Automated Structure- and Sequence-Based Design of Proteins for High Bacterial Expression and Stability

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As a result of an author oversight in the originally published version of this article, the Rosetta script file design.xml provided in Data S1 was missing the res_type_constraint value, causing code crash. The value appears in the script filterscan.xml that is used just before design.xml. We updated the file design.xml, which now includes the res_type_constraint value reported in the main text (0.4). In addition, a file called design_text had no use and was omitted in this update. The authors apologize for the error and any inconvenience that may have resulted.

SUPPLEMENTAL INFORMATION

Supplemental Information includes a data file containing RosettaScripts, flags, and command lines and can be found with this article online at https://doi.org/10.1016/j.molcel.2018.03.035.