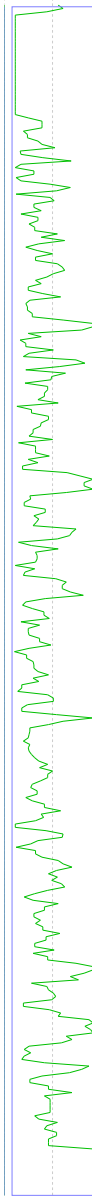


% Identity



(0-100%)

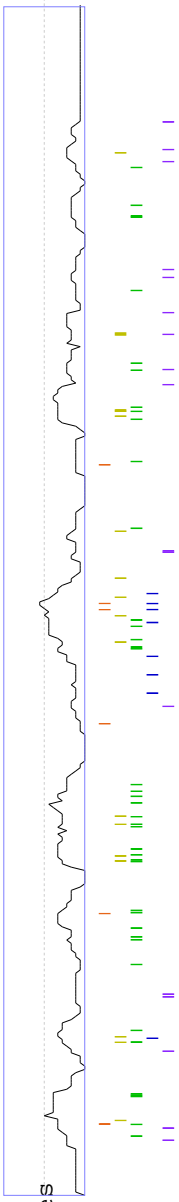
alignment gaps (>80bp)



gray - D.mel gap

site density - all D.mel sites

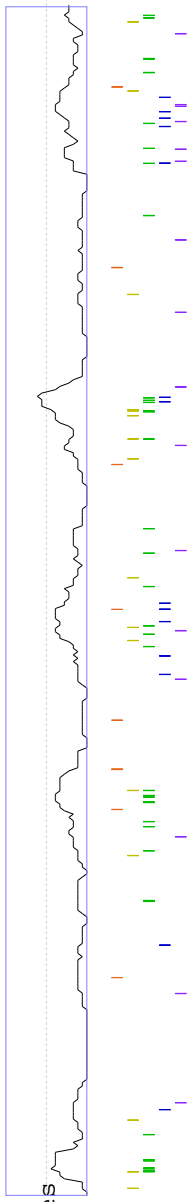
Bcd  
Cad  
Hb  
Kr  
Kni



(0-18 sites)

site density - all D.pse sites

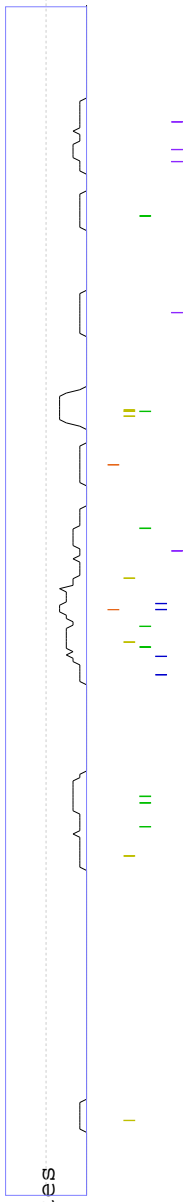
Bcd  
Cad  
Hb  
Kr  
Kni



(0-18 sites)

site density - all aligned sites

Bcd  
Cad  
Hb  
Kr  
Kni



(0-12 sites)