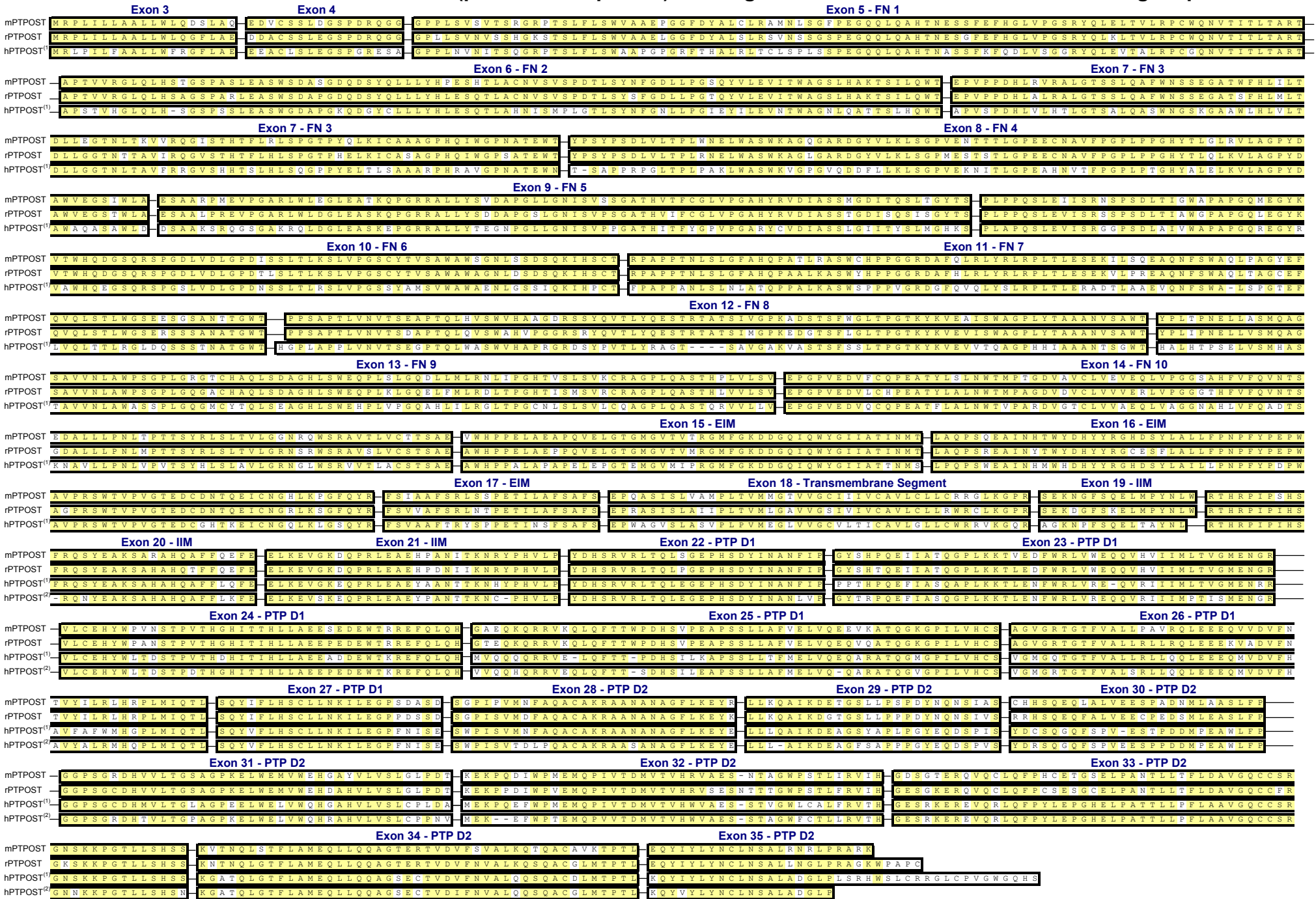


Exon structure of human PTP-OST (predicted sequence) and alignment with the mouse and rat ortholog sequences



⁽¹⁾ This hOST-PTP sequence is predicted from genomic clones AL592300 and AL356953. These clones map to 1q32 - a region with synteny to the rat and mouse loci for OST-PTP.

⁽²⁾ This hOST-PTP sequence is a fragment and is predicted from genomic clone AL354751, which in the current assembly (Build 31) has been aligned to Chromosome 9

FN - Fibronectin type III repeat; EIM - Extracellular domain inter-motif; IIM - Intracellular domain inter-motif; D1 - Domain 1; D2 - Domain 2 (Nomenclature according to Morrison & Mauro, Gene 257 (2000) p.195-208)