art and `ParameterSet` validation:

Technology preview

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FHiCL and *art*

- *art* processes are configured using FHiCL files, that can look like this.

```
#include "minimalMessageService.fcl"
#include "standardProducers.fcl"
#include "standardServices.fcl"

process_name : PbarS3

# Start form an empty source
source :
{
    module_type : EmptyEvent
    maxEvents : 1000
}

services :
{
   message               : @local::default_message
   TFileService          : { fileName : "hist_pbar_s3.root" }
   RandomNumberGenerator : { }

   GeometryService   : { inputFile : "JobConfig/TDR/geom_pbar_s3.txt" }
   ConditionsService : { conditionsfile : "Mu2eG4/test/conditions_01.txt" }
   GlobalConstantsService : { inputFile : "Mu2eG4/test/globalConstants_01.txt" }
}

physics : {
   producers: {

      g4run : {
         module_type          : G4
         inputSimParticles    : "g4filter:s0"
         generatorModuleLabel : rotatetarget
         SDConfig : {
            sensitiveVolumes: [ TS1Vacuum, Coll11, Coll12 ]
         }
      }
   }

  p1 : [ g4run ]
  trigger_paths : [p1]

}
```

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                sensitiveVolumes: [ TS1Vacuum, Coll11, Coll12 ]
            }
        }
    }

   p1 : [ g4run ]
   trigger_paths : [p1]

}
```

Atom: no underlying structure

```
module_type : EmptyEvent
```

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   ConditionsService : { conditionsfile : "Mu2eG4/test/conditions_01.txt" }
                                                          tants_01.txt" }
```

Sequence: list whose elements are unnamed objects

```
sensitiveVolumes: [ TS1Vacuum, Coll11, Coll12 ]
```

```
            sensitiveVolumes: [ TS1Vacuum, Coll11, Coll12 ]
        }
      }
   }

  p1 : [ g4run ]
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}
```

FHiCL and *art*

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   ConditionsService : { conditionsfile : "Mu2eG4/test/conditions_01.txt" }
                                          est/globalConstants_01.txt" }
                                          oll12 ]
}
```

Table: Object with underlying name-value pairs.

```
source :
{
    module_type : EmptyEvent
    maxEvents : 1000
}
```

FHiCL is very flexible

- The user decides how complicated the job configuration should be
  - Simpler configurations are better.
  - However, it can support very complicated nested structures.

- Retrieving parameter values in your source code is fairly straightforward (e.g.)

```
pset.get<int>("someInt")
pset.get<double>("someTable.someDouble");
```

- But the current system has limitations …

Some limitations

- Parameter misspellings are not noticed for parameters that have defaults.
- It is not possible to know which parameters are supported for a given module
  - i.e. no FHiCL description for a given module
- Parameter retrievals (i.e. pset.get<T>) can be awkward for structures more complicated than atoms.

ParameterSet validation goals

Validation

- Provide a means of validating configuration files against a specified reference for a given module.
  - Notifies user of parameters in *.fcl files that are not supported (e.g. misspellings fall into this category)
  - Notifies user of parameters that are ***missing*** from their FHiCL.

Description

- The specified reference must serve as a description so that users do not need to look at source code to determine the allowed configuration.

Ease of use

- User interface must be straightforward to understand and use.

Show and tell

- Today I'm using a toy module.
- This is very similar to what you would see when *art* constructing an *art* module.

```
#include "fhiclcpp/static_types/Atom.h"
#include "fhiclcpp/static_types/Sequence.h"
#include "fhiclcpp/static_types/Table.h"
#include "test/static_types/macros.h"

#include <iostream>
#include <string>

using namespace fhicl::static_types;

namespace {

  //================================================================================
  // Job configuration
  //
  struct Parameters {
  };

  //================================================================================
  // Module declaration
  //
  class MyModule : public art::EDProducer {
  public:

    using Parameters = ::Parameters;

    MyModule(Table<Parameters> const &) {
    }

  };

}

DEFINE_TEST(MyModule)
```

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No longer uses `fhicl::ParameterSet`

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#include <string>

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  //====================================
  // Job configuration
  //
  struct Parameters {
  };

  //================================================================================
  // Module declaration
  //
  class MyModule : public art::EDProducer {
  public:

    using Parameters = ::Parameters;

    MyModule(Table<Parameters> const &) {
    }

  };

}

DEFINE_TEST(MyModule)
```

Allowed parameters be specified here.

No longer uses `fhicl::ParameterSet`

The simple case – one atom

```
#include "fhiclcpp/static_types/Atom.h"
#include "fhiclcpp/static_types/Sequence.h"
#include "fhiclcpp/static_types/Table.h"
#include "test/static_types/macros.h"

#include <iostream>
#include <string>

using namespace fhicl::static_types;

namespace {

  //======================================================================
  // Job configuration
  //
  struct Parameters {
  };

  //======================================================================
  // Module declaration
  //
  class MyModule : public art::EDProducer {
  public:

    using Parameters = ::Parameters;

    MyModule(Table<Parameters> const &) {
    }

  };

}

DEFINE_TEST(MyModule)
```

```
pset: {

   oneAtom : "g-2"

}
```

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#include "fhiclcpp/static_types/Atom.h"
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#include <iostream>
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using namespace fhicl::static_types;

namespace {

  //================================================================================
  // Job configuration
  //
  struct Parameters {
  };

  //================================================================================
  // Module declaration
  //
  class MyModule : public art::EDProducer {
  public:

    using Parameters = ::Parameters;

    MyModule(Table<Parameters> const &) {
    }

  };

}

DEFINE_TEST(MyModule)
```

```
pset: {

   oneAtom : "g-2"

}
```

Loading module:

```
The following keys are present in your FHiCL configuration but not supported:
+ pset.oneAtom
```

The simple case – one atom

```
namespace {

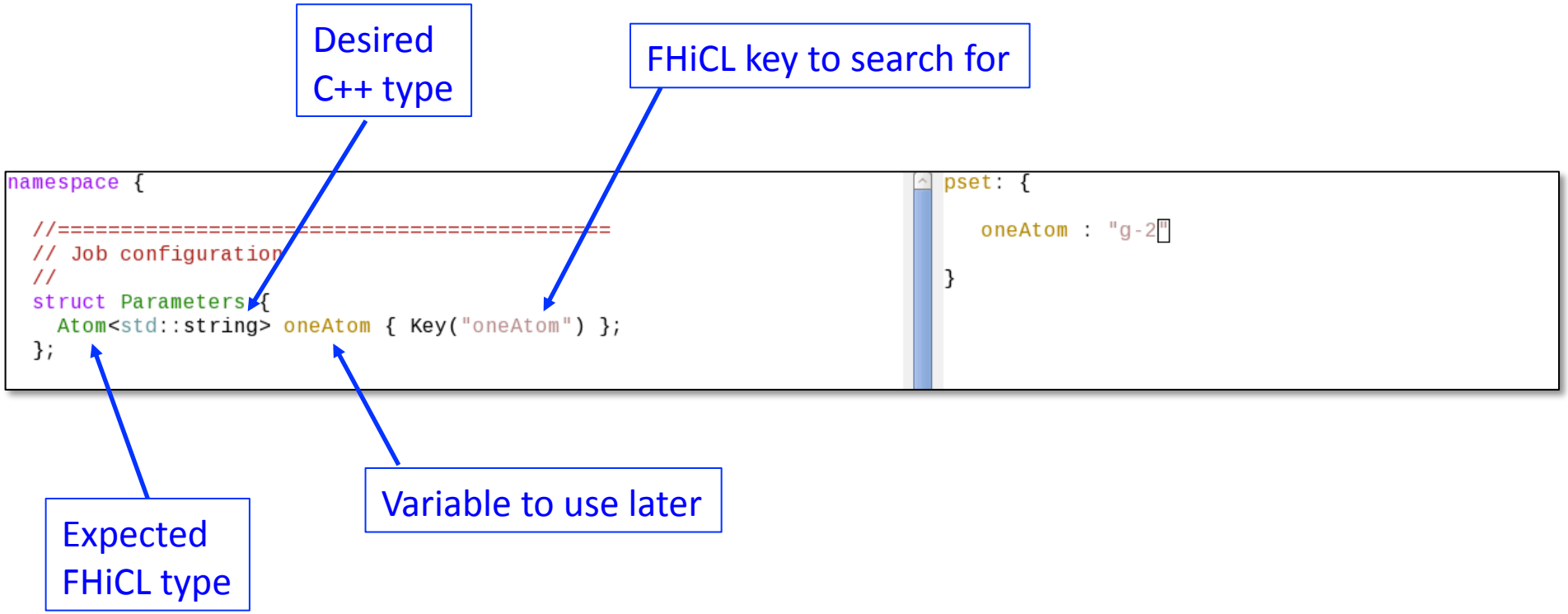
  //==========================================
  // Job configuration
  //
  struct Parameters {
    Atom<std::string> oneAtom { Key("oneAtom") };
  };
```

```
pset: {

   oneAtom : "g-2"

}
```

The simple case – one atom



Adding some structure

```
namespace {

  //=============================================
  // Job configuration
  //
  struct Parameters {
    Atom<std::string> oneAtom { Key("oneAtom") };

     What goes here?
  };
```

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

}
```

Adding some structure

```
namespace {

  //============================================
  // Job configuration
  //
  struct Parameters {
    Atom<std::string>  oneAtom  { Key("oneAtom")  };
    Atom<int>          value    { Key("value"), 8 };
    Atom<std::vector<int> > /*?*/ list { Key("list") };
  };
```

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

}
```

If you compile you get …

Adding some structure

```
namespace {

  //============================================
  // Job configuration
  //
  struct Parameters {
    Atom<std::string>  oneAtom  { Key("oneAtom")  };
    Atom<int>          value    { Key("value"), 8 };
    Ato
  };
```

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

}
```

```
fhiclcpp error: Cannot create an 'Atom' with any of the following types

                .. std::array
                .. std::pair
                .. std::vector
                .. std::tuple

                Please use one of the 'Sequence' options:

                .. Sequence<int>              ===> std::vector<int>
                .. Sequence<int,4>            ===> std::array <int,4u>
                .. Tuple<int,double,bool>     ===> std::tuple <int,double,bool>
                .. Sequence<Sequence<int>,4> ===> std::array <std::vector<int>,4u>
                .. etc.
```

Compile-time (!) error

Including sequences

```
namespace {

  //===============================================
  // Job configuration
  //
  struct Parameters {
    Atom<std::string> oneAtom { Key("oneAtom") };
    Atom<int> value { Key("value") };
    Sequence<int> list { Key("list") };
  };
```

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

}
```

Including sequences

```
namespace {

  //================================================
  // Job configuration
  //
  struct Parameters {
    Atom<std::string> oneAtom { Key("oneAtom") };
    Atom<int> value { Key("value") };
    Sequence<int> list { Key("list") };
  };
```

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

}
```

- Three kinds of containers that support sequences (e.g.):

fhiclcpp type	**Underlying `std::` type**
`Sequence<int>`	`std::vector<int>`
`Sequence<int,4>`	`std::array<int,4>`
`Tuple<int,double>`	`std::tuple<int,double>`

Introducing defaults

Attempt to override default

```
namespace {

  //===========================================
  // Job configuration
  //
  struct Parameters {
    Atom<std::string> oneAtom { Key("oneAtom") };
    Atom<int> value { Key("value"), 8 };
    Sequence<int> list { Key("list") };
  };
```

```
pset: {

   oneAtom : "g-2"
   valu    : 7
   list    : [1,3,17]

}
```

Introducing defaults

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```
namespace {

  //================================================
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  struct Parameters {
    Atom<std::string> oneAtom { Key("oneAtom") };
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    Sequence<int> list { Key("list") };
  };
```

```
pset: {

   oneAtom : "g-2"
   valu    : 7
   list    : [1,3,17]

}
```

```
The following keys are present in your FHiCL configuration but not supported:
+ pset.valu
```

Introducing nested tables

```
namespace {

  //===========================================
  // Job configuration
  //
  struct G4Config {
    Atom<double> energyCutoff { Key("energyCutoff") };
  };

  struct Parameters {
    Atom<std::string> oneAtom  { Key("oneAtom")  };
    Atom<int>         value    { Key("value"), 8 };
    Sequence<int>     list     { Key("list")     };
    Table<G4Config>   g4config { Key("g4Config") };
  };
```

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

   g4Config : {
      energyCutoff : 27
   }

}
```

Introducing nested tables

```
namespace {

  //============================================
  // Job configuration
  //
  struct G4Config {
    Atom<double> energyCutoff { Key("energyCutoff") };
  };

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    Atom<int>         value    { Key("value"), 8 };
    Sequence<int>     list     { Key("list")     };
    Table<G4Config>   g4config { Key("g4Config") };
  };
```

```
pset: {

   oneAtom  : "g-2"
   value    : 7
   list     : [1,3,17]

   g4Config : {
       energyCutoff : 27
     }

}
```

Introducing nested tables

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  // Job configuration
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  struct G4Config {
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  };

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  };
```

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

   g4Config : {
      energyCutoff : 27
   }

}
```

Introducing nested tables

Omitting key that's expected

```
namespace {

  //=============================================
  // Job configuration
  //
  struct G4Config {
    Atom<double> energyCutoff { Key("energyCutoff") };
  };

  struct Parameters {
    Atom<std::string> oneAtom  { Key("oneAtom")  };
    Atom<int>         value    { Key("value"), 8 };
    Sequence<int>     list     { Key("list")     };
    Table<G4Config>   g4config { Key("g4Config") };

  };
```

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

   g4Config : {
      # energyCutoff   27
   }

}
```

```
The following keys are missing from your FHiCL configuration:
- pset.g4Config.energyCutoff
```

Introducing nested tables

```
namespace {

  //============================================
  // Job configuration
  //
  struct G4Config {
    Atom<double> energyCutoff { Key("energyCutoff") };
  };

  struct Parameters {
    Atom<std::string> oneAtom  { Key("oneAtom")  };
    Atom<int>         value    { Key("value"), 8 };
    Sequence<int>     list     { Key("list")     };
    Table<G4Config>   g4config { Key("g4Config") };
  };
```

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

   g4Config : {
      energyCutoff : 27
   }

}
```

- From these tools, the C++ source can describe any FHiCL parameter set.

Allowed types (e.g.)

```
Atom<T>
Sequence<T>
Sequence<T,SZ>
Tuple<T…>
Tuple< Sequence<T>, U… >
Tuple< Sequence<T,SZ>, U… >
Tuple< Tuple<T…>,U…>
Sequence< Tuple<T…> >
Sequence< Tuple<T…>, SZ >
Sequence< Sequence<T> >
Sequence< Sequence<T,SZ> >
Sequence< Sequence<T>, SZ >
Sequence< Sequence<T,SZ>, SZ >
Table<S>
Sequence< Table<S> >
Sequence< Table<S>, SZ >
Tuple< Table<S>, U… >
Tuple< Sequence< Table<S> >, U… >
Tuple< Sequence< Table<S>, SZ>, U… >
Sequence< Tuple< Table<S>, U… > >
Sequence< Tuple< Table<S>, U… >, SZ>
```

N.B. List is meant to illustrate flexibility of system,
not to encourage the use of complicated types.

Print description of expected parameters

```
art --module-description MyModule
```

```
//================================================
// Job configuration
//
struct G4Config {
  Atom<double> energyCutoff { Key("energyCutoff") };
};

struct Parameters {
  Atom<std::string> oneAtom  { Key("oneAtom")  };
  Atom<int>         value    { Key("value"), 8 };
  Sequence<int>     list     { Key("list")     };
  Table<G4Config>   g4config { Key("g4Config") };

};
```

Print description of expected parameters

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art --module-description MyModule
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//================================================
// Job configuration
//
struct G4Config {
  Atom<double> energyCutoff { Key("energyCutoff") };
};

struct Parameters {
  Atom<std::string> oneAtom  { Key("oneAtom")  };
  Atom<int>         value    { Key("value"), 8 };
  Sequence<int>     list     { Key("list")     };
  Table<G4Config>   g4config { Key("g4Config") };

};
```

```
pset : {

   oneAtom : <string>
   value : 8  # dflt
   list : [ <int>, <int>, ... ]
   g4Config : {

      energyCutoff : <double>
   }
}
```

- This is a significant improvement from where things stand now; but I still don't know what the (e.g.) "energyCutoff" is.

Introducing Comment

```
//================================================
// Job configuration
//
struct G4Config {
  Atom<double> energyCutoff { Key("energyCutoff"), Comment("This is a number in units of GeV.\n"
                                                           "Geant4 uses it to interpolate between\n"
                                                           "different physics lists.") };

};

struct Parameters {
  Atom<std::string> oneAtom  { Key("oneAtom")  };
  Atom<int>         value    { Key("value"), 8 };
  Sequence<int>     list     { Key("list")     };
  Table<G4Config>   g4config { Key("g4Config") };

};
```

`art --module-description MyModule` yields …

Introducing Comment

```
//================================================
// Job configuration
//
struct G4Config {
  Atom<double> energyCutoff { Key("energyCutoff"), Comment("This is a number in units of GeV.\n"
                                                           "Geant4 uses it to interpolate between\n"
                                                           "different physics lists.") };

};

struct Parameters {
  Atom<std::string> oneAtom  { Key("oneAt
  Atom<int>         value    { Key("value"
  Sequence<int>     list     { Key("list"
  Table<G4Config>   g4config { Key("g4Con

};
```

```
art --module-c
```

```
pset : {

   oneAtom : <string>
   value : 8  # dflt
   list : [ <int>, <int>, ... ]
   g4Config : {


      # This is a number in units of GeV.
      # Geant4 uses it to interpolate between
      # different physics lists.
      energyCutoff : <double>
   }
}
```

Accessing the elements

- After the *.fcl file has been validated, the module is constructed and users can access the elements.

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- Reminder of our module:

```
namespace {

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  // Job configuration
  //
  struct G4Config {
    Atom<double> energyCutoff { Key("energyCutoff") };
  };

  struct Parameters {
    Atom<std::string> oneAtom  { Key("oneAtom")  };
    Atom<int>         value    { Key("value"), 8 };
    Sequence<int>     list     { Key("list")     };
    Table<G4Config>   g4config { Key("g4Config") };

  };

  //================================================
  // Module declaration
  //
  class MyModule : public art::EDProducer {
  public:

    using Parameters = ::Parameters;

    MyModule(Table<Parameters> const & pset) {

    }

  };

}
```

Accessing the elements

- No more `pset.get<>`.
  - Accessing elements is now done using syntax very similar to FHiCL syntax.

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

   g4Config : {
      energyCutoff : 27
   }

}
```

```
MyModule(Table<Parameters> const & pset) {
  std::string str = pset().oneAtom();
  int     some_int = pset().value();
  double    cutoff = pset().g4config().energyCutoff();



}
```

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  - Accessing elements is now done using syntax very similar to FHiCL syntax.

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

   g4Config : {
      energyCutoff : 27
   }

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```

```
MyModule(Table<Parameters> const & pset) {
  std::string str = pset().oneAtom();
  int     some_int = pset().value();
  double    cutoff = pset().g4config().energyCutoff();

  std::cout << str      << std::endl;
  std::cout << some_int << std::endl;
  std::cout << cutoff   << std::endl;
}
```

```
g-2
7
27
```

What about sequences?

- No more `pset.get<>`.
  - Accessing elements is now done using syntax very similar to FHiCL syntax.

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

   g4Config : {
      energyCutoff : 27
   }

}
```

```
MyModule(Table<Parameters> const & pset) {
  std::string str = pset().oneAtom();
  int    some_int = pset().value();
  double   cutoff = pset().g4config().energyCutoff();

  std::vector<int> nums = pset().list();

  // Second element of list
  int okay   = pset().list()[1];
  int better = pset().list(1);
}
```

What about sequences?

- No more `pset.get<>`.
  - Accessing elements is now done using syntax very similar to FHiCL syntax.

```
pset: {

   oneAtom : "g-2"
   value   : 7
   list    : [1,3,17]

   g4Config : {
      energyCutoff : 27
   }

}
```

```
MyModule(Table<Parameters> const & pset) {
  std::string str = pset().oneAtom();
  int    some_int = pset().value();
  double   cutoff = pset().g4config().energyCutoff();

  std::vector<int> nums = pset().list();

  // Second element of list
  int okay   = pset().list()[1]; std::cout << okay   << std::endl; 3
  int better = pset().list(1);   std::cout << better << std::endl; 3
}
```

Features not yet implemented

- Specialized conversions
  - Sometimes you want to read in (e.g.) a sequence as a different kind of structure:

```
pset.get<CLHEP::HepLorentzVector>("lvec");
```

  - Takes 4-element sequence of doubles.

- Will implement specialized versions (e.g.):

```
SequenceAs<CLHEP::HepLorentzVector,double,4>
TupleAs<SomeClass,double,string>
```

Features not yet implemented

- Conditional configuration (e.g.)

```
std::string const& shape = pset.get<string>("shape");

if ( shape == "box") {
  makeBox( pset.get< array<double,3> >("halfLengths") );
{
else if (shape == "sphere") {
  makeSphere( pset.get<double>("radius") );
}
```

- Are working to implement something to support this. May not be in place by Aug. 1.

Please give us your input!

- To be released Aug. 1, 2015.
- This is meant to be a help for users of *art*.
- We've made significant progress, but we want your thoughts.
  - knoepfel@fnal.gov , or
  - artists@fnal.gov
- Thanks!