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Materials List for

Computational Prediction of Amino Acid Preferences of Potentially Multispecific Peptide-Binding Domains Involved in Protein-Protein Interactions

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Materials

Name	Company	Catalog Number	Comments
BUDE Alanine Scan Server	University of Edinburgh	https://pragmaticproteindesign.bio.ed.ac.uk/balas/	doi: 10.1021/acschembio.9b00560
Rosetta Modeling Software	Rosetta Commons	https://www.rosettacommons.org/software	doi: 10.1002/prot.22851
UCSF Chimera	University of California San Francisco	https://www.cgl.ucsf.edu/chimera/	doi: 10.1002/jcc.20084