

# Supporting Information for “A Highly Dissociative and Concerted Mechanism for the Nicotinamide Cleavage Reaction in Sir2Tm Enzyme Suggested by *ab initio* QM/MM Molecular Dynamics Simulations”

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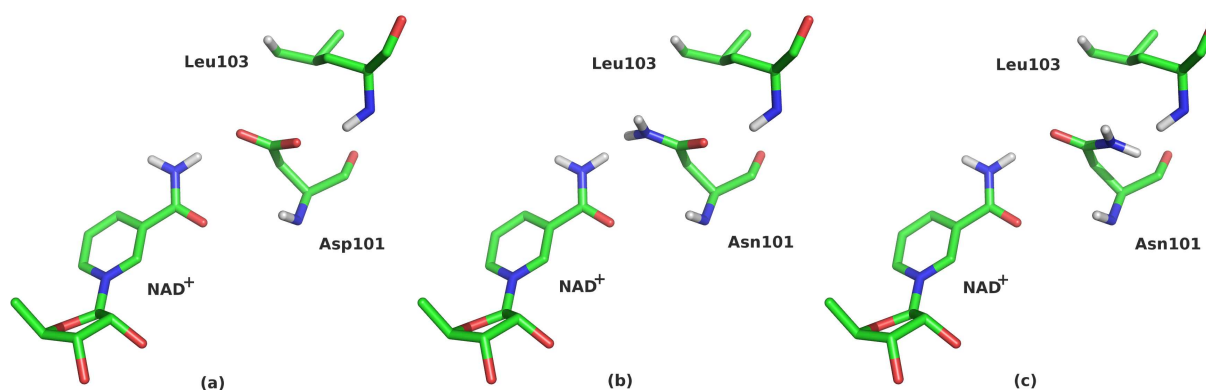


Figure S1: Illustration of two sets of Sir2Tm D101N mutations. (a) Wild-type Sir2Tm enzyme; (b) The first D101N mutation, where the carboxyl oxygen hydrogen-bonding to the NAD<sup>+</sup> was replaced with an amide group; (c) The second D101N mutation, where the carboxyl oxygen hydrogen-bonding to the Leu103 was replaced with an amide group.

## Complete list of references 10, 20, 66 and 67.

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