



Additional file 1. Number and length distributions of predicted open reading frames. (a) Number and length distributions of predicted open reading frames (ORFs) in 53,861 non-redundant transcripts in sweet potato (Ib53861) and 51,184 in *I. trifida* (It51184). (b,c) Number and length distributions of characteristic components of 34,963 non-redundant full-length transcripts in sweet potato (b) and 33,637 in *I. trifida* (c), UTR, untranslated region; ORF, open reading frame.