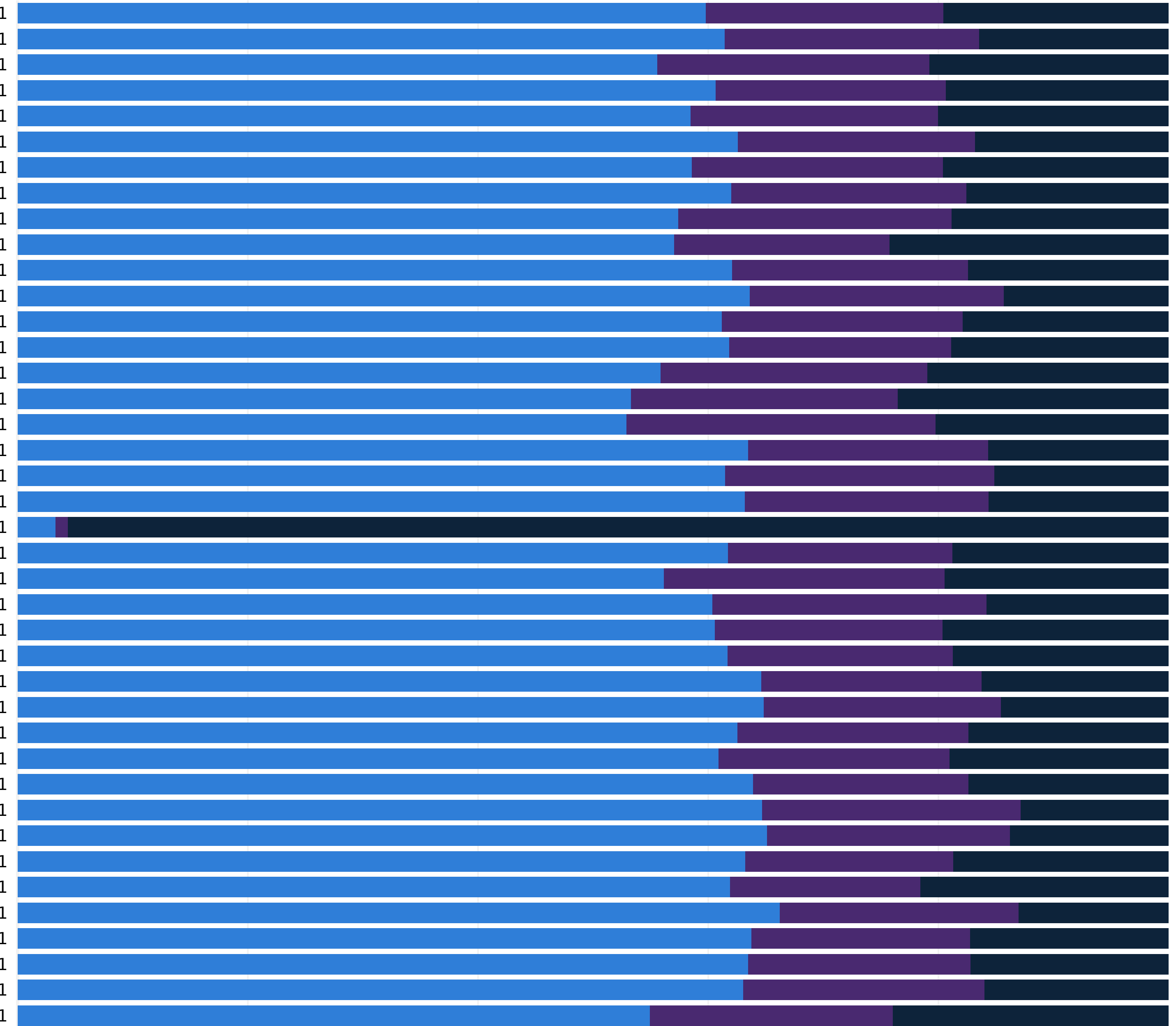


Bismark: Alignment Scores

40 samples

ACR-139-TP1.ACR-139-TP1_R1_001
ACR-139-TP2.ACR-139-TP2_R1_001
ACR-139-TP3.ACR-139-TP3_R1_001
ACR-139-TP4.ACR-139-TP4_R1_001
ACR-145-TP1.ACR-145-TP1_R1_001
ACR-145-TP2.ACR-145-TP2_R1_001
ACR-145-TP3.ACR-145-TP3_R1_001
ACR-145-TP4.ACR-145-TP4_R1_001
ACR-150-TP1.ACR-150-TP1_R1_001
ACR-150-TP2.ACR-150-TP2_R1_001
ACR-150-TP3.ACR-150-TP3_R1_001
ACR-150-TP4.ACR-150-TP4_R1_001
ACR-173-TP1.ACR-173-TP1_R1_001
ACR-173-TP2.ACR-173-TP2_R1_001
ACR-173-TP3.ACR-173-TP3_R1_001
ACR-173-TP4.ACR-173-TP4_R1_001
ACR-186-TP1.ACR-186-TP1_R1_001
ACR-186-TP2.ACR-186-TP2_R1_001
ACR-186-TP3.ACR-186-TP3_R1_001
ACR-186-TP4.ACR-186-TP4_R1_001
ACR-225-TP1.ACR-225-TP1_R1_001
ACR-225-TP2.ACR-225-TP2_R1_001
ACR-225-TP3.ACR-225-TP3_R1_001
ACR-225-TP4.ACR-225-TP4_R1_001
ACR-229-TP1.ACR-229-TP1_R1_001
ACR-229-TP2.ACR-229-TP2_R1_001
ACR-229-TP3.ACR-229-TP3_R1_001
ACR-229-TP4.ACR-229-TP4_R1_001
ACR-237-TP1.ACR-237-TP1_R1_001
ACR-237-TP2.ACR-237-TP2_R1_001
ACR-237-TP3.ACR-237-TP3_R1_001
ACR-237-TP4.ACR-237-TP4_R1_001
ACR-244-TP1.ACR-244-TP1_R1_001
ACR-244-TP2.ACR-244-TP2_R1_001
ACR-244-TP3.ACR-244-TP3_R1_001
ACR-244-TP4.ACR-244-TP4_R1_001
ACR-265-TP1.ACR-265-TP1_R1_001
ACR-265-TP2.ACR-265-TP2_R1_001
ACR-265-TP3.ACR-265-TP3_R1_001
ACR-265-TP4.ACR-265-TP4_R1_001

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



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

Reads