Figure S5. Alignment of amino acid sequences from mouse Cwc22R2d1, Cwc22R2d2 and retro-Cwc22, plus Cwc22 orthologs from 19 other placental mammals plus opossum, platypus and chicken as outgroups. Residues are colored according to biochemical properties and gaps are shown in grey. Information content of each column in the alignment, measured as the Jenson-Shannon divergence, is plotted in the lower panel.