Gromacs - Bug #1218
vrescale_resamplekin assumes an integer, but an integer isn't always passed

04/09/2013 04:27 AM - Michael Shirts

Description
vrescale_resamplekin expects an integer:

```c
static real vrescale_resamplekin(real kk, real sigma, int ndeg, real taut, gmx_rng_t rng)
```

But the value that is passed in to ndeg is `opts->nrdf[i]`, which, due to subtraction of COM degrees of freedom which are then partitioned among the temperature coupling groups, may not be an integer. In my case, I was trying every atom as its own tcoupling group, which lead to `opts->nrdf[i] = 2.99` or so, and ndeg becomes 2. It might make more sense to round?

I've tried without COM removal with one thermostat per atom, so that there is now exactly 3 DOF per temperature group, to see if it gives the correct temperature, but it gives a temperature too high. (306 vs 300) I wonder if there is some other approximation that fails when nrdf becomes low?

I've attached a .tpr, though it will fail without increasing the size of several of the temperature group variables.

Getting vrescale to work right for many temperature groups is probably not a super high priority, but I'm worried that this behavior may be indicative of subtle errors with in more normal situations.

Associated revisions

Revision 1f8a4f94 - 04/29/2014 03:22 PM - Berk Hess

Removed truncation of nrdf in v-rescale thermostat

The resampling function for the v-rescale thermostat expected an integer value for nrdf, but a real was passed, which was truncated. With a single coupling coupling group nrdf is analytically an int, but could be off by a bit. The could lead to incorrect kinetic energy fluctuations (averages were correct).

Now fractional nrdf's are properly handled for nrdf > 3.

For nrdf < 3 a check is added for integer values with a small margin for rounding.

Fixes #1218

Change-Id: I4c60c337f9874d0bff51220ad09429140be2a056

History

#1 - 04/09/2013 11:46 AM - Berk Hess

11/25/2015
We should change ndeg to a real. Have you tried that with your many group case? It could be that some atoms get a ndeg of 3 - one bit.

#2 - 04/09/2013 08:26 PM - Michael Shirts
I'm don't think making it ndeg a real would work, since some of the algorithms that vrescale_resamplekin uses require integers.

I didn't pursue this fully, because when I used COM = none, so I got full integers everywhere, then the temperature was too high -- 306, statistically significantly (errors was about 0.5 K, over about 10 ps or so). So the thermostat has additional problems beyond this.

Note that for, say, 900 argon molecules with one temperature group, the differences are statistically insignificant (see the checkensemble paper), so I don't know how vital it is to fix this -- but it should be fixed eventually.

#3 - 04/10/2013 12:26 PM - Berk Hess
Indeed, for small ndeg we need integers.
I'm still not sure if there are no rounding issues. FP calculations are not that simple, there could still be rounding issues due to the ratio of sums being off by one bit. To check that you would need to round ndeg before passing it.
Your tpr format changed, so I can't check your input.
I ran a 20 system LJ particle system and didn't observe any issues.
What tau_t are you using compared to the LJ time?
It could be that the v-rescale algorithm "simply" is not valid with multiple/many t-coupl groups.
Indeed, not a critical issue, but we need to understand what is going on.

#4 - 04/10/2013 02:57 PM - Michael Shirts
FP calculations are not that simple, there could still be rounding issues due to the ratio of sums being off by one bit. To check that you would need to round ndeg before passing it.

Agreed. Though being off by one bit is better than being off by 0.99.
Your tpr format changed, so I can't check your input.

Ah, sorry. Let me try to construct a system that runs under unmodified gromacs. Might take a day or two to get a chance.

I ran a 20 system LJ particle system and didn't observe any issues.
What tau_t are you using compared to the LJ time?

The .tpr is with water (not constrained), not LJ. tau_t is 0.1, timestep was 0.0005.

It could be that the v-rescale algorithm "simply" is not valid with multiple/many t-coupl groups.
Could be, will need to investigate this a bit more.

Indeed, not a critical issue, but we need to understand what is going on.

Right.

#5 - 04/11/2013 06:23 AM - Michael Shirts
Still working on constructing a small enough problem to fit in the unmodified 4.6 to have it show up statistically. Perhaps for < 256 groups it ends up not being statistically significant? Certainly the ndeg error should be fixed, but I can't yet determine if the other issues beyond that are relevant or not.

#6 - 05/22/2013 05:00 AM - Mark Abraham
- Target version changed from 4.6.2 to future
- Affected version set to 4.6.1

#7 - 05/22/2013 05:38 AM - Mark Abraham
- Target version changed from future to 4.6.x

#8 - 10/02/2013 04:47 AM - Michael Shirts
Was there any final solution here?

#9 - 10/02/2013 03:00 PM - Mark Abraham
Not that I know of

#10 - 04/23/2014 02:08 PM - Berk Hess
I don't see an easy fully accurate solution for fractional degrees of freedom.
I would suggest that we add rounding for nrdf, which will always be correct for a single coupling group and at least better for multiple groups, which you should anyhow only use with large groups and weak coupling, where the error will be small.
Should I put this in 4.6, which should be merged into 5.0?

#11 - 04/23/2014 03:29 PM - Gerrit Code Review Bot
Gerrit received a related patchset '1' for Issue #1218.
Uploader: Berk Hess (hess@kth.se)
Change-Id: I4c60c337f9874d0bff51220ad09429140be2a056
Gerrit URL: https://gerrit.gromacs.org/3392

#12 - 04/29/2014 03:30 PM - Berk Hess
- Status changed from New to Resolved
- % Done changed from 0 to 100

Applied in changeset 1f8a4f94581bb4ade86901208580bc6f01ee2abe.

#13 - 05/12/2014 10:29 AM - Rossen Apostolov
- Status changed from Resolved to Closed

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<td>04/09/2013</td>
<td>Michael Shirts</td>
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