Hall A Analysis Software & Computing Update

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Jefferson Lab

Hall A Collaboration Meeting February 10, 2022

Podd Status

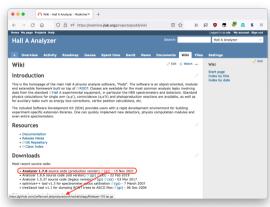
- Current release: 1.7.0 (16 Nov 2021)
 - Many new features (already presented at previous meetings)
 - Additional improvements and bugfixes based on early SBS data taking
 - Significant speedup, primarily in decoder and database
 - Improved CODA 3 support
 - Dynamic raw data event buffer size
 - PID calculation based on Bayesian likelihoods
 - ▶ Requires C++11 compiler and ROOT 6. Installed in counting house and on the farm.
- Priority development: 2.0-devel (Summer 2022, delayed because of SBS work)
 - Multithreading
 - Will benefit SBS and Hall C, primarily for online replay
 - Requires C++17 (e.g. gcc 9+, available on ifarm)
 - Existing code will need minor modifications
- Auxiliary development: 1.8-devel (if time permits)
 - Add small new features missed in 1.7
 - Maintain system requirements and API of version 1.7 as much as possible

Podd: Profile-Based Code Optimization

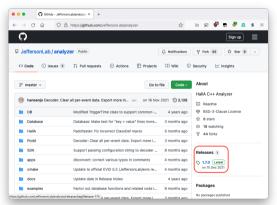
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Podd Source Code & Documentation

JLab Redmine



GitHub



Podd: Building with CMake

Prerequisites:

- Install ROOT (ensure root-config is in PATH, or set \$ROOTSYS)
 - Farm: run setroot_CUE.csh. RHEL: install from EPEL. macOS: install from Homebrew.
 - See also https://redmine.jlab.org/projects/podd/wiki/ROOT_Installation_Guide
- Ensure you have CMake ≥ 3.5 (cmake --version. cmake3 on RedHat)

Building & Installing Podd with CMake \geq 3.15

```
$ git clone https://github.com/JeffersonLab/analyzer.git
$ cmake -S analyzer -B analyzer-build [-DCMAKE_INSTALL_PREFIX=/some/dir]
$ cmake --build analyzer-build [-j4]
$ ./analyzer-build/apps/analyzer
$ [cmake --install analyzer-build]
$ [/some/dir/bin/analyzer]
```

Notes:

- Installing recommended (cmake --install): Set CMAKE_INSTALL_PREFIX
- Will phase out aging SCons build system (too many limitations)

Pre-Installed Podd

farm/ifarm (works in Counting House, too)

\$ module use /group/halla/modulefiles \$ module load analyzer \$ analyzer --version Podd 1.7.0 Linux-3.10.0-1160.31.1.el7.x86_64-x86_64 git @e26c21d ROOT 6.22/06

Counting House (local installation, faster, safer)

```
$ module use /adaqfs/apps/modulefiles
$ module load analyzer
$ analyzer --version
Podd 1.7.0 Linux-3.10.0-1160.31.1.el7.x86_64-x86_64 git @e26c21d ROOT 6.24/06
```

The SDK is located in \$ANALYZER/../src/SDK/

Podd 2.0

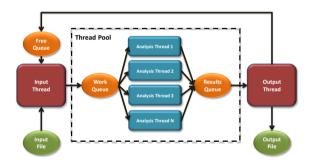
• Event-based parallelization/multithreading

- Important for online replay
- Reduced memory footprint compared to multiple individual jobs
- Requires thread safe user code (\rightarrow only const or protected globals, statics)
- $\bullet~I/O~improvements$
 - ▶ Output system upgrade (full set of data types, object variables) largely complete
 - TBD: HIPO or PODIO output file format support
 - ► TBD: EVIO 6 input format support (HIPO-like raw data files)
 - ► Goal: Make output easily usable with Python and Julia tools (*e.g.* uproot, UnROOT)

ETA: This summer. Delayed because of work on SBS.

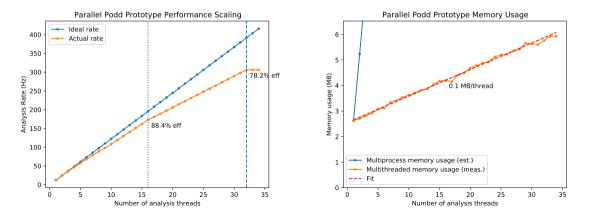
Podd Parallel Processing Prototype

- https://github.com/hansenjo/parallel
- Small standalone toy analyzer with hand-crafted multithreading (std::thread)
- Mimics main components of Podd (*e.g.* decoder, analysis variables, output)
- A few example "detectors" included whose processing is intended to burn CPU cycles
- Exploring migration to TBB (Intel Thread Building Blocks)



Parallel Podd Performance Scaling Benchmark

- Benchmark processing rate as function of number of analysis threads
- Run on aonl1 (16 hyperthreaded cores, Intel Xeon E5-2650 v2 @ 2.60GHz), RHEL 7.9, idle
- Admittedly extreme example: maximally CPU-bound (negligible I/O & memory use)



Remaining Podd Limitations



Separated Data and Algorithms

- Algorithms and Data are closely coupled
 - More work to add new algorithms
 - Difficult to stream event data only
- $\bullet\,$ No native event data I/O and API
 - Podd cannot take its own output as input
 - One-pass analysis only:

EVIO raw data \rightarrow ROOT trees + histograms

Major limitation with large data sets

• Addressing these would require complete re-write



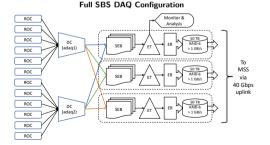
Ole Hansen (Jefferson Lab)

SBS Online Computing

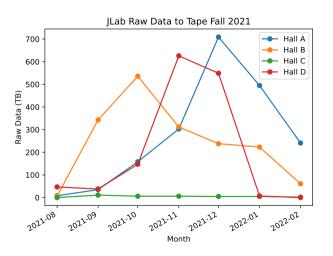
- Traditional CODA3 DAQ for GMn: Single Event Builder host (new high-performance server), demonstrated ≥ 1 GB/s peak raw data rate
- Plan to use CODA's scalable event stream parallelization to achieve up to \approx 3 GB/s
- Online replay on aonIX systems (128 threads), 2014-vintage servers (to be upgraded)
- SBS is the first experiment to take full advantage of these systems, running 100 automated parallel analysis jobs.
- Online replay typically able to keep up with incoming data.

New CODA Event Builder Machines





SBS-GMn Data Volume in Comparison



Hall	Total (TB)
А	1,949
В	1,719
С	35
D	1,414
Sum	5,117

Fun fact #1: SBS-GMn took \approx 5 times more data in these 6 months than all prior Hall A experiments combined in 25 years

Fun fact #2: The entire SBS program expects to accumulate \approx 25 PB raw data through 2024

Scientific Computing Resources

- Farm/ifarm upgraded to CentOS 7.9. RHEL 8 clones being evaluated.
- Farm batch system has been transitioned to slurm and swif2. Legacy Auger/swif commands will stop working March 1, 2022. See https://scicomp.jlab.org/docs/FarmUsersGuide.
- Current farm resources
 - ► Disk: Lustre: 4.1 PB, Work: 1.4 PB (recent upgrade).
 - ► CPU: 14192 cores / 28384 threads. Total capacity 249 M-core-hours/year
 - Almost half the capacity is on AMD EPYC 7502 64C/128T systems (speed demons!)
 - 6 nodes with Nvidia TitanRTX GPUs dedicated for ML applications
- Mass storage system (as of Feb 2022)
 - Throughput $\approx 8 \text{ GB/s}$ (20 LTO-8 drives, uncompressed, theoretical)
 - ightarrow = 150 PB capacity (LTO-8, uncompressed), pprox 85 PB used (23.4 raw, 26.7 rawdup).
 - Significant capacity headroom (more frames, LTO-9) with current silo, up to pprox 325 PB.

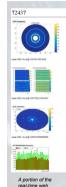
New Counting House Desktop Systems

- Clean separation of desktops and servers. \rightarrow increased reliability and stability.
- Platform for browsers, editors, slow controls (some), remote logins
- Extensive use of VNC servers/clients, very successful
- No significant issues. Small updates planned (EPICS etc.)
- Feedback welcome (ole@jlab.org)



AI-Assisted Online Monitoring (Hydra)

- EPSCI group has offered support to deploy the Hall D Hydra system in Hall A for automated data quality monitoring.
- Will tap into online histograms generated by panguin.
- Currently being set up. Test version expected \approx March–April.
- One-time human review ("labeling") required. Volunteers welcome.



dashboard of Hydra

Hydra A.I. Data Quality monitoring

- Traditionally, scientists working shifts must frequently scan dozens of plots to ensure the quality of incoming data
- Plots are themselves just pictures. A.I.'s are now very good at classifying pictures.
- This is applied A.I. since it uses models already designed for image classification such as Google's Inception_v3 network
- Between 93 and 99% accurate when compared to expert labeling

 Has found mislabeling by human experts indicating an irreducible error that is expert dependent

- Currently capable of analyzing an image in under 200ms
 - This equates to a throughput in excess of 10,000 images a day when running. (far more than a human)

(slide from David Lawrence, Jan 2021)





- "Podd" analysis software continues to be actively maintained and used by current experiments in Halls A & C.
- Significant modernization work (multithreading etc.) underway.
- The large data volumes from SBS are putting Hall A in the same league as Halls B & D in terms of computing resource needs. This will require careful planning going forward.
- Experience with the upcoming SBS mass replays on the farm will inform future direction of the Hall A software.