# How to add a tree prior to BEAST2?

This tutorial follows the same wording and structure as BEAST2: how to write a BEAST2 package, but for a tree prior instead.

### **Empty Package Explorer**

When starting Eclipse in a new workspace, the Package Explorer is still empty. In this example, my workspace is ~/GitHubs/Java.

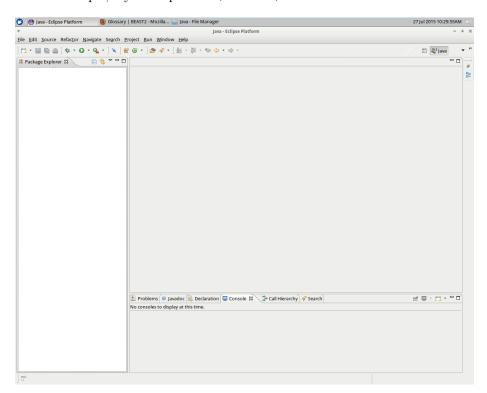


Figure 1: Eclipse with an empty package explorer

### Adding BEAST2 to the Package Explorer

To add BEAST2 to the Package Explorer, it must first be downloaded, then put into a Java project.

#### Download BEAST2

In this example, I download BEAST2 from the SVN repository from withing the '~/GitHubs/Java' folder:

cd ~/GitHubs/Java svn checkout http://beast2.googlecode.com/svn/trunk/beast2 This will create a folder called beast2.

```
Terminal
                                                                         - + ×
File Edit View Terminal Tabs Help
p230198@fwn-biol-132-102:~$ cd ~/GitHubs/Java/
p230198@fwn-biol-132-102:~/GitHubs/Java$ svn checkout http://beast2.googlecode.c
om/svn/trunk/ beast2
    beast2/release
beast2/release/Linux
    beast2/release/Linux/bin
    beast2/release/Linux/bin/logcombiner
    beast2/release/Linux/bin/addonmanager
    beast2/release/Linux/bin/densitree
    beast2/release/Linux/bin/treeannotator
    beast2/release/Linux/bin/beauti
    beast2/release/Linux/bin/beast
     beast2/release/Mac
     beast2/release/Mac/createdmg.sh
    beast2/release/Mac/install.png
     beast2/release/common
     beast2/release/common/LICENSE.txt
     beast2/release/common/tools
```

Figure 2: Start downloading BEAST2

#### Put BEAST2 into a Java project

First, we will add beast2 to the Package Explorer.

Select File | New | Java Project:

This is how the 'Create a Java Project dialog' looks like:

Just type in the word beast. It will detect there is a folder called beast2 and the dialog will change:

Just press Finish.

Now, the Eclipse Package Explorer shows the project beast2. Well done!

```
Terminal
File Edit View Terminal Tabs Help
     beast2/examples/beast2vs1/testStrictClock2.xml
beast2/examples/beast2vs1/testSRD06CP12_3.xml
beast2/examples/beast2vs1/testRandomLocalClock.xml
     beast2/examples/beast2vs1/testCoalescentTipDatesSampling.xml
     beast2/examples/beast2vs1/testCoalescentNoClock1.xml beast2/examples/beast2vs1/testBirthDeathModel_10taxa.xml
      beast2/examples/beast2vs1/testTipDates.xml
      beast2/examples/beast2vs1/testStrictClockTipDatesSampling.xml
     beast2/examples/beast2vs1/testYuleModel_10taxa.xml
beast2/examples/beast2vs1/testCoalescentTipDates.xml
      beast2/examples/testSliceHKY.xml
     beast2/examples/starbeastinit
      beast2/examples/starbeastinit/sbi-01.xml
     beast2/examples/starbeastinit/sbi-02.xml
      beast2/examples/starbeastinit/sbi-03.xml
     beast2/examples/testTipDates.xml
beast2/templates
     beast2/templates/Standard.xml
     beast2/templates/SubstModels.xml
     beast2/templates/StarBeast.xml
beast2/test.xml
     beast2
Checked out revision 1068.
p230198@fwn-biol-132-102:~/GitHubs/Java$
```

Figure 3: End of downloading BEAST2

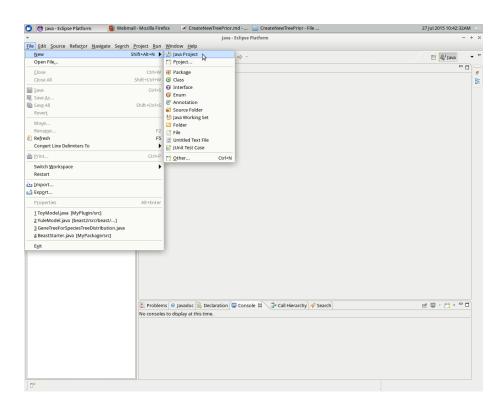


Figure 4: Start a new Java Project

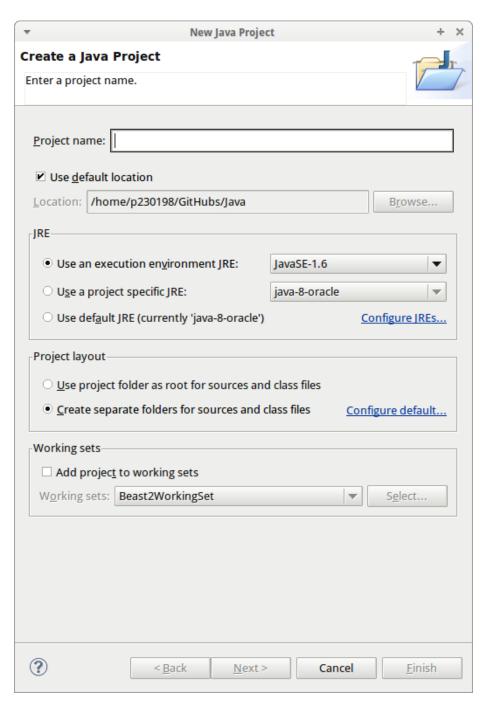


Figure 5: Create a Java Project

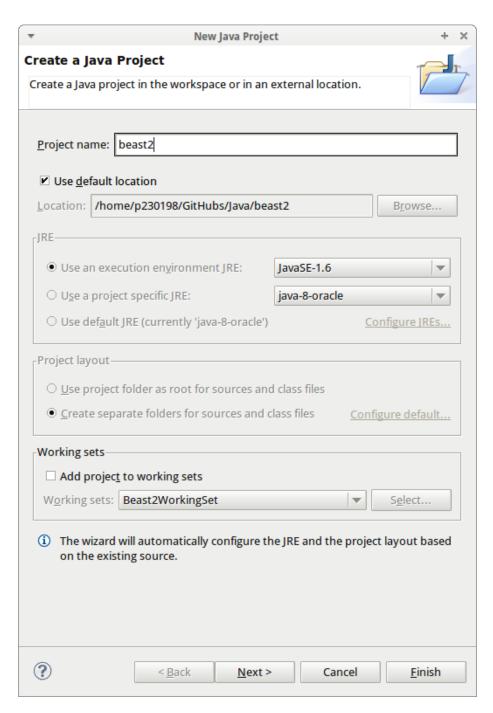


Figure 6: Create a BEAST2 Java Project

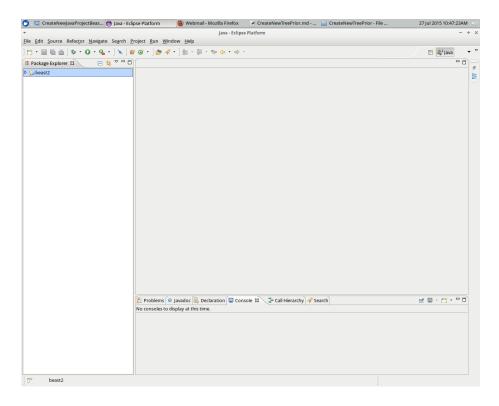


Figure 7: Eclipse with BEAST2 in the package explorer

### Creating your own project

Now you need to create an Eclipse project for your package and make it depend on the beast2 project.

Select File | New | Java Project:

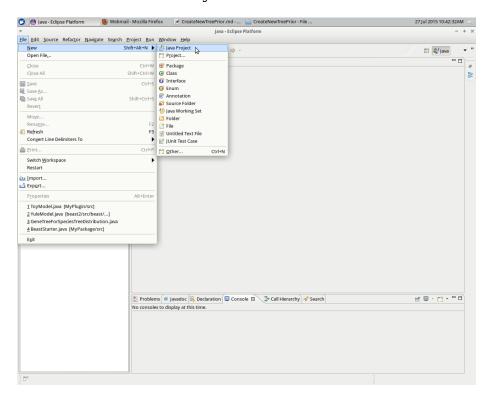


Figure 8: Start a new Java Project

In Project name, type in the name, for example MyProject. For the JRE, be sure JavaSE-1.6 or something higher is checked. Click Finish.

Now you should see two projects in the Package Explorer pane: beast2 and MyProject.

## Add BEAST2 to your own project

Right-click MyProject and choose Properties from the pop-up menu that appears

Go to Java build path | Projects and click Add.

Add beast2 to your project build path:

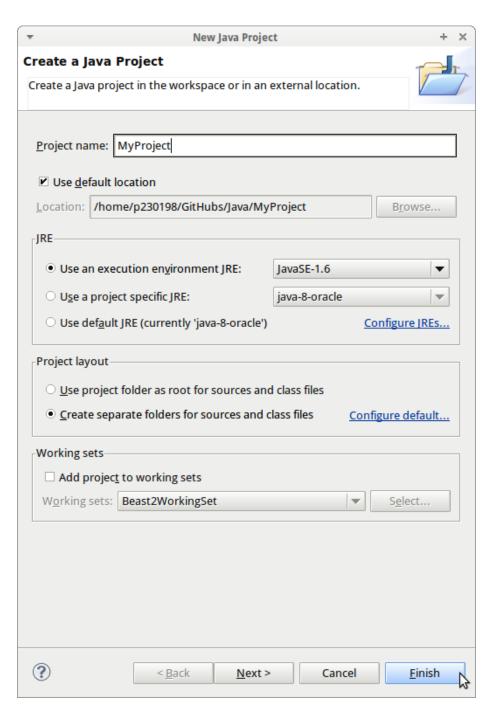


Figure 9: Create a Java project: MyProject

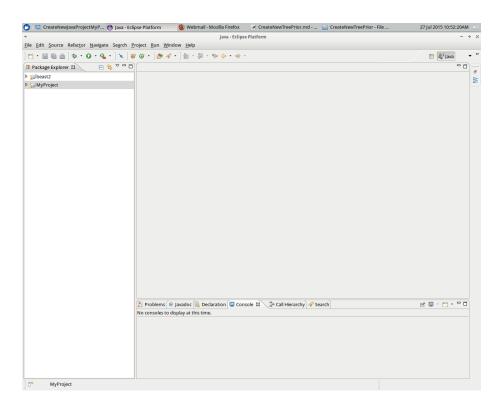


Figure 10: Package Explorer with BEAST2 and MyProject

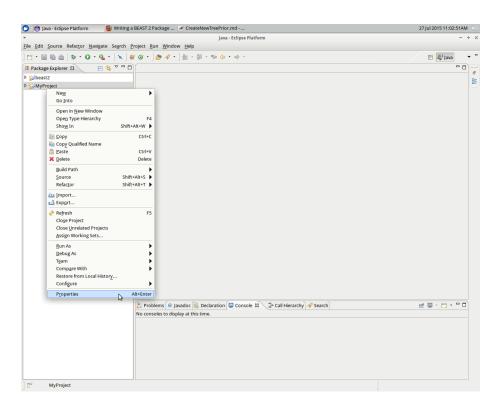


Figure 11: Modify the project properties

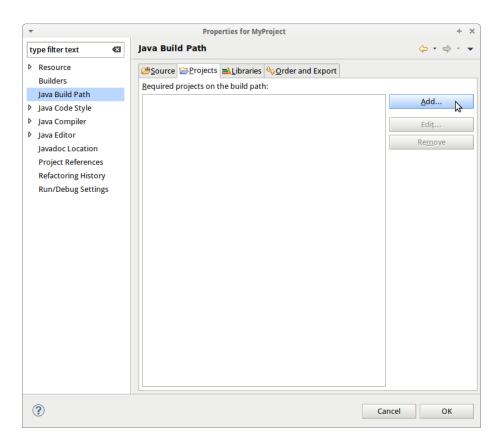


Figure 12: Adding a build path to MyProject

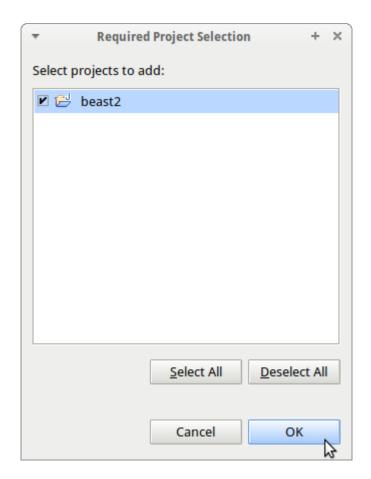


Figure 13: Adding BEAST2 to your project build path

Properties for MyProject + × type filter text Java Build Path ▶ Resource <u>B</u>ource <u>Projects</u> <u>Libraries</u> <u>Order and Export</u> Builders Required projects on the build path: Java Build Path ▷ 🗁 beast2 Add. ▶ Java Code Style Java Compiler Edit ... Java Editor Re<u>m</u>ove Javadoc Location Project References Refactoring History Run/Debug Settings ? OK N Cancel

Now BEAST2 is in your project build path, click OK.

Figure 14: Added BEAST2 to your project build path

This takes you back to the Package Explorer pane.

#### Create a new tree prior

To create a new tree prior, I will use the Yule model already present in BEAST2.

Select MyProject in the Package Explorer, right-click and select 'New | Class'

Change the class name to MyTreePrior. Ignore all the warnings (as we won't use the generated code anyways) and click Finish.

Now you will see the initial code of MyTreePrior:

This initial code, however, will be replaced soon.

I will replace the initial code by the Yule tree prior. In the Package Explorer, find the Yule tree prior code. It is under  $beast2 \mid src \mid beast.evolution.speciation$ .

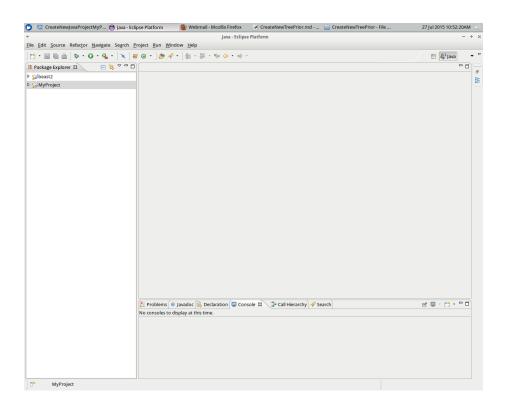


Figure 15: Package Explorer with BEAST2 and MyProject

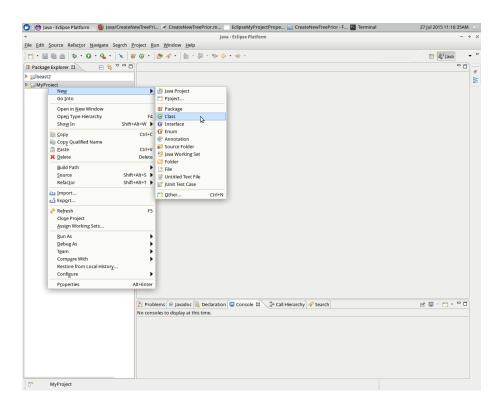


Figure 16: Create a new class

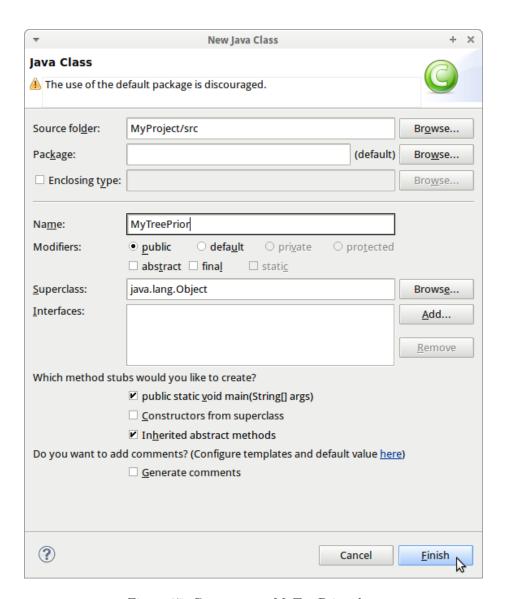


Figure 17: Create a new MyTreePrior class

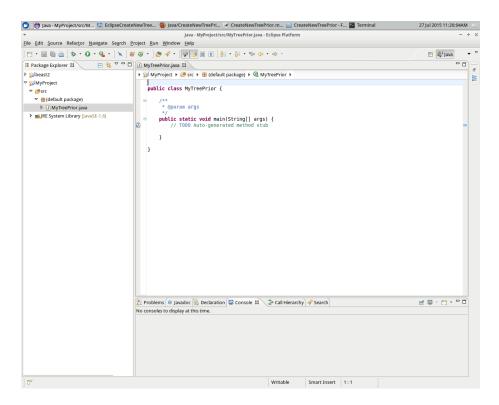


Figure 18: Initial code of MyTreePrior

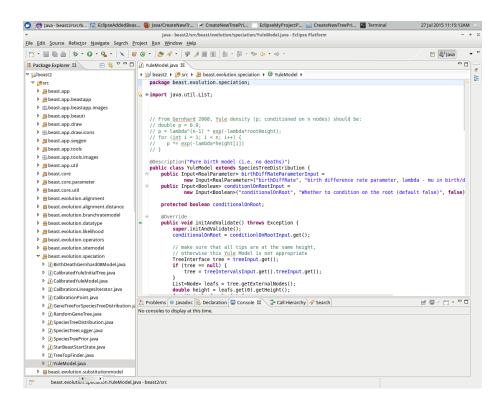


Figure 19: Yule model code

Copy-paste the source code into the MyTreePrior code. A lot of errors pop up that we'll fix now.

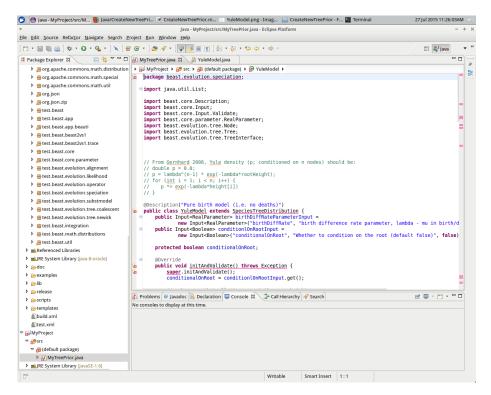


Figure 20: Copied Yule model code

The first error suggests to put the class in the appropriate package.

Double-click this fix.

The next error is that YuleModel is not the correct type and Eclipse suggest to rename it to MyTreePrior instead.

Double-click this fix. Our class now has the correct (new) class name.

#### Add the tree prior to BEAUti

To allow BEAUti to find the new tree prior, a template XML files needs to be created. Because MyTreePrior is a copy of YuleModel, we will use the XML code of that model as a starting point.

In the MyProject folder, you will need to create a templates folder. I just copied ~/GitHubs/Java/beast2/templates to ~/GitHubs/Java/MyProject/templates.

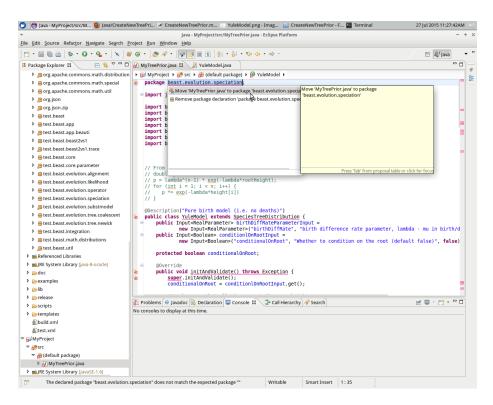


Figure 21: Suggest to move to package

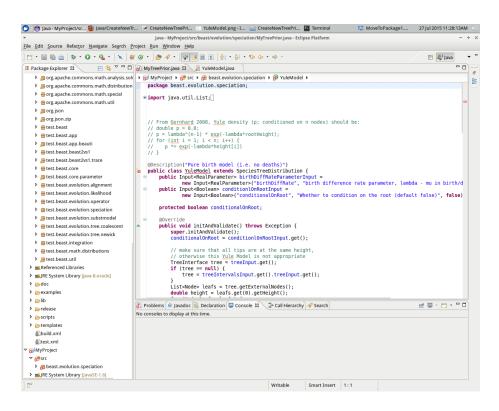


Figure 22: Yule model copy in correct package

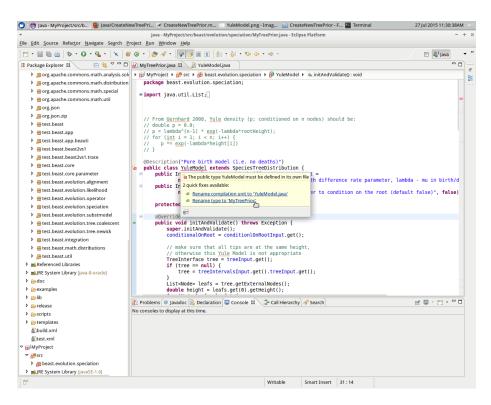


Figure 23: Eclipse suggests to rename the type

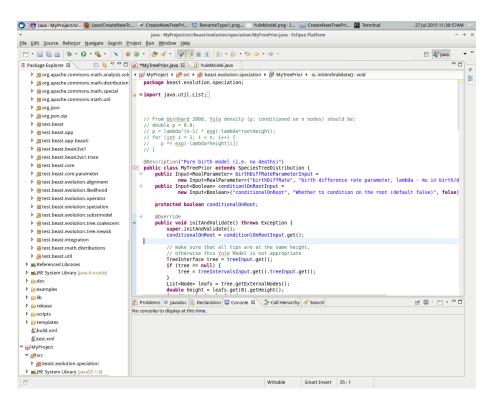


Figure 24: Eclipse renamed the type

In that folder, only keep SubstModels.xml and rename it to MyTreePrior.xml. Then edit MyTreePrior.xml. You can find YuleModel somewhere in that file. It will be the only one to keep.

```
🐧 😝 Java - MyProjec... 🧕 Java/CreateNe... 🎅 CreateNewTree... 🌠 MyTreePrior.x... 🕍 RenameType1.... 🗆 YuleModel.png ... 🗀 templates - File ... 🖪 Terminal
<u>File Edit View Text Document Navigation Help</u>
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           <!-- tree priors -->
<mergewith point='treePriorTemplates'>
               <sintemplate id='YuleModel' class='beast.evolution.speciation.YuleModel' mainid='YuleModel.t:$(n)'>
    <![CDATA[</pre>
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                    <prior id='YuleBirthRatePrior.t:$(n)' x='@birthRate.t:$(n)'><distr spec="beast.math.distributions.U"</pre>
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596 ]]>
                  <scale id='YuleBirthRateScaler.t:$(n)' spec='ScaleOperator' scaleFactor="0.75" weight="3" parameter='</pre>
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                     </p
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                     <connect srcID='YuleBirthRatePrior.t:$(n)' targetID='prior' inputName='distribution'
    if='inposterior(YuleModel.t:$(n)) and inposterior(Tree.t:$(n)) and birthRate.t:$(n)/estima
    Yule speciation process birth rate of partition t:$(n)</pre>
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                    <connect srcID='YuleModel.t:$(n)' targetID='tracelog' inputName='log'
    if='inposterior(YuleModel.t:$(n)) and inposterior(Tree.t:$(n)) and Tree.t:$(n)/estimate=tr
<connect srcID='birthRate.t:$(n)' targetID='tracelog' inputName='log'
    if='inposterior(YuleModel.t:$(n)) and inposterior(Tree.t:$(n)) and birthRate.t:$(n)/estima</pre>
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                </subtemplate>
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               <!-- Calibrated Yule -->
<cubtemplate_id='CalibratedYuleModel'_class='heast_evolution_speciation_CalibratedYuleModel</pre>
× Fi<u>n</u>d: Yule
                         ♦ Next ♠ Previous 📳 Highlight All 🗷 Match Case
                                                                                                      Filetype; XML Line: 617 Column: 22 Selection: 2223 OVR
```

Figure 25: MyTreePrior.xml before edit

Remove all other things, except for YuleModel. Be sure to keep the closing XML tags. Or just copy paste this code:

```
<connect srcID='YuleModel.t:$(n)' targetID='prior' inputName='distribution'</pre>
                      if='inposterior(YuleModel.t:$(n)) and inposterior(Tree.t:$(n)) and Tree
                 speciation prior on tree t:$(n)
            </connect>
            <connect srcID='birthRate.t:$(n)' targetID='state' inputName='stateNode'</pre>
                      if='inposterior(YuleModel.t:$(n)) and inposterior(Tree.t:$(n)) and inposterior
            <connect srcID='YuleBirthRatePrior.t:$(n)' targetID='prior' inputName='distribut
</pre>
                      if='inposterior(YuleModel.t:$(n)) and inposterior(Tree.t:$(n)) and bir
                Yule speciation process birth rate of partition t:$(n)
            </connect>
            <connect srcID='YuleBirthRateScaler.t:$(n)' targetID='mcmc' inputName='operator</pre>
                      if='inposterior(YuleModel.t:$(n)) and inposterior(Tree.t:$(n)) and bir
                Scale birth rate of Yule prior of tree t:$(n)
            </connect>
            <connect srcID='YuleModel.t:$(n)' targetID='tracelog' inputName='log'</pre>
                      if='inposterior(YuleModel.t:$(n)) and inposterior(Tree.t:$(n)) and Tree
            <connect srcID='birthRate.t:$(n)' targetID='tracelog' inputName='log'</pre>
                      if='inposterior(YuleModel.t:$(n)) and inposterior(Tree.t:$(n)) and bir
        </subtemplate>
    </mergewith>
</beast>
Some YuleModel occurrances must be replaced by MyTreePrior.
<beast version='2.0'</pre>
       namespace='beast.app.beauti:beast.core:beast.evolution.branchratemodel:beast.evolutio
    <!-- tree priors -->
    <mergewith point='treePriorTemplates'>
        <!-- Yule -->
        <subtemplate id='MyTreePrior' class='beast.evolution.speciation.MyTreePrior' mainid=</pre>
            <! [CDATA [
            <plugin spec='MyTreePrior' id="MyTreePrior.t:$(n)" tree='@Tree.t:$(n)'>
                 <parameter name='birthDiffRate' id="birthRate.t:$(n)" value='1.0'/>
            </plugin>
            <pri>or id='YuleBirthRatePrior.t:$(n)' x='@birthRate.t:$(n)'><distr spec="beast.r"</pre>
```

<scale id='YuleBirthRateScaler.t:\$(n)' spec='ScaleOperator' scaleFactor="0.75" v</pre>

]]>

```
<scale id='YuleBirthRateScaler.t:$(n)' spec='ScaleOperator' scaleFactor="0.75" v</pre>
]]>
            <connect srcID='MyTreePrior.t:$(n)' targetID='prior' inputName='distribution'</pre>
                      if='inposterior(MyTreePrior.t:$(n)) and inposterior(Tree.t:$(n)) and Tr
            </connect>
            <connect srcID='birthRate.t:$(n)' targetID='state' inputName='stateNode'</pre>
                      if='inposterior(MyTreePrior.t:$(n)) and inposterior(Tree.t:$(n)) and in
            <connect srcID='YuleBirthRatePrior.t:$(n)' targetID='prior' inputName='distribut
</pre>
                      if='inposterior(MyTreePrior.t:$(n)) and inposterior(Tree.t:$(n)) and b
                Yule speciation process birth rate of partition t:$(n)
            </connect>
            <connect srcID='YuleBirthRateScaler.t:$(n)' targetID='mcmc' inputName='operator</pre>
                      if='inposterior(MyTreePrior.t:$(n)) and inposterior(Tree.t:$(n)) and b
                Scale birth rate of Yule prior of tree t:$(n)
            </connect>
            <connect srcID='MyTreePrior.t:$(n)' targetID='tracelog' inputName='log'</pre>
                      if='inposterior(MyTreePrior.t:$(n)) and inposterior(Tree.t:$(n)) and Tr
            <connect srcID='birthRate.t:$(n)' targetID='tracelog' inputName='log'</pre>
                      if='inposterior(MyTreePrior.t:$(n)) and inposterior(Tree.t:$(n)) and be
        </subtemplate>
    </mergewith>
```

#### Run BEAUti

</beast>

To create a debug configuration that runs BEAUti, choose  ${\tt Run}$  |  ${\tt Debug}$   ${\tt Configurations}$ .

In the Debug Configurations dialog, select Java Application, right-click, then click New.

- In the Name field at the top of the dialog box, type BEAUti
- In the Project field of the Main tab, type MyProject
- For the Main class, type beast.app.beauti.Beauti

Click Apply.

Click Debug to start BEAUti.

Use File | Import Alignment or click the + button at the lower left, then navigate to the ~/GitHubs/Java/beast2/examples/nexus folder and select the anolis.nex file. Note: you can avoid doing this step if you specify -nex

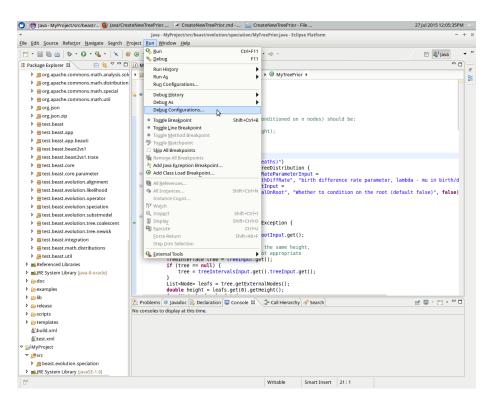


Figure 26: Debug configurations

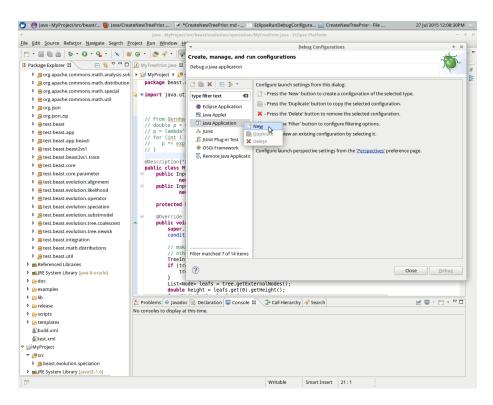


Figure 27: Debug configuration: add new java application

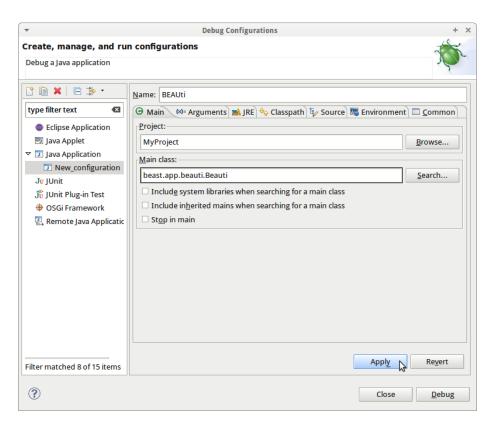


Figure 28: Debug configuration done

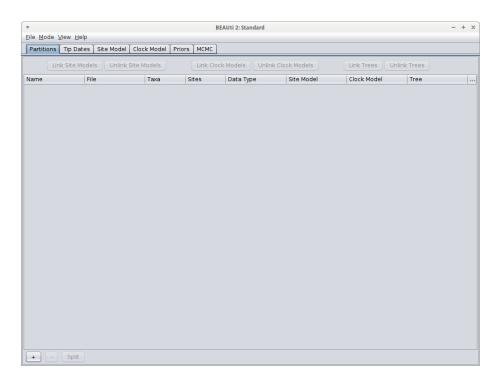


Figure 29: BEAUti2 still empty

../beast2/examples/nexus/anolis.nex in the Program arguments field of the Arguments tab in the BEAUti debug configuration.

Now the alignment is used in BEAUti.

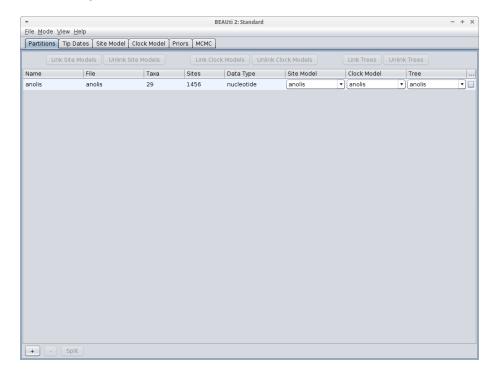


Figure 30: BEAUti2 Anolis

Under the tab Priors you can select the My Tree Prior.

And there it is!

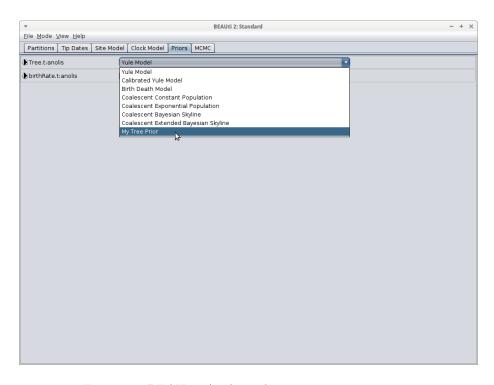


Figure 31: BEAUti2 Anolis and new tree prior is present

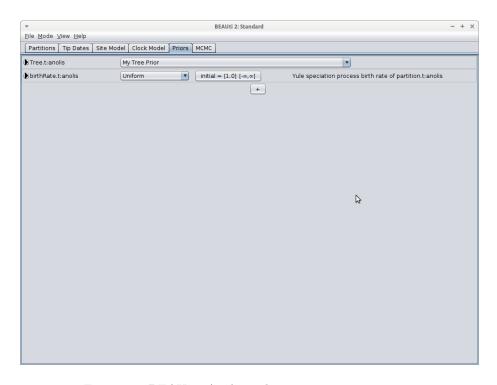


Figure 32: BEAUti2 Anolis and new tree prior is present