

Characterization and catalytic-site-analysis of an aldo-keto reductase with excellent solvent tolerance

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Figure legends:

Fig.S1. Alignment of the sequence of AKR3-2-9 with the sequence of 1og6.1.

Fig.S2. Summary of the structure of other aldo-keto reductases in the PDB.

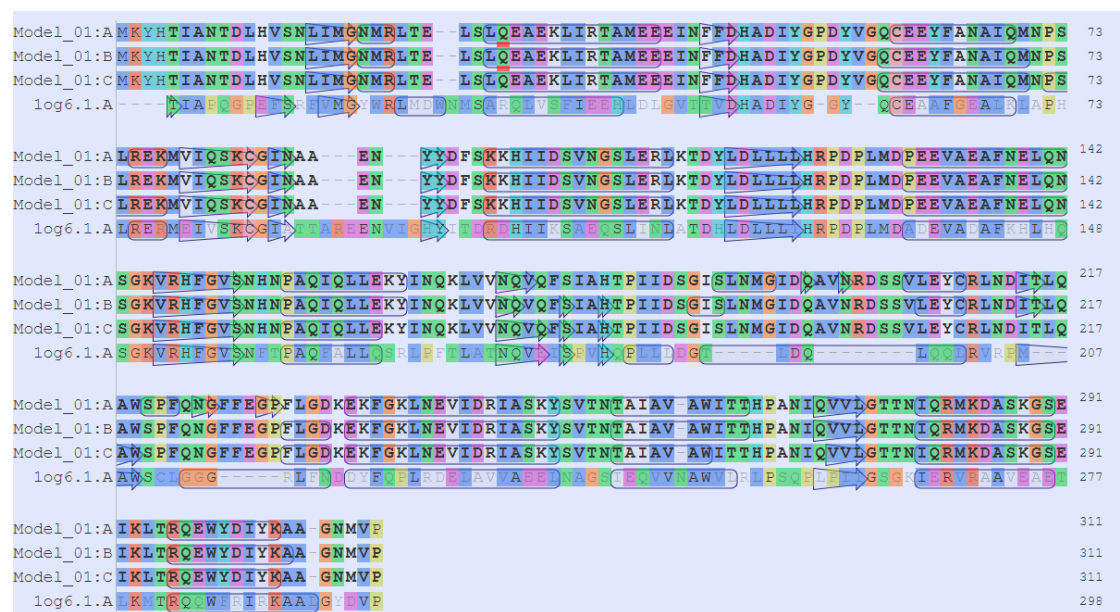


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