# RNTuple: latest developments & the RNTupleProcessor

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Based on 25+ years of **TTree** experience, **RNTuple** is a redesigned columnar I/O subsystem aiming at:

- Less disk and CPU usage through smaller files and higher throughput;
- Systematic use of data checksums and runtime exceptions to prevent silent I/O errors;
- Efficient support for modern hardware;
- Native support for local and remote ROOT files and object stores;
- Coverage of all of today's TTree use cases (integration in both Athena and CMSSW);
- A binary format defined in a dedicated specification.

The first production version of the **RNTuple** on-disk binary format is available in ROOT 6.34. The API will start moving out of ROOT::Experimental from ROOT 6.36 onward.

#### Recent performance results I





CMS I/O Throughput vs Threads



CHEP '24: RNTuple: A CMS perspective

### Recent performance results II





CHEP '24: ROOT RNTuple and EOS: The Next Generation of Event Data I/O

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## Combining data sets with the **RNTupleProcessor**

Why do we want to combine data sets?



A data set is typically stored across multiple files (*samples*), but we want **seamless** event processing.

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#### Why do we want to combine data sets?



A data set is typically stored across multiple files (*samples*), but we want **seamless** event processing.

- → need to be able to **vertically concatenate** samples.
- 1. Analysis may require objects not present in the compact data format;
- 2. Analyses could be sped up by storing and reusing (expensive) intermediate computation results.
- → need to be able to **horizontally concatenate** samples.

#### Combining HEP data sets



**TTree** has the ability to concatenate data sets in two directions:

- 1. Vertically through the TChain interface;
  - → comparable to a SQL UNION ALL operation (*but not exactly*).
- 2. *Horizontally* through the TTree::AddFriend interface, possibly using a TTreeIndex for unaligned entries
  - → comparable to a SQL JOIN operation (*but not exactly*).

They can be combined using TChain::AddFriend.

Similar functionality is desired for **RNTuple**. We want to provide additional composition flexibility and above all, prevent users from accidentally getting erroneous data.

#### Data set joins: the ideal case





#### Data set joins: a realistic scenario





#### The caveats of unaligned data set joins



- Which events belong together?
  - Both false positives and negatives are unacceptable!
- What if the right-hand side event data is missing?
- What if my events are scattered across multiple files?
- What if want to distribute my analysis?



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- What if want to distribute my analysis?
- + How to express all of this nicely?



### Handling unaligned joins



When events between two data sets don't align on their entry numbers, we need a join table:

- Mapping between values of one or multiple *join columns* and corresponding entry numbers;
  - Support for up to 4 integral-type join columns;
  - Multiple column values are combined into a single hash.
- Built for the *auxiliary data set*;
- Probed using values from the *primary data set*.



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```
std::vector<RNTupleSourceSpec> ntuples{
    {"myElectrons", "electrons1.root"}, {"myElectrons", "electrons2.root"}};
auto processor = RNTupleProcessor::CreateChain(ntuples);
for (const auto &entry : *processor) {
    std::cout << "pt = " << *entry.GetPtr<float>("pt") << std::endl;
}</pre>
```

#### → See the ntpl012\_processor\_chain.C tutorial

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```
std::vector<RNTupleSourceSpec> ntuples{
    {"myElectrons", "electrons.root"}, {"myMuons", "muons.root"}};
auto processor = RNTupleProcessor::CreateJoin(ntuples, {"run", "event"});
for (const auto &entry : *processor) {
    std::cout << "electron pt = " << *entry.GetPtr<float>("pt") << std::endl;
    std::cout << "muon pt = " << *entry.GetPtr<float>("myMuons.pt") << std::endl;
}</pre>
```

#### → See the ntpl015\_processor\_join.C tutorial

## Composability of the RNTupleProcessor





Each processor implements the same interface for loading entries, allowing for arbitrary composition ordering.

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### Chain-first approach





```
auto electrons = {Create({"myElectrons", "electrons1.root"}), Create({"myElectrons", "electrons2.root"})};
auto muons = {Create({"myMuons", "muons1.root"}), Create({"myMuons", "muons2.root"})};
auto electronChain = CreateChain(electrons);
auto muonChain = CreateChain(muons);
auto processor = CreateJoin({electronChain, muonChain}, {"run", "event"});
```

#### Join-first approach





```
auto emPair1 = {Create{{"myElectrons", "electrons1.root"}), Create{{"myMuons", "muons1.root"})};
auto emPair2 = {Create({"myElectrons", "electrons2.root"}), Create({"myMuons", "muons2.root"})};
auto electronMuonJoin1 = CreateJoin(emPair1, {"run", "event"});
auto electronMuonJoin2 = CreateJoin(emPair2, {"run", "event"});
auto processor = CreateChain({electronMuonJoin1, electronMuonJoin2});
```

#### Foreseen integration with RDataFrame

- The RNTupleProcessor will become the under-the-hood processing engine in RDataFrame;
  - → Completely transparent to users!
- Opportunity to further evolve **RDataSetSpec**.
  - → This will become the interface for "building" the RNTupleProcessor.



```
"samples": [
    "identifier": "electrons",
    "name": "mvElectrons".
    "files": ["electrons1.root",
              "electrons2.root"].
    "joinWith": {
      "sample": "muons".
      "joinOn": ["run", "event"].
      "eventAlignment": "file"
   Ъ.
    "identifier": "muons".
    "name": "mvMuons".
    "files": ["muons1.root".
              "muons2.root"]
                              spec.ison
```

#### Current status and outlook



- RNTupleChainProcessor and RNTupleJoinProcessor are currently available in ROOT master;
- Work towards enabling composing RNTupleProcessors is well underway;
- Integration with **RDataFrame** is planned for 2025;
- Performance profiling and optimization for *unaligned joins* is planned for 2025;
- Support for the **RNTupleProcessor** in *distributed settings* (through **RDataFrame**) is foreseen.