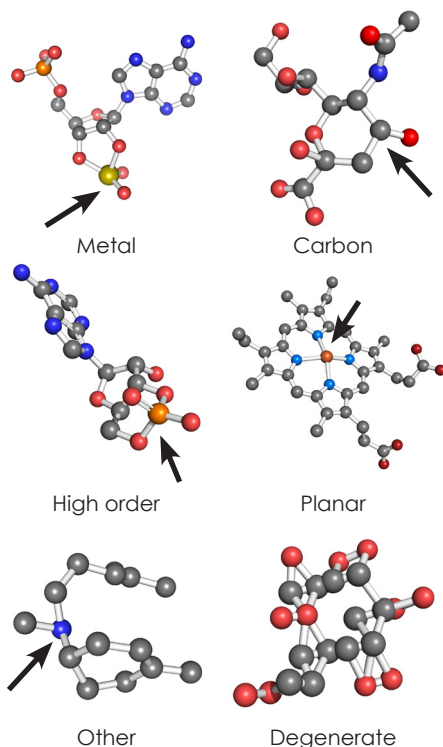
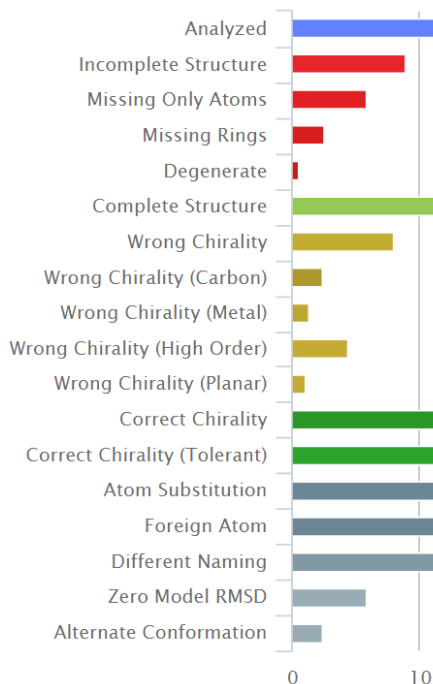


ValidatorDB

database of up-to-date and comprehensive validation results for ligands and non-standard residues from the PDB.

- Up-to-date validation of annotation of individual entries so as non-standard residues in the whole PDB.
- Custom validation for user specified classes of structures.
- Detection of missing atoms.
- Classification of variety chirality issues.

Issues detected:



<http://ncbr.muni.cz/ValidatorDB>

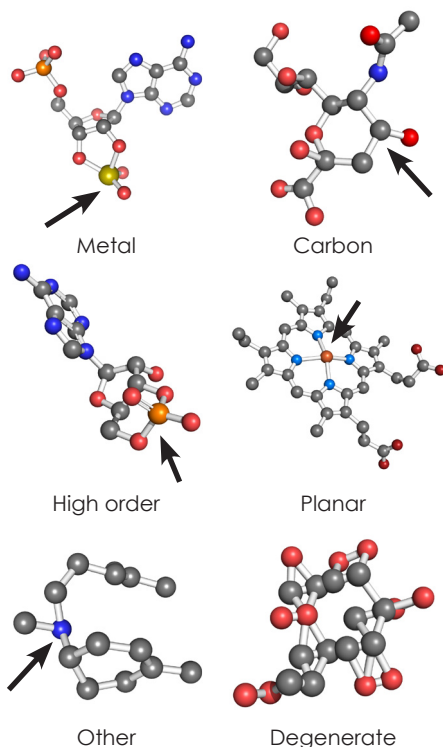
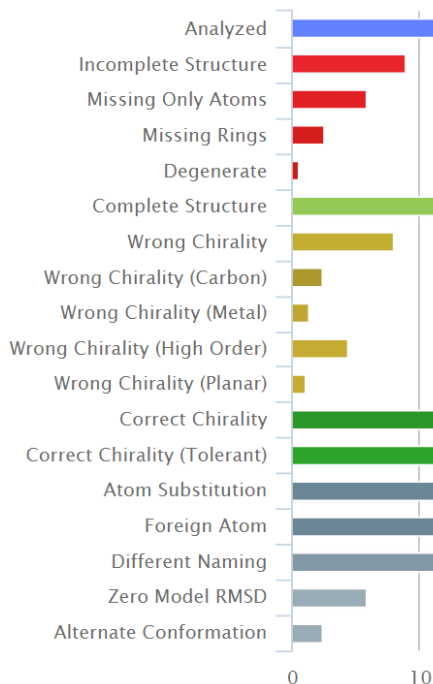
[1] Sehnal, D., et al., (2014) ValidatorDB: database of up-to-date and comprehensive validation results for ligands and non-standard residues from the Protein Data Bank. *Nucleic Acids Res.*, submitted.

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