

GROMACS - Bug #2472

confusing error message when OMP_NUM_THREADS is used with GPUs

03/29/2018 01:56 PM - Szilárd Páll

Status:	Closed	
Priority:	Normal	
Assignee:	Szilárd Páll	
Category:	mdrun	
Target version:	2018.2	
Affected version - extra info:	2018.x	Difficulty: uncategorized
Affected version:	2018	

Description

The r2018 code does not allow setting only the OpenMP thread count in a GPU run (in tMPI builds), but as the OpenMP thread count handling was changed and part of the reporting seems short-circuited (the env var-related reporting from the omp_nthreads module does not happen), this leads to potentially confusing error messages that lack context.

```
$ OMP_NUM_THREADS=2 gmx mdrun -nsteps 0  
[...]
```

```
GROMACS:      gmx mdrun, version 2018  
Executable:   /opt/tcbsys/gromacs/2018/AVX2_256/bin/gmx  
Data prefix:  /opt/tcbsys/gromacs/2018/AVX2_256  
Working dir:  /home/pszilard/projects/gromacs/testing/water-048k  
Command line:  
  gmx mdrun -nsteps 0
```

```
Back Off! I just backed up md.log to ./#md.log.95#  
Reading file topol.tpr, VERSION 4.6-beta3-dev-20121222-492378e (single precision)  
Note: file tpx version 82, software tpx version 112  
The number of OpenMP threads was set by environment variable OMP_NUM_THREADS to 2
```

```
-----  
Program:      gmx mdrun, version 2018  
Source file:  src/gromacs/taskassignment/resourcedivision.cpp (line 224)
```

```
Fatal error:  
When using GPUs, setting the number of OpenMP threads without specifying the  
number of ranks can lead to conflicting demands. Please specify the number of  
thread-MPI ranks as well (option -ntmpi).
```

```
For more information and tips for troubleshooting, please check the GROMACS  
website at http://www.gromacs.org/Documentation/Errors
```

```
-----  
In contrast, in r2016, besides there being no error, it is pretty clear that the environment variable's value is used (that may have not  
been set by the user / at the time of mdrun invocation):
```

```
GROMACS:      gmx mdrun, version 2016  
Executable:   /opt/tcbsys/gromacs/2016/AVX2_256/bin/gmx  
Data prefix:  /opt/tcbsys/gromacs/2016/AVX2_256  
Working dir:  /home/pszilard/projects/gromacs/testing/water-048k  
Command line:  
  gmx mdrun -nsteps 0
```

```
Running on 1 node with total 4 cores, 8 logical cores, 2 compatible GPUs  
Hardware detected:  
CPU info:  
  Vendor: Intel  
  Brand: Intel(R) Core(TM) i7-4790K CPU @ 4.00GHz  
  SIMD instructions most likely to fit this hardware: AVX2_256
```

```
SIMD instructions selected at GROMACS compile time: AVX2_256
```

```
Hardware topology: Basic
```

```
GPU info:
```

```
Number of GPUs detected: 2
```

```
#0: NVIDIA GeForce GTX 1080, compute cap.: 6.1, ECC: no, stat: compatible
```

```
#1: NVIDIA GeForce GTX 960, compute cap.: 5.2, ECC: no, stat: compatible
```

```
Reading file topol.tpr, VERSION 4.6-beta3-dev-20121222-492378e (single precision)
```

```
Note: file tpx version 82, software tpx version 110
```

```
Changing nstlist from 10 to 40, rlist from 1 to 1.101
```

```
The number of OpenMP threads was set by environment variable OMP_NUM_THREADS to 2
```

```
Overriding nsteps with value passed on the command line: 0 steps, 0 ps
```

```
Using 2 MPI threads
```

```
Using 2 OpenMP threads per tMPI thread
```

```
2 compatible GPUs are present, with IDs 0,1
```

```
2 GPUs auto-selected for this run.
```

```
Mapping of GPU IDs to the 2 PP ranks in this node: 0,1
```

Associated revisions

Revision f0c98f46 - 06/12/2018 03:09 PM - Szilárd Páll

Also issue OMP_NUM_THREADS reading note to the log

The note that was meant to inform users that OMP_NUM_THREADS was setting the number of threads in their run (as this value can be inherited by the env) has not been logged. It was also printed right after the tpx reading status making it hard to notice. Removed stderr output now that this is no longer required.

This change makes the note easier to notice prepending a newline and issues it to the log file too.

Refs #2472

Change-Id: I73fc9de5e9d747f9d7a094c6678ffc1547481b94

History

#1 - 03/29/2018 01:57 PM - Szilárd Páll

- Description updated

#2 - 03/29/2018 02:31 PM - Szilárd Páll

- Description updated

- Status changed from New to In Progress

Note that as the hw_opt reads the env var in check_and_update_hw_opt_1() by calling gmx_omp_nthreads_read_env() and the report that's not issued anymore with 2018 should be in fact printed, so something else is wrong.

#3 - 03/29/2018 02:45 PM - Szilárd Páll

- Status changed from In Progress to Rejected

Wow, I'm blind it seems. Only the newline went missing -- and we'd be better off logging this as well as printing a note.

#4 - 03/29/2018 02:57 PM - Szilárd Páll

Ref user report:

https://mailman-1.sys.kth.se/pipermail/gromacs.org_gmx-users/2018-March/119506.html

Will try to add the message in question to the log.

#5 - 03/29/2018 03:02 PM - Gerrit Code Review Bot

Gerrit received a related patchset '1' for Issue [#2472](#).

Uploader: Szilárd Páll (pall.szilard@gmail.com)

Change-Id: gromacs~release-2018~173fc9de5e9d747f9d7a094c6678ffc1547481b94

Gerrit URL: <https://gerrit.gromacs.org/7741>

#6 - 03/29/2018 04:26 PM - Mark Abraham

Putting the full context in the error message requires that `hw_opt` keep track of the reason why `nthreads_omp` has the value that it does. That would be wise to do in general, but probably not in the release branch.

#7 - 06/12/2018 03:09 PM - Mark Abraham

- Status changed from *Rejected* to *Fix uploaded*

- Assignee set to *Szilárd Páll*

#8 - 06/13/2018 06:09 PM - Mark Abraham

- Status changed from *Fix uploaded* to *Resolved*

#9 - 06/14/2018 10:50 AM - Mark Abraham

- Status changed from *Resolved* to *Closed*