

Supplementary data:

Novel mutation in ATP-binding domain of *ABCD1* gene in adrenoleucodystrophy

Neeraj Kumar, Krishna K. Taneja, Atul Kumar, Deepti Nayar, Bhupesh Taneja, Satindra Aneja, Madhuri Behari, Veena Kalra and Surendra K. Bansal

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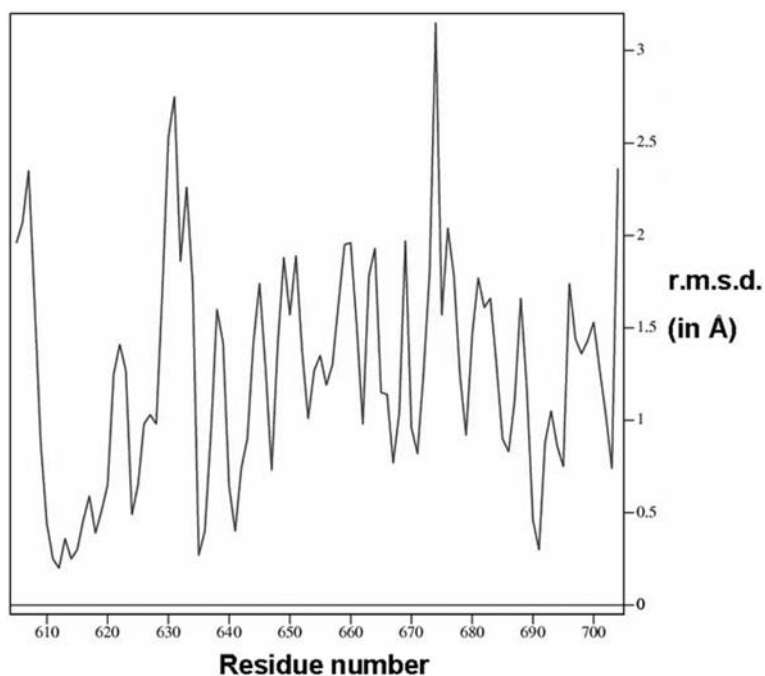


Figure 1. Rmsd plot of native and Arg617Ser substituted models. A plot of rmsd per residue for the two models after equal number of minimization cycles using AMBER. Largest deviations are clearly seen at the termini and at loops encompassing residues 629–642 and residues 674–681. Deviations at other regions or loops are much less than these regions.