

PÁRHAZAMOS AGYI BIOLEKTROMOS JELFELDOLGOZÁS MULTI-GPU KÖRNYEZETBEN A SLICES IMEC INFRASTRUKTÚRA TRANSZNACIONÁLIS HASZNÁLATÁVAL

PARALLEL MULTI-GPU BIOELECTRICAL SIGNAL PROCESSING BY THE TRANS-NATIONAL USE OF THE SLICES IMEC INFRASTRUCTURE

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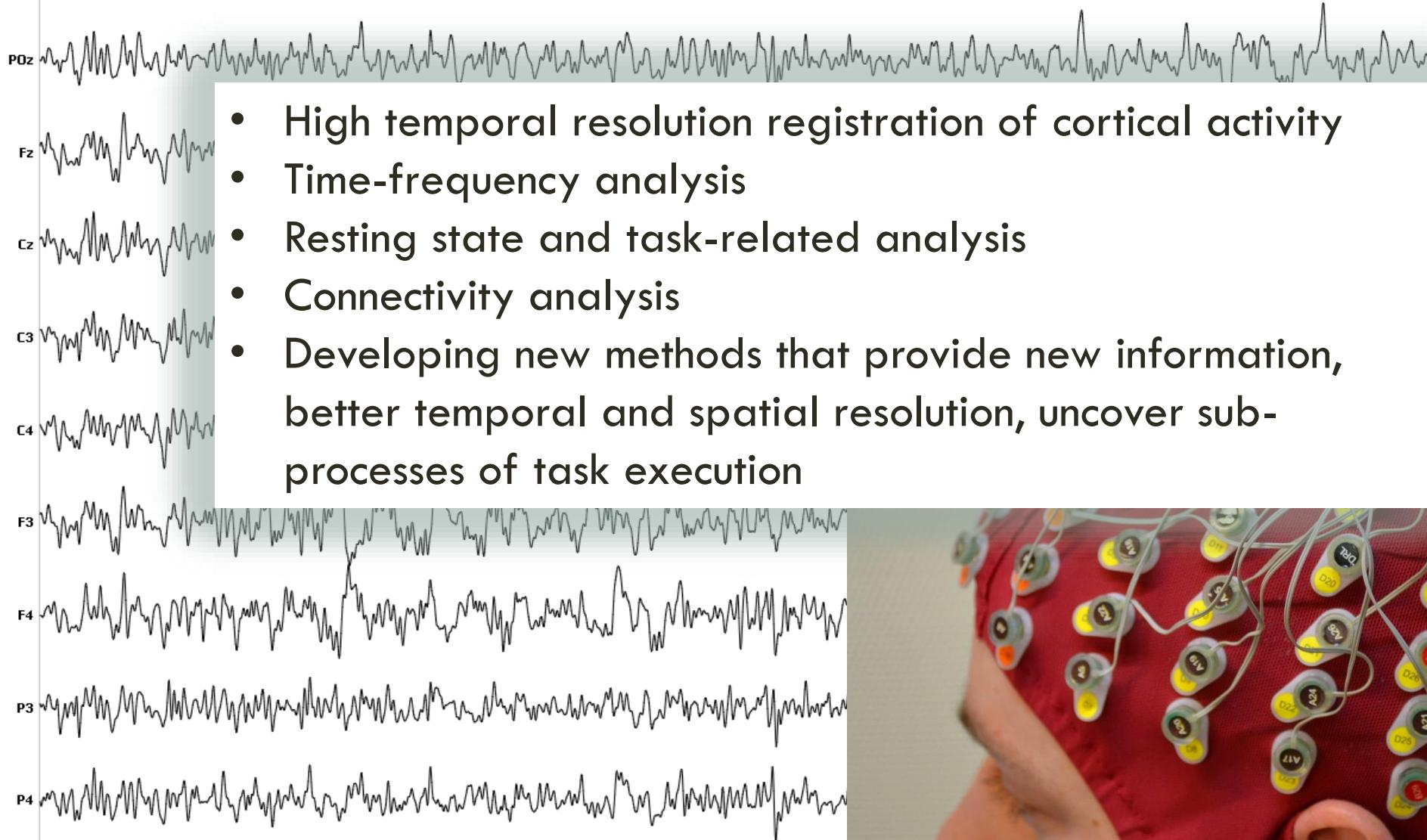
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SETTING THE SCENE

Bioelectrical Brain Imaging Laboratory

1. What we do: parallel biomedical signal processing and analysis
2. Why we need high-performance computational resources
3. Problems we normally face: financial, administrative and technical
4. Why we chose the SLICES-SC resources
5. Benefits and results

BIOMEDICAL (EEG) SIGNAL PROCESSING



NEED FOR HIGH-PERFORMANCE COMPUTATIONAL RESOURCES

- EEG research is mainly conducted using MATLAB scripts
- tendency to use **limited number of channels, low sampling frequency, short measurements**, to keep execution time within practical limits
- Most sophisticated experiments and analysis methods are excluded!
 - 2k samples/channel/second → 512k samples for 256 channels/sec → 10^9 samples for 30 min measurement
 - at 1k ops per sample → 10^{12} ops (1 Top/s)
- MATLAB execution times are in the order of **hours per subject**
- limited parallelism support, mainly clusters

Research goal:

- Massively parallel GPU algorithms to reduce time from hours/days to seconds
- single-GPU and multi-GPU systems, price/performance ratio

TYPICAL HPC-RELATED DIFFICULTIES

Options for increasing computational capacity:

- in-house cluster (CPU and/or GPU) – very high upfront cost, only viable if used for 24/7 production use
- Cloud – cost, still limited HPC and GPU support, mainly single node architecture
- Supercomputers – fine if access is granted, high threshold to entry
 - current situation in Hungary: there is no suitable system available to us
 - existing systems (Leo) are outdated (GPU model no longer supported)
 - Komondor (A100 GPUs, NVLink!) is still in test mode
 - PRACE resources: possible, but mainly for production runs
- SLICES-SC project – fast access to resources for **experimentation / development**

EEG PROCESSING FUNDAMENTALS

Typical operations

- high-pass, low-pass filtering, mean removal, segmenting into trials
- trial averaging and quantification, statistical tests
- spectrum estimation (global or time-varying); FFT, wavelets
- artefact removal: Independent Component Analysis
- source localisation
- connectivity estimation, calculation of graph metrics

Typically performed for subject/patient populations of size 20-100

OPPORTUNITIES FOR PARALLEL EXECUTION

Fortunately, EEG processing lends itself to parallel execution

- subjects: “embarrassingly parallel”
- channels:
- samples:
- spatial ops: one time point but all electrodes
- algorithmic level:

High degree of parallelism, ideal for GPU execution

One GPU is a huge improvement: 200-2000x speedup over MATLAB

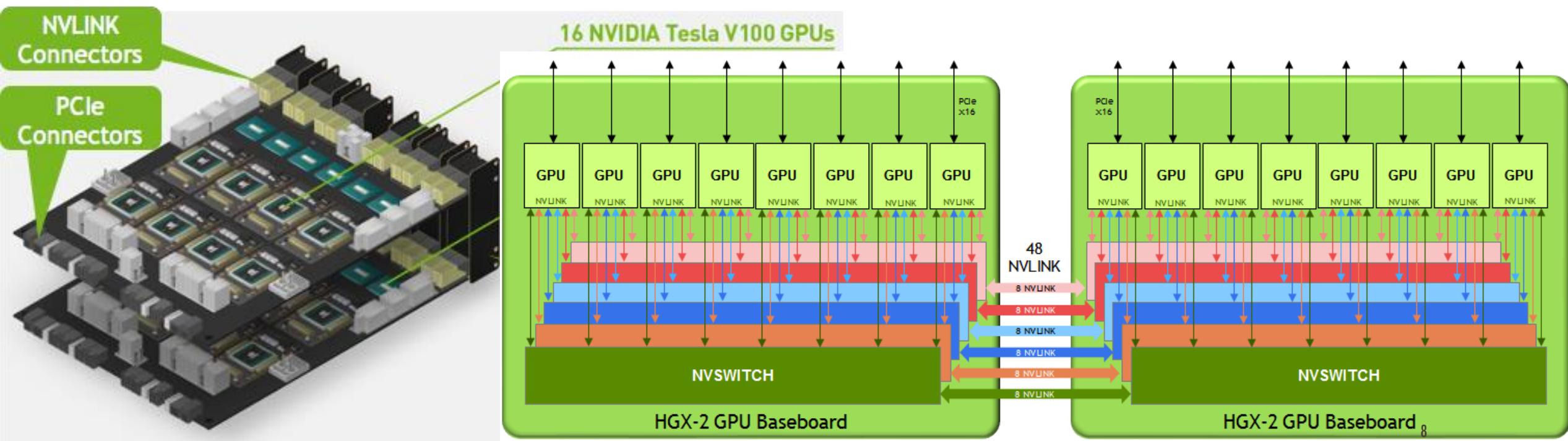
What about multiple GPUs? Can we utilise them effectively?

GPULAB RESOURCES

Several GPU clusters with different types of GPU cards

What we needed was Cluster 6:

- NVIDIA HGX-2 with **16 Tesla V100 GPUs**, 96 2.7Ghz vCPU cores with 1.5TB RAM
- NVLink switch connection fabric



ACCESS AND USAGE

We used the GPU Lab system both in interactive and CLI batch mode

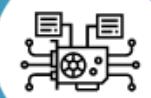
Interactive: mainly for checking resources, job status and uploading files -- JupyterHub

Batch: program execution using a batch execution system

The system is based on the Docker container system

Many predefined Docker images are available

We created our own image for HPC workloads



Dashboard

Jobs

All Jobs

My Jobs

+ Create Job

Live

Available Resources

Available Resources Per Cluster

Cluster 6

GPU Model

Tesla V100-SXM3-32GB
with 31 GB RAM

Storage

/project_ghent/
/project_scratch/

Nodes

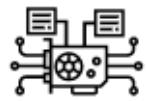
gpulab6A

ONLINE

4 / 16 GPUs available

24 / 96 CPUs available

4 / 1583 GB CPU RAM available



Jobs

List

+ Create job

Live

Clusters

Slaves

History

Summary

Users

Slaves

Jobs

Running

Queued

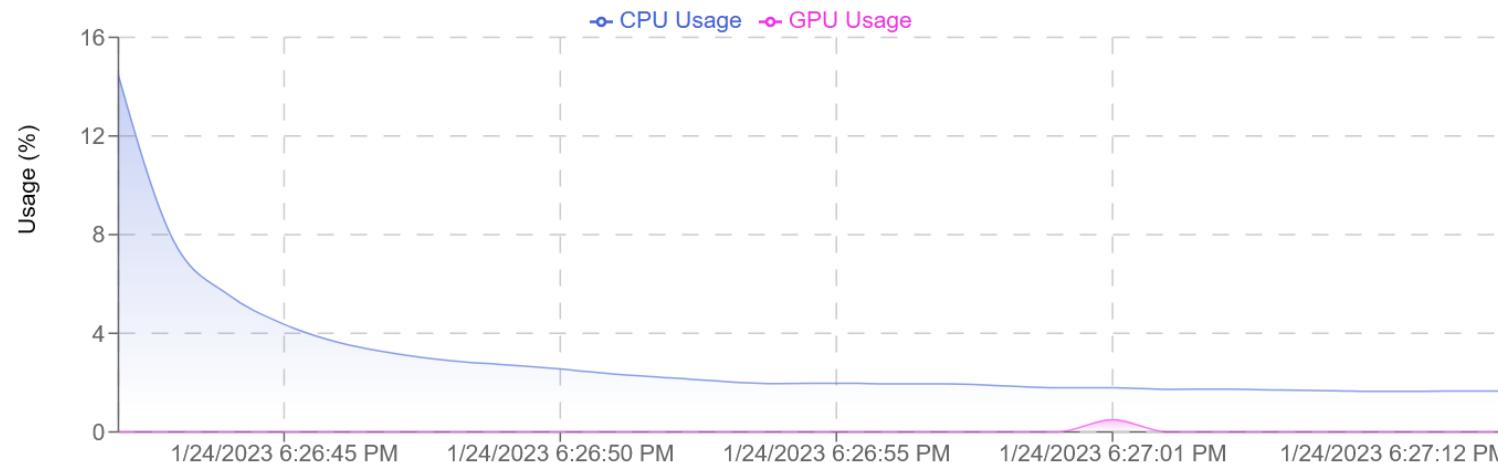
5 jobs

0 jobs

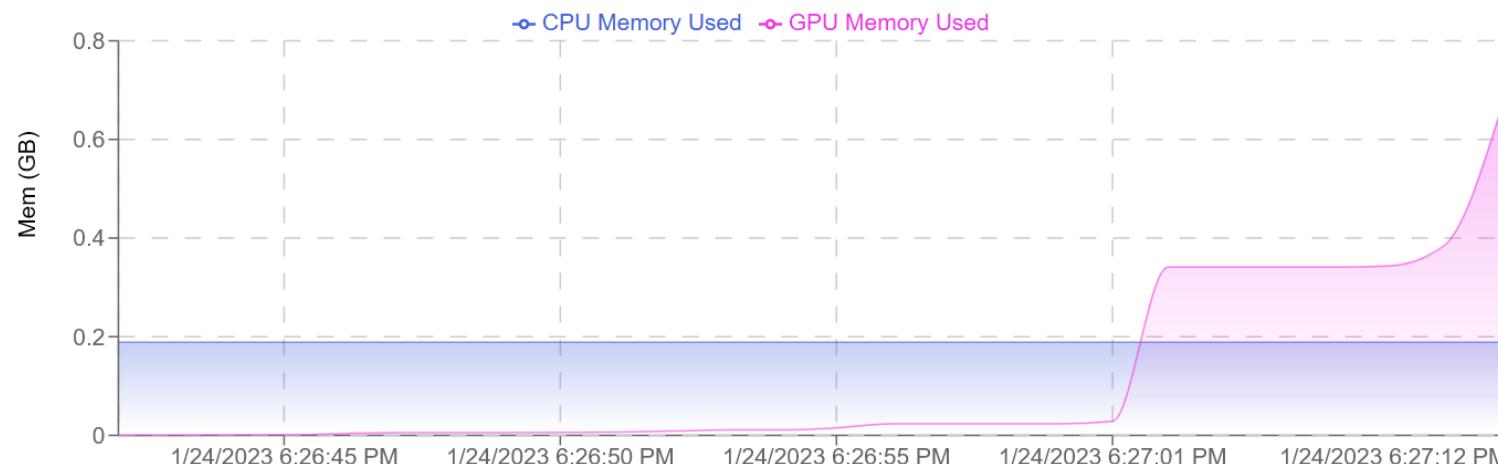
ID	Project	Username	Name	State	Created	Host
7eaec7	Orca	pbonte	JupyterHub-singleuser	RUNNING	July 22, 2019 8:25 AM	4B
ca38f0	DeepBeamforming	ykuno	JupyterHub-singleuser	RUNNING	July 22, 2019 3:55 AM	4A

[General Info](#)[Logs](#)[Debugging Logs](#)[Usage Graphs](#)[Raw Job JSON](#)

CPU/GPU Utilization

[CPU](#) [GPU](#) [Network](#) [Other](#)

Memory Utilization

[CPU](#) [GPU](#) [Network](#) [Other](#)

Sample job definition file

```
{  
  "name": "CUDA_MEMD",  
  "description": "Test_CUDA_MEMD",  
  "request": {  
    "docker": {  
      "image": "gitlab.ilabt.imec.be:4567/ilabt/gpu-docker-stacks/code-server-notebook:latest",  
      "command": "/project_ghent/startScript.sh",  
      "storage": [  
        {  
          "hostPath": "/project_ghent",  
          "containerPath": "/project_ghent"  
        }  
      ]  
    },  
    "resources": {  
      "clusterId": 6,  
      "cpus": 4,  
      "gpus": 2,  
      "cpuMemoryGb": 16,  
      "minCudaVersion": 11  
    },  
    "scheduling": {  
      "maxDuration": "1 min",  
      "interactive": false  
    }  
  }  
}
```

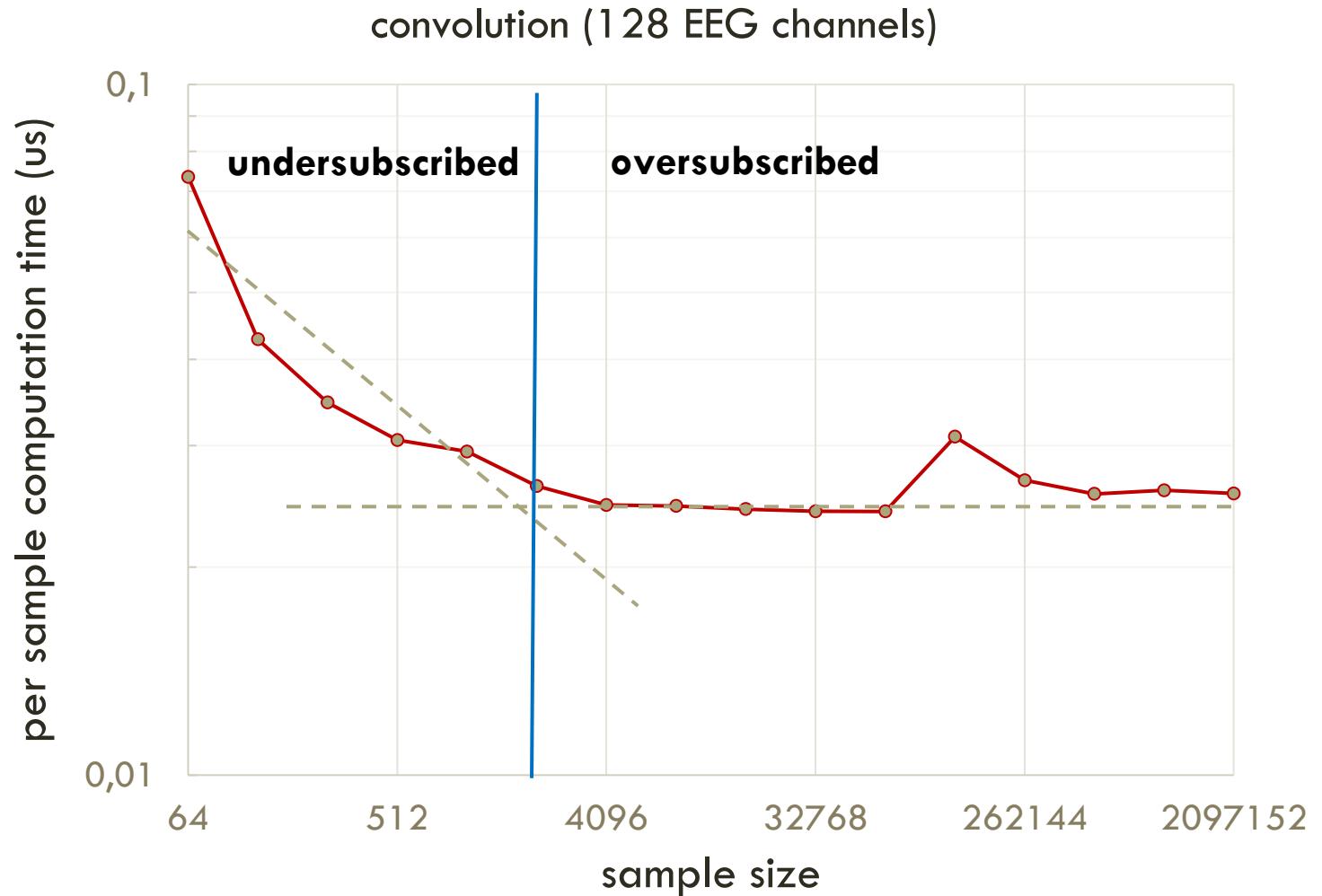
GPU EXECUTION PROPERTIES

GPU uses a fixed number of cores

throughput, not latency optimised

inefficient if there is not enough work

will not run faster if fully loaded

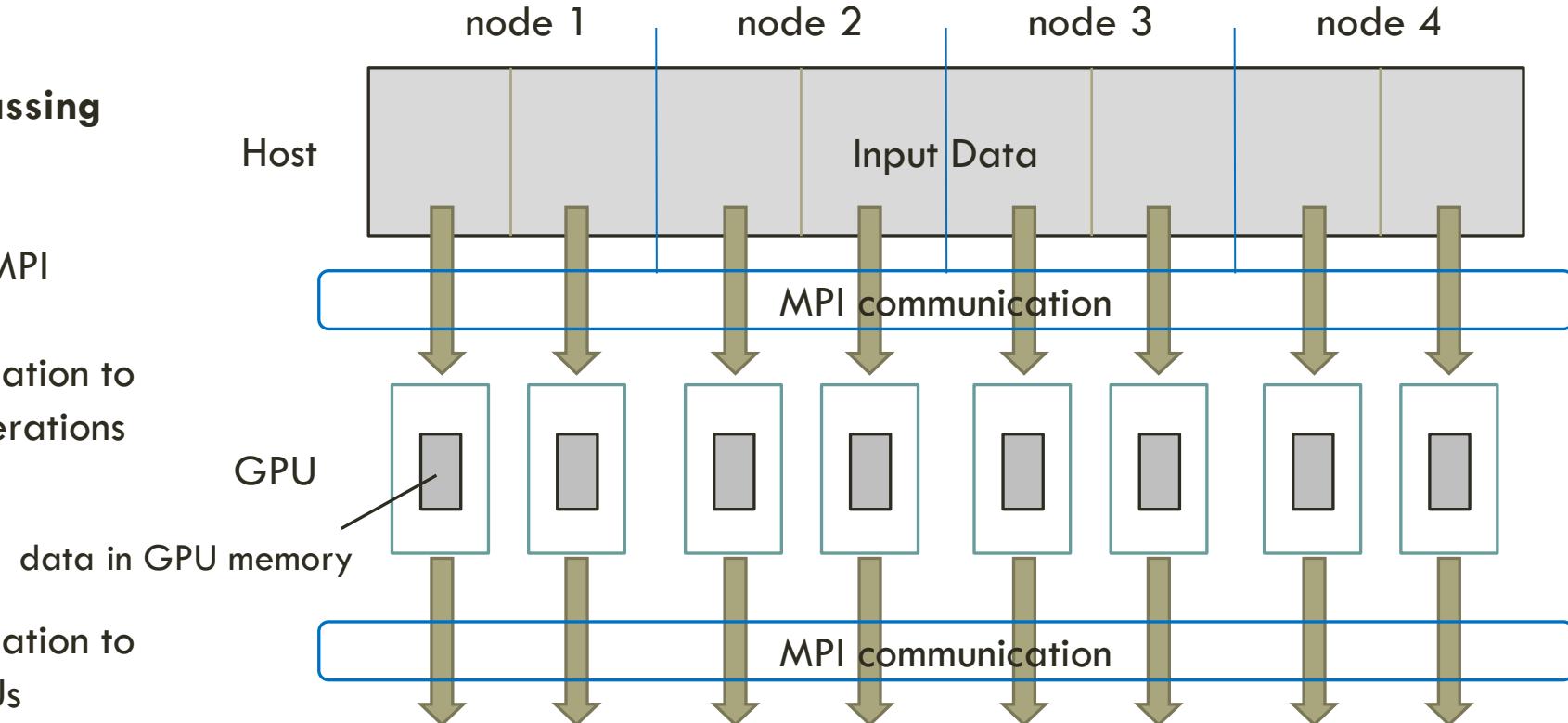


TRADITIONAL MULTI-GPU APPROACH

Distributed, message-passing
programming model

1. Distribute data using MPI
2. Perform MPI communication to collect data for GPU operations
3. Execute GPU code
4. Perform MPI communication to send results to other GPUs
5. If necessary, collect data on host and re-distribute results among nodes

...



DIRECT MULTI-GPU APPROACH

Shared-memory programming model, asynchronous in-kernel communication

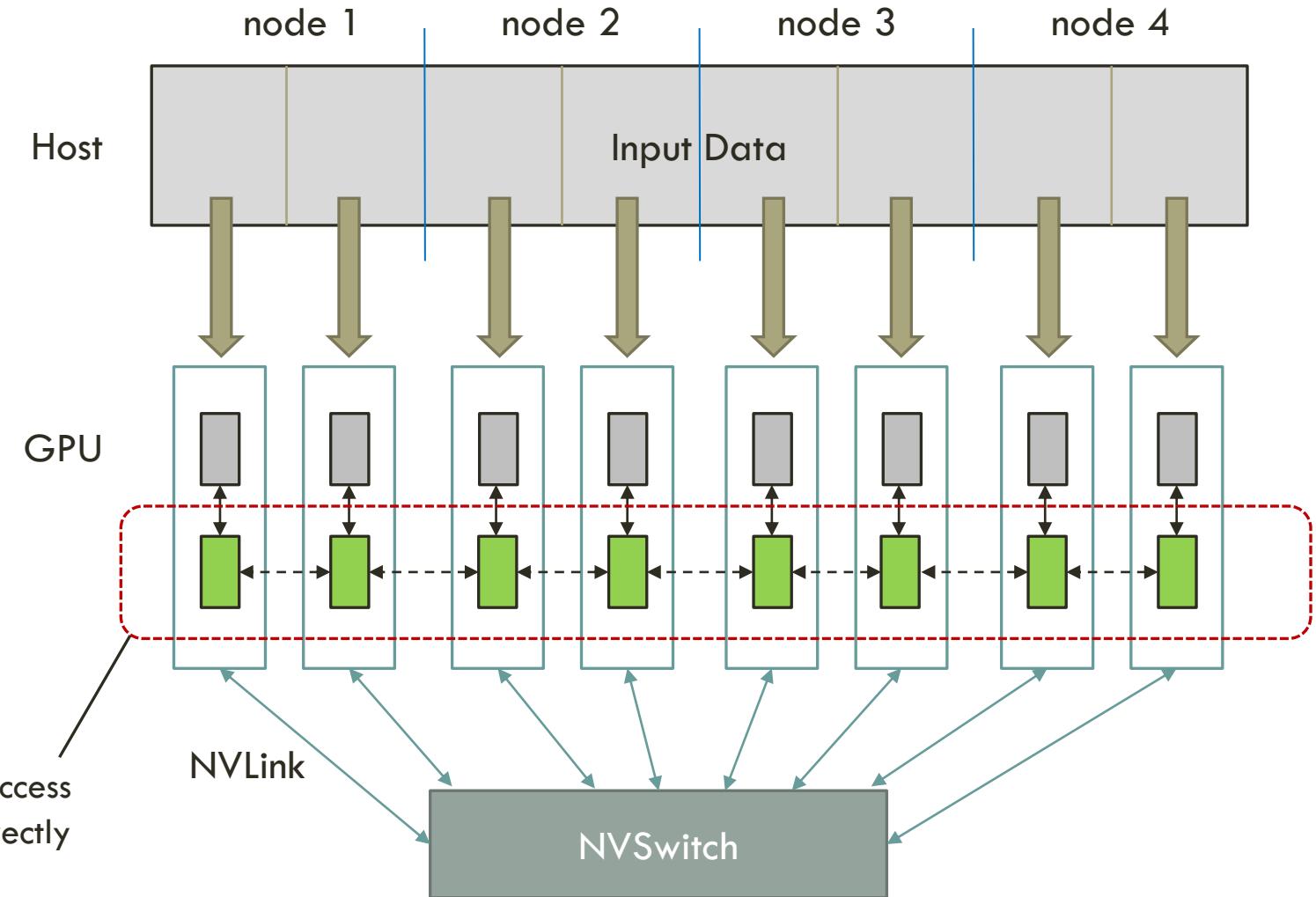
1. Distribute data using MPI

2. Execute GPU code, access other GPU memory directly for data (load/store)

3. If necessary, collect data on host and re-distribute results among nodes

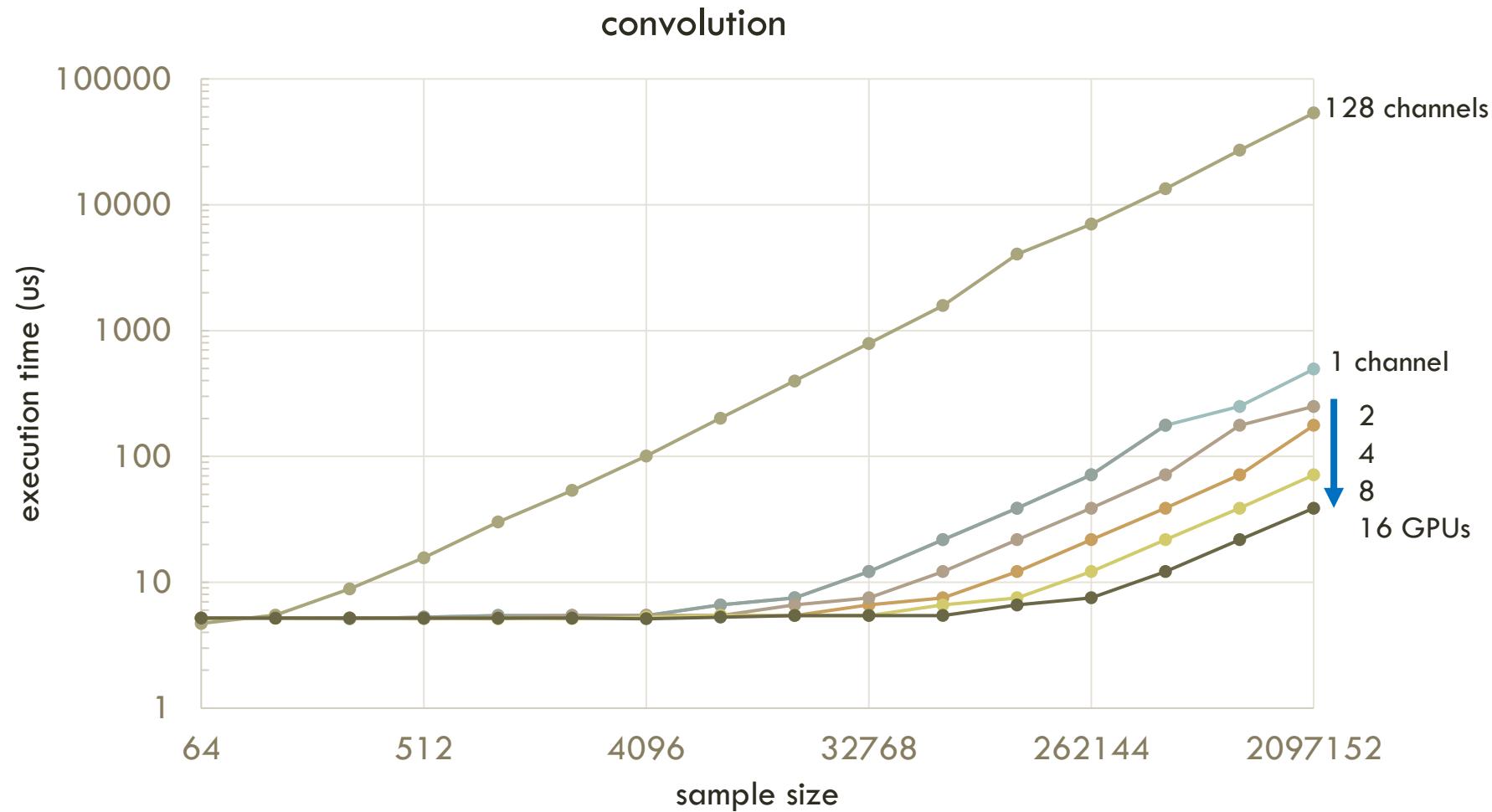
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GPU kernels access peer memory directly



RESULTS

- Single-GPU algorithms tested on the V100 GPU
- simple multi-GPU test programs developed and executed
- multi-GPU EEG algorithms designed, some are tested for functionality
- full scale scalability and performance tests in the coming weeks



CONCLUSIONS

Several multi-GPU EEG processing algorithms have been developed

Full-scale (16 GPU) scalability test is yet to be done

The SLICES-SC project and the GPULab infrastructure gave us state-of-the-art resources for the development

Allowed us to progress with our work until other HPC resources become available