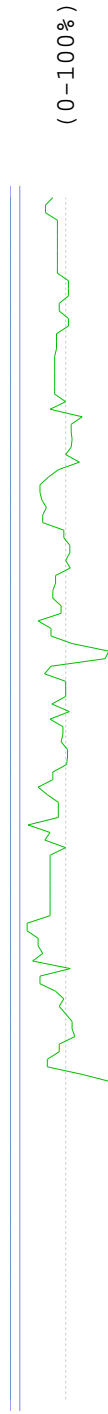


PCE8662_ZOOM_5927_18647



% Identity



alignment gaps (>80bp)



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



Cad
Hb

