



Supplementary Figure 5 : Phylogenetic tree of *jsd* sequences. Unrooted tree generated by neighbor-joining/UPGMA method version 3.572c. All *jsd* orthologs (processed pseudogenes) are represented by the name of the species that they were sequenced from, the X-linked progenitor genes are indicated by species name followed by an X, and the only identified pseudogene (found in mouse) is indicated by mouseP.