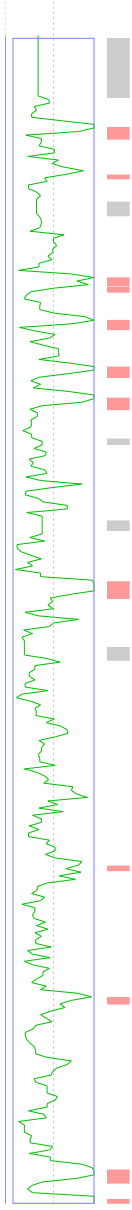


% Identity

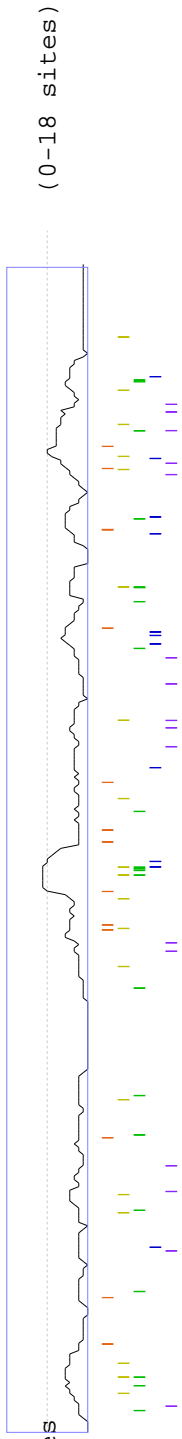


alignment gaps (>80bp)

gray - D.mel gap

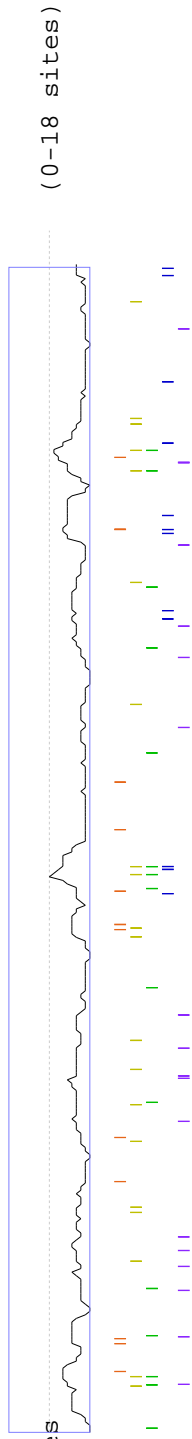
site density - all D.mel sites

Bcd
Cad
Hb
Kr
Kni



site density - all D.pse sites

Bcd
Cad
Hb
Kr
Kni



site density - all aligned sites

Bcd
Cad
Hb
Kr
Kni

