Molecular characterization of *OsCURT1A* from upland rice in response to osmotic stress

Bernadette Toni, Nurulhikma Md Isa, Cheng Seng Tan, Ismanizan Ismail, Zamri Zainal*

Supplementary information

Sequence data used for phylogenetic analysis are available in the NCBI database (www.ncbi.nlm.nih.gov) under the following accession numbers:
AtCURT1A(XP_020180454.1), SbCURT1A (XP_021315547.1), OslCURT1A (EAZ00433.1), DoCURT1A (OEL26938.1), ZmCURT1B (ONM56978.1), SiCURT1B (XP_004957648.1), BdCURT1B (XP_003563146.1), OsjapCURT1B (XP_015645395.1), OsjapCURT1C (XP_015621702.1), ObraCURT1C (XP_006646347.1), AcCURT1C (XP_020099268.1), ZmCURT1D (XP_008646297.2), OsjapCURT1D (XP_015623062.1), PdCURT1D (XP_008802300.1), AcCURT1D (OAY85311.1), GmCURT1A (XP_003548597.1), VaCURT1A (XP_017410228.1), CcCURT1A (XP_020233680.1), AiCURT1A (XP_016180476.1), LaCURT1A (XP_019456941.1), CaCURT1A (XP_004494779.1), ApCURT1A (KFM26712.1), Chlorella1 (EFN53210.1), Micromonas1 (XP_003058644.1), Monoraphidium1 (XP_013892548.1), Oscillatoria1 (CBN54640.1), Arthrospira1 (EDZ94246.1), Raphidiopsis1 (EFA71519.1), Microcystis1 (CAO88457.1) and Strepto aaRSs (AKI57929.1).