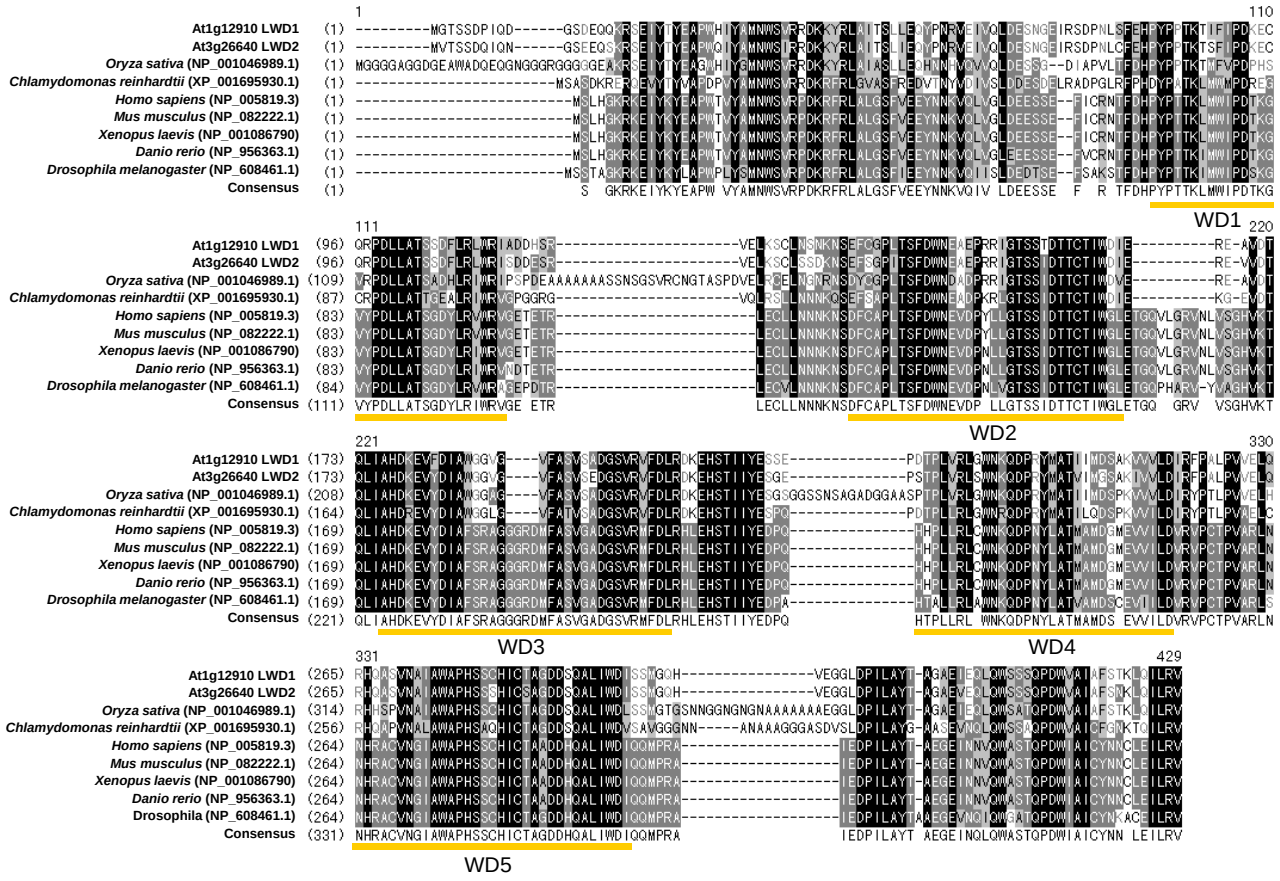

Supplemental Figure S7. Amino acid sequence alignment of LWD1 orthologous proteins.

Amino acid sequences of LWD1, LWD2 and their orthologs are aligned by use of Vector NTI AlignX. Identical amino acids are shaded in black. The dark grey shading represents positions with conserved amino acids. The light grey shading represents positions with similar amino acids. The 5 WD repeats were marked according to positions in LWD1 and LWD2. The alignment shows two distinct groups for LWD orthologs originating from plant or non-plant species.