

Geometry Access in DANA

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Issue: How does one access geometry information in the reconstruction code without hardwiring it?

DGeometry class

- Provides an interface to the underlying *JGeometry* object
- A single *DGeometry* object is used for all threads/factories needing to access the geometry it represents
- Multiple *DGeometry* objects can exist in a single process if it is analyzing events corresponding to different geometries.
- Provides both a ***low-level*** and a ***high-level*** API for accessing the geometry info.

Telling *JANA* where the geometry information is stored

- Set the *JANA_GEOMETRY_URL* environment variable to the location of the geometry:

```
setenv JANA_GEOMETRY_URL xmlfile:///${HALLD_HOME}/src/programs/Simulation/hdds/main_HDDS.xml
```

- URL specifies top-level XML file. Other XML files that are referenced in it are automatically read in by the underlying parser (Xerces)
- Using a URL in this way allows the possibility to change the mechanism through which the geometry is obtained in the future (e.g. http, database, ...)

Getting a pointer to *DGeometry*

- A pointer is obtained through the *DApplication* object:

```
47 jerror_t DX_factory::brun(JEventLoop *loop, int runnumber)
48 {
49     DApplication *dapp = dynamic_cast<DApplication*>(loop->GetJApplication());
50     const DGeometry *dgeom = dapp->GetDGeometry(runnumber);
51
52     // ... use dgeom ...
```

Low-level API: using *xpath**

- *xpath* is a commonly used syntax for specifying a specific tag or attribute in an XML string.

```
<A>
  <B type="3">
    <C id="1"></C>
  </B>
  <B type="4">
    <C id="2"></C>
  </B>
</A>
```

The following *xpath* can be used to specify the “type” attribute of the tag=“B” node that has a subtag “C” with an “id” attribute equal to “2” in the XML snippet on the left.

```
/A/B/@type/C[@id="2"]
```

*Full *xpath 1.0* specification not supported. Only an *xpath*-like syntax for now

A more practical example

xpath for extracting radiation length of Oxygen gas from the XML below

```
//materials/element[@name='Oxygen']/real[@name='radlen']/@value
```

```
<materials version="3.0"
  date="2006-11-22"
  author="R.T. Jones"
  specification="v1.0" >

  <element name="Oxygen" symbol="O" z="8" a="15.9995">
    <real name="density" value="0.00133" unit="g/cm^3" />
    <real name="radlen" value="34.24" unit="g/cm^2" />
    <real name="collen" value="63.2" unit="g/cm^2" />
    <real name="abslen" value="91.0" unit="g/cm^2" />
    <real name="dedx" value="1.801" unit="MeV/g/cm^2" />
  </element>

</materials>
```

Constructing an appropriate *xpath*

- The *JANA* tool *jgeomread* can be used to both 1.)test *xpaths* and 2.)provide guidance as to what *xpaths* are available in the current XML string.
- To get a list of **all** *xpaths*, use the “-L” argument.
- To include **all** attributes in the listing, use the “-a” argument

```
fwing-dhcp13:~> jgeomread -L -a | grep "Oxygen" | grep "radlen"  
/HDDS[@specification='v1.0' and @xmlns='http://www.gluex.org/hdds' and @xmlns:xsi='http://www.w3.org/  
2001/XMLSchema-instance' and @xsi:schemaLocation='http://www.gluex.org/hdds HDDS-1_0.xsd']/  
Material_s/materials[@author='R.T. Jones' and @date='2006-11-22' and @specification='v1.0' and  
@version='3.0']/element[@a='15.9995' and @name='Oxygen' and @symbol='O' and @z='8']/real  
[@name='radlen' and @unit='g/cm^2' and @value='34.24']
```

```
//materials/element[@name='Oxygen']/real[@name='radlen']/@value
```


Low-level methods in *DGeometry*

- Access is similar to other *JANA* access methods, using a *Get(...)* method
- First argument is an *xpath* specifying the desired attribute(s).
- Second argument is reference to container to hold result

```
56 bool Get(string xpath, string &sval) const;  
57 bool Get(string xpath, map<string, string> &svals) const;  
58 template<class T> bool Get(string xpath, T &sval) const;  
59 template<class T> bool Get(string xpath, vector<T> &svals, string delimiter=" ") const;  
60 template<class T> bool Get(string xpath, map<string,T> &svals) const;
```

Example using low-level API

To get the atomic mass of Antimony, one could do the following:

```
63 double a;  
64 dgeom->Get(" '//hdds:element[@name='Antimony']/@a", a);  
65 // a now equals 121.75
```

To get all of the attributes of Antimony, one could do the following:

```
67 map<string,string> attr;  
68 dgeom->Get(" '//hdds:element[@name='Antimony']", attr);  
69 // attr now contains the following entries:  
70 // a      121.75  
71 // name   Antimony  
72 // symbol Sb  
73 // z      51
```

High-level API using *DGeometry*

DGeometry contains several “convenience” methods. Most users will want to use these.

```
68 bool GetFDCZ(vector<double> &z_wires) const; ///< z-locations for each of the FDC wire planes in cm
69 bool GetFDCStereo(vector<double> &stereo_angles) const; ///< stereo angles of each of the FDC wire layers
70 bool GetFDCRmin(vector<double> &rmin_packages) const; ///< beam hole size for each FDC package in cm
71 bool GetFDCRmax(double &rmax_active_fdc) const; ///< outer radius of FDC active area in cm
72
73 bool GetCDCOption(string &cdc_option) const; ///< get the centralDC_option-X string
74 bool GetCDCCenterZ(double &cdc_center_z) const; ///< z-location of center of CDC wires in cm
75 bool GetCDCAxialLength(double &cdc_axial_length) const; ///< length of CDC axial wires in cm
76 bool GetCDCStereo(vector<double> &cdc_stereo) const; ///< stereo angle for each CDC layer in degrees
77 bool GetCDCRmid(vector<double> &cdc_rmid) const; ///< Distance of the center of CDC wire from beamline for
78 bool GetCDCNwires(vector<int> &cdc_nwires) const; ///< Number of wires for each CDC layer
79 bool GetCDCEndplate(double &z,double &dz,double &rmin,double &rmax) const;
80
81 bool GetBCALRmin(double &bc_al_rmin) const; ///< minimum distance of BCAL module from beam line
82 bool GetBCALNmodules(unsigned int &bc_al_nmodules) const; ///< Number of BCAL modules
83 bool GetBCALCenterZ(double &bc_al_center_z) const; ///< z-location of center of BCAL module in cm
84 bool GetBCALLength(double &bc_al_length) const; ///< length of BCAL module in cm
85 bool GetBCALDepth(double &bc_al_depth) const; ///< depth (or height) of BCAL module in cm
```

Documentation on the wiki

A HOWTO describing all of this in more detail
can be found on the web at:

[http://www.jlab.org/Hall-D/software/wiki/index.php/
HOWTO_Access_Geometry_Information_from_JANA/
DANA](http://www.jlab.org/Hall-D/software/wiki/index.php/HOWTO_Access_Geometry_Information_from_JANA/DANA)

i.e.

GlueX wiki: Offline Software->HOWTO Access Geometry Information from JANA/DANA