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heavy

AMD Ryzen 5 3400G testing with a ASUS PRIME B450M-A (2006 BIOS) and ASUS AMD Picasso 2GB on Ubuntu 19.10 via the Phoronix Test Suite.

Test Systems:

1

2

3

Processor: AMD Ryzen 5 3400G @ 3.70GHz (4 Cores / 8 Threads), Motherboard: ASUS PRIME B450M-A (2006 BIOS), Chipset: AMD Raven/Raven2, Memory: 2 x 8192 MB DDR4-3000MT/s CRUCIAL, Disk: 29GB INTEL MEMPEK1W032GA + 4 x 6001GB Seagate ST6000VN0033-2EE, Graphics: ASUS AMD Picasso 2GB (1400/1500MHz), Audio: AMD Raven/Raven2/Fenghuang, Monitor: SyncMaster, Network: Realtek RTL8111/8168/8411

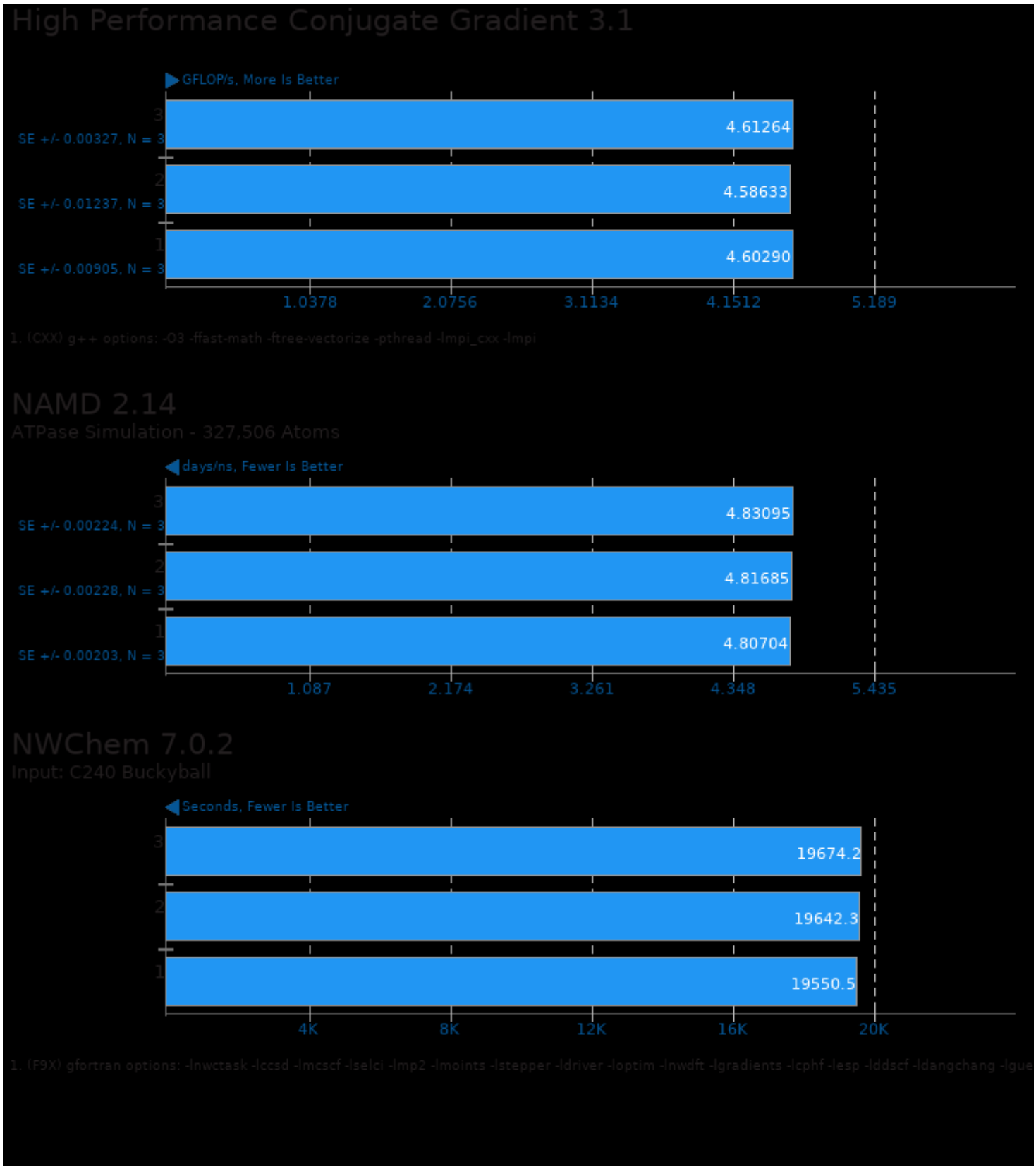
OS: Ubuntu 19.10, Kernel: 5.3.0-46-generic (x86_64), Desktop: GNOME Shell 3.34.1, Display Server: X Server, Display

heavy

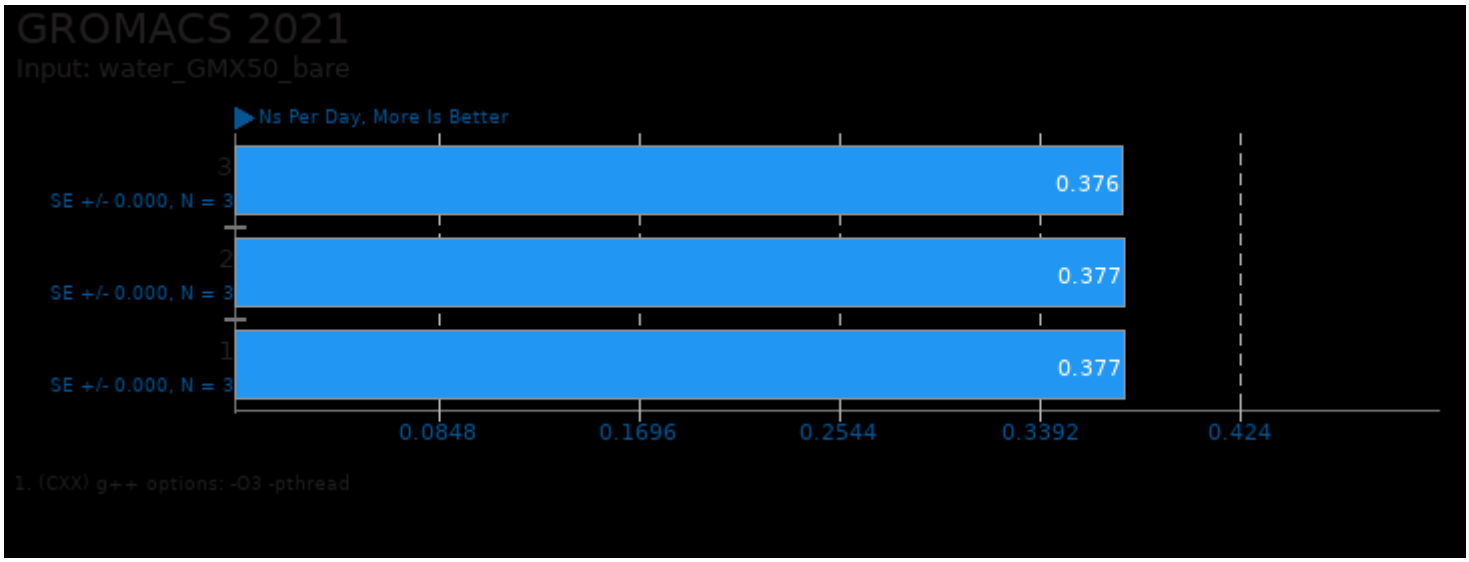
Driver: asus, Vulkan: 1.1.107, Compiler: GCC 9.2.1 20191008, File-System: ext4, Screen Resolution: 1280x800

Kernel Notes: Transparent Huge Pages: madvise
Compiler Notes: --build=x86_64-linux-gnu --disable-vtable-verify --disable-werror --enable-bootstrap --enable-checking=release --enable-clocale=gnu --enable-default-pie --enable-gnu-unique-object --enable-languages=c,ada,c++,go,brig,d,fortran,objc,obj-c++,gm2 --enable-libstdcxx-debug --enable-libstdcxx-time=yes --enable-multiarch --enable-multilib --enable-nls --enable-offload-targets=nvptx-none,hsa --enable-plugin --enable-shared --enable-threads=posix --host=x86_64-linux-gnu --program-prefix=x86_64-linux-gnu- --target=x86_64-linux-gnu --with-abi=m64 --with-arch-32=i686 --with-default-libstdcxx-abi=new --with-gcc-major-version-only --with-multilib-list=m32,m64,mx32 --with-target-system-zlib=auto --with-tune=generic --without-cuda-driver -v
Processor Notes: Scaling Governor: acpi-cpufreq ondemand (Boost: Enabled) - CPU Microcode: 0x8108109
Security Notes: itlb_multihit: Not affected + l1tf: Not affected + mds: Not affected + meltdown: Not affected + spec_store_bypass: Mitigation of SSB disabled via prctl and seccomp + spectre_v1: Mitigation of usercopy/swapgs barriers and __user pointer sanitization + spectre_v2: Mitigation of Full AMD retpoline IBPB: conditional STIBP: disabled RSB filling + tsx_async_abort: Not affected

	1	2	3
High Performance Conjugate Gradient (GFLOP/s)	4.60290	**4.58633**	**4.61264**
Normalized	99.79%	99.43%	100%
Standard Deviation	0.3%	0.5%	0.1%
NAMD - ATPase Simulation - 327,506 Atoms (days/ns)	**4.80704**	4.81685	**4.83095**
Normalized	100%	99.8%	99.51%
Standard Deviation	0.1%	0.1%	0.1%
NWChem - C240 Buckyball (sec)	**19551**	19642	**19674**
Normalized	100%	99.53%	99.37%
GROMACS - water_GMX50_bare (Ns/Day)	**0.377**	**0.377**	**0.376**
Normalized	100%	100%	99.73%
Standard Deviation	0%	0.2%	0.2%


High Performance Conjugate Gradient 3.1
GFLOP/s, More Is Better
3
SE +/- 0.00327, N = 3
4.61264
2
SE +/- 0.01237, N = 3
4.58633
1
SE +/- 0.00905, N = 3
4.60290
1.0378
2.0756
3.1134
4.1512
5.189
1. (CXX) g++ options: -O3 -ffast-math -ftree-vectorize -pthread -lmpi_cxx -lmpi
NAMD 2.14
ATPase Simulation - 327,506 Atoms
days/ns, Fewer Is Better
3
SE +/- 0.00224, N = 3
4.83095
2
SE +/- 0.00228, N = 3
4.81685
1
SE +/- 0.00203, N = 3
4.80704
1.087
2.174
3.261
4.348
5.435
NWChem 7.0.2
Input: C240 Buckyball
Seconds, Fewer Is Better
3
19674.2
2
19642.3
1
19550.5
4K
8K
12K
16K
20K
1. (F9X) gfortran options: -lnwctask -lccsd -lmcscf -lselci -lmp2 -lmoints -lstepper -ldriver -loptim -lnwdft -lgradients -lcphf -lesp -lddscf -ldangchang -lgue

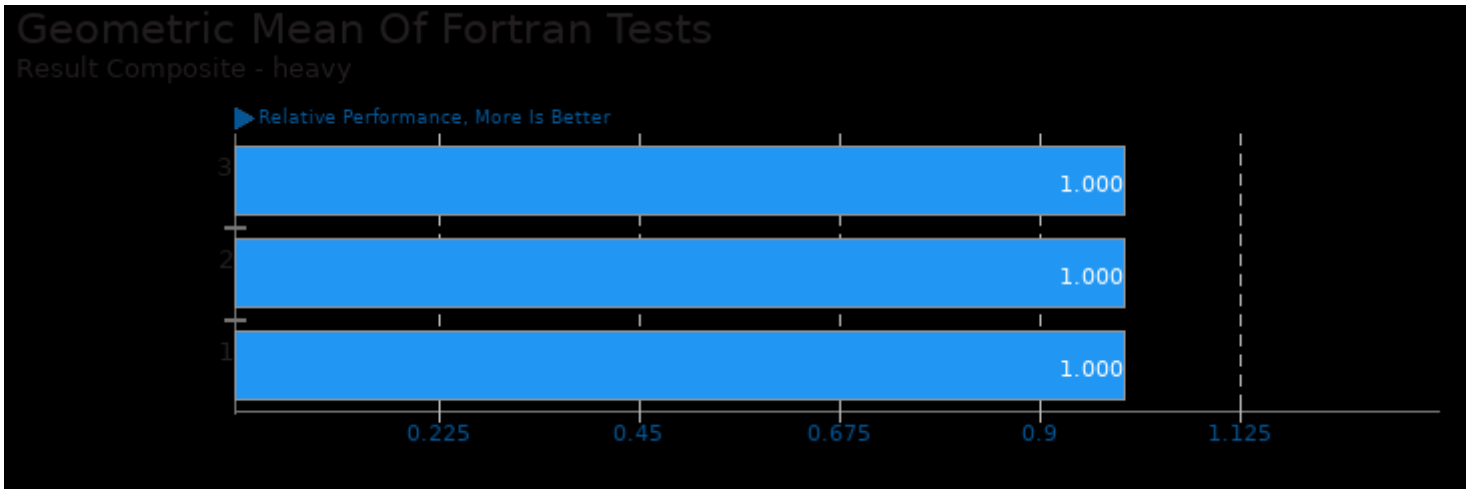
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GROMACS 2021
Input: water_GMX50_bare
Ns Per Day, More Is Better
3
SE +/- 0.000, N = 3
0.376
2
SE +/- 0.000, N = 3
0.377
1
SE +/- 0.000, N = 3
0.377
0.0848
0.1696
0.2544
0.3392
0.424
1. (CXX) g++ options: -O3 -pthread

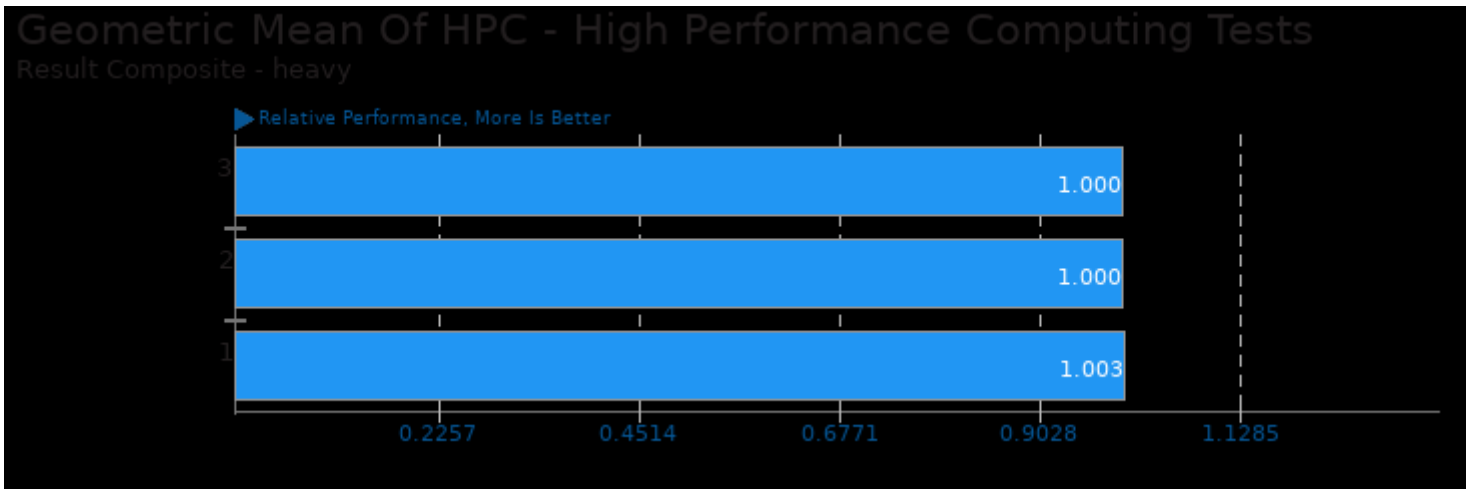
These geometric means are based upon test groupings / test suites for this result file.



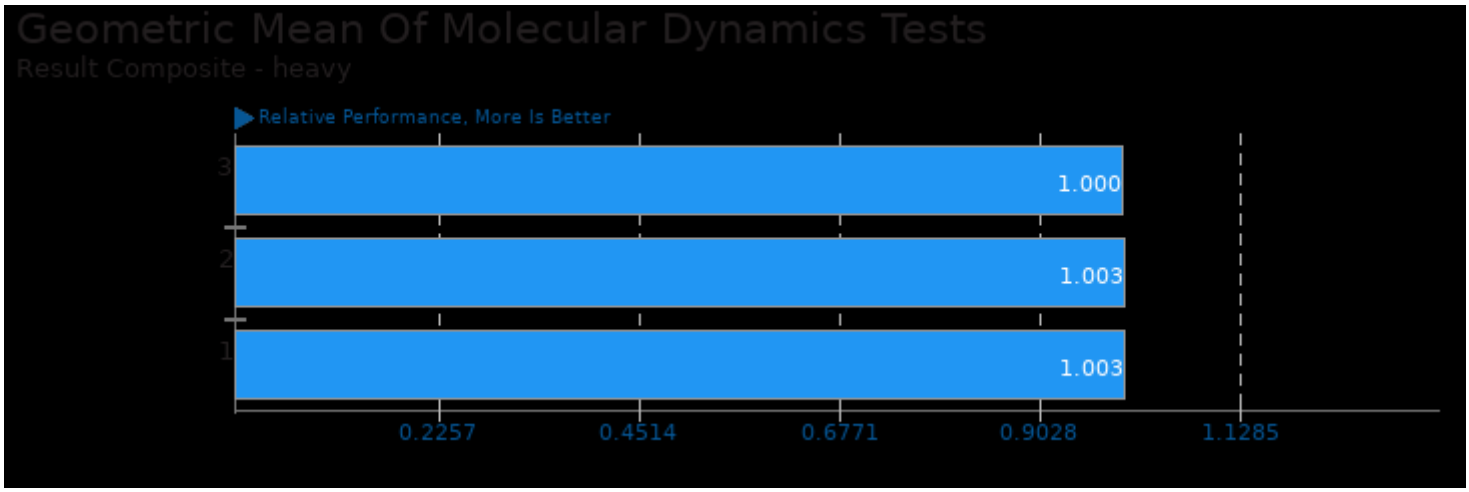
Geometric mean based upon tests: pts/hpcg and pts/namd



Geometric mean based upon tests: pts/hpcg and pts/nwchem



Geometric mean based upon tests: pts/hpcg, pts/namd, pts/gromacs and pts/nwchem



Geometric mean based upon tests: pts/namd, pts/gromacs and pts/nwchem

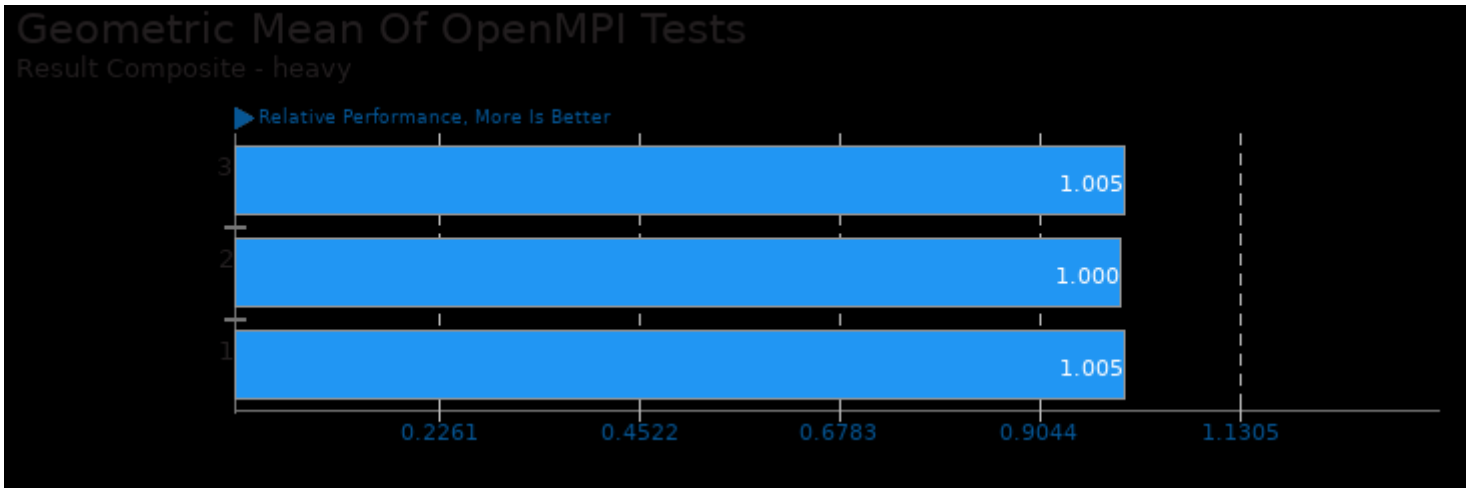


Geometric mean based upon tests: pts/gromacs and pts/hpcg

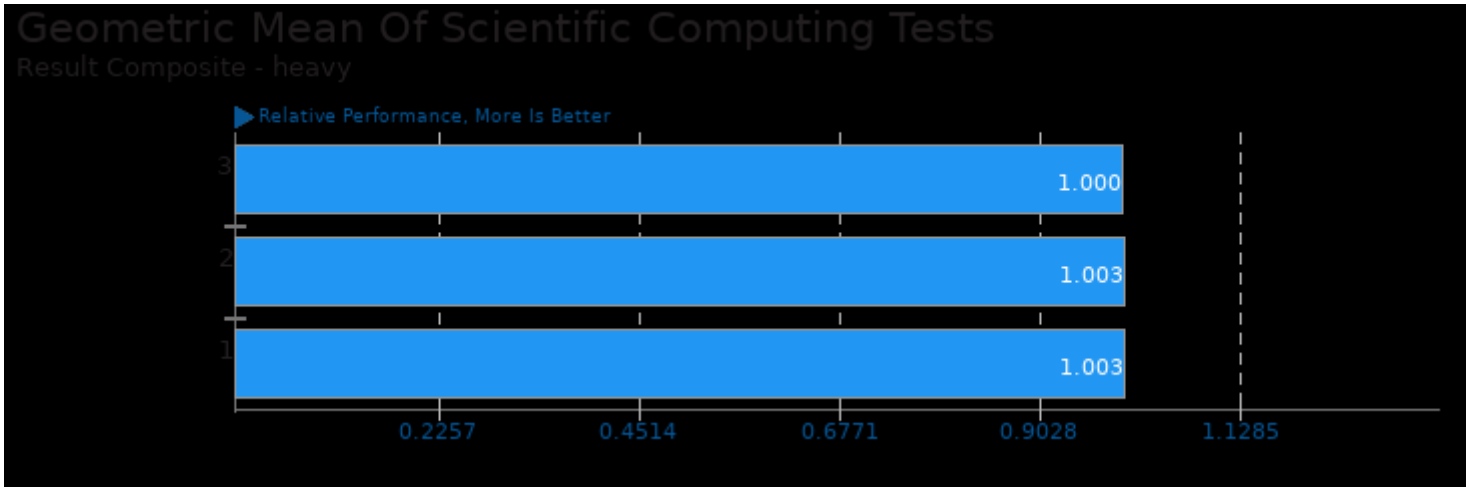


Geometric mean based upon tests: pts/namd, pts/gromacs and pts/hpcg

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Geometric mean based upon tests: pts/hpcg, pts/nwchem and pts/gromacs



Geometric mean based upon tests: pts/namd, pts/gromacs and pts/nwchem

This file was automatically generated via the Phoronix Test Suite benchmarking software on Friday, 29 March 2024 08:24.