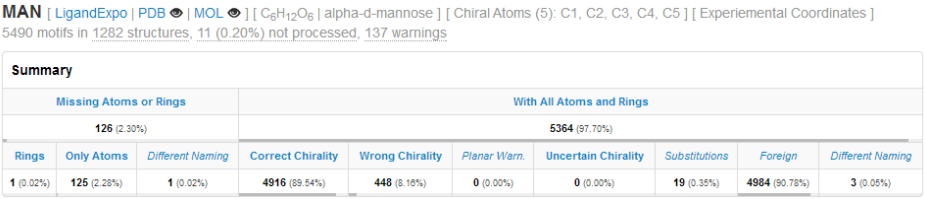
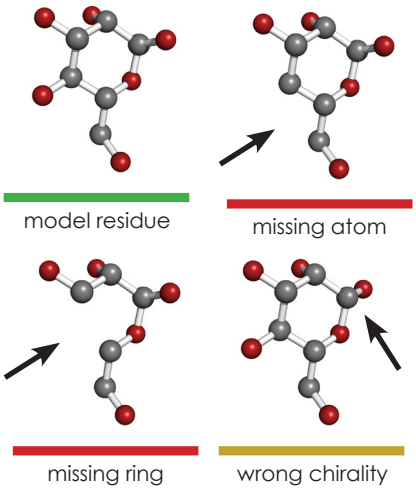
MotiveValidator

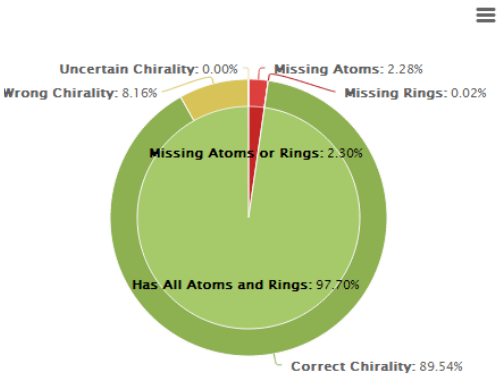
interactive web-based validation of ligand and residue structure in biomolecular complexes

Description:

- Fast and accurate detection of inconsistencies in *ligand* structures in PDB files against wwPDB CCD models.
- Expert *sugar* structure validation.
- User defined *motif/fragment* validation.
- Intuitive and user friendly interface.
- Supports pdb, mol & cif file formats.







http://ncbr.muni.cz/MotiveValidator

[1] Vařeková, R.S., et al., (2014) MotiveValidator: interactive web-based validation of ligand and residue structure in biomolecular complexes. Nucleic Acids Res., 42, W227–33, doi: 10.1093/nar/gku426.

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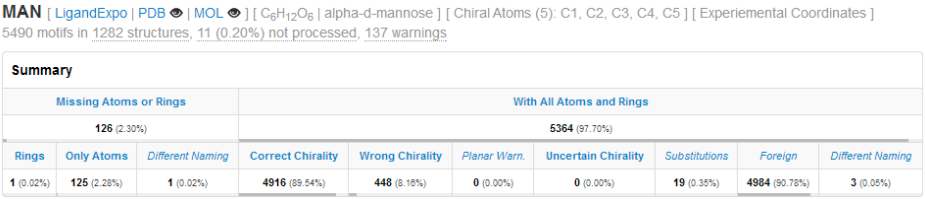

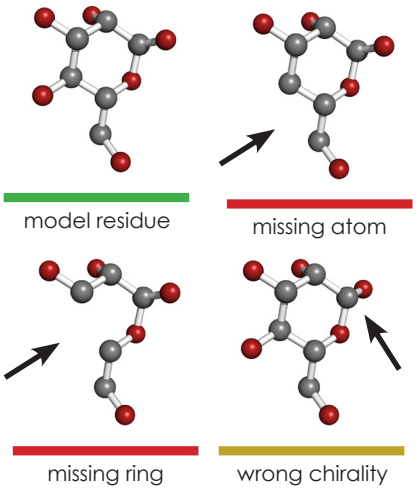

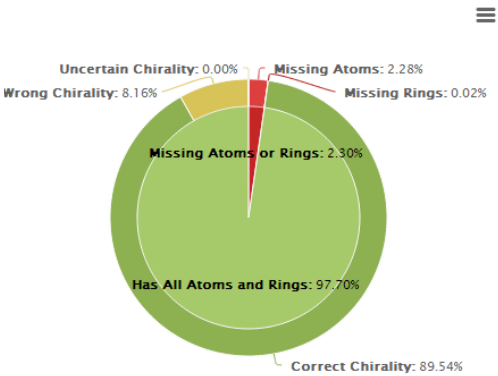

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