

# GPU Cluster Monitoring and Management

November 15<sup>th</sup>, 2012



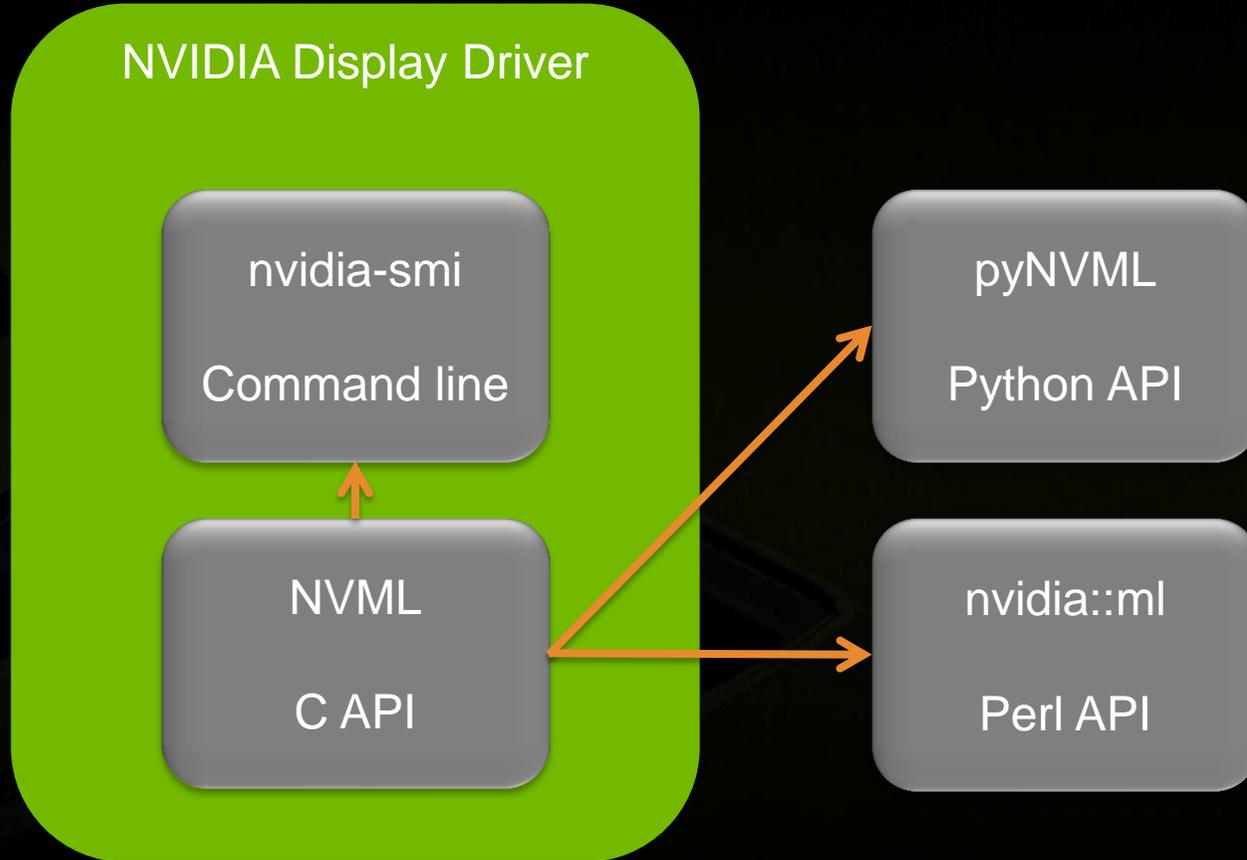
# Introductions

- **Robert Alexander**
  - **CUDA Tools Software Engineer at NVIDIA**
  - **Tesla Software Group**

# Overview

- **NVIDIA Interfaces**
- **Third Party Interfaces**
  - GPU Monitoring
  - GPU Management
  - GPU Scheduling
- **nvidia-healthmon**
- **Links and contact info**

# NVIDIA Interfaces



# GPU Monitoring Ecosystem



Third Party Tools

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NVIDIA APIs

nvidia-smi

Command line

NVML

C API

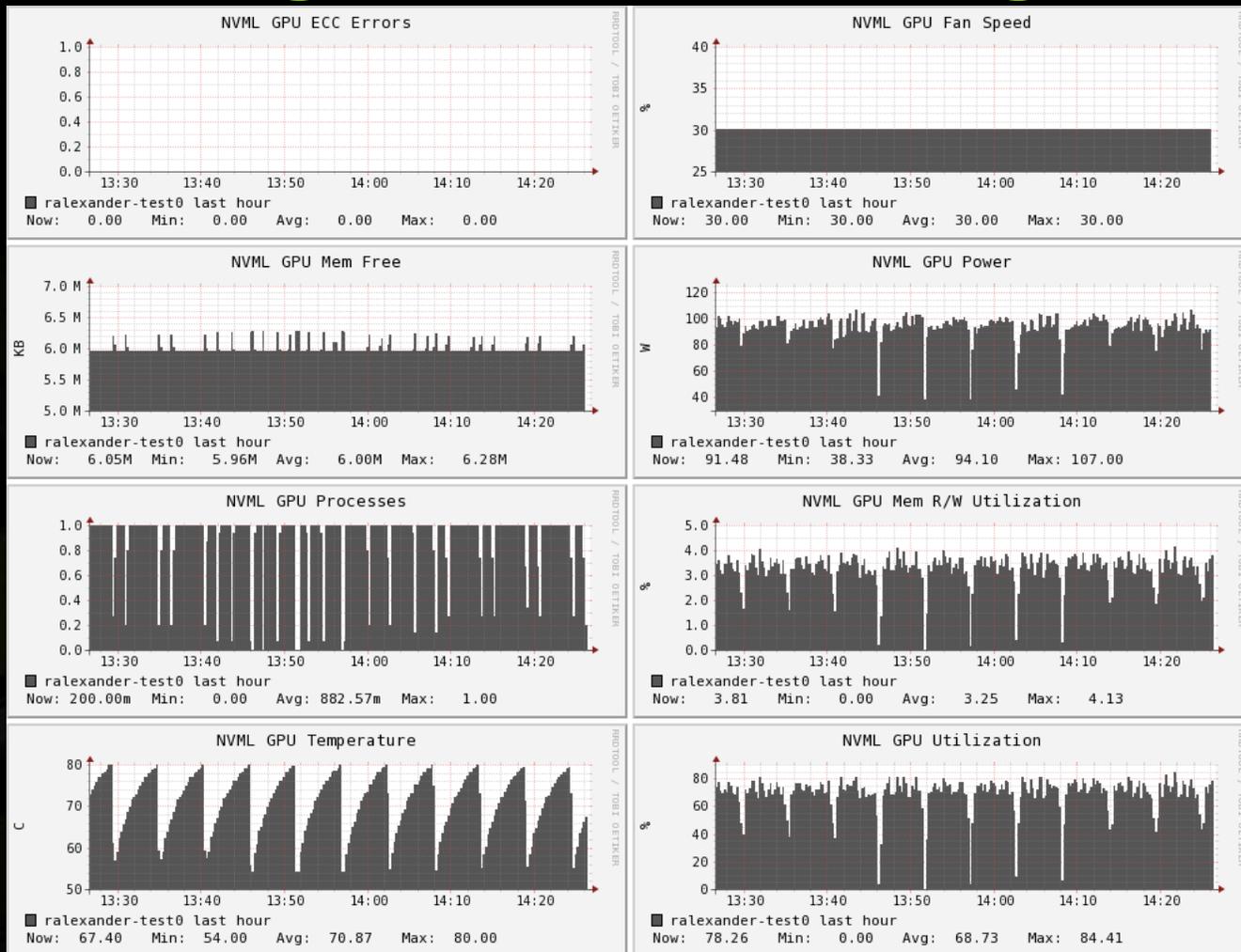
pyNVML

Python API

nvidia::ml

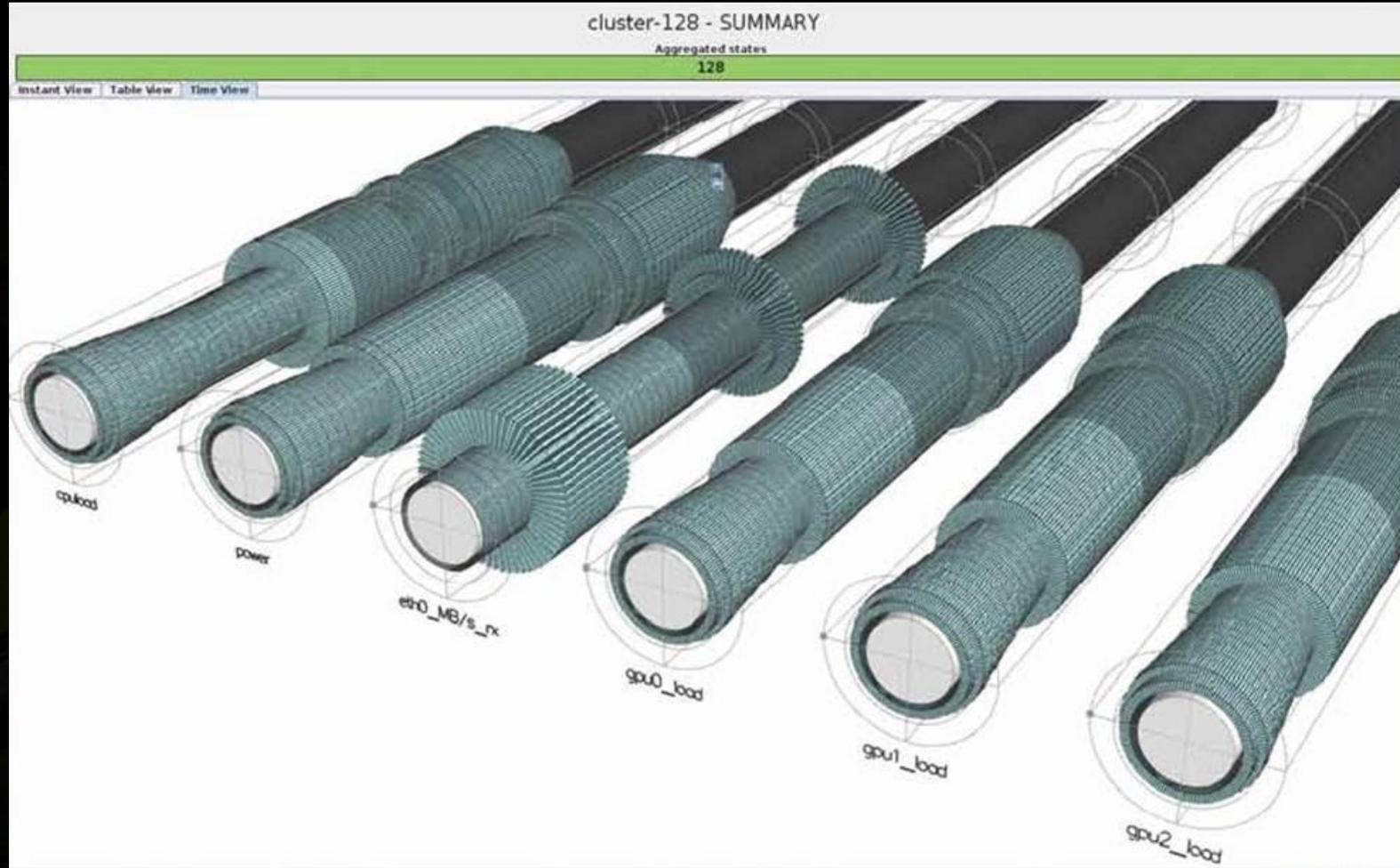
Perl API

# sFlow + Ganglia GPU Monitoring



From <http://blog.sflow.com>

# HP Insight Cluster Management Utility



From <http://hp.com>

# GPU Management Ecosystem



Third Party Tools

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Perl API

# Managing GPUs with Bright Computing



The screenshot shows the Bright Cluster Manager interface. The main window is titled 'Monitoring Configuration' and displays a table of monitoring rules. A 'Monitoring Rules Wizard' dialog box is open in the foreground, showing a list of categories to select from.

Category	Metric/Healthcheck	Parameter	Condition	Action	Action Parameter
All GPU units	gputemp	1	> 70 C	Power off	
All GPU units	gputemp	2	> 70 C	Power off	
All GPU units	gputemp	3	> 70 C	Power off	
All GPU units	gputemp	4	> 70 C	Power off	
All GPU units	ECC Errors	gpu1	> 10	SendEmail	gpu_dev@mycomp.com
All GPU units	ECC Errors	gpu2	> 10	SendEmail	gpu_dev@mycomp.com
All GPU units	gpufanspeed	1	< 1.95 KiRPM	SendEmail	gpu_dev@mycomp.com
All GPU units	gpufanspeed	2	< 1.95 KiRPM	SendEmail	gpu_dev@mycomp.com
All GPU units	gpufanspeed	3	< 1.95 KiRPM	SendEmail	gpu_dev@mycomp.com
All GPU units	gpufanspeed	4	< 1.95 KiRPM	SendEmail	gpu_dev@mycomp.com
All GPU units	gpufanspeed	5	< 1.95 KiRPM	SendEmail	gpu_dev@mycomp.com
All GPU units	gpufanspeed	6	< 1.95 KiRPM	SendEmail	gpu_dev@mycomp.com
All GPU units	gpufanspeed	7	< 1.95 KiRPM	SendEmail	gpu_dev@mycomp.com
All GPU units	gpufanspeed	8	< 1.95 KiRPM	SendEmail	gpu_dev@mycomp.com

The 'Monitoring Rules Wizard' dialog box is titled 'Monitoring Rules Wizard' and contains a list of categories to select from. The 'All GPU units' category is highlighted in green.

- All Chassis
- All Ethernet Switches
- All GPU units
- All Generic Devices
- All IB Switches
- All Master Nodes
- All Myrinet Switches
- All Power Distribution Units
- All Rack Sensors
- slave

Buttons: Cancel, Previous, Next

Buttons: Edit, Add (Wizard), Remove, Refresh, Save

MonConf 'All Chassis' refreshed, MonConf 'slave' refreshed

# GPU Scheduling Ecosystem



Third Party Tools

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nvidia::ml

Perl API

# GPU Job Scheduling with TORQUE



```
$ ./configure --with-nvml-lib=<...> --with-nvml-include=<...>
```

```
$ qsub -l nodes=1:gpus=1 myjob.sh
```

```
$ qsub -l nodes=1:gpus=2:exclusive_process myjob.sh
```

# GPU Job Scheduling with IBM PCM



The screenshot displays the IBM PCM job scheduling interface. On the left, a 'Jobs' table lists several jobs, with Job ID 104 highlighted in green and its status 'Running'. Below the table, the details for Job ID 104 are shown, including its name 'test' and status 'Running'. On the right, the 'Submit Job: GPU Job' configuration window is open, showing various options for job submission. The 'Basic Job Options' section includes fields for 'Command to run' (mm-se), 'Job Name' (Modeling -GPU), and 'Notify me when job status changes' (On). The 'Advanced' section includes 'Requirements' for processors and GPUs, and 'Limits' for maximum run time (10 hours, 0 minutes). The 'Data' section includes buttons for 'Add Local File' and 'Add Server File'.

Job ID	Job Name	Job Status	App
104	test	Running	gene
103	Is	Done	-
102	Modeling 2 - GPU	Exited	gene
101	Modeling - GPU	Exited	gene

**Submit Job: GPU Job**

Submit

**Basic Job Options**

Command to run \* mm-se

Job Name Modeling -GPU

Notify me when job status changes On

**Advanced**

Requirements

--Number of processors required to run job

Total number of CPUs required \* 2

Number of processors to use on each host 2

Total number of GPUs required \* 4

Number of GPUs to use on each host

Limits

--Maximum run time

Hour 10

Minute 0

**Data**

Input file

Specifying CPU and GPU topology in job submission

# nvidia-healthmon



- **Quick health check**
- **Not a full diagnostic**
- **Suggest remedies to SW and system configuration problems**

# nvidia-healthmon – Use Cases



- **Cluster scheduler's prologue / epilogue script**
- **Health and diagnostic suites**
  - **Designed to integrate into third party tools**
  - **Support in Warewulf NHC tool**
  - **<http://warewulf.lbl.gov/trac/wiki/Node%20Health%20Check>**
- **After provisioning cluster node**
- **Run directly, manually**

# nvidia-healthmon – Feature Set



- **Basic CUDA and NVML sanity check**
- **Diagnosis of GPU failure-to-initialize problems**
- **Check for conflicting drivers (I.E. VESA)**
- **InfoROM validation**
- **Poorly seated GPU detection**
- **Check for disconnected power cables**
- **ECC error detection and reporting**
- **Bandwidth test**

# Downloads



- **CUDA 5.0 Driver**
  - <http://developer.nvidia.com/cuda/cuda-toolkit>
- **NVML SDK + nvidia-healthmon**
  - <http://developer.nvidia.com/tesla-deployment-kit>
- **Python NVML Bindings**
  - <http://pypi.python.org/pypi/nvidia-ml-py/>
- **Perl NVML Bindings**
  - <http://search.cpan.org/~nvbinding/nvidia-ml-pl/>

# Contact Info and Links



- **Robert Alexander**
  - [ralexander@nvidia.com](mailto:ralexander@nvidia.com)
- **Developer Information**
  - <http://www.developer.nvidia.com/nvidia-management-library-nvml>

# Third Party Links



- <http://ganglia.sourceforge.net/>
- <http://sflow.org/>
- <http://www.hp.com>
- <http://www.ibm.com>
- <http://www.adaptivecomputing.com/>
- <http://www.brightcomputing.com/>
- <http://www.penguincomputing.com/>
- <http://www.stackiq.com/>
- <http://www.univa.com/>
- <http://gridscheduler.sourceforge.net/>
- <https://computing.llnl.gov/linux/slurm/>
- <http://www.pbsworks.com/>

Thanks!



# Questions?

