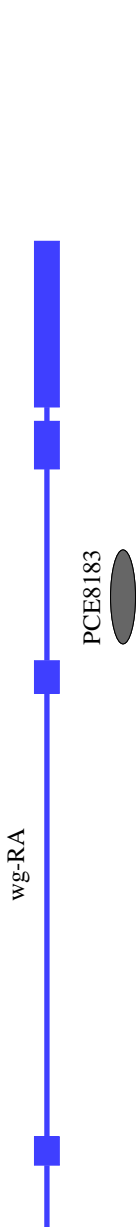
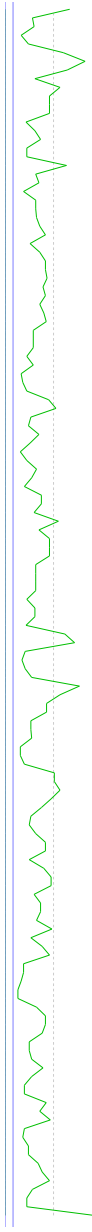


PCE8183_ZOOM_10057_21137



% Identity (0-100%)



alignment gaps (>80bp)

gray - D.mel gap

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)

