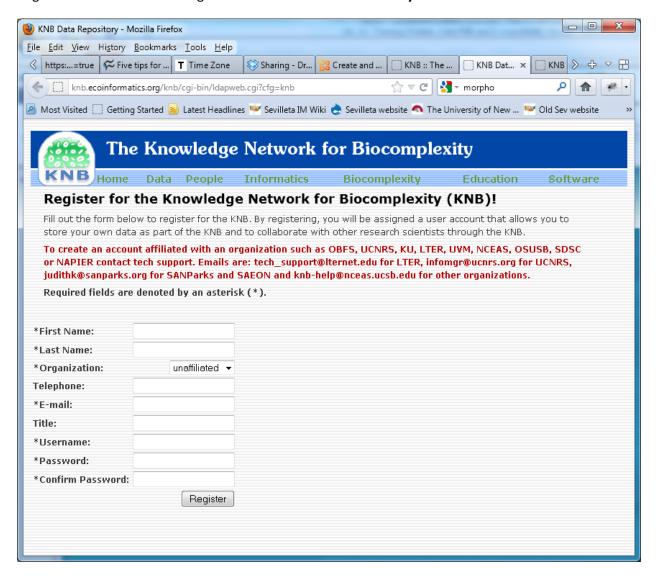
In order to use Morpho, you have to register with the Knowledge Network for Complexity. They created this software.

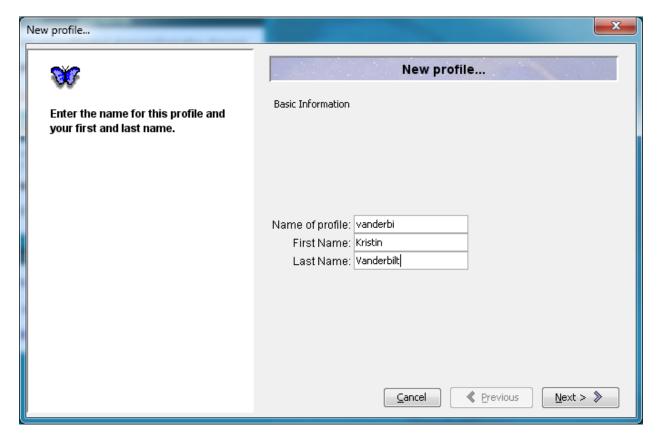
Go to: http://knb.ecoinformatics.org/index.jsp and scroll down to "Login and Registration". Select 'Create a New Account':

Register as an 'unaffiliated' Organization. Be sure to write down your Username and Password!!!

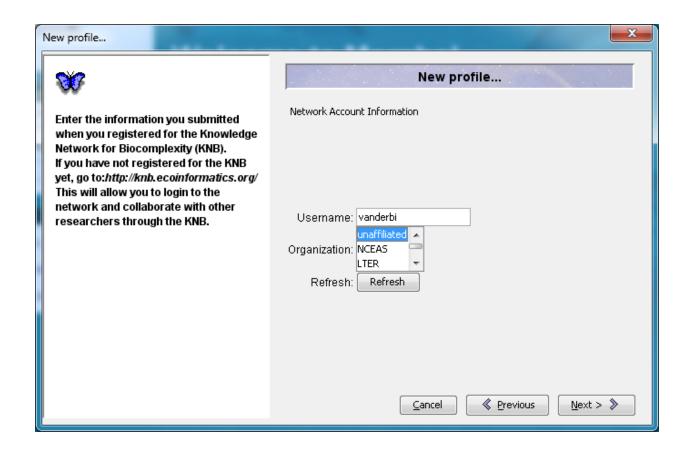


Then, open Morpho.

When you first open Morpho, you will be asked to create a user profile. This profile is used on your local machine to manage your data packages (which can consist of metadata, or data plus metadata). Enter a 'Name of Profile' of your choice (it can be anything), and your First Name and Last Name:



On the next screen, enter the 'Username' you chose when registering for the KNB and 'LTER' for your organization.

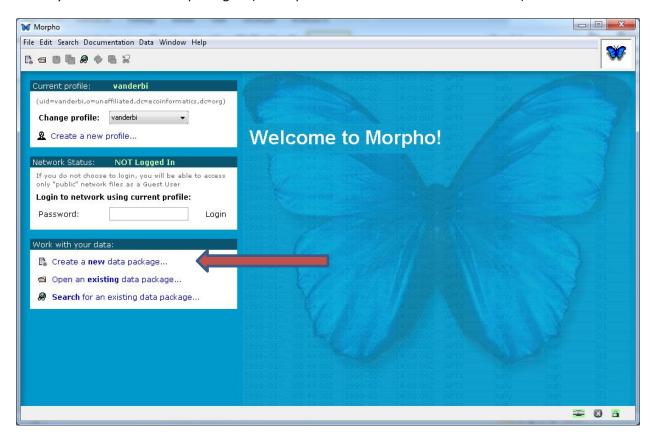


On the next screen, enter an Identifier Prefix for your data packages. This can be anything, and will be used to help name your data packages. You can use your profile name, for instance.

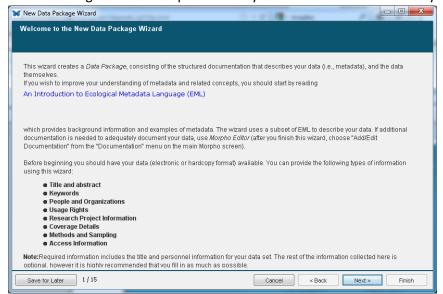


Then click 'Finish', and you will be taken back to this screen:

At this time, we will be working offline, and not connected to the KNB database, so click on the entry that says 'Create a new data package'. (Arrow points to this link in above screenshot).

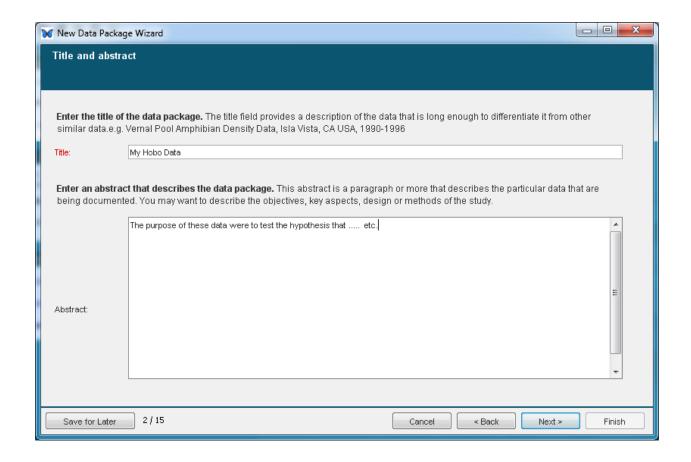


A Data Package wizard will open and tell you what kind of information you will be entering. Click 'Next'.



On this screen, you will enter the title for your data package. The abstract is not a 'required' element, but you should enter an abstract briefly describing your Hobo study.

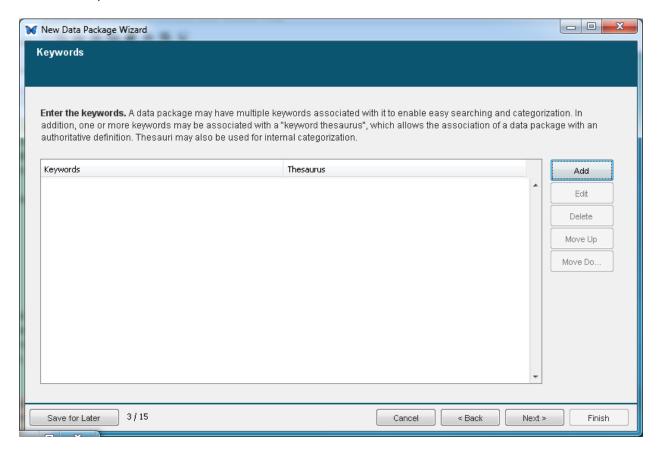
Click 'next'.



Next you will add keywords. Enter a few terms that would help other users of your data find it in a data catalog. Go to the NBII's Biocomplexity Thesaurus and find at least one term that applies:

http://www.usgs.gov/core science systems/csas/biocomplexity thesaurus/index.html

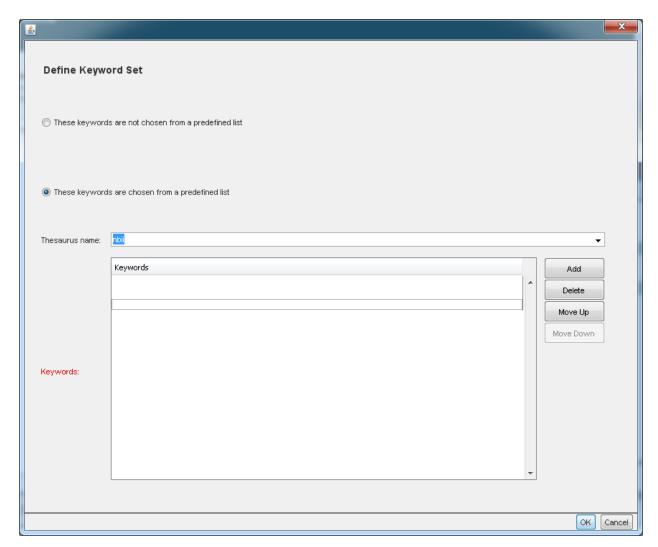
To add terms, click 'Add' on this screen:



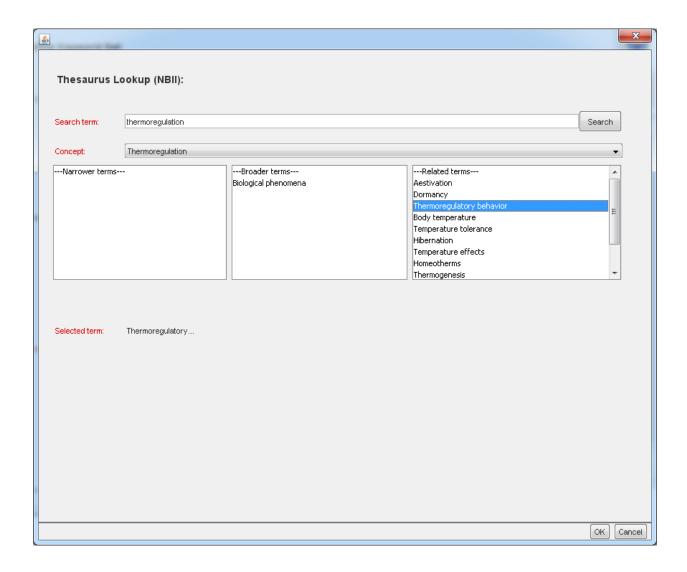
Then this screen will appear:

Select These keywords are chosen from a predefined list. From the dropdown list next to Thesaurus name, choose "nbii"

Thank click 'Add'

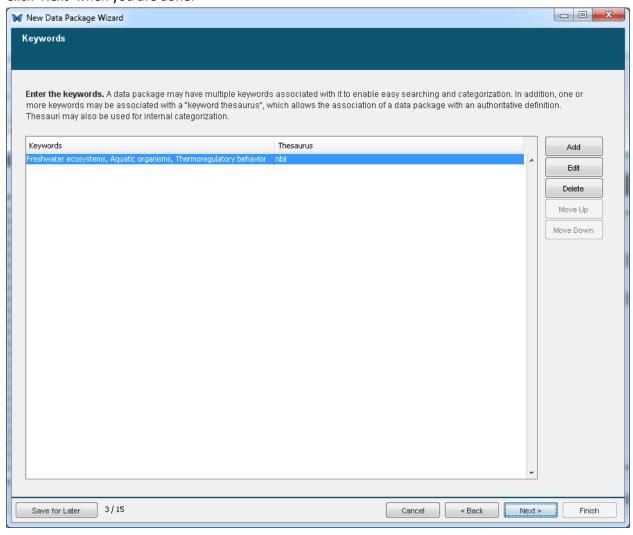


Add three appropriate keywords, and click OK when you are done.

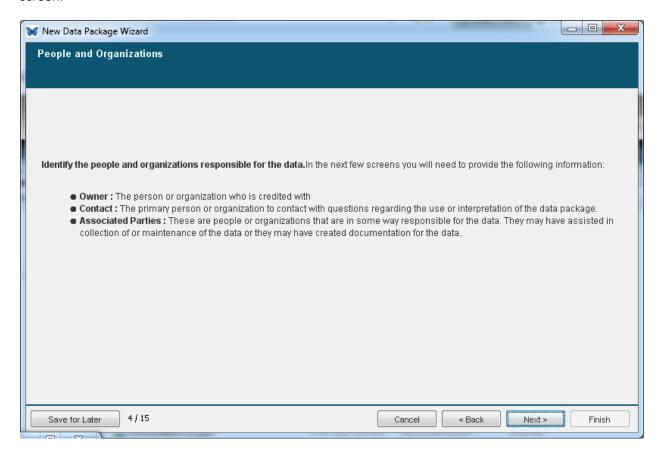


Here you can review or change what keywords you entered:

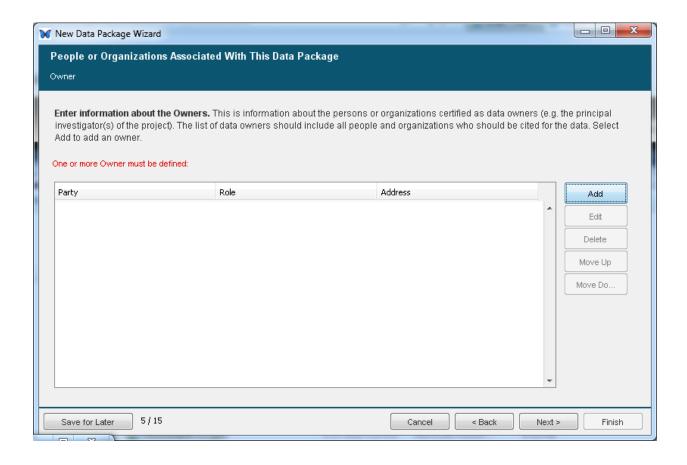
Click 'Next' when you are done:



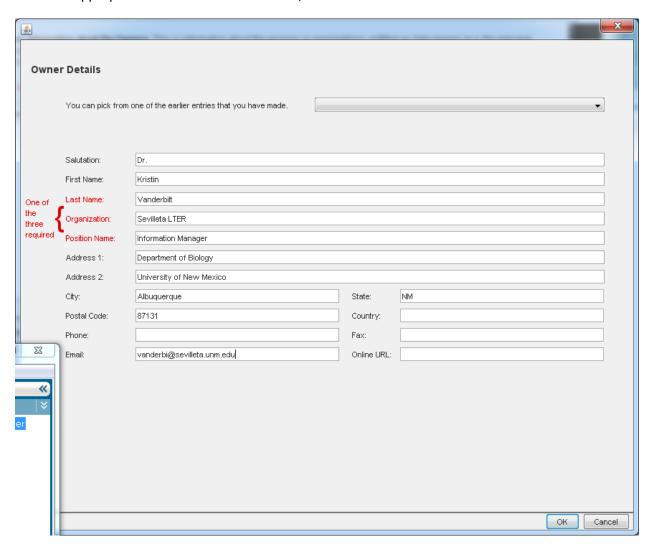
Next, you will enter information about the people associated with your dataset. Click 'Next' on this screen.



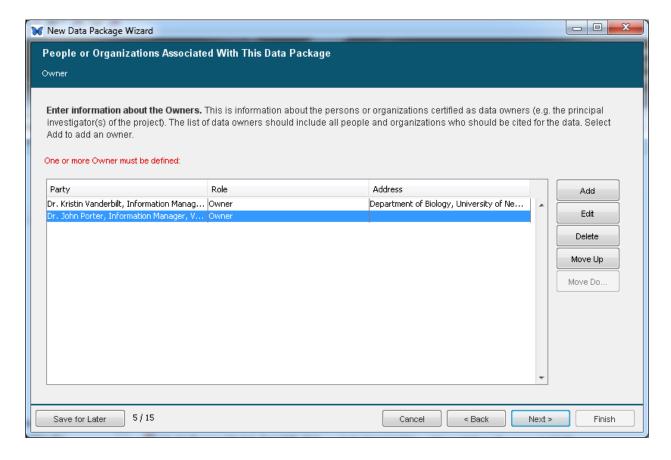
You should be considered Owner (principal investigator) on the project. Click 'Add'



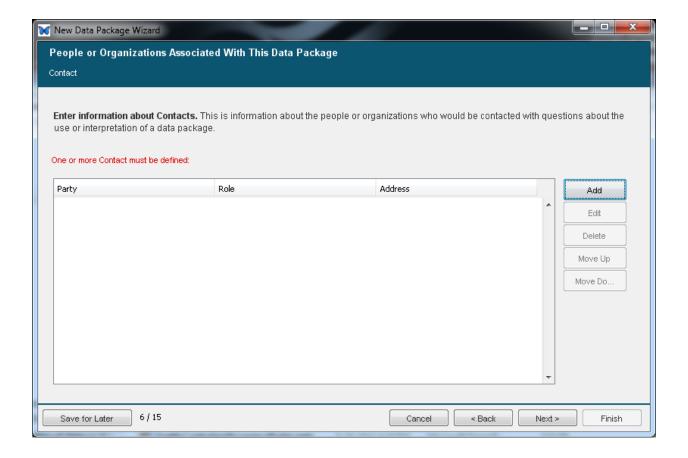
Enter the appropriate information in the screen, and click 'OK'.



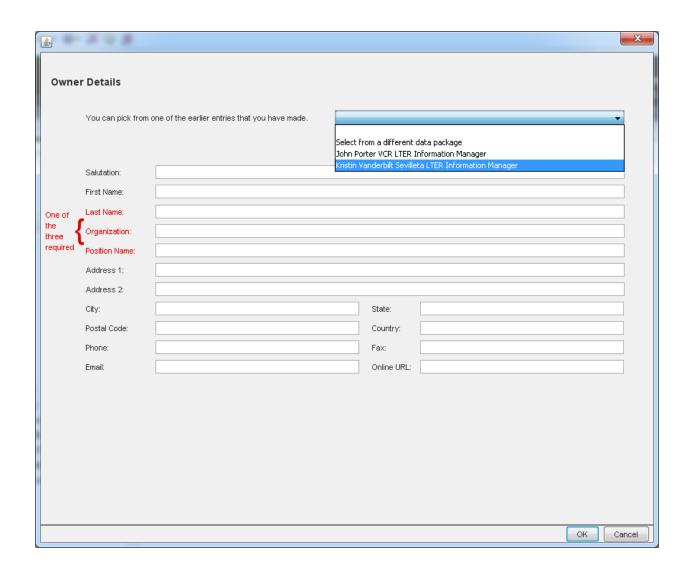
If you worked with other people, you can add them as Owners. Click 'Next' once you are done adding owners:



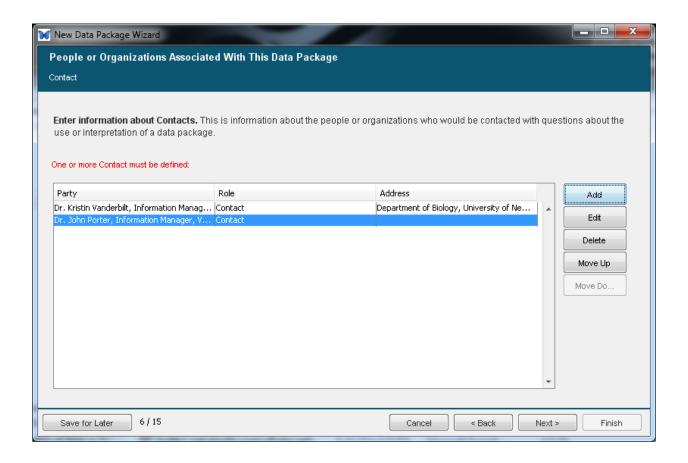
Now you will add contact information, to designate who should be contacted with questions about this dataset. Click 'Add' on this screen.



You are also probably the contact for this data set, so you can use the dropdown list in the upper right corner to select people who have already been entered. Add yourself and anyone else relevant:

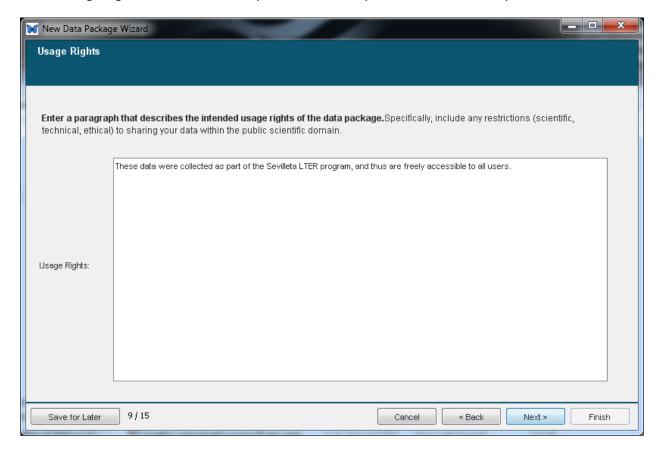


When you are done entering contacts, select 'Next' on this screen:

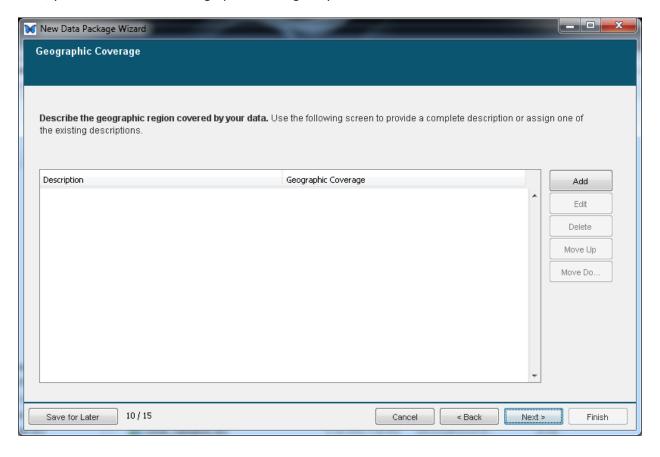


We are not going to enter Associated Parties or Research Project Information, so click 'Next' on the next two screens.

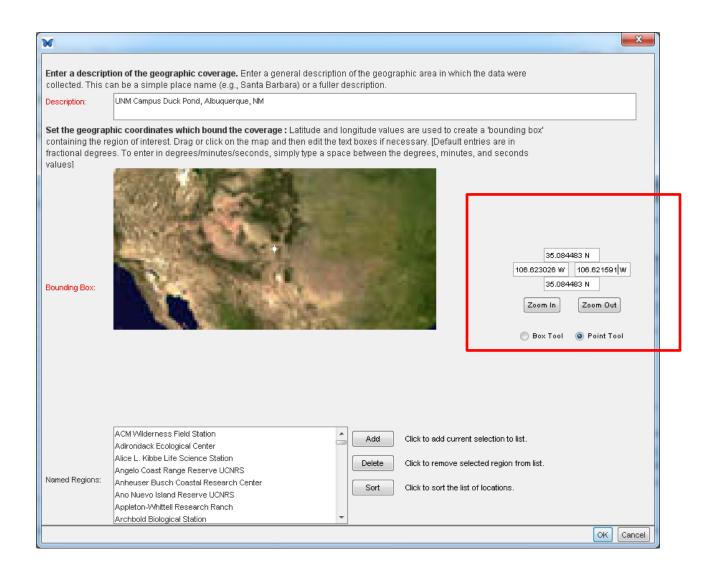
On the Usage Rights screen, indicate any restrictions that you want on the use of your data.



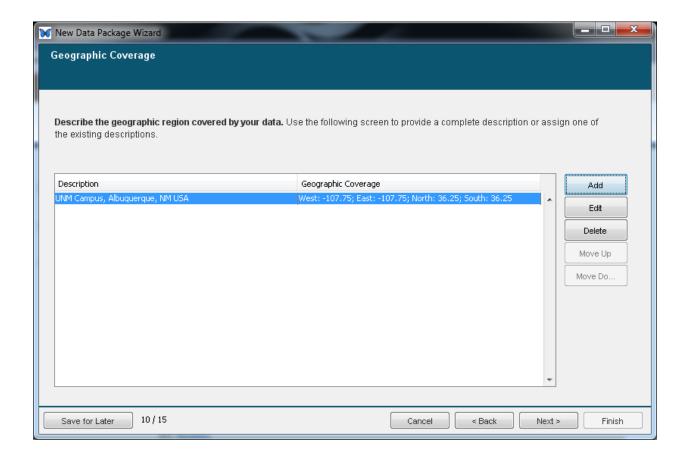
Next, you can describe the Geographic Coverage of your data. Click 'Add' on this screen:



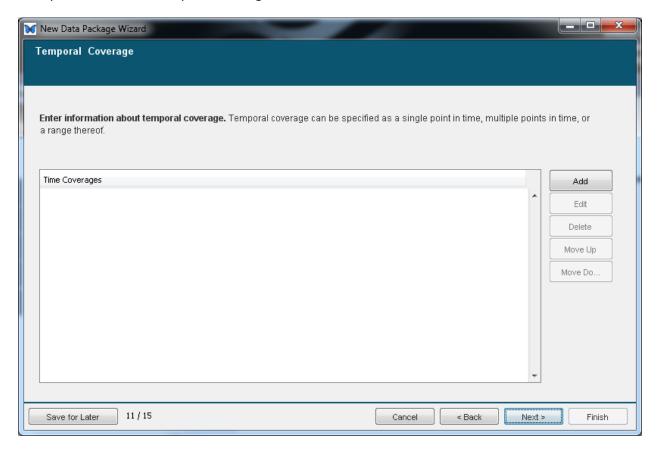
On this screen, enter a description of the study location. A bounding box for the duck pond area is entered below. Click 'OK' when you have entered the coordinates.



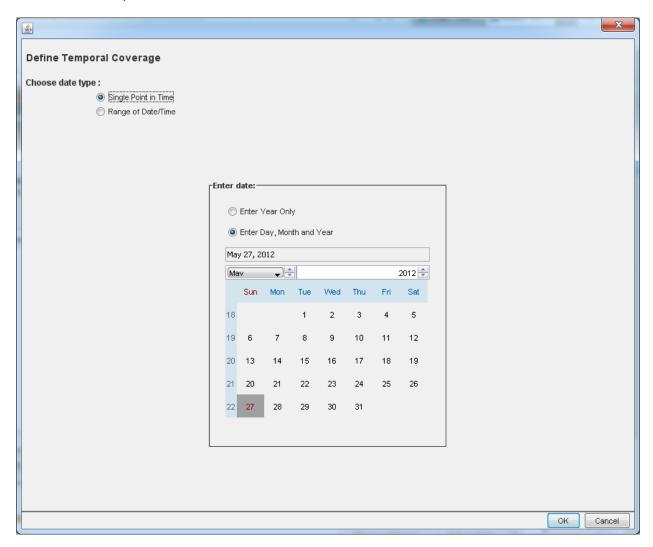
Enter 'Next' on this screen:



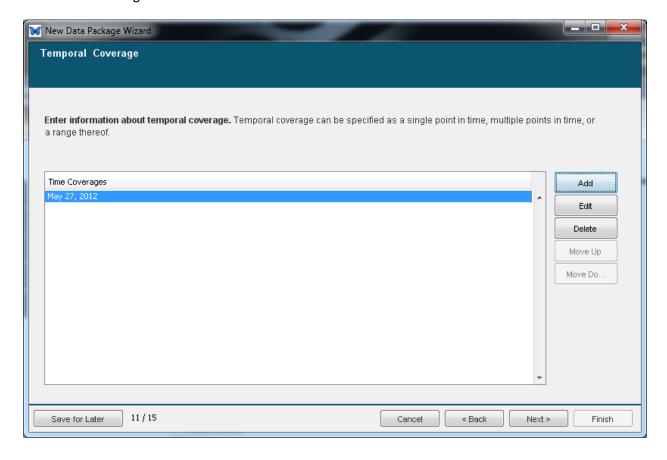
Now you will enter the temporal coverage. Click 'Add' on this screen:



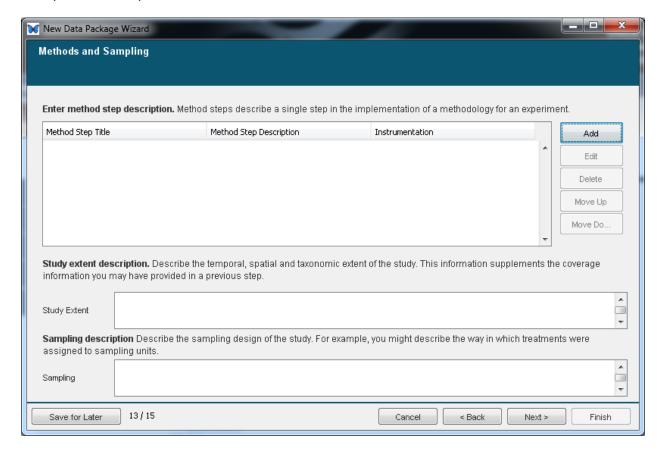
Enter the Date of your data Collection and select 'OK'.



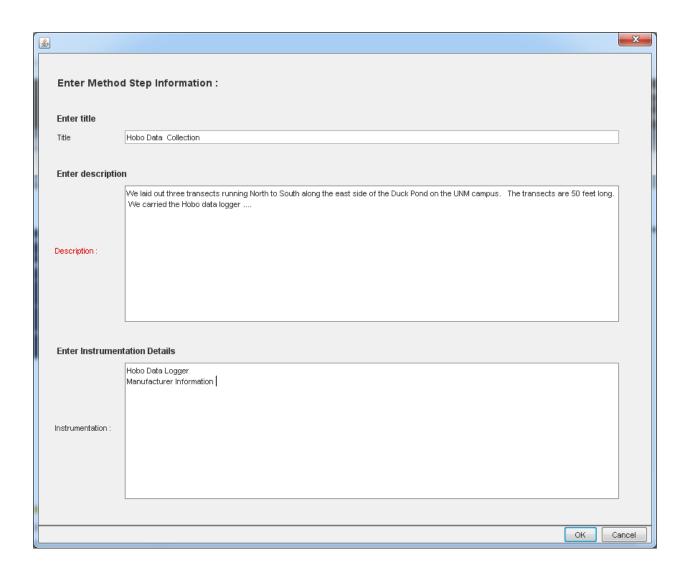
Click 'Next' on this screen. We are not going to enter taxonomic Coverage data, so click 'Next' on the Taxonomic Coverage screen.



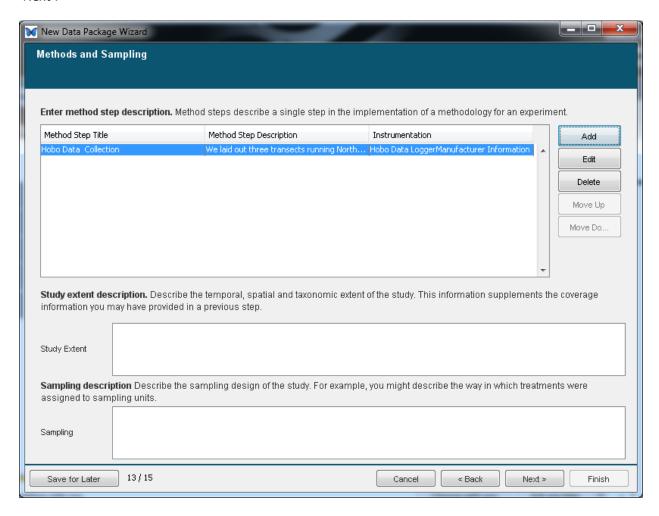
Now you will enter your methods. Click 'Add' on this screen:



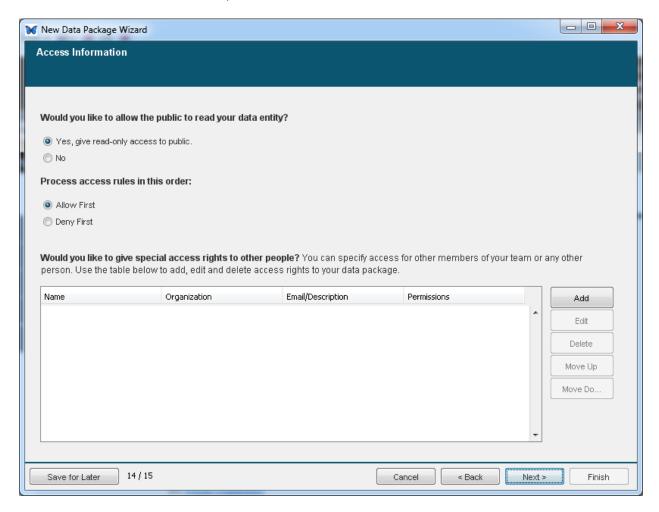
You can have several method steps. Use as many as you need to describe what you did. It would be a good idea to enter information about the Instrument you are using. Click 'OK' after entering each Method Step.



For our purposes, there is no need to enter Study Extent Description and Sampling Description. Click 'Next':

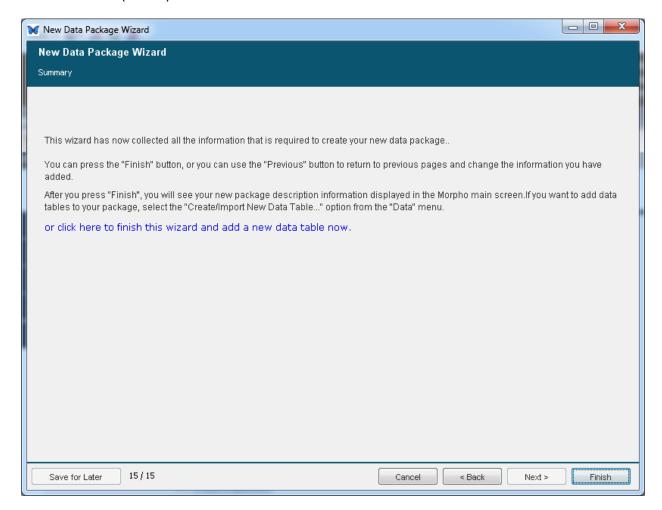


Now, decide if you want your data to be publicly accessible if it was added to a Metacat, such as the KNB metacat. Yes is the default. Then, choose 'Next':



Now you will be able to add your data table to the metadata package.

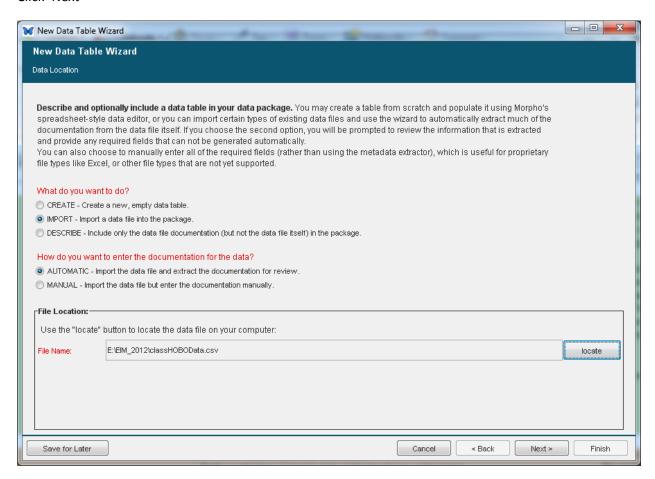
Click on the Link (in blue) to "finish this wizard and add a new data table now."



Select 'Import' and 'Automatic' on this screen.

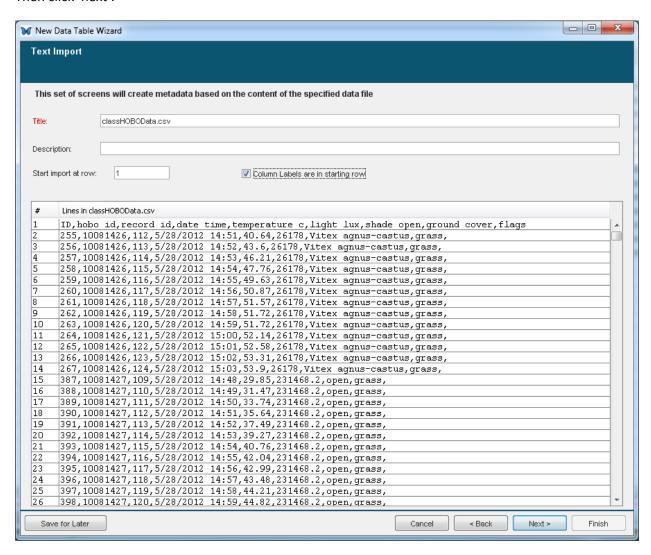
Click on Locate, and find the file classHOBOData.csv.

Click 'Next'



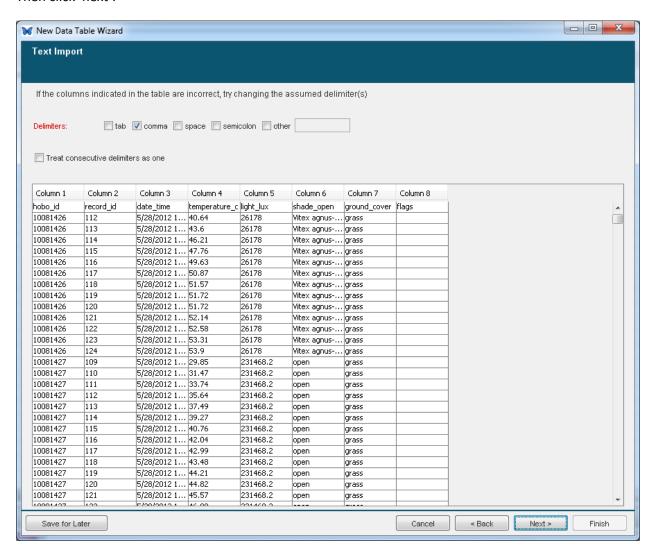
Check the "Column Labels are in starting row" box on this screen:

Then click 'next'.



Ensure Delimiters = Comma is checked on this screen:

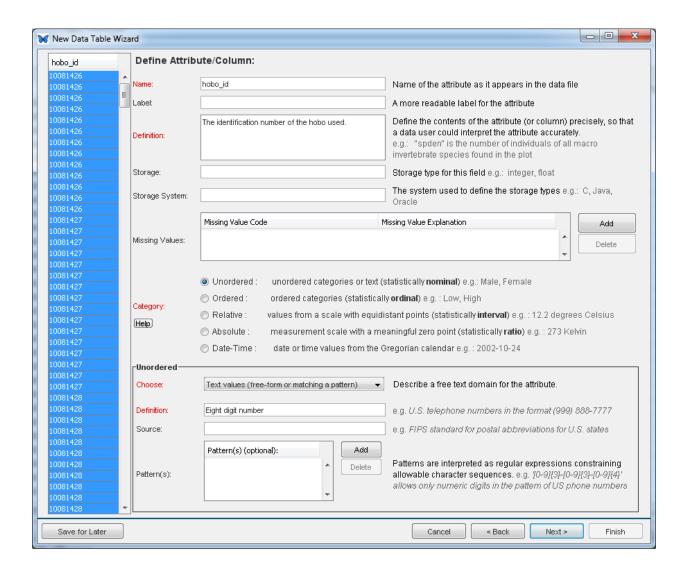
Then click 'next':



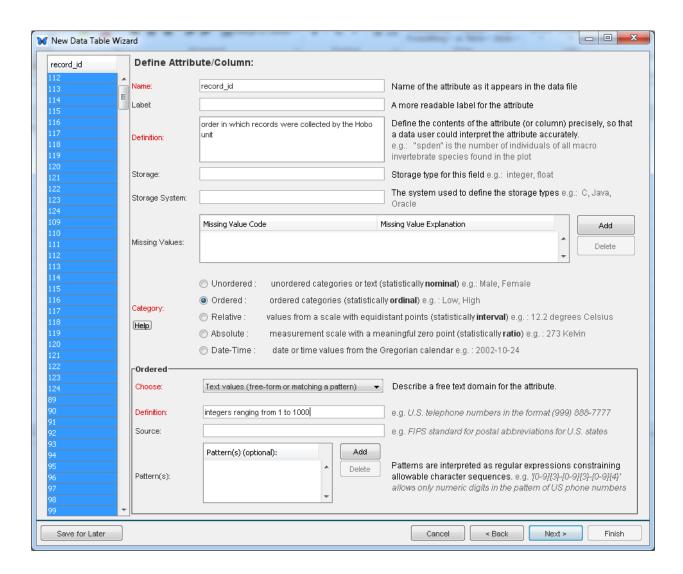
Now, you will define Metadata for Each Attribute/Column. You will need to enter something for every field marked in red.

Below is what I would enter for hobo_id. The id is just a label, which is why I chose 'nominal' as the category of variable.

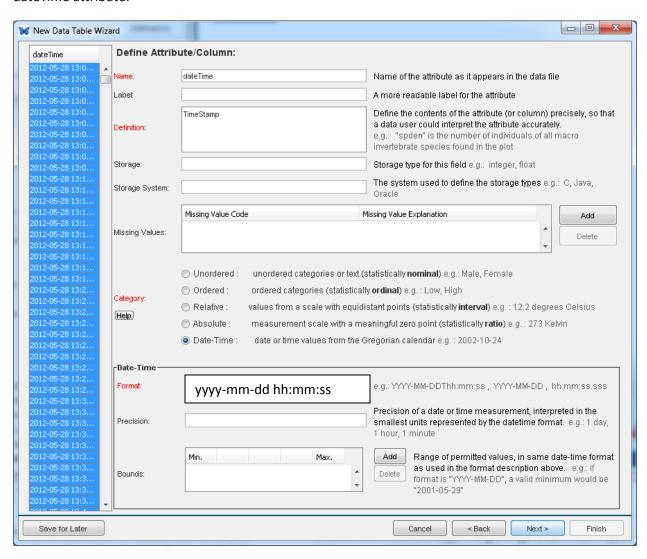
Click next when you have finished entering the metadata for hobo id.



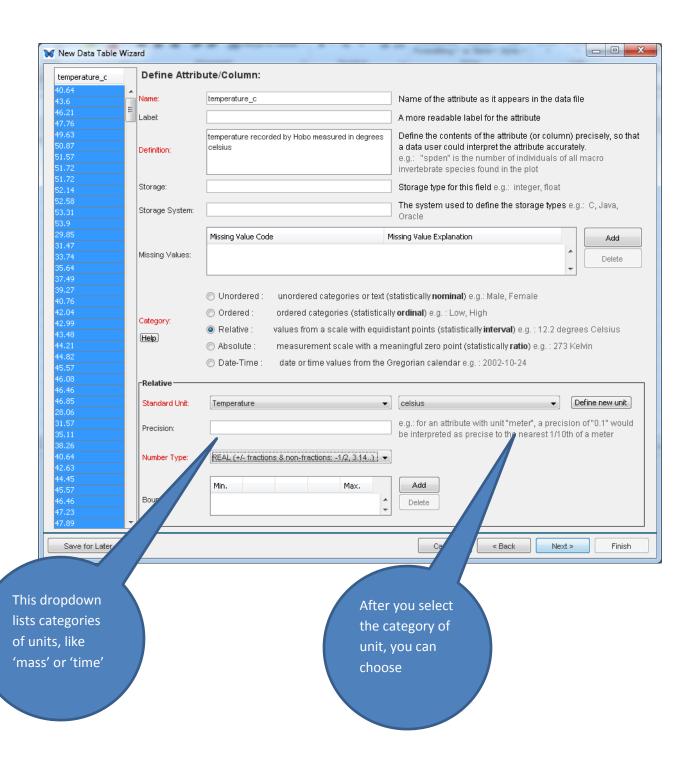
Record_id



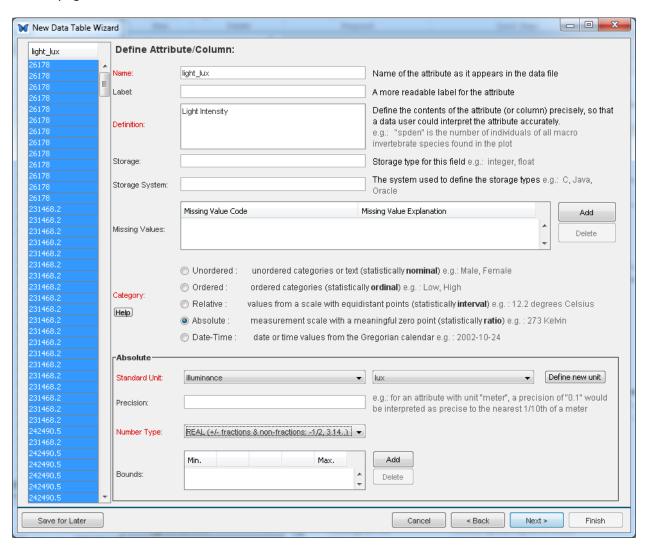
dateTime attribute:



temperature attribute: For this attribute, you have to choose the category of Unit, and then the unit itself:



intensityLight attribute: StandardUnit = Illuminance; Unit = lux.



Shade_open attribute:

If this were an EML document you were going to keep, you would define the codes for the trees. I recommend that you make this easy and define the codes as:

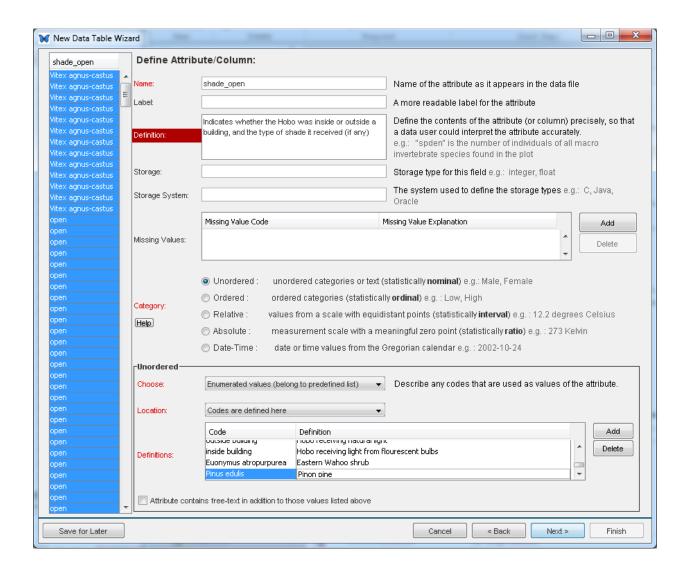
Vitex: shrub Open: Unshaded

Celtis occidentalis: shrub Platanus hybrida: tree Styphnolobium: tree

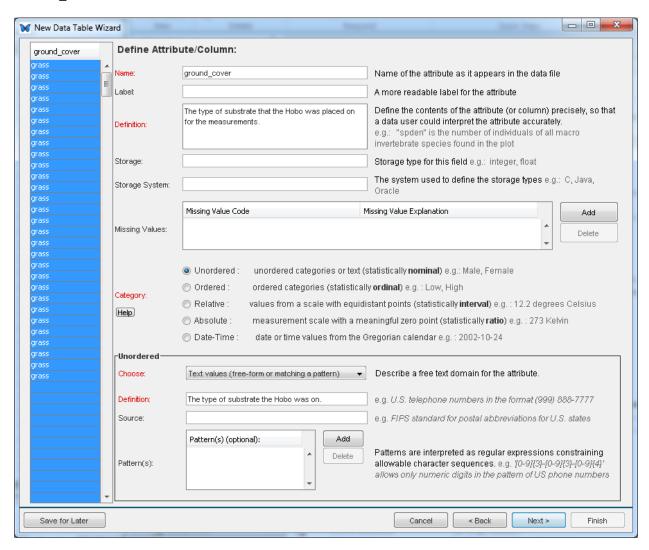
Outside building: Hobo receiving natural light

Inside building: Hobo receiving light from fluorescent bulbs

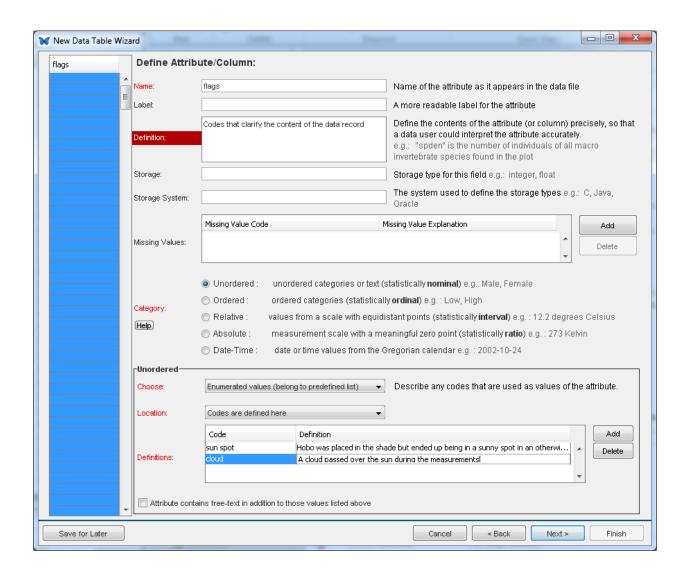
Euonymus: shrub Pinus edulis: tree



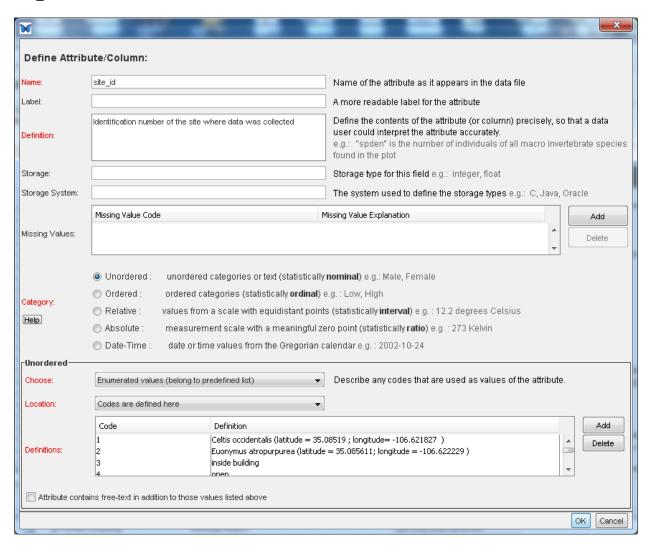
Ground_cover



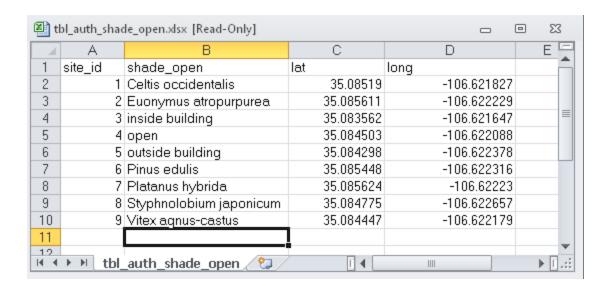
Flags:



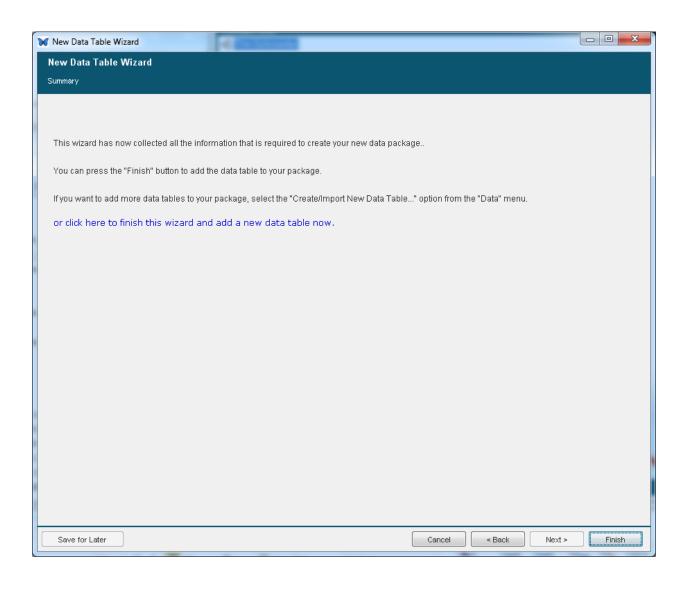
Site_id:



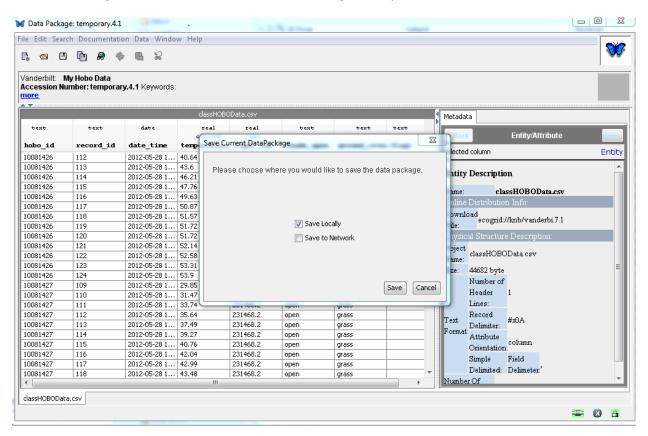
Here is a table of Site_id and definitions. You don't need to enter all the latitudes and longitudes.



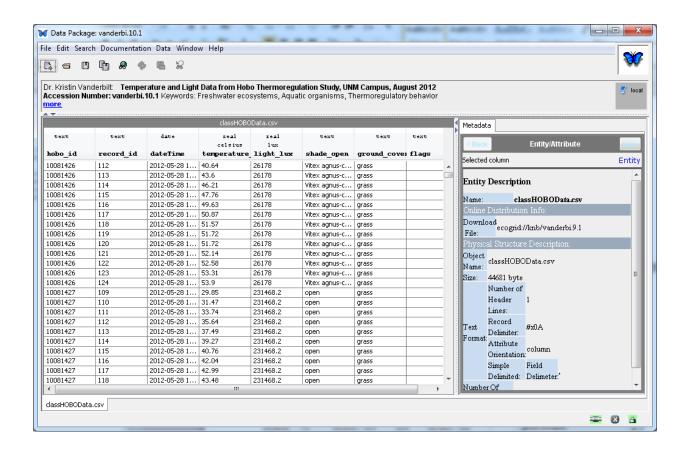
Yay!!! You are done. Click Finish on this screen:



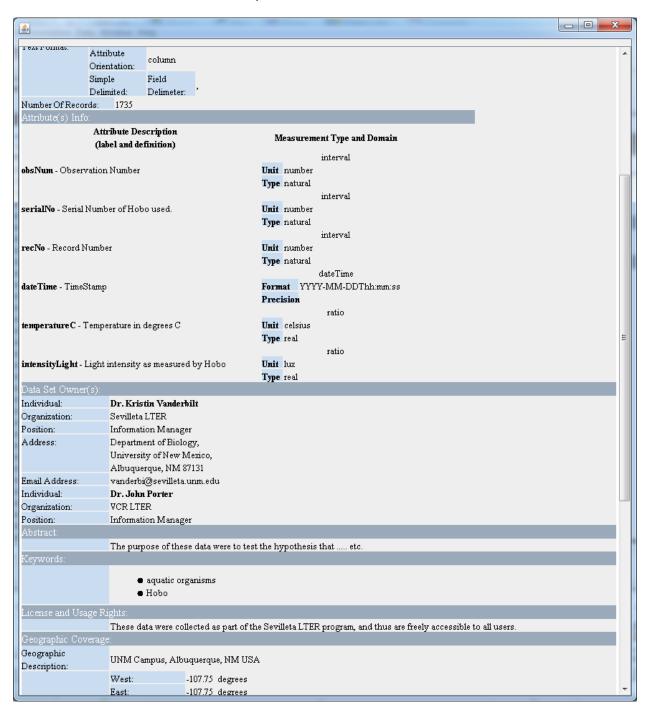
On this screen, go to File -> SAVE. Save the DataPackage Locally.



On this screen, select Documentation -> View Documentation. Then you can look at all the metadata you entered.



Scroll down to see all the metadata that you entered:



The actual EML file is stored in:

If you double-click on it, it should open in Notepad and you can see the XML structure.

