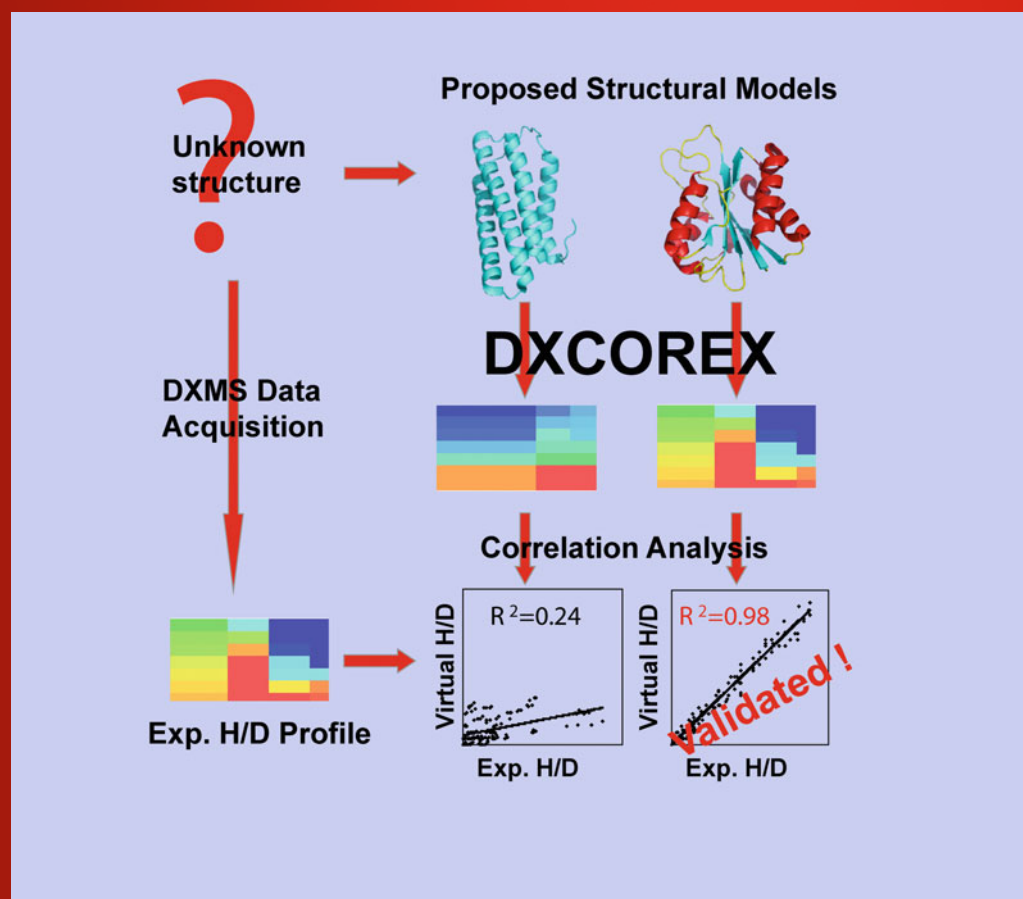


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Assessment of Protein Structural Models by Comparison of H/D Exchange MS Data with DXCOREX-predicted Exchange Behavior, see page 43.

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