

PCE8362_ZOOM_13455_31234

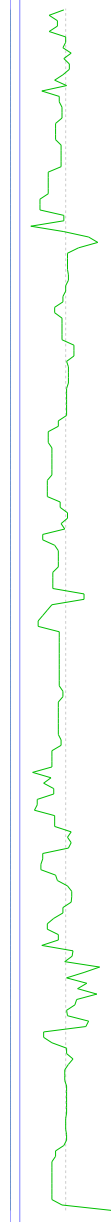


PCE8362



% Identity

(0-100%)



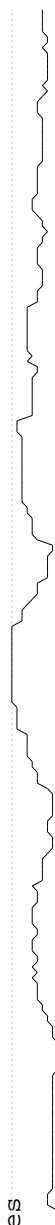
alignment gaps (>80bp)

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)

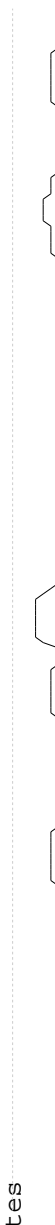


Bcd
Cad
Hb
Kr
Kni



site density - all aligned sites

(0-12 sites)



Cad
Hb

