

Supporting Information

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Engineering the Fullerene-protein Interface by Computational Design: The Sum is More than its Parts

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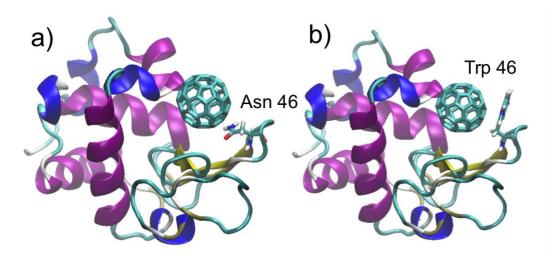


Figure S1. a) Wild-Type Lysozyme in complex with C_{60} - Residue Asn 46, b) Mutated Lysozyme N46W

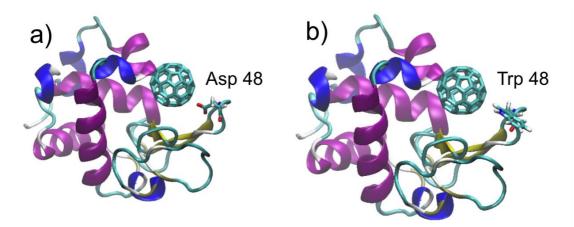


Figure S2. a) Wild-Type Lysozyme in complex with C_{60} - Residue Asp 48, b) Mutated Lysozyme D48W

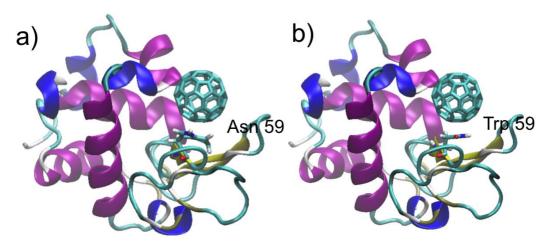


Figure S3. a) Wild-Type Lysozyme in complex with C_{60} - Residue Asn 59, b) Mutated Lysozyme N59W.

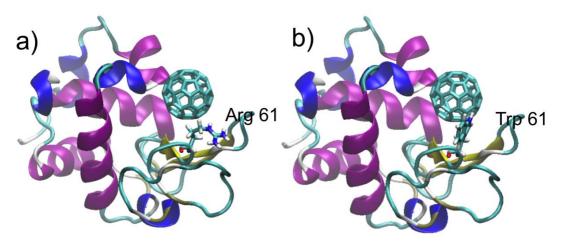


Figure S4. a) Wild-Type Lysozyme in complex with C_{60} - Residue Arg 61, b) Mutated Lysozyme R61W.

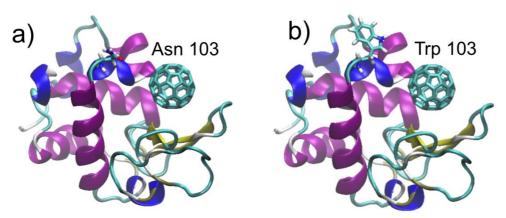


Figure S5. a) Wild-Type Lysozyme in complex with C_{60} - Residue Asn 103, b) Mutated Lysozyme N103W.

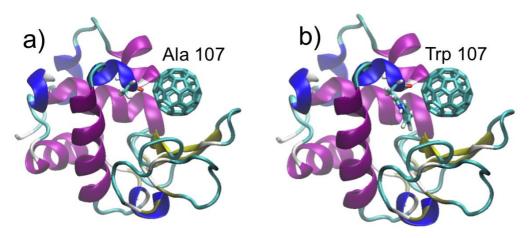


Figure S6. a) Wild-Type Lysozyme in complex with C_{60} - Residue Ala 107, b) Mutated Lysozyme A107W.

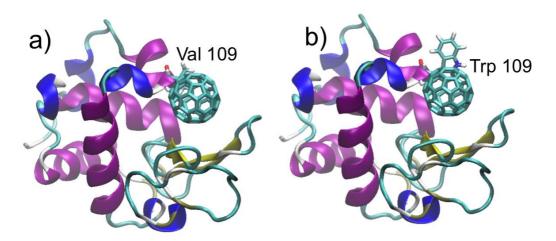


Figure S7. a) Wild-Type Lysozyme in complex with C_{60} - Residue Val 109, b) Mutated Lysozyme V109W.