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Supporting information for article:

Crystal structure of the pseudoenzyme PDX1.2 in complex with its cognate enzyme PDX1.3: A total eclipse

Graham R. Robinson, Markus Kaufmann, Céline Roux, Jacobo Martinez-Font, Michael Hothorn, Stéphane Thore and Teresa B. Fitzpatrick **Supplementary Figure S1** Intensity distributions of crystals of the PDX1.2/PDX1.3 complexes. (A) and (C), L-test and NZ-test, respectively, for the dataset of the PDX1.2/PDX1-3 complex. (B) and (D), L-test and NZ-test, respectively, for the PDX1.2/PDX1.3-adduct dataset. Intensity analyses were performed with XTRIAGE (<u>Adams *et al.*</u>, 2010).

