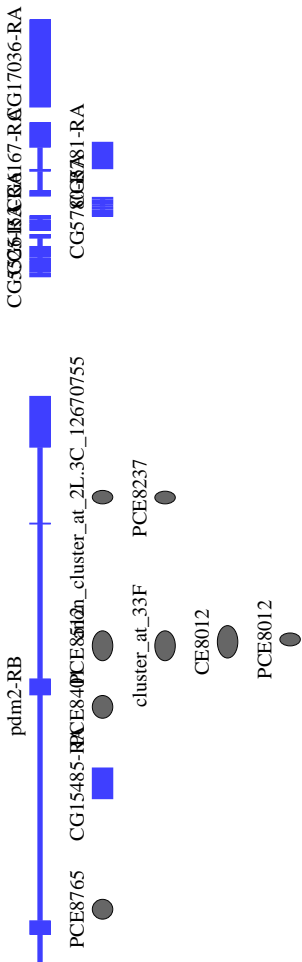
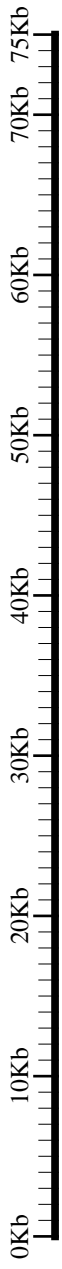
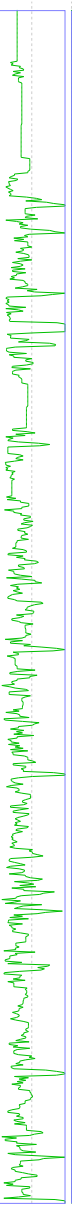


PCE8512_ZOOM_26016_37975



% Identity



(0-100%)

alignment gaps (>80bp)



gray - D.mel gap

site density - all D.mel sites

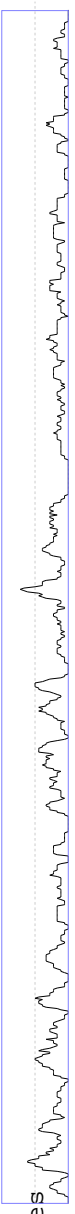


(0-18 sites)

Bcd
Ctfd
Hb
kr
Knl



site density - all D.pse sites



(0-18 sites)

Bcd
Ctfd
Hb
kr
Knl



site density - all aligned sites



(0-12 sites)

Bcd
Ctfd
Hb
kr
Knl

