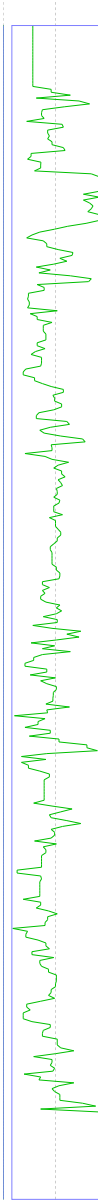




PCE8742



% Identity

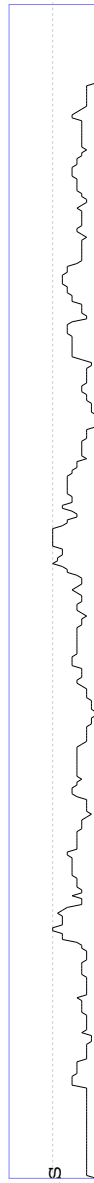


alignment gaps (>80bp)



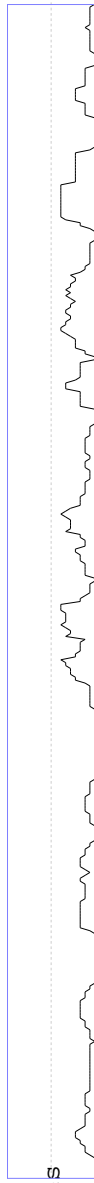
gray - D.mel gap

site density - all D.mel sites



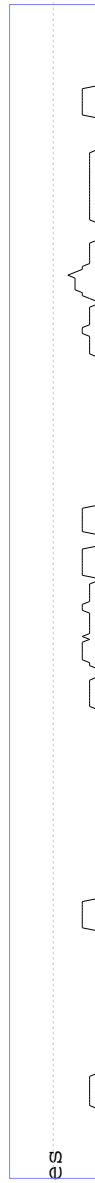
Bcd
Cad
Hb
Kr
Kni

site density - all D.pse sites



Bcd
Cad
Hb
Kr
Kni

site density - all aligned sites



Bcd
Cad
Hb
Kr
Kni