

Supporting information

Crystal structure of a glycoside hydrolase family 68 β -fructosyltransferase from *Beijerinckia indica* subsp. *indica* in complex with fructose

Takashi Tonozuka, Junichi Kitamura, Mika Nagaya, Reika Kawai, Atsushi Nishikawa, Katsuaki Hirano, Keisuke Tamura, Tadashi Fujii, Takumi Tochio

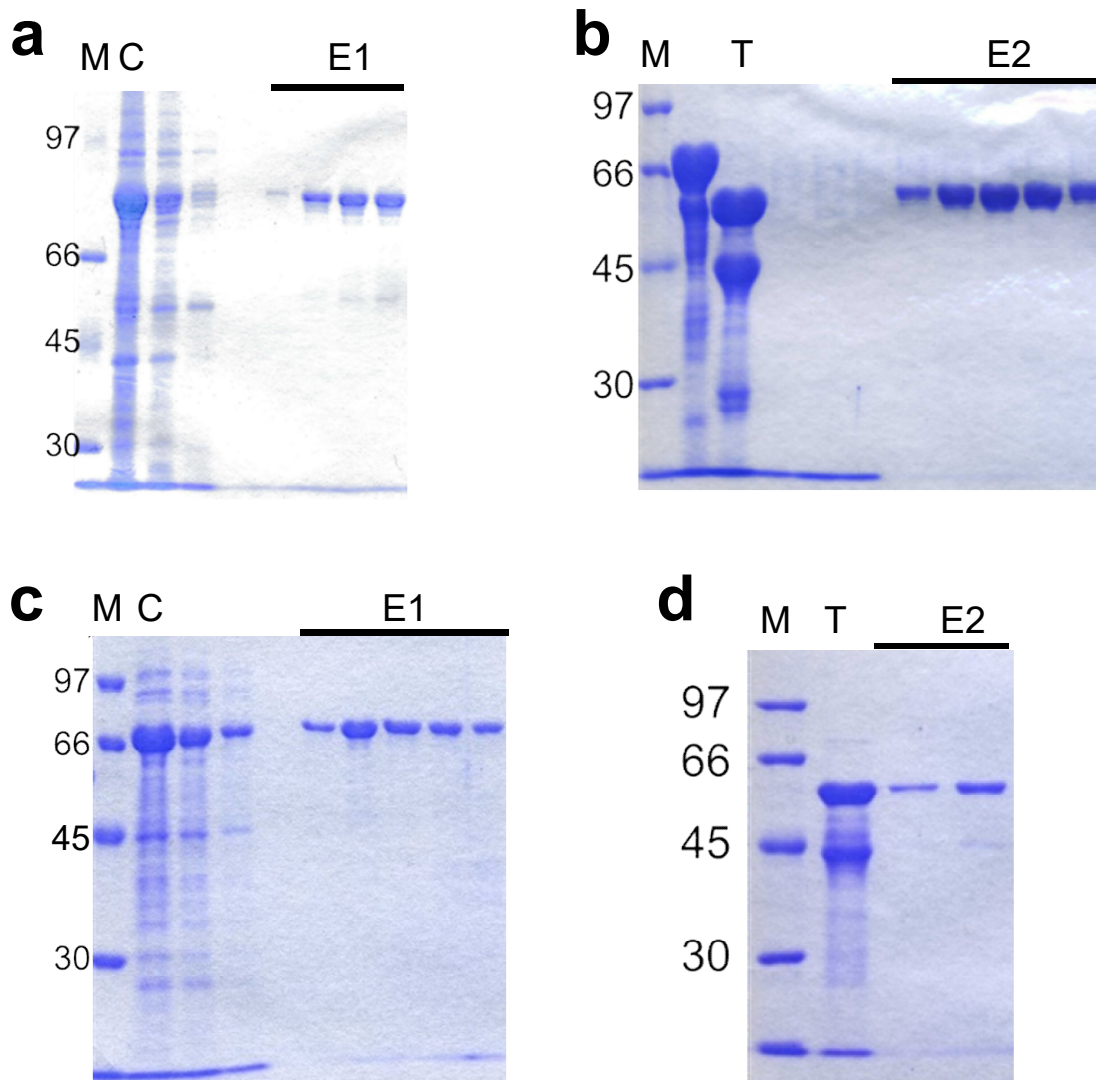


Figure S1. SDS-PAGE showing the purification of BiBftA and BiBftA $\Delta A\Delta C$. (a) Purification of thioredoxin-His-tagged full-length BiBftA using Ni-NTA agarose. (b) Purification of full-length BiBftA using HiTrap Q HP column. (c) Purification of thioredoxin-His-tagged BiBftA $\Delta A\Delta C$ using Ni-NTA agarose. (d) Purification of BiBftA $\Delta A\Delta C$ using HiTrap Q HP column. Lanes: M, molecular weight marker, numbers indicate molecular sizes in kDa; C, crude enzyme; E1, elution from Ni-NTA agarose; T, after treatment with thrombin; E2, elution from HiTrap Q HP column.