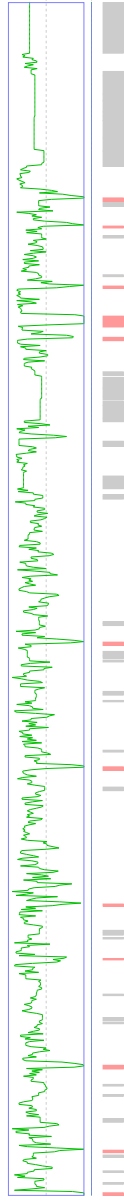


% Identity

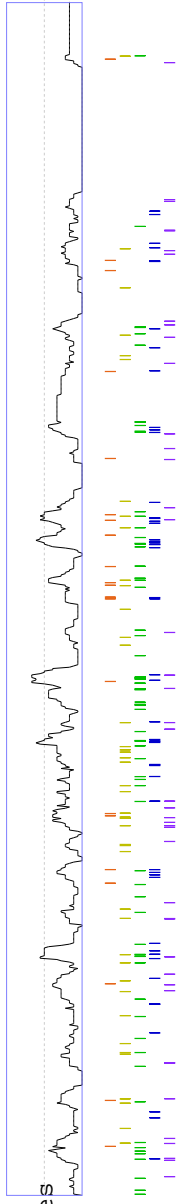


(0-100%)

alignment gaps (>80bp)

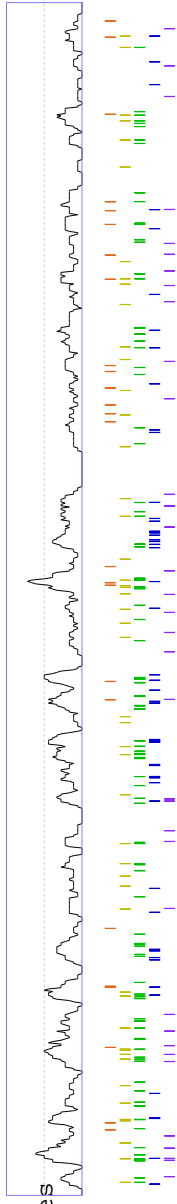
gray - D.mel gap

site density - all D.mel sites



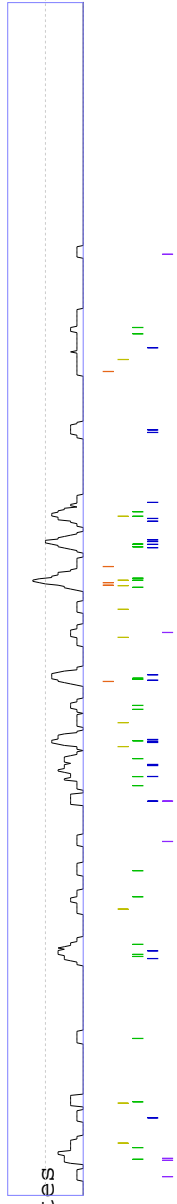
(0-18 sites)

site density - all D.pse sites



(0-18 sites)

site density - all aligned sites



(0-12 sites)