Additional file 1. Representative ITC data of the interactions of CuaA and CuaD with cellodextrins

Interactions of CuaA (A) and CuaD (B) with cellobiose are shown. The top half of each panel represents the raw binding heats and the lower half represents the integrated binding heats minus the dilution control heats fitted to a single-site binding model. Insert presents the binding parameters, with n designating the number of binding sites on the protein.