

### Quantum Espresso GPU on Marconi100

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#### GPU enabled version of QuantumESPRESSO

#### The last release of QE-GPU is available at

#### https://gitlab.com/QEF/q-e-gpu

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• Tasks ported to GPO		Tasks ported to GPU					

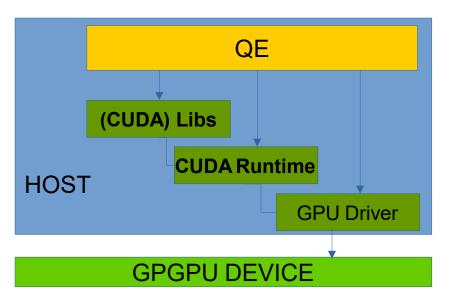
#### Compiling QE GPU

#### Compiling is as simple as...

./configure --with-cuda=XX --with-cuda-runtime=Y.y --with-cuda-cc=ZZ --enable-openmp [ --with-scalapack=no ]

where XX is the location of the CUDA Toolkit (in HPC environments it is generally \$CUDA\_HOME, be sure that this variable is not empty with a simple echo \$CUDA\_HOME), Y.y is the version of the CUDA Toolkit (Y and y are the two numbers identifying major and minor release, e.g. 9.0) and ZZ is the compute capability (cc) of the card. This information can be found on the internet using the model name of the GPU card or by using pgaccelinfo command.

Openmp is required in order to successfully compile the accelerated version.





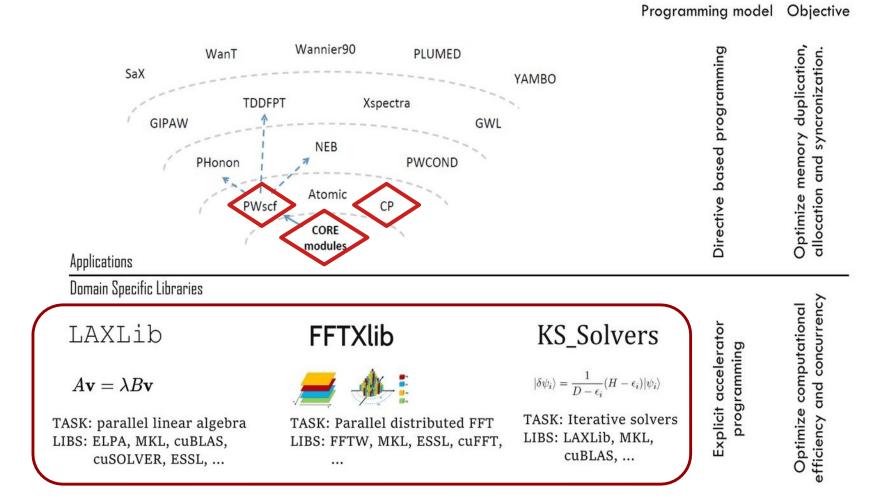
Cuda Toolkit is generally in \$CUDA\_HOME.

The *compute capabilities* codify the features and specifications of the target device.

<pre>[pbonfa00@login02 scf]\$</pre>	pgaccelinfo   grep cc
PGI Default Target:	-ta=tesla: <mark>cc</mark> 70
PGI Default Target:	-ta=tesla: <mark>cc</mark> 70
PGI Default Target:	-ta=tesla: <mark>cc</mark> 70
PGI Default Target:	ta=tesla:cc70



#### What does QE GPU provide





#### What does QE GPU provide

#### What can be done with the accelerated version of pw.x

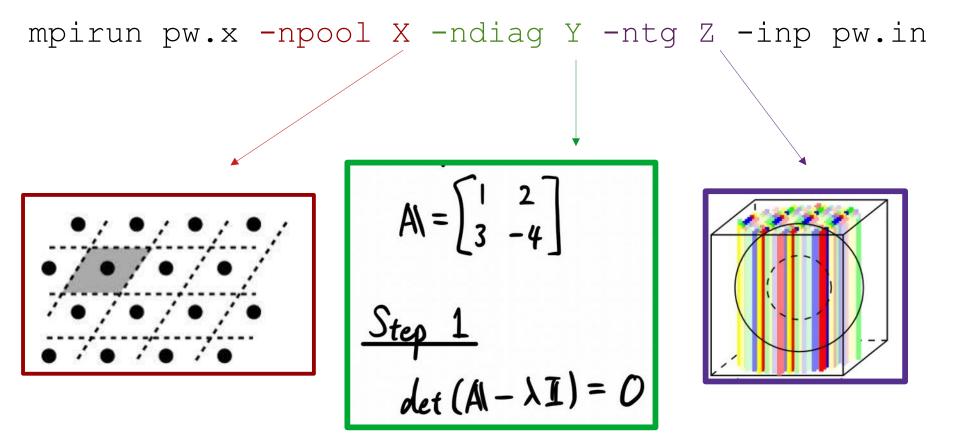
GPU version	Total Energy (K points)	Forces	Stress	Collinear Magnetism	Non- collinear magnetism	Gamma trick	US PP	PAW PP	DFT+U	All other functions
v5.4	A	W	W	B (?)	U	A	A	?	W (?)	W (?)
v6.1	A	A	A	A	U	W (*)	A	A (*)	U (?)	U (?)
v6.4	Α	W	W	Α	А	A	Α	A (*)	W	W
V6.5a1	Α	A	W	Α	А	A	Α	Α	W	W
V6.5a2	А	Α	Α	Α	А	А	А	А	W	W

#### Accelerated, Working, Unavailable, Broken

\* Acceleration obtained from other parts of the code.



You know how to run QE efficiently on a HPC machine:

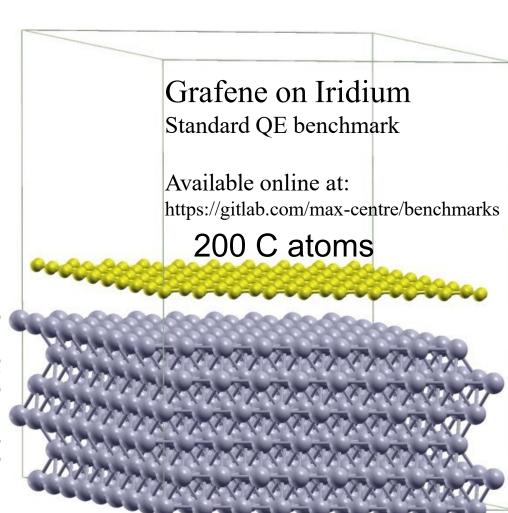




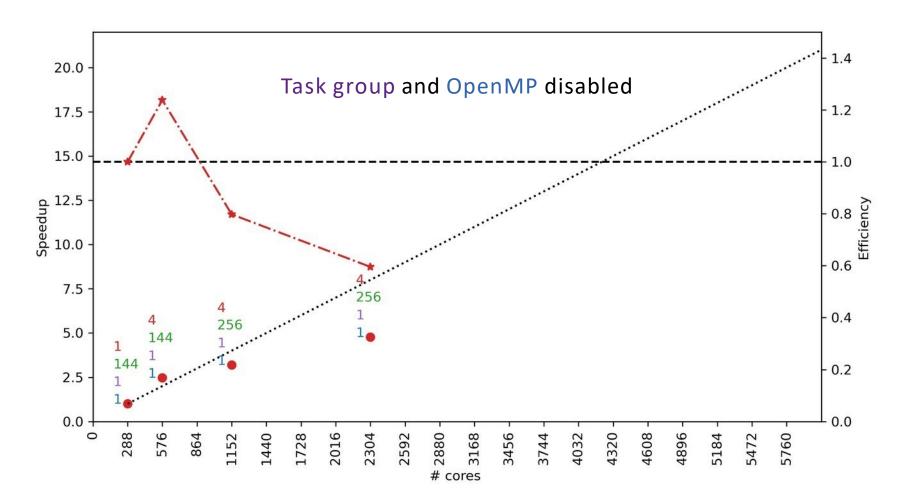
#### mpirun pw.x -npool X -ndiag Y -ntg Z -inp pw.in

```
&control
    calculation = 'scf'
   prefix='GRIR'
   restart mode='from scratch'
   pseudo dir='./',
 &system
    ibrav= 4
    celldm(1) = 46.5334237988185d0
    celldm(3) = 1.274596
    nat=686
   ntyp= 2,
    ecutwfc=30
   occupations = 'smearing'
    smearing='mv'
    degauss=0.025d0
    nspin = 2
    starting magnetization(1) = +.00
    starting magnetization(2) = +.00
 &electrons
   conv thr = 1.0d-5
   mixing beta=0.3d0
   mixing mode='local-TF'
    startingwfc='atomic'
    diagonalization='david'
    electron maxstep = 1
ATOMIC SPECIES
     12.010 C.pbe-paw kj-x.UPF
 С
Ir 192.22
             Ir.pbe-paw kj.UPF
K POINTS {automatic}
2 2 2 0 0 0
```

# 486 Ir atoms

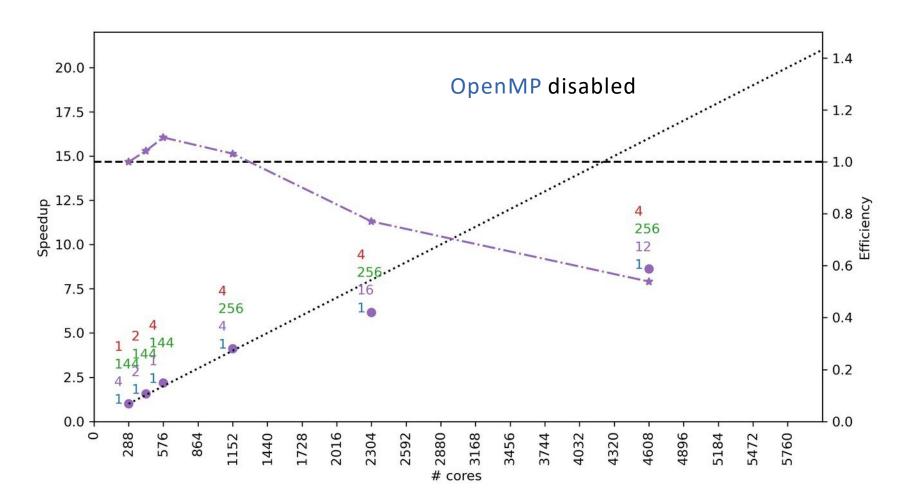


mpirun pw.x -npool X -ndiag Y -ntg Z -inp pw.in



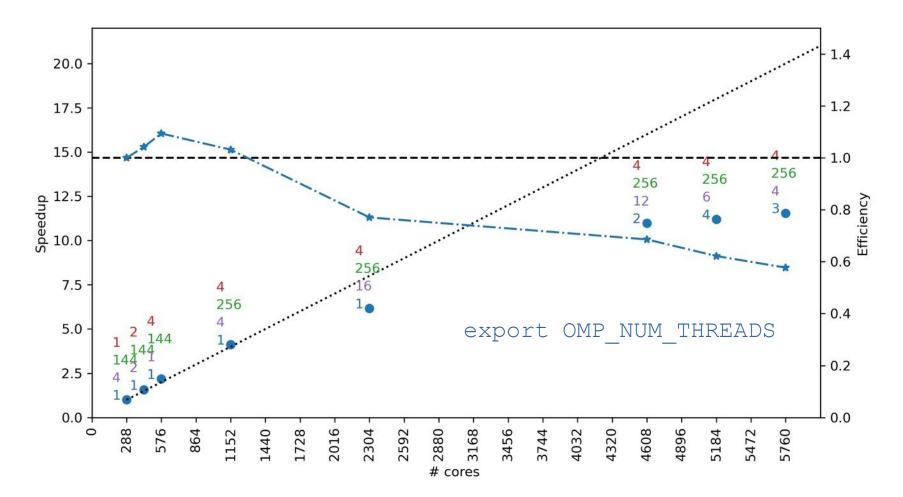


mpirun pw.x -npool X -ndiag Y -ntg Z -inp pw.in





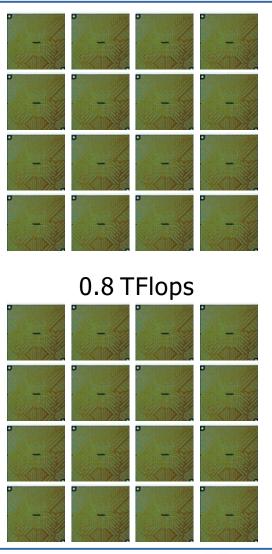
mpirun pw.x -npool X -ndiag Y -ntg Z -inp pw.in



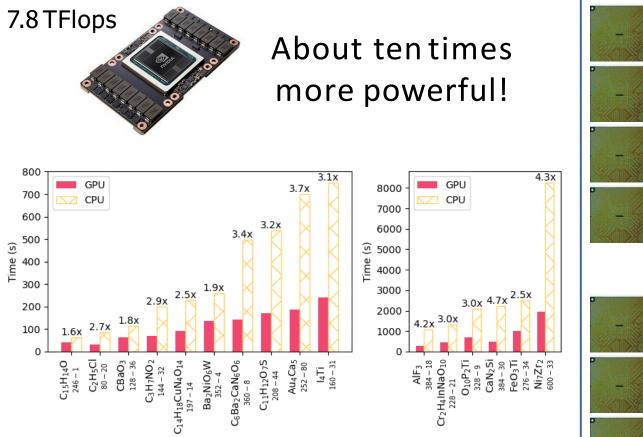




# About tentimes more powerful!



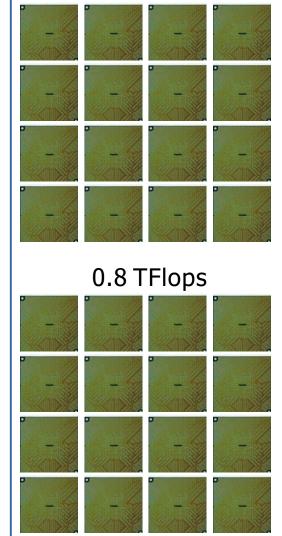




Quantum ESPRESSO toward the exascale

P. Giannozzi *et al.* 

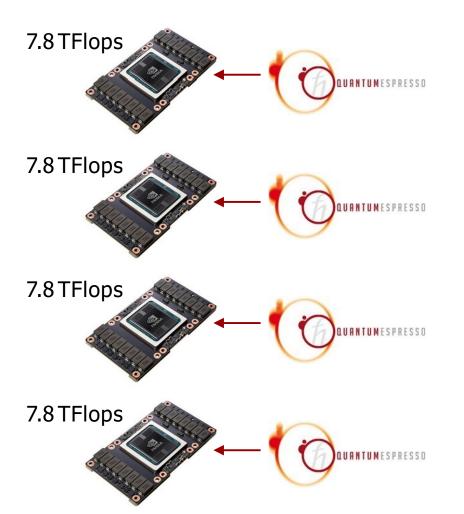
J. Chem. Phys. 152, 154105 (2020); DOI: 10.1063/5.0005082

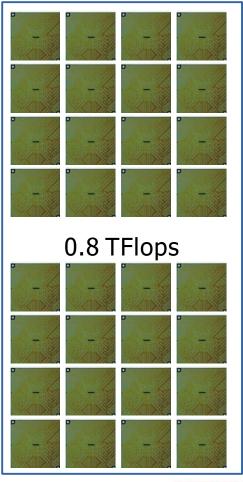




There are 4 GPUs per node on Marconi100!

mpirun -np 4 pw.x

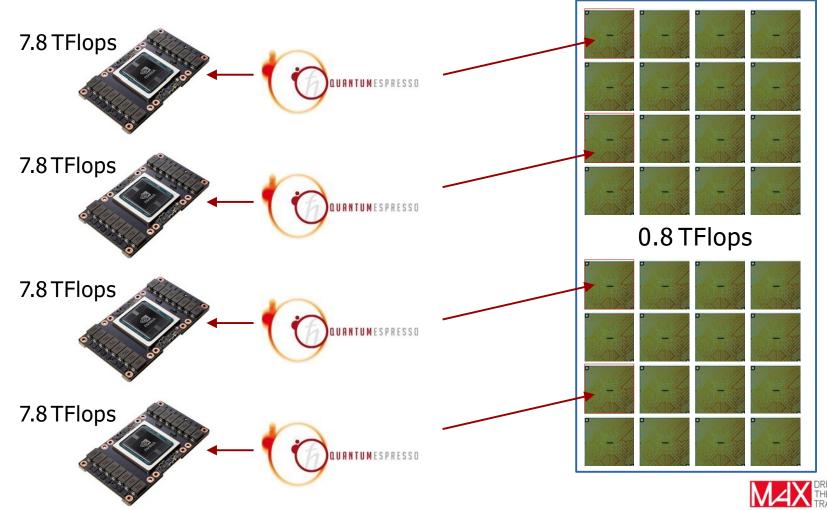


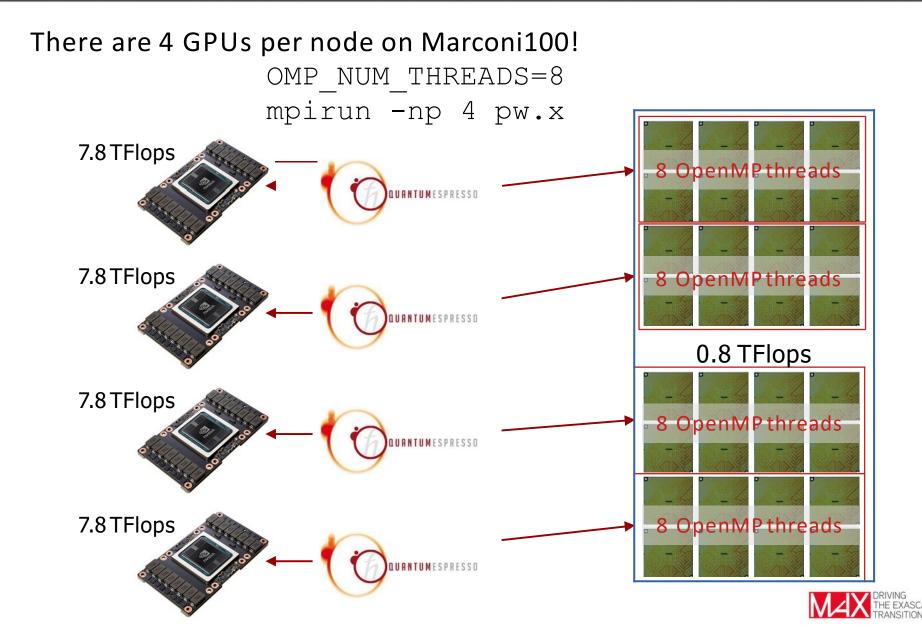




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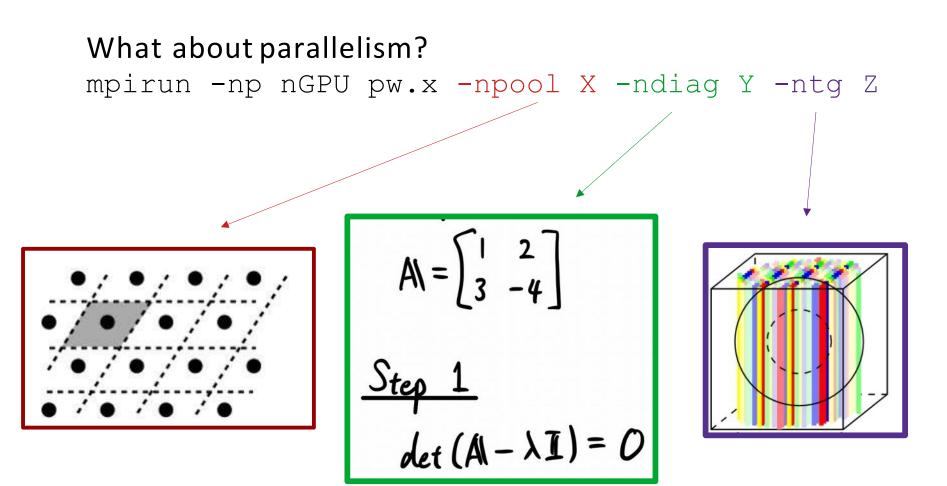


**One MPI process per GPU!** mpirun -np nGPU pw.x ...

What about parallelism? mpirun -np nGPU pw.x -npool X -ndiag Y -ntg Z

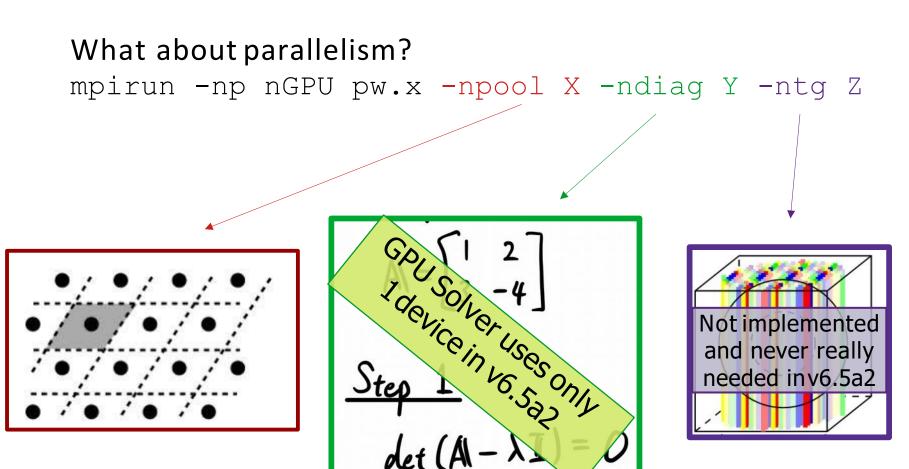


One MPI process per GPU! mpirun -np nGPU pw.x ...





**One MPI process per GPU!** mpirun -np nGPU pw.x ...







One MPI process per GPU! mpirun -np nGPU pw.x ...

```
What about parallelism?
mpirun -np nGPU pw.x -npool X -ndiag 1 -ntg 1
```

```
Subspace diagonalization in iterative solution of the eigenvalue problem:
a serial algorithm will be used 
[...]
GPU acceleration is ACTIVE.
```



What about memory...



#### mpirun -np nGPU pw.x -npool X -ndiag 1 -ntg 1

Check memory estimator!

X=4 Estimated max dynamical RAM per process > 14.72 GB X=1 Estimated max dynamical RAM per process > 2.97 GB

Choose the largest value for X that can fit available memory.



#### Tips & Tricks

You run out of memory ... what to do?

- use more GPUs...
- <u>reduce subspace dimension in</u> <u>Davidson algorithm</u>
- Change diagonalization method

&electrons	
$conv_thr = 1.0d-9$	
mixing_beta=0.3d0	
startingwfc='atomic'	
diago david ndim='2'	
/	

```
&electrons
    conv_thr = 1.0d-9
    mixing_beta=0.3d0
    startingwfc='atomic'
    diagonalization='cg'
/
```

Feature X is slow!

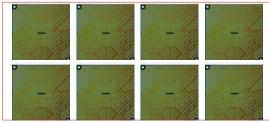
→Open an issue athttps://gitlab.com/QEF/q-e-gpu/-/issues



A few things you should remember when running the GPU version of the code:

- 1MPI process per GPU,
- CPU cores can (must!) be exploited with *OpenMP parallelism*
- Pool parallelism is very effective, but requires memory
- The dense eigenvalue problem is (as of v6.5a2) solved on 1 GPU, use the serial eigensolver.
- Check the Wiki, it's updated with a collaborative effort!
- More details: P. Giannozzi *et al.* J. Chem. Phys. 152, 154105 (2020)









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## THANKS



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