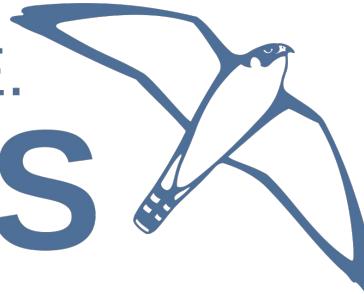


FAST. FLEXIBLE. FREE.

GROMACS



Testing and testing infrastructure

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Testing



- “In simple terms, it’s a process of checking something if it does what it intended to do”

<https://keencoder.dev/unit-testing-for-absolute-beginners>

Testing



- Unit tests: verify individual methods and functions
- Integration tests: verify that different modules work well together
- Functional tests: verify the final output of the application
- End-to-end tests: replicate a user behavior
- Acceptance testing: verify entire application for end goals
- Performance testing: verify reliability, speed, scalability
- Smoke testing: basic checks of major features

One of the possible nomenclatures; many others exists

Testing



- CI: practice of merging all developers' working copies to a shared mainline several times a day
- CD: software engineering approach in which teams produce software in short cycles, ensuring that the software can be reliably released at any time and, following an automated testing pipeline
 - + Less conflicts between team members
 - + Less need for manual testing
 - + No need to follow later with (external) contributors

Testing



- CI: practice of merging all developers' working copies to a shared mainline several times a day
- CD: software engineering approach in which teams ~~produce software in short cycles~~, ensuring that the software can be reliably released at any time and, following an automated testing pipeline
- Manual testing: not everything can be (easily) automated

GROMACS CI pipeline



Report NBNXM GPU supercluster dimensions in the log

Open

Szilárd Páll requested to merge [sz_allow_setting_and_report...](#) into [main](#) 1 day ago

Overview 8

Commits 2

Pipelines 2

Changes 2

2 unreso

Also allow setting DGMX_GPU_NB_NUM_CLUSTER_PER_CELL_X/Y/Z=1 with other backends too not just SYCL. The only case failing tests is GMX_GPU_NB_CLUSTER_SIZE=4 on Intel (not on NVIDIA), but since we can't check at build-time for this, we forbid setting the cluster per cell in all cases when cluster size is 4.



0



0



Merge request pipeline #994237396 passed

Merge request pipeline passed for [68e00fc2](#) 22 hours ago



8

Revoke approval

Requires 1 approval from GMX Developer Main. Approved by you



Test summary: no changed test results, 499 total tests

Full report



GROMACS CI pipeline



Pipeline Needs Jobs 55 Tests 499

Group jobs by [Stage](#) [Job dependencies](#)

pre-build	configure-build	build	test	documentation	source-check	post-test
clang-format	clang-tidy:configure-mr	gromacs:clang-9:build	gmxaapi:clang-9:py-3.10	docs:build	check-source	webpage:build
copyright-check	docs:configure	gromacs:clang-13-mpi:build	gmxaapi:clang-13-mpi:py-3.7		clang-tidy:test	
python-format	gromacs:clang-9:configure	gromacs:clang-ASAN:build	gromacs:clang-9:regressiontest			
simple-build	gromacs:clang-13-mpi:configure	gromacs:clang-TSAN:build	gromacs:clang-9:test			
	gromacs:clang-ASAN:configure	gromacs:clang-UBSAN:build	gromacs:clang-13-mpi:test			
	gromacs:clang-TSAN:configure	gromacs:clang-static-analyzer:build	gromacs:clang-13:regressiontest			
	gromacs:clang-UBSAN:configure	gromacs:gcc-9-cuda-11.0.3:build	gromacs:clang-ASAN:regressiontest			
	gromacs:clang-static-analyzer:configure	gromacs:gcc-9-cuda-11.0.3:buildMPI	gromacs:clang-ASAN:test			

Parts of GROMACS automated testing

- Pre-build: code formatting, copyrights, simple build
- Configure-build: CMake in different configurations
- Build: Build the configurations above
- Test: Run the tests for the builds above
- Documentation: Check that documentation is building
- Source-check: Static analysis checks
- Post-test: Manual webpage can be built

Code style



- Clang-format: C++ code style
 - Very sensitive to clang-format version, needs to be 11.0.1
 - `sudo apt install clang-format-11` (Ubuntu 22.04)
 - `pipx install clang-format==11.0.1` (Ubuntu 24.04)
- Python-format: Python code style (`black`)
- Copyright-check: Copyright headers

Code style



GROMACS > GROMACS > Jobs > #4985233672

```
at.sh --rev=$REV diff | tee clang-format.patch ; exit 1 ; fi
42 clang-format.sh found issues!
43 Patch below (can also be downloaded from artifacts)
44 diff --git org/api/nplib/util/setup.cpp new/api/nplib/util/setup.cpp
45 index b68d75a..03c5ea8 100644
46 --- org/api/nplib/util/setup.cpp
47 +++ new/api/nplib/util/setup.cpp
48 @@ -125,8 +125,7 @@ std::vector<Vec3> generateVelocity(real tempi, unsigned int seed, std::vector<re
49 //! (so the contrary of isfinite, see https://en.cppreference.com/w/cpp/numeric/math/isfinite)
50     bool isRealValued(gmx::ArrayRef<const Vec3> values)
51     {
52         return std::all_of(values.begin(), values.end(), [](const Vec3& val)
53         {
54             return std::all_of(values.begin(), values.end(), [](const Vec3& val) {
55                 for (int m = 0; m < dimSize; ++m)
56                 {
57                     if (!std::isfinite(val[m]))
58                         Uploading artifacts for failed job
59                         Uploading artifacts...
60                         clang-format.log: found 1 matching files and directories
61                         clang-format.patch: found 1 matching files and directories
```

clang-format



Duration: 28 seconds

Finished: 1 week ago

Queued: 2 seconds

Timeout: 1h (from project) [?](#)

Runner: #22213073 (Coj35txY9)
gitlab-runner-gitlab-runner-c94b4dbff-cg7lj

Job artifacts [?](#)

The artifacts will be removed in 3 weeks [?](#)

Commit 171f228c [!](#) in !3782

Micro-optim isRealValued by using std::all_of + isfinite

✖ Pipeline #987039987 for !3782 with micro_optim [!](#)

Configure + Build + Test



```
cmake .. && make && make check
```

- Compiler: Clang/GCC
 - MPI: on/off
 - GPU: off/CUDA/OpenCL/SYCL
 - Sanitizers: none/TSAN/ASAN/UBSAN
-
- “-test” and “-regressiontests”: two different test harnesses
 - A few gmxapi-specific jobs

GROMACS CI pipeline



Pipeline Needs Jobs 55 Tests 499

Group jobs by [Stage](#) [Job dependencies](#)

pre-build	configure-build	build	test	documentation	source-check	post-test
clang-format	clang-tidy:configure-mr	gromacs:clang-9:build	gmxaapi:clang-9:py-3.10	docs:build	check-source	webpage:build
copyright-check	docs:configure	gromacs:clang-13-mpi:build	gmxaapi:clang-13-mpi:py-3.7		clang-tidy:test	
python-format	gromacs:clang-9:configure	gromacs:clang-ASAN:build	gromacs:clang-9:regressiontest			
simple-build	gromacs:clang-13-mpi:configure	gromacs:clang-TSAN:build	gromacs:clang-9:test			
	gromacs:clang-ASAN:configure	gromacs:clang-UBSAN:build	gromacs:clang-13-mpi:test			
	gromacs:clang-TSAN:configure	gromacs:clang-static-analyzer:build	gromacs:clang-13:regressiontest			
	gromacs:clang-UBSAN:configure	gromacs:gcc-9-cuda-11.0.3:build	gromacs:clang-ASAN:regressiontest			
	gromacs:clang-static-analyzer:configure	gromacs:gcc-9-cuda-11.0.3:buildMPI	gromacs:clang-ASAN:test			

Documentation + webpage: preview

The screenshot shows a build job summary for 'webpage:build'. It includes details like duration (12 minutes 12 seconds), finish time (55 minutes ago), queue time (5 seconds), timeout (1h from project), and runner information (#22213073). A section for 'Job artifacts' indicates they will be removed in 6 days, with options to 'Keep', 'Download' (which is highlighted with an orange box), or 'Browse'. Below this, it shows a commit (d7c51bc2) in pipeline #3722, applying 1 suggestion(s) to 1 file(s). It also shows a checked pipeline entry for 'Pipeline #995466495 for !3722 with gyrate-taf'. A dropdown menu shows 'post-test' and a final step arrow pointing to 'webpage:build'.

Artifacts / build-docs / docs / html:

- index.html
- manual-2024-dev.pdf

Source-check



- check-source: basic documentation format checks
- clang-tidy: advanced code static analysis

Source-check



- check-source: basic documentation format checks
- clang-tidy: advanced code static analysis

```
if (i == 1 && i == 2) { // correct?  
}
```

Even more automated tests



- Post-merge (multi-GPU tests)
 - Heavy tests not to run on every minor change to the MR
- Nightly (exotic hardware)
 - <https://gitlab.com/gromacs/gromacs/-/pipelines>
- GitHub actions (Windows, macOS)
 - <https://github.com/gromacs/gromacs/actions>

Not covered by automated testing

- Performance testing
- Many analysis tools
- Large-scale runs (> 2 GPUs, > 4 ranks)
- Rare devices (POWER9, ARM, high-end GPUs, etc)
- Long-running physical validation tests

Test frameworks



- GoogleTest for most tests
 - Modern and convenient
 - We call them “unit tests”, but that’s not always the case
- Old Perl scripts for regression tests
 - `cmake -DREGRESSIONTEST_DOWNLOAD=ON`
 - Don’t touch it unless you’re changing `mdrun` behavior :)
 - <https://gitlab.com/gromacs/gromacs-regressiontests/>

Running tests



- All tests:
 - make tests && make check
 - ctest .
- Specific test set:
 - ctest -R MdrunIOTests
 - ./bin/mdrun-io-test
- Specific test case:
 - ./bin/mdrun-io-test --gtest_filter=GromppTest.*
 - Some tests might also need to have -ntomp N or -ntmpi N set
- Run multiple times (flaky test):
 - ctest --repeat-until-fail 100 --output-on-failure -R MdrunIOTests

Finding tests



- File: `src/gromacs/utility/logger.cpp`
- Tests: `src/gromacs/utility/tests/logger.cpp`
- Reference data: `src/gromacs/utility/tests/refdata/`
- Also in: `src/programs/*/tests/`

Understanding tests



```
TEST(EnergyTermTest, AddFrameWorks)
{
    EnergyTerm term(0, true, "test", "test");
    term.addFrame(2, 1000, 10, 50, 5, 255);
    term.addFrame(4, 2000, 10, 100, 10, 155);
    EXPECT_EQ(term.numFrames(), 2);
    auto errorEstimate = term.errorEstimate(1);
    ASSERT_TRUE(errorEstimate.has_value());
    EXPECT_REAL_EQ(errorEstimate.value(), 0);
}
```

Parametrized tests



```
TEST_P(HbondModuleTest, Works)
{
    const auto params = GetParam();
    std::string name = std::get<0>(params);
    int startingValue = std::get<1>(params);
    // Do something with name and startingValue
}

INSTANTIATE_TEST_SUITE_P( //...
    ::testing::Combine(::testing::Values("name1", "name2"),
                      ::testing::Range(4, 8, 1)), // ...
```

A lot of boilerplate code omitted

<https://google.github.io/googletest/>

Reference data



```
TEST_F(WrapperTest, WrapsCorrectly)
{
    std::vector<std::string> wrapped = doThings();
    checker().checkSequence(wrapped.begin(), wrapped.end(), "Wrapped");
}
```

checker() will compare with data in
src/gromacs/*/tests/**refdata/WrapperTest_WrapsCorrectly.xml**

```
$ ./bin/utility-test -ref-data update-all
```

Input data



If you need some input data (trajectories, etc):

src/testutils/simulationdatabase/

```
class TrjconvWithDifferentInputs: public gmx::test::CommandLineTestBase{};  
  
TEST_F(TrjconvWithDifferentInputs, WithIndexGroupSubset) {  
    auto& cmdline = commandLine();  
    setInputFile("-s", "spc2.gro");  
    ASSERT_EQ(0, gmx_trjconv(cmdline(argc()), cmdline.argv()));  
}
```

Writing tests



- When fixing a bug: write a test first
 - Helps replicating
 - Helps verifying that your fix works
 - Helps prevent it from happening again
- When adding a new feature: think first about how to test it
 - Test behavior, not implementation
- Some things are very hard to test. Such is life.

Further information



- Testing in GROMACS:
 - <https://manual.gromacs.org/current/dev-manual/testutils.html>
- More on reference data:
 - https://manual.gromacs.org/current/doxygen/html-lib/page_refdata.xhtml
- GoogleTest Primer:
 - <https://google.github.io/googletest/primer.html>