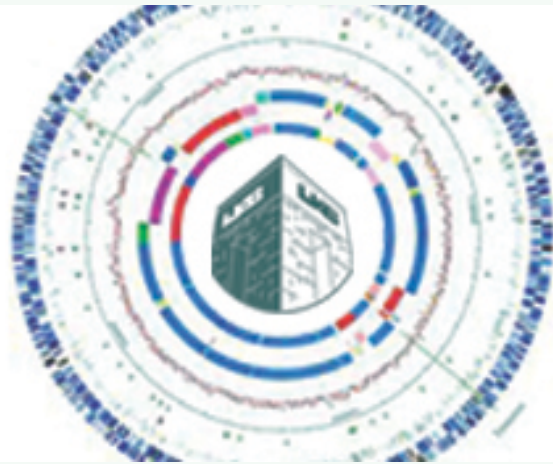


CB²



COMPUTATIONAL BIOLOGY
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CB2 Journal Club and Seminars

Shivani Viradia

Research Assistant
Department of Pathology

Combining structural modeling with ensemble machine learning to accurately predict protein fold stability and binding affinity effects upon mutation.

PLOS ONE. PMID: 25243403

Thursday, April 30, 2015 at 10:00 AM
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