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Supporting information for article:

LAHMA: structure analysis through local annotation of homology-matched amino acids

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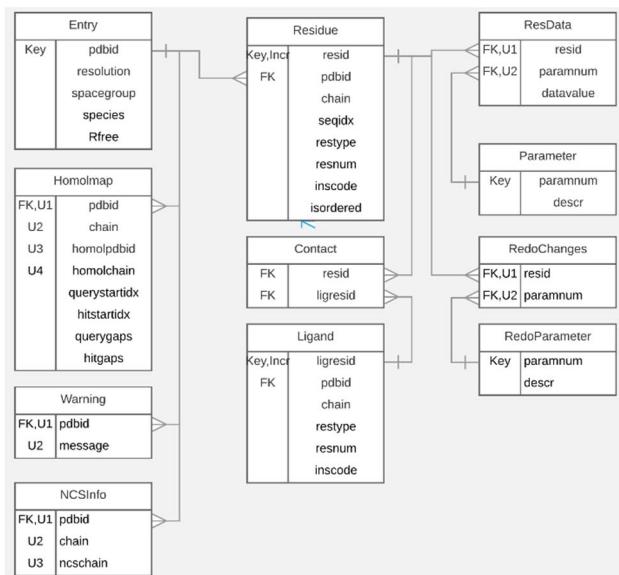


Figure S1 Scheme of the database. The following information describes the tables: key, primary key of the table; FK, foreign key in a different table; Incr, this variable is auto-incremented; U1, U2, etc., parameters that together are constrained to be unique.