

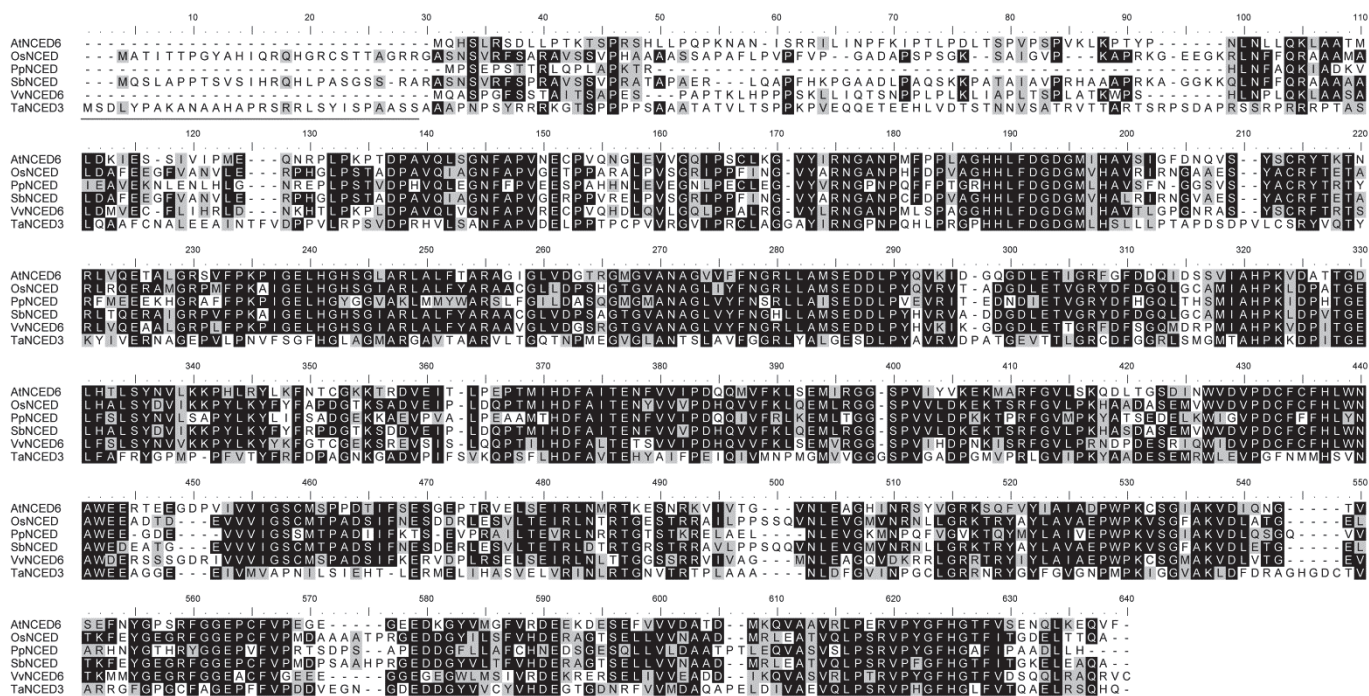


Fig. 1 Suppl. The multiple alignment of deduced amino acid sequences of TaNCED and NCED proteins from other species. Identical amino acids are shaded in *black* and similar amino acids are shaded in *gray*. The region *underlined* indicates the putative chloroplast transit peptide. Accession numbers for the proteins are as follows: OsNCED (NP_001050765.1), SbnCED (XP_002466753.1), AtNCED6 (NP_189064.1), VvNCED6 (XP_002283185.), and PpNCED (XP_001784596.1).

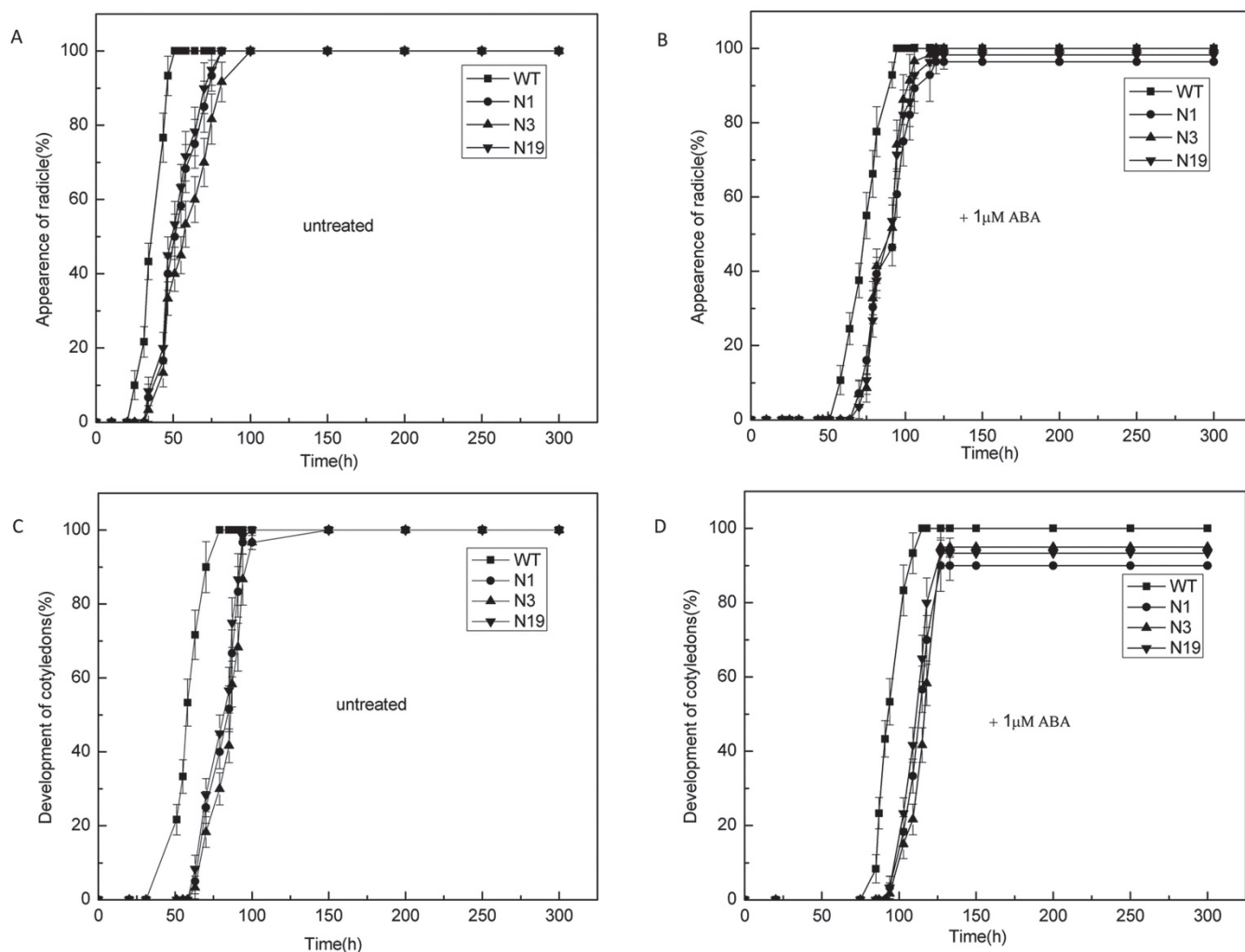


Fig. 2 Suppl. Seed germination assay of transgenic and wild type (WT) plants with or without ABA. *A* - Appearance time of a radicle without ABA. *B* - Appearance time of a radicle with 1 μ M ABA. *C* - Time for full development of a cotyledon without ABA. *D* - Time for full development of a cotyledon with 1 μ M ABA. The data represent means $\pm$ SEs ($n = 3$).