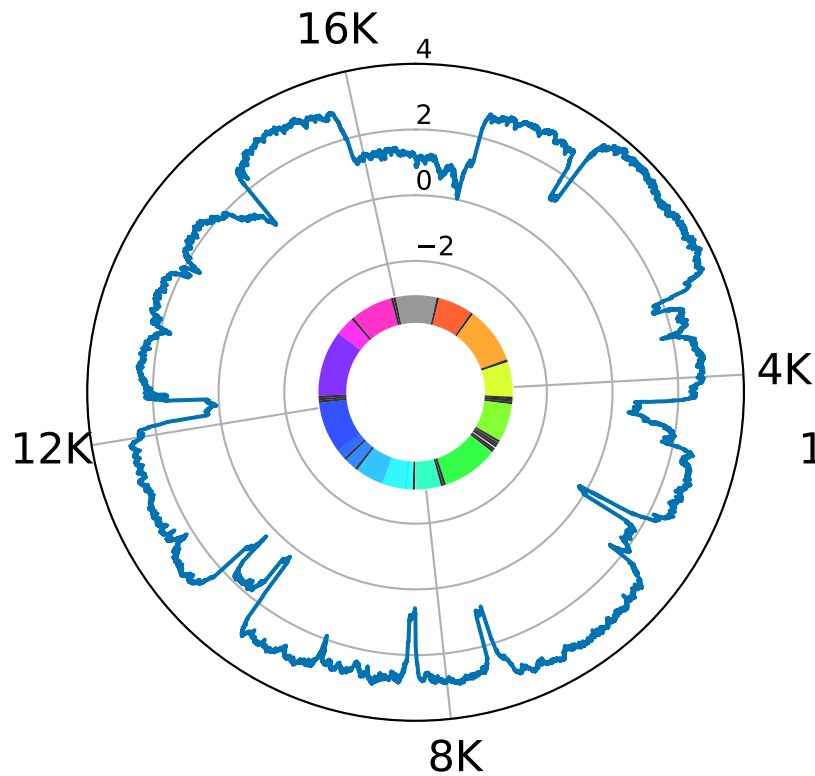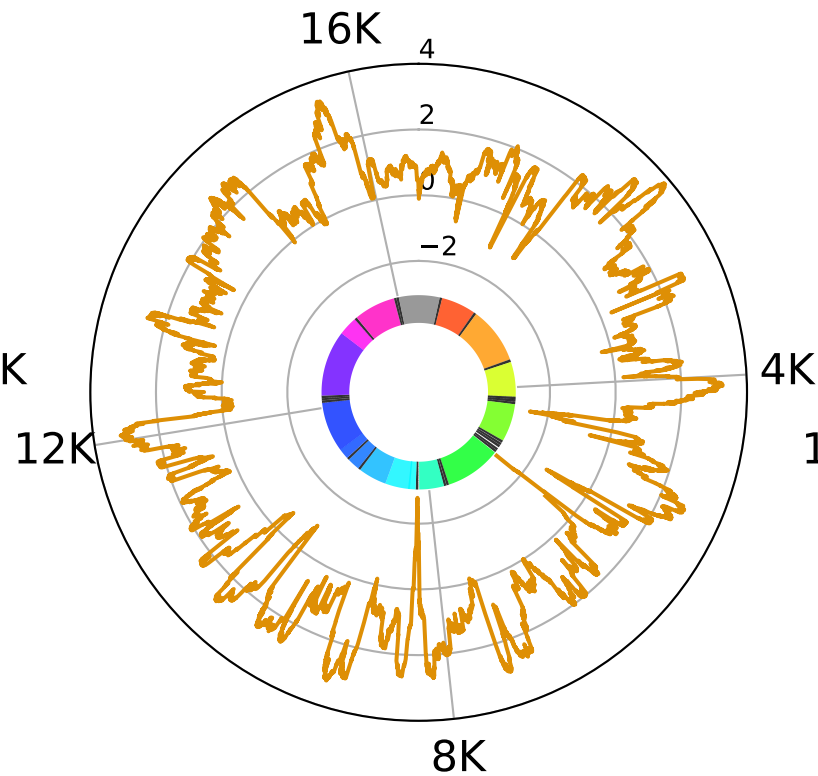# Average depth at base ($\log_{10}$)

smartseq2   CEL_SEQ2   10x

Legend: trna, RNR1, RNR2, ND1, ND2, CO1, CO2, ATP8, ATP6, CO3, ND3, ND4L, ND4, ND5, ND6, CYB, Control