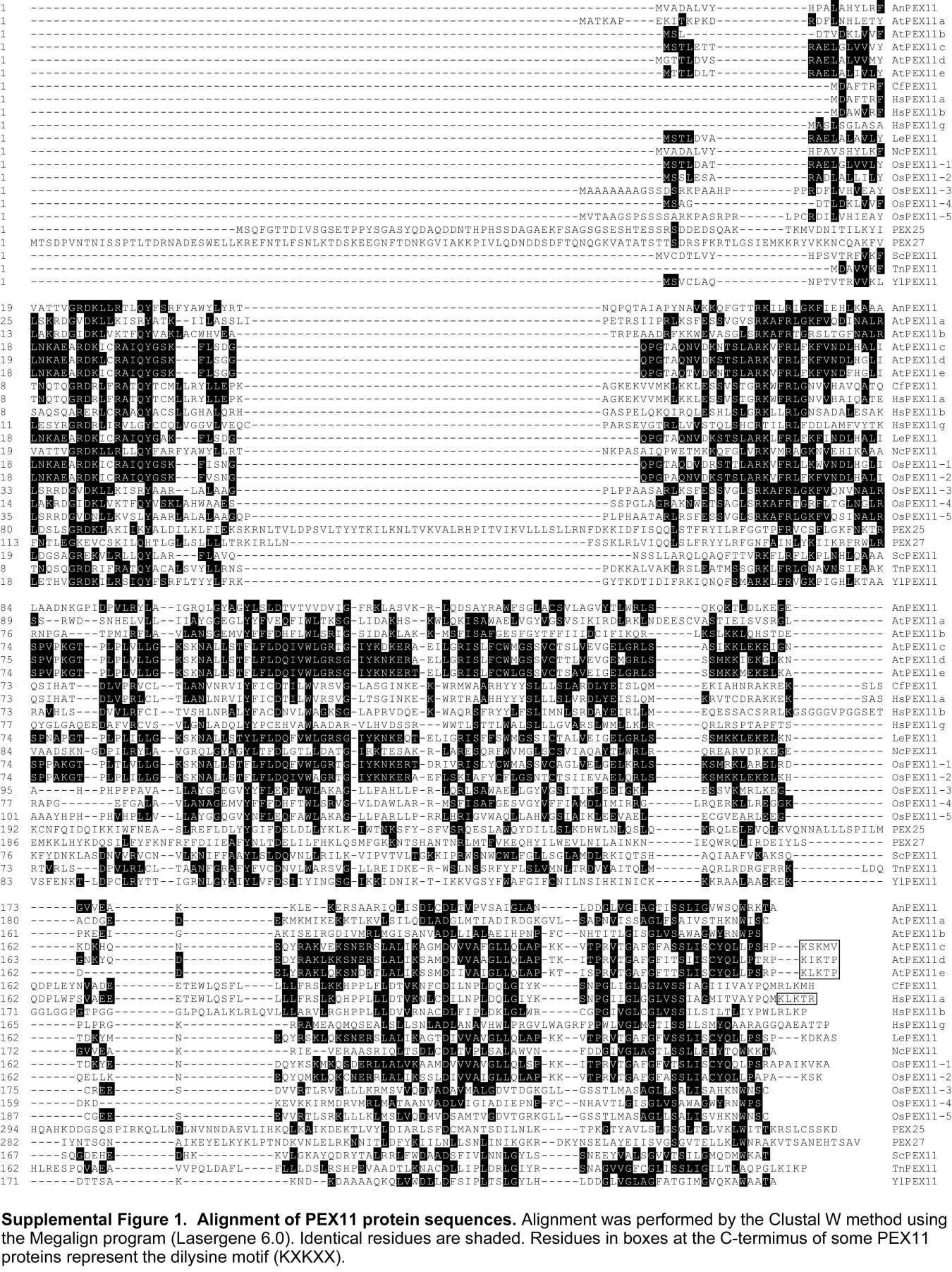

Supplemental Figure 1. Alignment of PEX11 protein sequences. Alignment was performed by the Clustal W method using the Megalign program (Lasergene 6.0). Identical residues are shaded. Residues in boxes at the C-termimus of some PEX11 proteins represent the dilysine motif (KXKXX).