

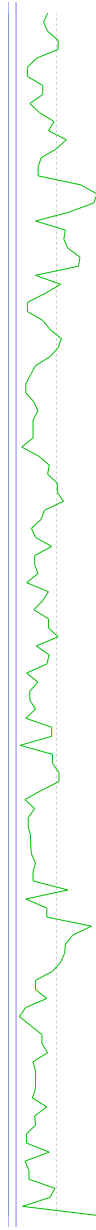
PCE8861_ZOOM_8800_19436



bru-3-RA

PCE8861

% Identity



alignment gaps (>80bp)

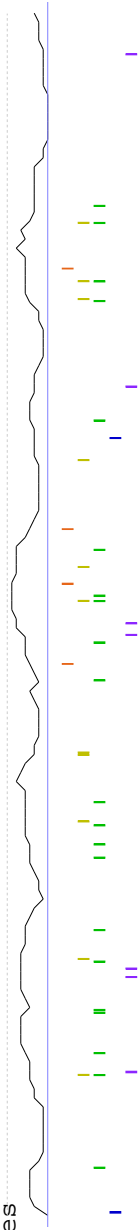
gray - D.mel gap



site density - all D.mel sites

(0-18 sites)

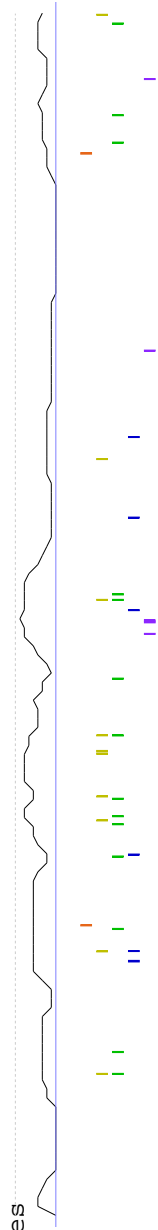
Bcd
Cad
Hb
Kr
Kni



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
Kni

