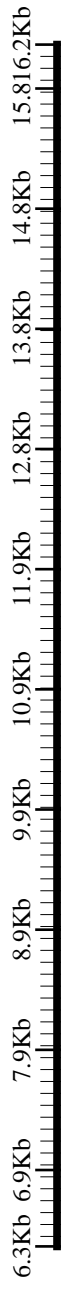
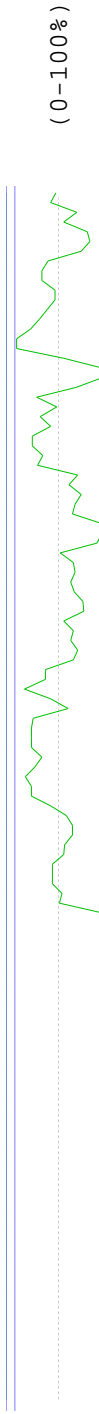


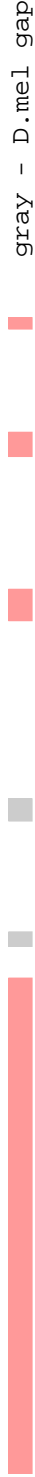
PCE8971_ZOOM_6288_16169



% Identity



alignment gaps (>80bp)



site density - all D.mel sites

(0-18 sites)



site density - all D.pse sites

(0-18 sites)



site density - all aligned sites

(0-12 sites)

