



Alignment Monitoring

Jim Pivarski, Dmitry Yakorev, Alexei Safonov

Texas A&M University

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Alignment monitoring: roughly four categories

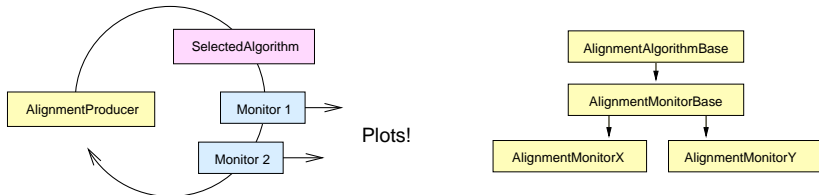
1. DQM-based monitoring upstream of alignment process
(reports an error if alignment used online is wrong)
2. Sanity checks in AlignmentProducer (convergence, improvement in residuals, overlap plots, p_T)
3. Geometry Validation— compare output geometries from different alignments: have the chambers moved?
4. Validation with reconstructed tracks: is it better?

Status of 2: Foundation for plotting modules in AlignmentProducer

Status of 3: Reading from multiple SQLite files, plotting difference

Plotting Modules in AlignmentProducer (#2)

- ▶ Extends existing plugin mechanism to add monitoring modules



- ▶ Lowers “potential barrier” to adding plots
- ▶ Independent of algorithm (though we can make **AlignmentMonitorHIP**, **AlignmentMonitorMillePede**, etc.)
- ▶ Aware of iteration number, selected alignables and parameters
- ▶ Collects and merges histograms/profiles from a distributed job



Plotting Modules in AlignmentProducer (#2)

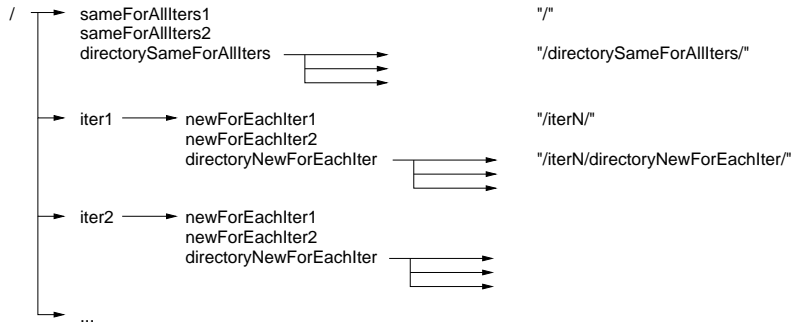
```
replace AlignmentProducer.monitorConfig = {  
    untracked vstring monitors = {"AlignmentMonitorHIP"}  
    PSet AlignmentMonitorHIP = {  
        string outpath = "./"  
        string outfile = "histograms.root"  
        bool collectorActive = false  
        int32 collectorNJobs = 0  
        string collectorPath = "./"  
    }  
}  
  
void AlignmentMonitorHIP::book() {  
    m_sameForAllIters = (TH1F*)(add("/", new TH1F(...)))  
    m_newForEachIter = (TH1F*)(add("/iterN/", new TH1F(...)))  
}
```

Plotting Modules in AlignmentProducer (#2)

```
m_sameForAllIters = (TH1F*)(add("/", new TH1F(...)))
m_newForEachIter = (TH1F*)(add("/iterN/", new TH1F(...)))
```

ROOT file structure

what to type in add()



Nothing more is needed for collection jobs



Plotting Modules in AlignmentProducer (#2)

What works (tested for 25 events):

- ▶ Loading an arbitrary number of modules
- ▶ Arbitrarily-deep ROOT directory structure
- ▶ Iteration (via `AlignmentProducer.maxLoops` and/or files)
- ▶ Merging histograms/profiles from a distributed job
- ▶ Generating histograms from selected Alignables

What's next:

- ▶ Add lots of plots to AlignmentMonitorHIP
- ▶ Test with lots of events
- ▶ Put into CVS?



Reading from multiple databases (#3, Dmitry)

Basic structure:

- ▶ Compiled C++ ROOT GUI
- ▶ Forks cmsRun processes which read SQLite files

What works:

- ▶ Loading two geometry files
- ▶ Calculating and displaying translation differences
- ▶ Tabs to switch between plots

What's next:

- ▶ Read from multiple databases— plot versus time
- ▶ Represent differences in rotation angles
- ▶ A different framework? DQM? Iguana? Compile database access into ROOT GUI?



Reading from multiple databases (#3, Dmitry)

vs.
 R 

$$(\vec{x}_1 - \vec{x}_2)$$

$$(\vec{x}_1 - \vec{x}_2) \cdot \hat{\phi}$$

$$(\vec{x}_1 - \vec{x}_2) \cdot \hat{Z}$$



Summary

2. Sanity checks in AlignmentProducer (convergence, improvement in residuals, overlap plots, p_T)

Laid a foundation that handles multiple modules, ROOT directory structure with iterations, merging histograms after parallel processing

3. Geometry Validation— compare output geometries from different alignments: have the chambers moved?

Measuring geometry differences across multiple SQLite files, beginnings of a GUI tool