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bioinfo-si-inserm-ubuntu-8B

VMware testing on Ubuntu 22.04 via the Phoronix Test Suite.

Test Systems:

bioinfo-si-inserm-ubuntu-8B

Processor: 8 x Intel Xeon Gold 6240R (15 Cores), Motherboard: Intel 440BX (6.00 BIOS), Chipset: Intel 440BX/ZX/DX, Memory: 32GB, Disk: 215GB Virtual disk, Graphics: VMware SVGA II, Network: 2 x VMware VMXNET3

OS: Ubuntu 22.04, Kernel: 5.15.0-43-generic (x86_64), Vulkan: 1.2.204, Compiler: GCC 11.3.0, File-System: ext4, Screen Resolution: 800x600, System Layer: VMware

Kernel Notes: Transparent Huge Pages: madvise
Compiler Notes: --build=x86_64-linux-gnu --disable-vtable-verify --disable-werror --enable-bootstrap --enable-cet --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++,m2 --enable-libphobos-checking=release
--enable-libstdcxx-debug --enable-libstdcxx-time=yes --enable-link-serialization=2 --enable-multiarch --enable-multilib --enable-nls --enable-objc-gc-auto
--enable-offload-targets=nvptx-none=/build/gcc-11-xKiWfi/gcc-11-11.3.0/debian/tmp-nvptx/usr,amdgcn-amdhsa=/build/gcc-11-xKiWfi/gcc-11-11.3.0/debian/tmp-gcn/usr
--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-build-config=bootstrap-lto-lean --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: CPU Microcode: 0x5003102

Python Notes: Python 3.10.6

Security Notes: `itlb_multihit`: KVM: Mitigation of VMX unsupported + `I1tf`: Mitigation of PTE Inversion + `mds`: Mitigation of Clear buffers; SMT Host state unknown + `meltdown`: Mitigation of PTI + `mmio_stale_data`: Mitigation of Clear buffers; SMT Host state unknown + `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 Retpolines `IBPB`: conditional `IBRS_FW` `STIBP`: disabled RSB filling + `srbds`: Not affected + `tsx_async_abort`: Not affected

bioinfo-si-inserm-ubuntu-8B**Himeno Benchmark - P.P.S (MFLOPS)** 2786

Standard Deviation 0.2%

Timed MAFFT Alignment - M.S.A - LSU RNA (sec) 12.205

Standard Deviation 1.6%

Timed HMMer Search - P.D.S (sec) 160.792

Standard Deviation 0%

QMCPACK - simple-H2O (Execution Time - sec) 37.981

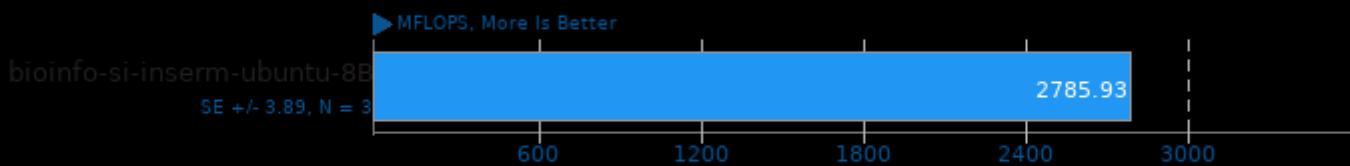
Standard Deviation 0.3%

Timed MrBayes Analysis - P.P.A (sec) 164.417

Standard Deviation 0.9%

Himeno Benchmark 3.0

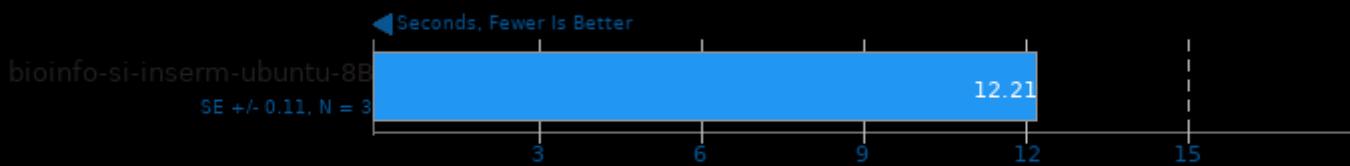
Poisson Pressure Solver



1. (CC) gcc options: -O3 -mavx2

Timed MAFFT Alignment 7.471

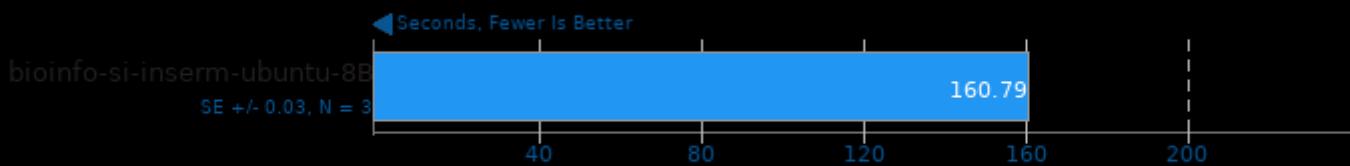
Multiple Sequence Alignment - LSU RNA



1. (CC) gcc options: -std=c99 -O3 -lm -lpthread

Timed HMMer Search 3.3.2

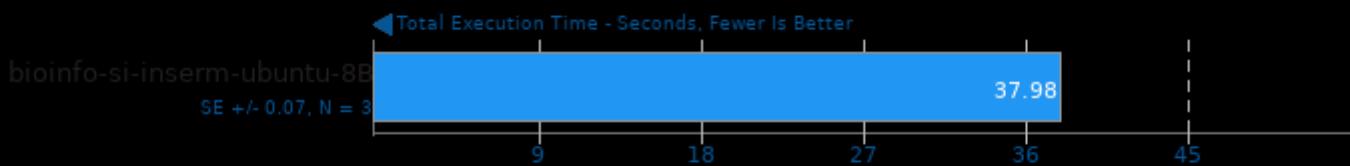
Pfam Database Search



1. (CC) gcc options: -O3 -pthread -lhmmer -leasel -lm -lmpi

QMCPACK 3.13

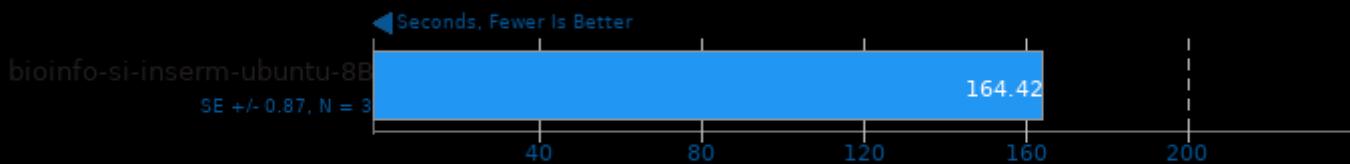
Input: simple-H2O



1. (CXX) g++ options: -finline-limit=1000 -fstrict-aliasing -funroll-all-loops -ffast-math -march=native -O3 -lm -ldl

Timed MrBayes Analysis 3.2.7

Primate Phylogeny Analysis



1. (CC) gcc options: -mmmx -msse -msse2 -msse3 -msse3 -msse4.1 -msse4.2 -maes -mavx -mfma -mavx2 -mrdrnd -mbmi -mbmi2 -madx -mabm -O3 -std

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