**Supplementary Figure 7.** Neighbor-joining phylogenetic tree from an amino acid alignment of Glyma.18G026900, the Arabidopsis homolog and paralogs AtCCR3, AtCCR4, and AtCR4, as well as paralogs from soybean (Wm82.a2.v1) built with Clustal Omega and EMBL Simple Phylogeny (Sievers and Higgins, 2014; Goujon et al., 2010).