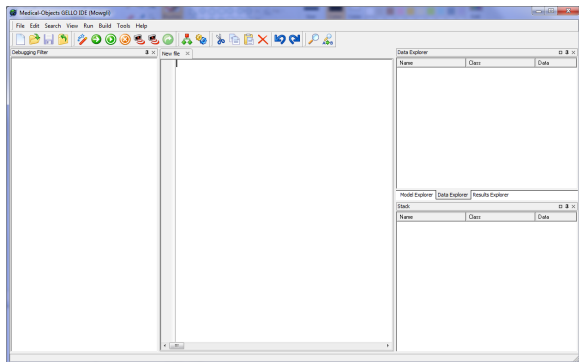


Introduction to the Medical Objects GELLO editor and first code generation

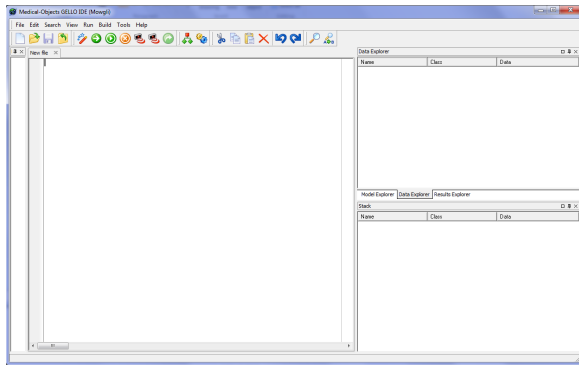
Introduction to the editor

- 1 [Introduction to the editor](#)
- 2 [Writing some code](#)

Put the downloaded file *mowgli_11.exe* in a folder called 'Gello' for example in C: drive. You can make a shortcut to it if you like, or pin it to the taskbar. Run it and you should have this:



On the left hand side (LHS) there is a debugging filter which we won't use so minimise up that window by adjusting the pane size, and make some more room for the RHS panes as well:



So, looking at the editor now we have a largish workspace area where we will write GELLO code, a top menu bar, and on the RHS, a Model Explorer, a Data Explorer and a Results Explorer (in the tabs of the top RHS pane), and a stack for debugging also. We won't use the Stack viewer very much however the pane above it is very important so give it some more vertical room.

GELLO is a query language for writing computer executable decision support in the health domain. It can access information out of a virtual medical record (VMR). A VMR is an information model that presents what is useful for clinical decision support out of an Electronic Health Record. So this will be things like patient age and gender, their past illnesses, operations, medications, allergies etc. GELLO also can perform logic against a reference terminology such as SNOMED CT which means it can do things like ask if the patient has a respiratory disease, without having to type in all possible codes for all possible respiratory diseases to find out.

Before we get started writing our first piece of code take a look at the Model Explorer tab on RHS. Unfold on **System Packages**:

Model Explorer		
Name	Kind	Detail
Application	package	package Application
System Packages	package	package System Packages
System	package	package System
XMLAny	package	package XMLAny
iso_21090_datatypes	package	package iso_21090_datatypes
HL7_v2_VMR_V1	package	package HL7_v2_VMR_V1

We'll take a look at the bottom two packages. These describe the ISO 21090 datatypes that GELLO uses (GELLO is strongly typed) and the HL7 v2 VMR which is the default VMR model - although other VMRs may be used. Unfold on **iso_21090_datatypes**, and scroll down and you can see a large number of types. We will use standard datatypes such as String, Real, Boolean (These are in fact in the **System** package further up in the last screenshot), and from iso_21090_datatypes - mostly we will use Concept Descriptor (CD), Physical Quantity (PQ), Timestamp (like a datetime) (TS) and the Sequence classes. Codes and code systems get used a lot in health IT and if you unfold the CD datatype class you can see the attributes and operations available to this type:

Model Explorer		
Name	Kind	Detail
BL	class	BL extends ANY
CD	class	CD extends ANY
code	attribute	code: String
codeSystem	attribute	codeSystem: String
codeSystemName	attribute	codeSystemName: String
codeSystemVersion	attribute	codeSystemVersion: String
codingRationale	attribute	codingRationale: String
displayName	attribute	displayName: ST
id	attribute	id: String
implies	operation	implies(other: CD): BL
implies	operation	implies(other: CS): BL
originalText	attribute	originalText: ED
source	attribute	source: XReference
translation	attribute	translation: Sequence(CD)
valueSet	attribute	valueSet: String
valueSetVersion	attribute	valueSetVersion: String

Now fold all that up and have a look at the **HL7_v2_VMR_V1** package. If you are a clinician you will recognise this as a fairly comprehensive summary of clinical information for a patient. Unfolding on Patient class you can see patients have both gender and date of birth, which are in turn of datatypes CD and TS respectively.

Writing some code

So lets write our first GELLO. It simply will be querying for a patients gender.

GELLO can be used in different contexts. Here we will be looking at a single patient at a time. So in the workspace write:

```
Context SinglePatient
```



and hit the **Compile** button.

What happens? Yes there should be a couple of error messages in a new bottom-most pane. The first message is saying it has no idea what you are talking about with the word "SinglePatient" and wants us to specify the model package that can define it. The best way to do this is to get in the habit of adding a line now *above* the one we have that says:

```
imports HL7_v2_VMR_V1
```

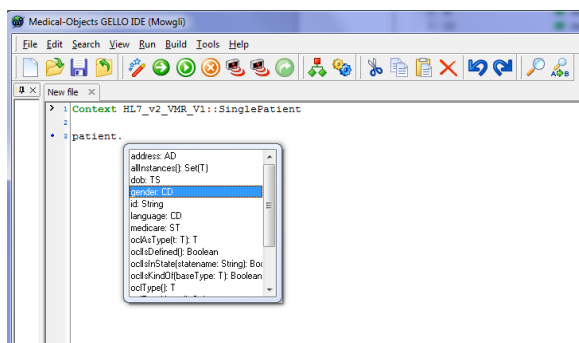
Compile that and we are left with the other error message that says in effect , "would you state what it is that you want the code to give you back" (GELLO is declarative). That's ok for now and this error does not stop things from working.

So what we want is the gender of the patient.

Below this put in a blank line and then type:

```
patient.
```

and pause - Code autocomplete will give us some hints:



Choose the gender attribute. [Notice anything else about the code in general in this screenshot ? - yes, we can collapse the top two lines into one single line that specifies the imported package and the context.]



Now compile - all good. Ok we choose the **Run** button . [NB - Not the one on the left of this that we can use to step through the code for debugging). The normal procedure is to compile code so that it is syntactically correct, and then run it against some test clinical data.

Ok, it wants some test data. Here is a small xml file for us to get started with. Cut and past the following text into a text editor such as notepad++ and save as *firstTest.xml* . Then when we run the GELLO code, navigate to where you have this file.

```
<?xml version="1.0" encoding="UTF-8"?>

<singlePatient xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
  xsi:noNamespaceSchemaLocation="/iso-21090-datatypes.xsd">

  <patient>

    <patientID

      root="1.2.36.174030967"

      extension="1234567892"

      reliability="VRF"

      scope="OBJ"/>

    <patientName>

      <part type="GIV" value="Alice" />

      <part type="GIV" value="A." />

      <part type="FAM" value="Someone" />

    </patientName>

    <dob value="19550621"/>

    <gender code="248152002"

      codeSystem="2.16.840.1.113883.6.96"

      codeSystemName="SNOMED-CT">

        <displayName value="Female" />

      </gender>

    <ssn value="987-65-4320"/>

    <address use="WP">

      <part type="AL" value="1050 W Wishard Blvd" />

    </address>

  </patient>

</singlePatient>
```

```
<part type="CTY" value="Indianapolis" />

<part type="STA" value="IN" />

<part type="ZIP" value="46240" />

</address>

</patient>

</singlePatient>
```

And here is the result - we are looking at top RHS of the editor screen:

Results Explorer		
Name	Class	Data
[-] _Context_	SinglePatient	<SinglePatient: TSinglePatient>
[-] Result_	CD	Female

Model Explorer | Data Explorer | Results Explorer

That's it for the first tutorial - well done!

Next time we will write more GELLO code using the date of birth; see how to set some test data in place for a GELLO editing session, and introduce the implies method to check for an allergy.