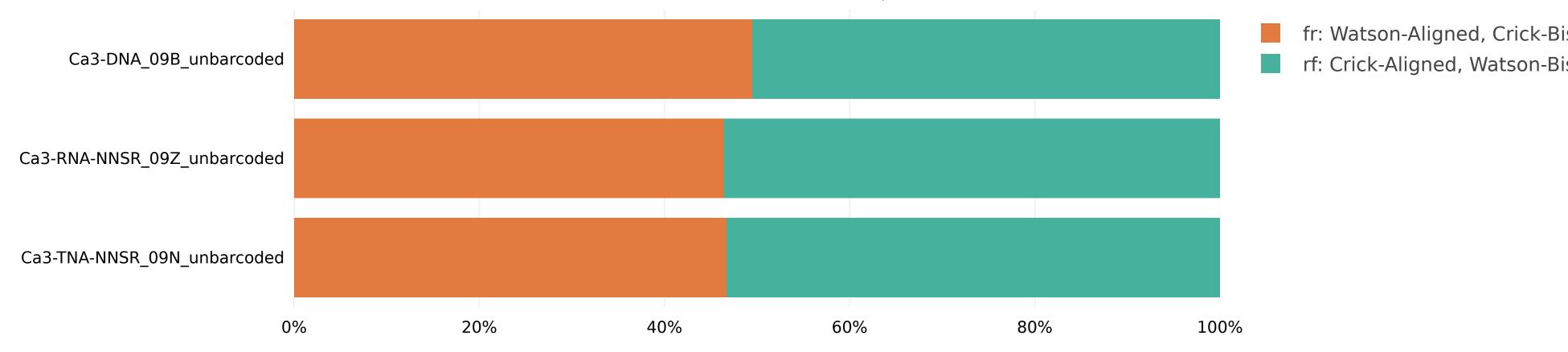
BISCUIT: Mapping Strand Distribution Read 2, 3 bars



fr: Watson-Aligned, Crick-Bisulfite Conversion

rf: Crick-Aligned, Watson-Bisulfite Conversion

Number of Reads