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26 Figure S2. Conservation of the protein structure of the MCR family.

27 A surface representation shows the conservation analysis calculated across 106 MCR family proteins,

28 in which red represents low homologous positions and green represents high conservation. The linker

29 region and the mutated residues are shown in cartoon and sticky representations, respectively. The

30 protein structure was analysed and plotted by PyMOL 2.6 software.

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