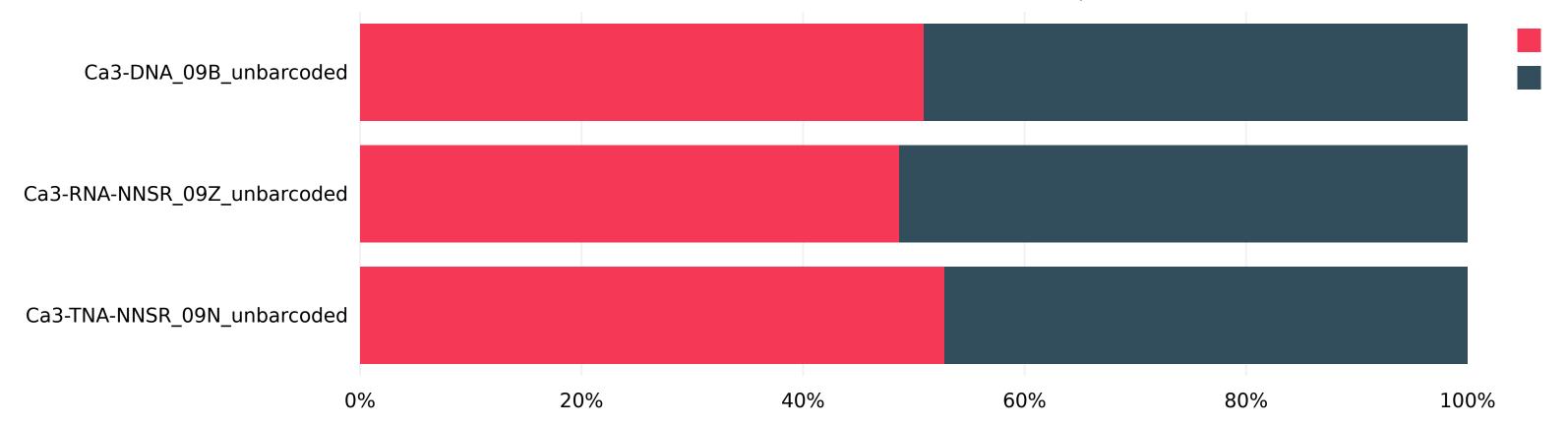
## BISCUIT: Mapping Strand Distribution Read 1, 3 bars



ff: Watson-Aligned, Watson-Bisulfite Conversion

rr: Crick-Aligned, Crick-Bisulfite Conversion

Number of Reads