

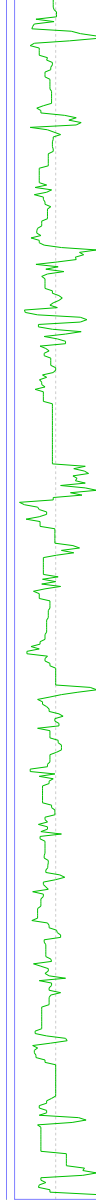
PCE8268_ZOOM_12398_29717



CG31760-RA

PCE8268

% Identity



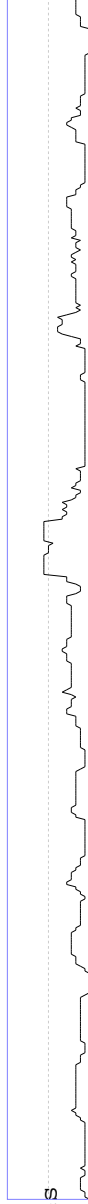
(0-100%)

alignment gaps (>80bp)

gray - D.mel gap



site density - all D.mel sites

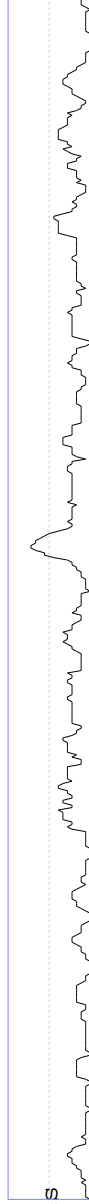


(0-18 sites)

Bcd
Cad
Hb
Kr
Kni

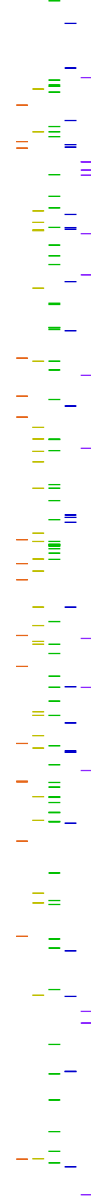


site density - all D.pse sites

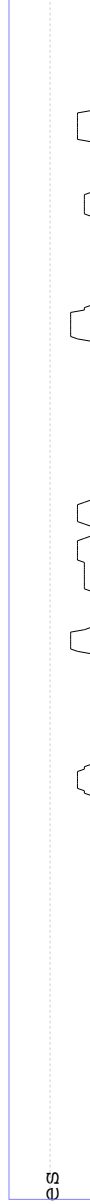


(0-18 sites)

Bcd
Cad
Hb
Kr
Kni



site density - all aligned sites



(0-12 sites)

Bcd
Cad
Hb
Kr

