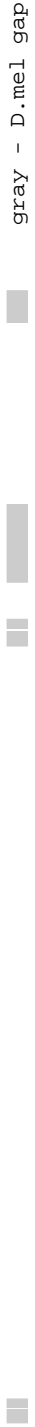


% Identity (0-100%)



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

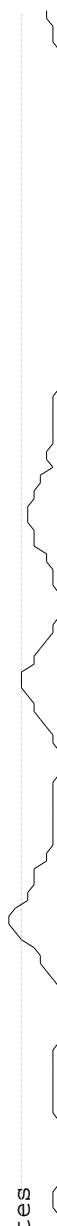


(0-18 sites)

Bcd
Cad
Hb
Kr
Kni



site density - all aligned sites



(0-12 sites)

Bcd
Cad
Hb
Kr
Kni

