# Scientific test: Protein Data Bank diagnostic

## ## AUTHOR AND DATE

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### **FAILURES**

The PDB diagnostic was run on 200,608 PDB files. 196,608 files passed the test and 4,000 failed!

## ## PURPOSE OF THE TEST

This test tries to load every PDB file in the PDB database and classifies the failures that occur. The command line below shows what was done; broadly all versions of this test examine load-time problems and more expensive versions (-PDB\_diagnostic::skip\_pack\_and\_min false) also check for errors during scoring, packing, and minimization.

"Hunting down these bugs is the most fun thing you can do on a Thursday morning" - Andy Watkins, probably.

An individual PDB passes or fails this test based on whether it errors out or completes the diagnostic. The test as a whole passes or fails based on a "reference results" system, like an expected result in a unit test. About 700 PDBs and 1200 CIFs fail at the time of this writing; the purpose of the test is to document the failures and watch for new ones, so PDBs failing in an expected manner does not constitute an overall test failure. The test will fail if PDBs pass or fail **UNEXPECTEDLY**, where the expectation is defined by the reference results (see below).

If you find that this page is telling you the test failed because there are **FEWER** errors: **GREAT**! You fixed some bugs! You can update the reference results following the instructions at the bottom of the page.

If you find that there are **MORE** failures than expected, especially non-timeout failures, consider it a warning that recent code changes may have introduced bugs into the PDB reading machinery. If you don't know what's going on: post to Slack or devel. **DO NOT** just update the reference results in this case.

If you want to know more about what this test does - pester Steven Lewis to write proper documentation.

Command line used: /home/benchmark/w/rosetta-2.hpc-c/rosetta/commits/main/source/test/timelimit.py 8 /home/benchmark/w/rosetta-2.hpc-c/rosetta/commits/main/source/bin/PDB\_diagnostic.default.linuxclangrelease -no\_color -out:file:score\_only /dev/null -jd2::delete\_old\_poses true -ignore\_unrecognized\_res false -load\_PDB\_components true -packing::pack\_missing\_sidechains false -packing::repack\_only true -s {input\_file} -ignore\_zero\_occupancy false -in:file:obey\_ENDMDL true -

#### **4,000** total PDBs failed with the following error codes:

```
2,950 Munrecognized residueM
           351 <a href="mailto:affill_missing_atoms">affill_missing_atoms</a>
           229 Emultiple_disulfides
           146 @missing_disulfide_partner@
                90 Bbad_patch
                54 Wunknown atom name
                35 Preroot_disconnected
                 33 <a href="mailto:adden.com">aduplicate</a> atom</a>_name
                17 🔣 ld_chirality
                17 Wexceed_timeout
                15 Mnu_conformer
                 14 Wunknown
                 11 Pro_usable_coords
                     5 @merge_with_next@
4 @insufficient_mainchain@
                      3 Whase of chim
                      3 <a href="mailto:apartly_recognized">apartly_recognized</a>
                      2 \( \text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\tinx{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\tinx{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\tinx{\tint{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\tinx{\tint{\text{\text{\text{\tint{\text{\text{\text{\text{\text{\tinit}}\\ \text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\tetx{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\tinit}}\\ \text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\texi}\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\texi}\text{\text{\texitile}}}\text{\text{\text{\text{\text{\texi}\text{\text{\text{\texit{\tet{\text{\text{\text{\text{\text{\texict{\texi{\texi{\texi{\texi{\
                       2 Wincompatible_w_polymer
                      1 Wzero_length_xyzVectorW
```

#### Test marked as **FAILED** due to following errors:

```
PDB 5NBX was passing test before but now failed with <code>@unrecognized_residue</code> error!
Its run-log could be found in $\mathbb{B}$5NBX$\mathbb{B}$
PDB 6E04 was passing test before but now failed with <code>Bunrecognized_residue</code> error!
Its run-log could be found in <a href="M6E048">M6E048</a>
PDB 6GSM was passing test before but now failed with <code>Bunrecognized_residue</code> error!
Its run-log could be found in M6GSMM
PDB 606E was passing test before but now failed with <code>Bunrecognized_residue</code> error!
Its run-log could be found in 8606E
PDB 6PWB was passing test before but now failed with Bexceed_timeout error! Its run-
log could be found in M6PWBM
PDB 6QMS was passing test before but now failed with @unrecognized_residue error!
Its run-log could be found in 260MS2
PDB 6YD3 was passing test before but now failed with <code>Bunrecognized_residue</code> error!
Its run-log could be found in M6YD3M
PDB 6YD4 was passing test before but now failed with <code>@unrecognized_residue</code> error!
Its run-log could be found in 186YD488
PDB 6YD7 was passing test before but now failed with Bunrecognized residue error!
Its run-log could be found in 26YD7
PDB 6Z6Y was passing test before but now failed with <code>Bunrecognized_residue</code> error!
Its run-log could be found in M6Z6YM
```

NOTE: 7412 PDB's passed the tests but was not listed in reference results.

To update reference results please copy the files below into the main repository:

reference-results.cif.new.json → main repository: as tests/benchmark/tests/scientific/
protein data bank diagnostic/reference-results.cif.json

blocklist.cif.new.json → main repository: as tests/benchmark/tests/scientific/
protein data bank diagnostic/blocklist.json (note the cif/pdb/fast mode is ignored: the blocklist is segfaults and huge PDBs and is shared)