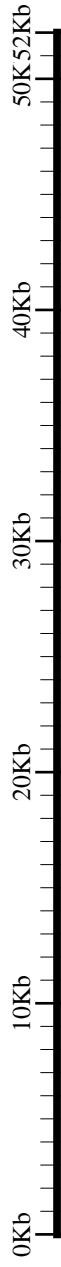


PCE8362_ZOOM_13455_31234

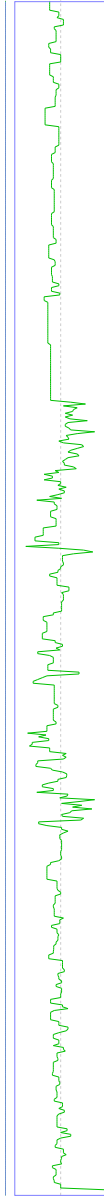


PCE8362

EG:BACR7A4.



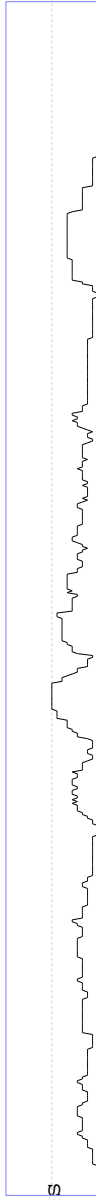
% Identity



alignment gaps (>80bp)

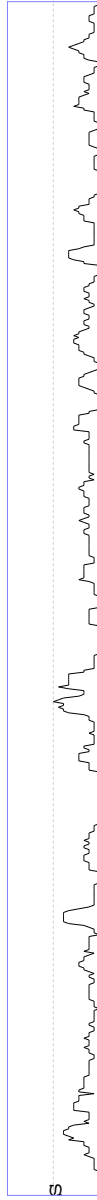


site density - all D.mel sites



Bcd
Cad
Hb
Kr
Kni

site density - all D.pse sites



Bcd
Cad
Hb
Kr
Kni

site density - all aligned sites



Cad
Hb