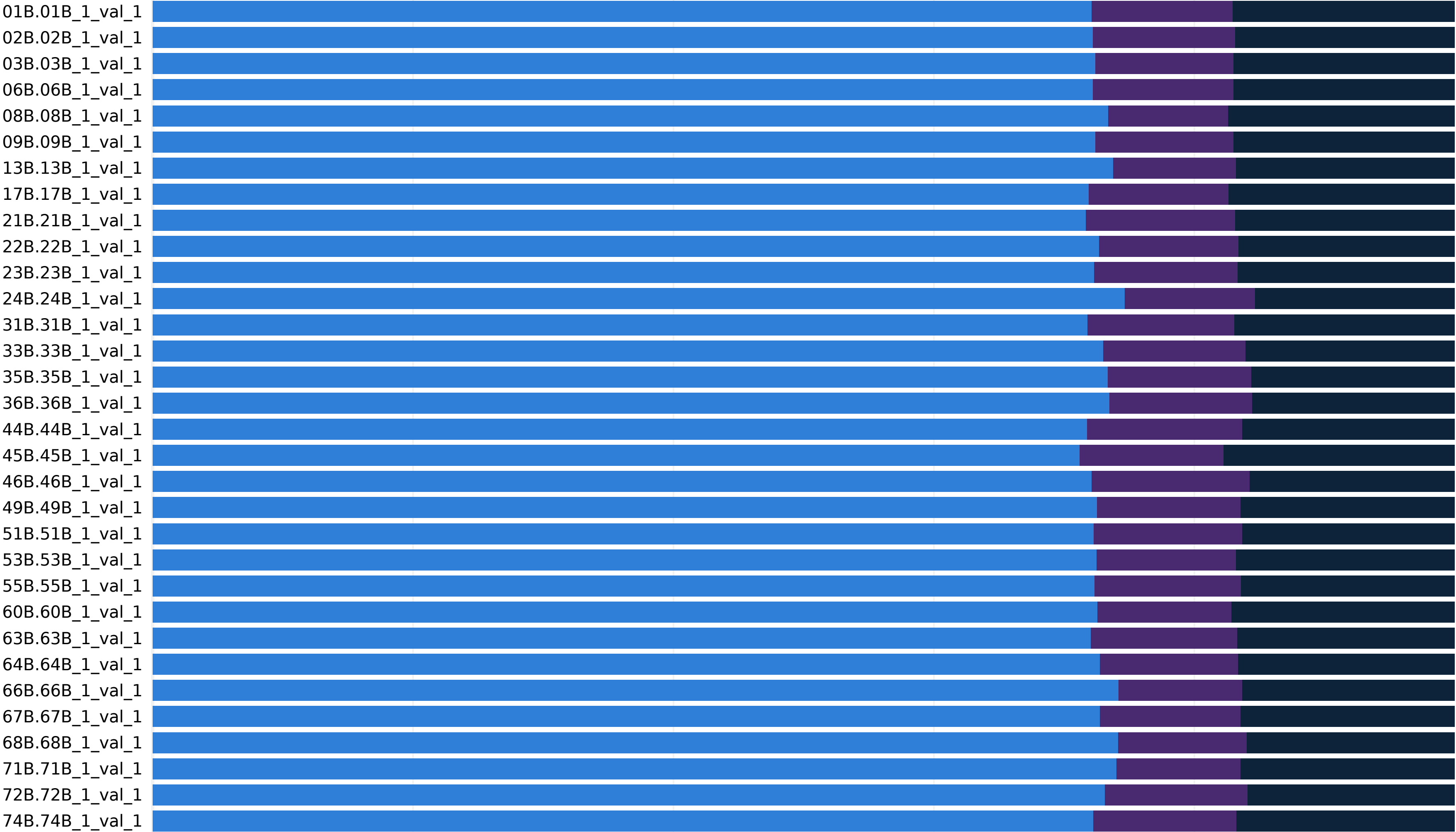


Bismark: Alignment Scores

32 samples

- Aligned Uniquely
- Aligned Ambiguously
- Did Not Align
- No Genomic Sequence



0% 20% 40% 60% 80% 100%

Reads