

## Supporting Information

# Application of enhanced sampling Monte Carlo methods for high-resolution protein-protein docking in Rosetta

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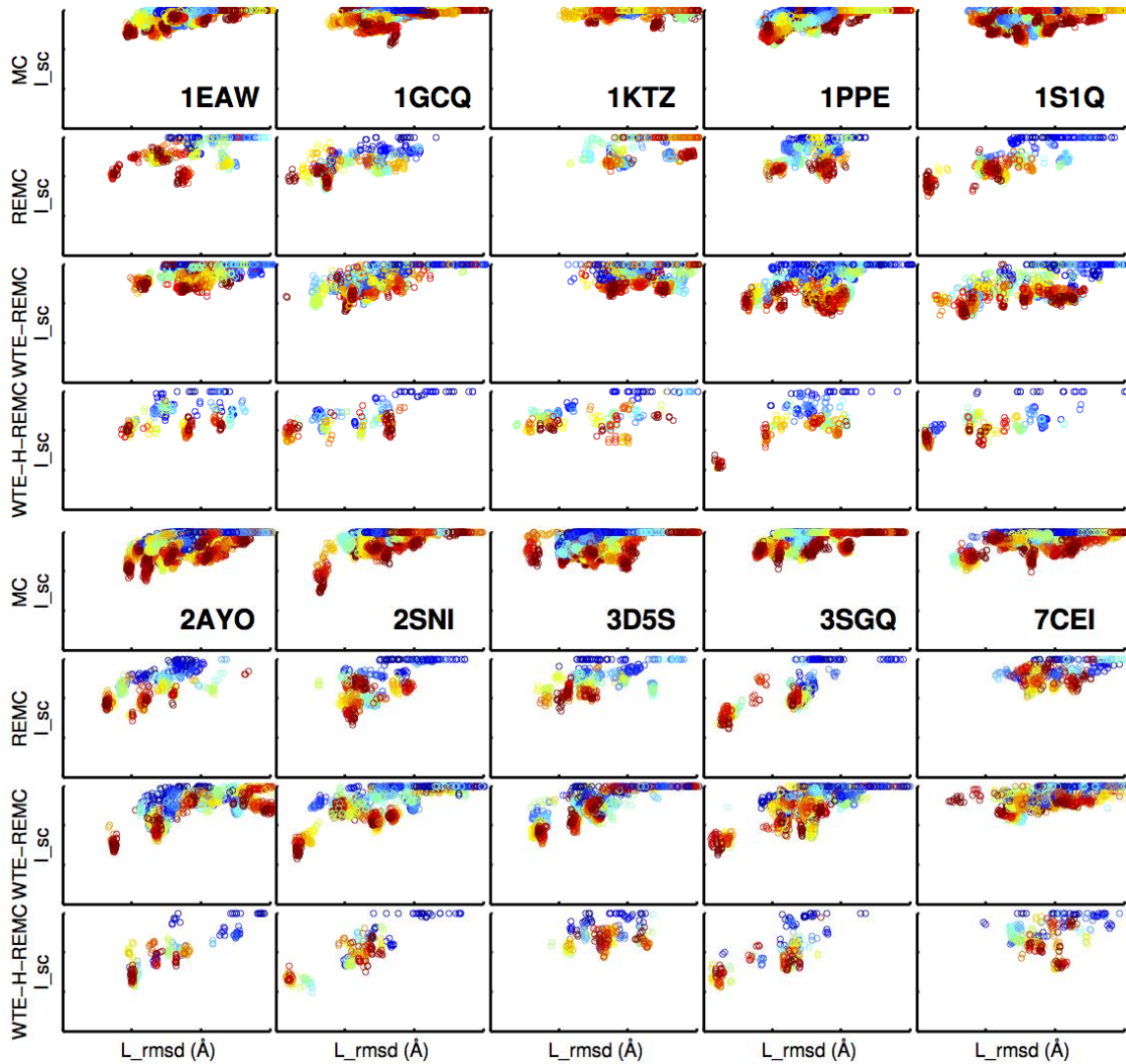
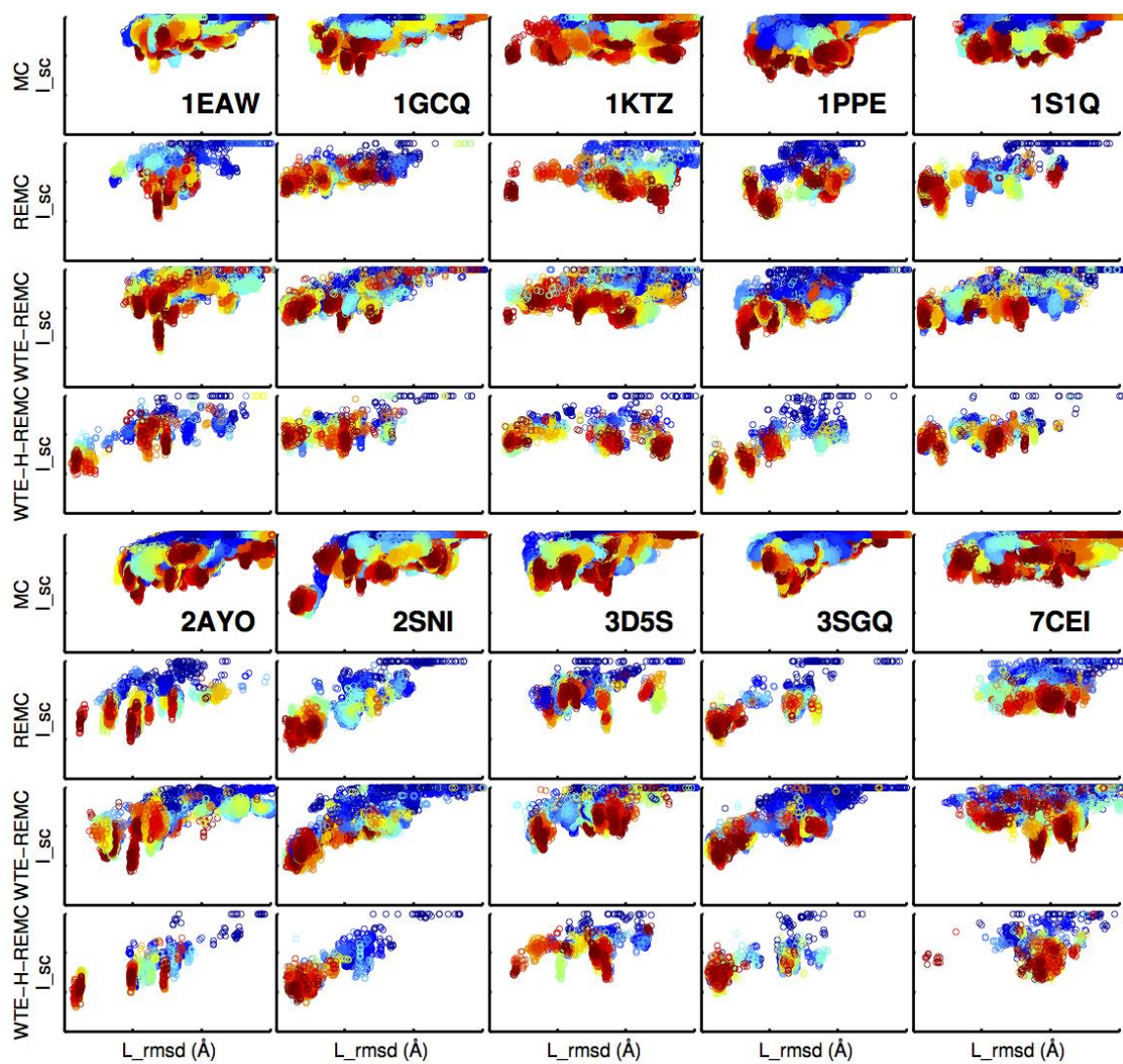
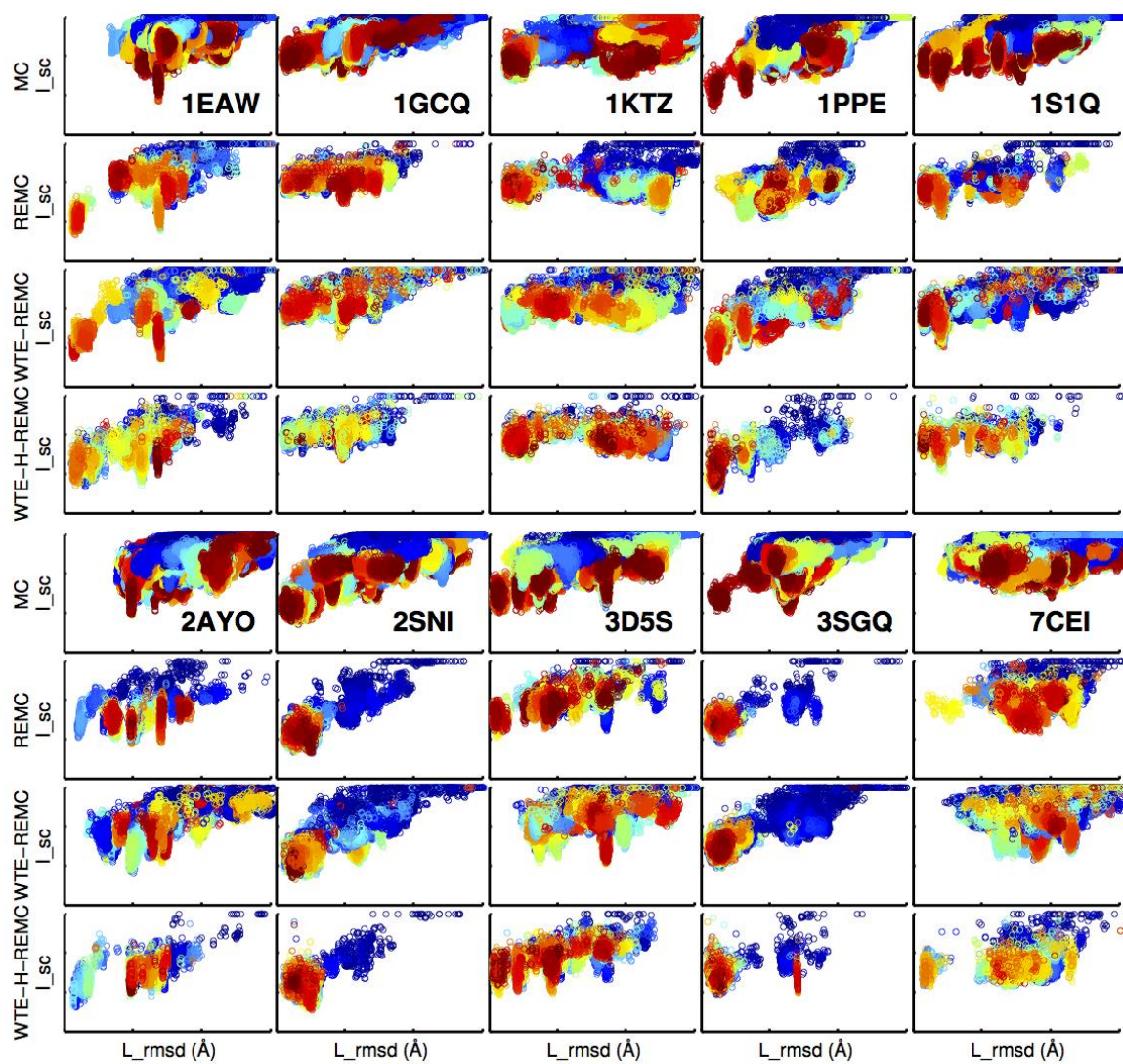


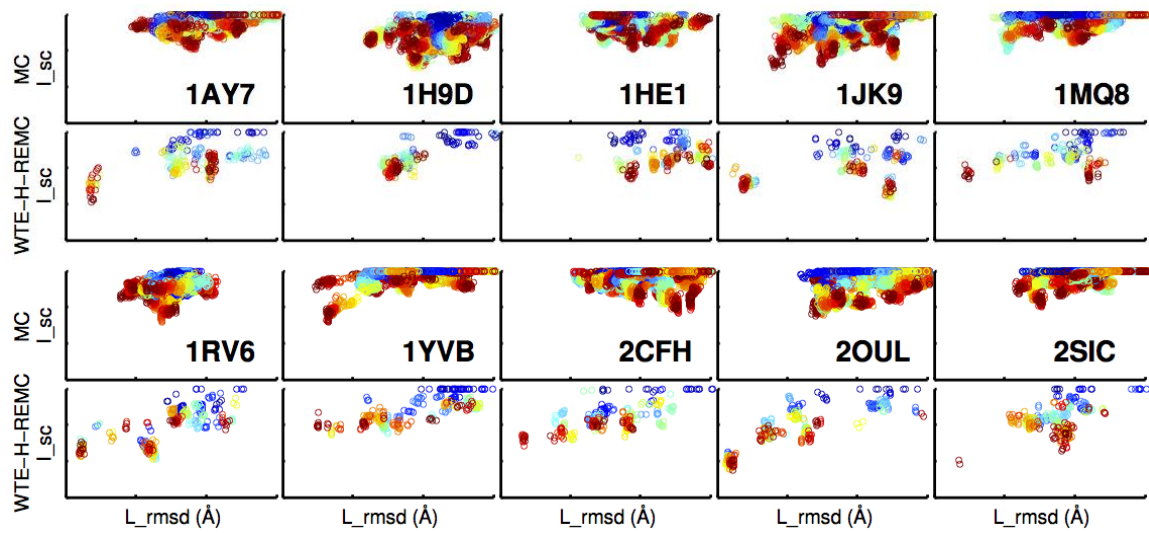
Figure A.



**Figure B.**

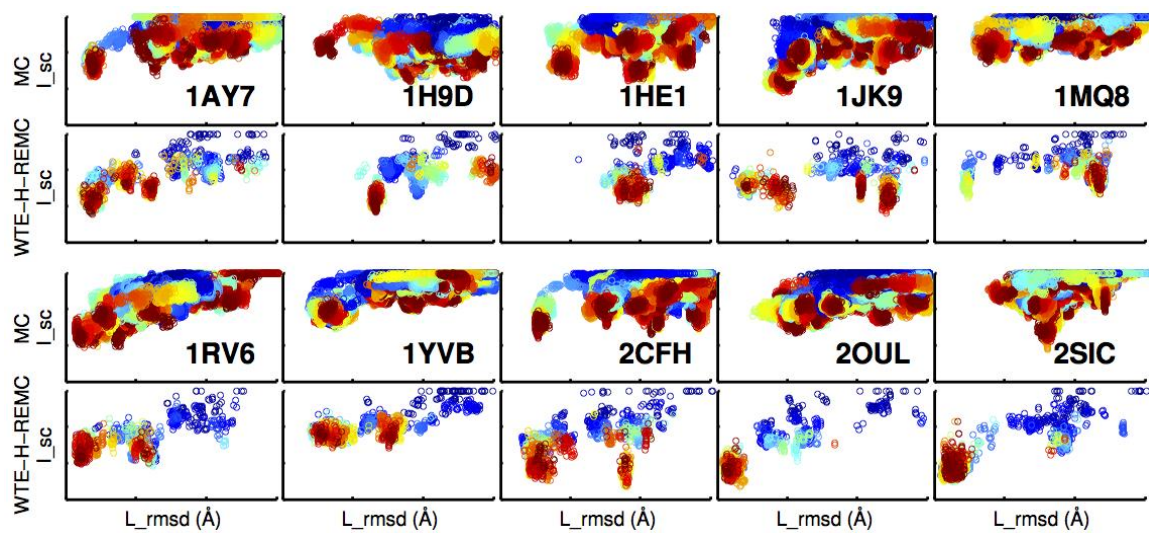


**Figure C.**

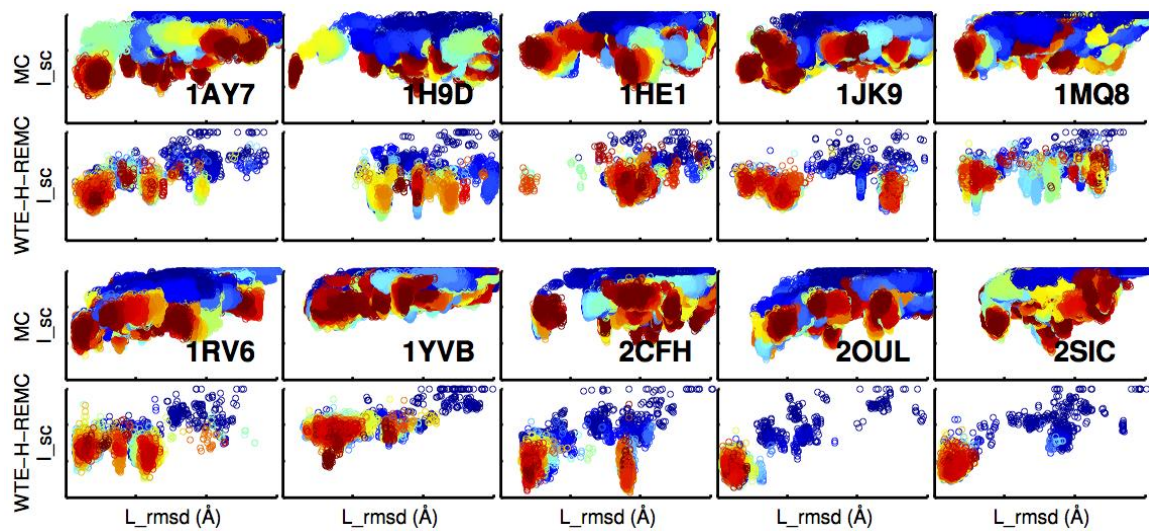


**Figure D.**





**Figure E.**



**Figure F.**