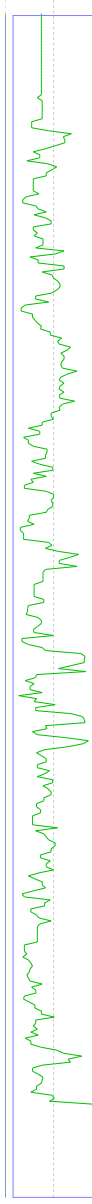




% Identity



(0-100%)

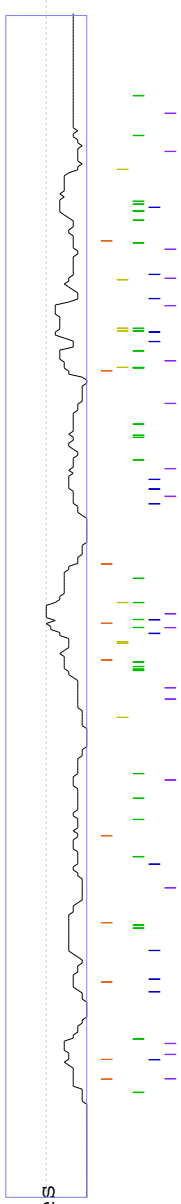
alignment gaps (>80bp)

gray - D.mel gap



site density - all D.mel sites

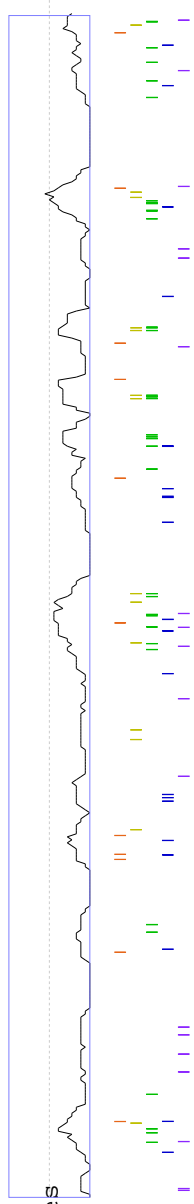
Bcd
Cad
Hb
Kr
Kni



(0-18 sites)

site density - all D.pse sites

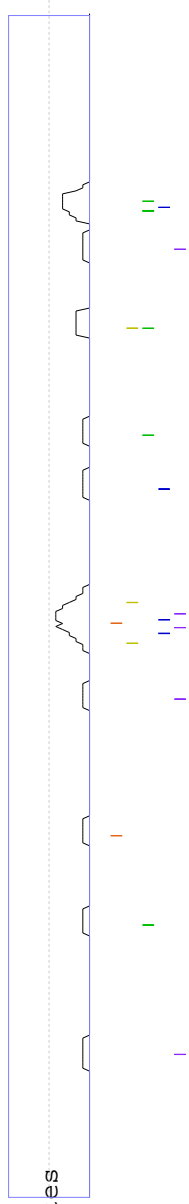
Bcd
Cad
Hb
Kr
Kni



(0-18 sites)

site density - all aligned sites

Bcd
Cad
Hb
Kr
Kni



(0-12 sites)