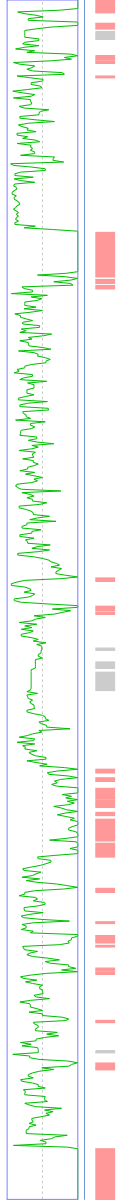


% Identity

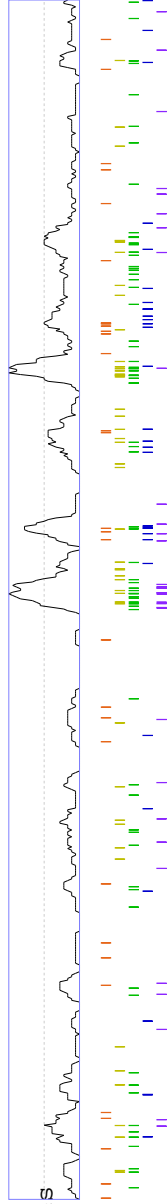


(0-100%)

alignment gaps (>80bp)

gray - D.mel gap

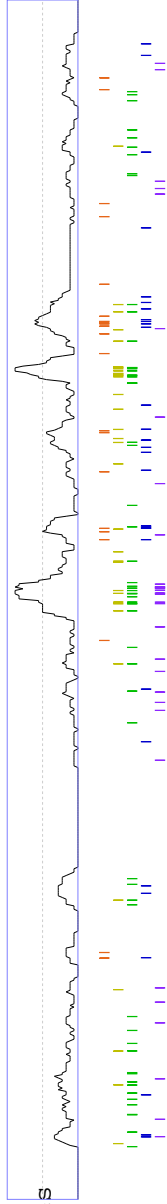
site density - all D.mel sites



(0-18 sites)

Bcd  
Cad  
Hb  
Kr  
Kni

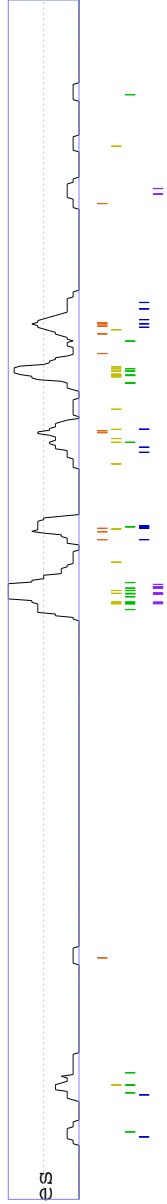
site density - all D.pse sites



(0-18 sites)

Bcd  
Cad  
Hb  
Kr  
Kni

site density - all aligned sites



(0-12 sites)

Bcd  
Cad  
Hb  
Kr  
Kni