



# WoRMS and OBIS services

Aleksandra Pawlik  
myGrid Team  
University of Manchester

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<http://www.taverna.org.uk/>



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# WoRMS services

- WoRMS is the World Register of Marine Species
- In this small exercise, we will use the WoRMS services to find the scientific names of species within a genus
- The naming tools are available as WSDL Web Services called the **AphiaNameService**
- Create a new workflow in Taverna
- In the **Service Catalogue** perspective, search for *AphiaNameService*



# AphiaNameService

- You should see 15 WSDL services
- The easiest way to use the Aphia services is by the AphiaID.
- So we will first get the AphiaID for the genus
- Scroll down to getAphiaID
- Right-click on it and select **Add to workflow**
- You will switch back to the design perspective



# AphiaNameService search

Using BiodiversityCatalogue at: <https://www.biodiversitycatalogue.org>

Clear AphiaNameService

WSDL services (15) REST services (0)

Tags  
 Tags (on Operations)  
 Tags (on Inputs)  
 Tags (on Outputs)

Search results for query "AphiaNameService"

getAphiaClassificationByID ✔  
**Part of:** AphiaNameService  
**WSDL location:** <http://www.marinespecies.org/aphia.php?p=soap&wsdl=1>  
 Get the complete classification for one taxon. This also includes any sub or super ranks. []  
**1 Input:** AphiaID  
**1 Output:** Please refer to this service's WSDL and schema documents to get details on outputs for this <br> operation.

getAphiaID ✔  
**Part of:** AphiaNameService  
**WSDL location:** <http://www.marinespecies.org/aphia.php?p=soap&wsdl=1>  
 Get the (first) exact matching AphiaID for a given name.  
 Parameters: `marine_only`: limit to marine taxa. Default=true. []  
**2 Inputs:** `marine_only`, `scientificname`  
**1 Output:** return

getAphiaNameByID ✔  
**Part of:** AphiaNameService  
**WSDL location:** <http://www.marinespecies.org/aphia.php?p=soap&wsdl=1>  
 Get the correct name for a given AphiaID. []

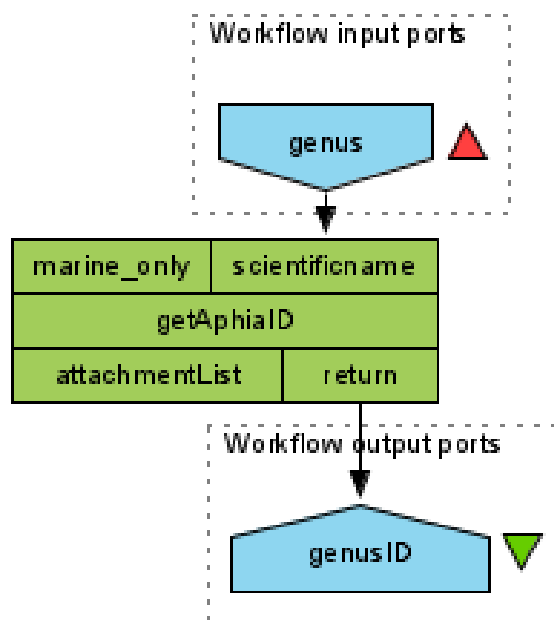


# Finding the Aphia ID of the genus

- Connect the input port *scientificname* of the service to a workflow input called *genus*
  - Connect the output port *return* of the service to a workflow output called *genusID*
  - Run the workflow with *Isurus*
  - You should get a genusID of *105743*
- Isurus is a type of mackerel shark



# Genus ID workflow



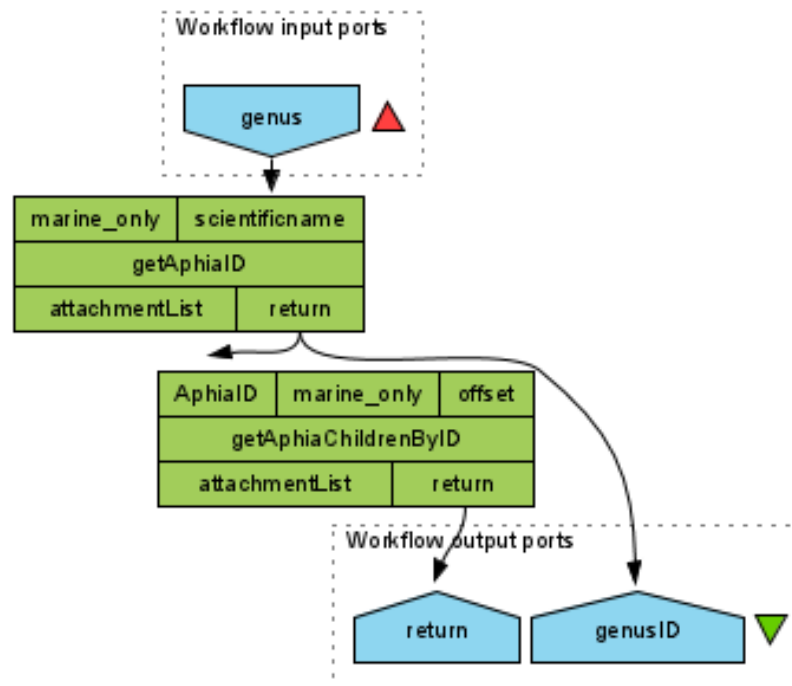


# Finding the genus's children

- We now want to find the “children” of the genus
- To do this, switch to the **Service Catalogue** perspective and find the service *getAphiaChildrenByID*
- Add it to the workflow
- Connect the *return* port of *getAphiaID* to the *AphiaID* port of *getAphiaChildrenByID*
- Connect the *return* port of *getAphiaChildrenByID* to a new workflow output port
- Run the workflow



# Genus children workflow







# Children return value

- If you look at the value returned by the *getAphiaChildrenByID*, it is an XML file - not very friendly
- We are just interested in the scientific names of the children
- There are two ways to extract this data, XML splitters or an XPath service – see the XPath service tutorial
- XML Splitters are used for WSDL services and they understand the XML documents produced/consumed by a service



# XML Splitters

- Delete the workflow *return* port
- Right-click on *getAphiaChildrenByID* and select **Add XML Output Splitter** – choose *return* to split
- If you connect the *return* port of *getAphiaChildrenByID\_return* to a workflow output and run the workflow, you can see it is still an XML document
- Right-click on *getAphiaChildrenByID\_return* and add an XML Output Splitter for its *return* port

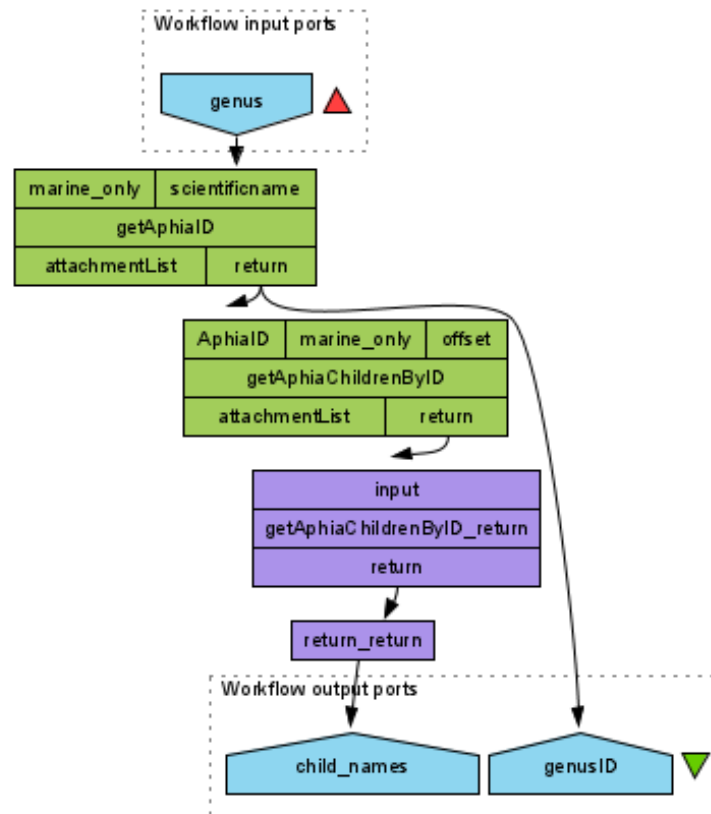


# Obtaining the children names

- Add a new workflow output port called *child\_names*
- Connect the *scientificname* port of *return\_return* to the *child\_names* port
- *Return\_return* has a lot of ports so it is easier to make the connection in the workflow explorer or to right-click on the service and choose **Link from output...** and pick *scientificname*
- Run the workflow
- *Child\_names* should now get a list of 15 scientific names



# Genus children names workflow





# WoRMS services

- Congratulations – you have now created a workflow using the WoRMS web services



# OBIS Services

- OBIS provides a set of Open Geospatial Consortium (OGC) Web Services at <http://www.iobis.org/geoserver>
- Access given to
  - Maps
  - Features – “things” on maps, including observations
  - Information about features
  - Legends to maps



# Getting a map

- The OBIS services are registered in BiodiversityCatalogue
- We will get a map of the counties in the world
- In Taverna, create a new workflow
- In the Service Catalogue perspective, search for *OBIS*
- Click on **REST services**



# GetMap

- Scroll down to *GetMap*
- Right-click and choose **Add to workflow**
- We need to set values for the ports of the *GetMap* service





# GetMap parameters

- Set the parameter values for the service
- Right-click on the port name and choose **Constant value**, enter the text value in the dialog.
- Make sure you do not have a newline in the value

Port name	Text value
bbox	-180,-90,180,90
format	image/jpeg
height	330
layers	OBIS:country
styles	country
width	660



# What do the parameters mean?

- *bbox* is the minimum longitude, minimum latitude, maximum longitude and maximum latitude as a comma separated list
- *format* says how we want the results returned – image/jpeg means as a JPEG image
- *height* and *width* specify the size of the JPEG image
- *layers* is a comma separated list of layers known to OBIS – OBIS:country is a map of countries
- You can find the possible layers and their formats at <http://www.iobis.org/geoserver/web/> under Layer Preview
- *styles* says for each layer how we want it displayed in the result

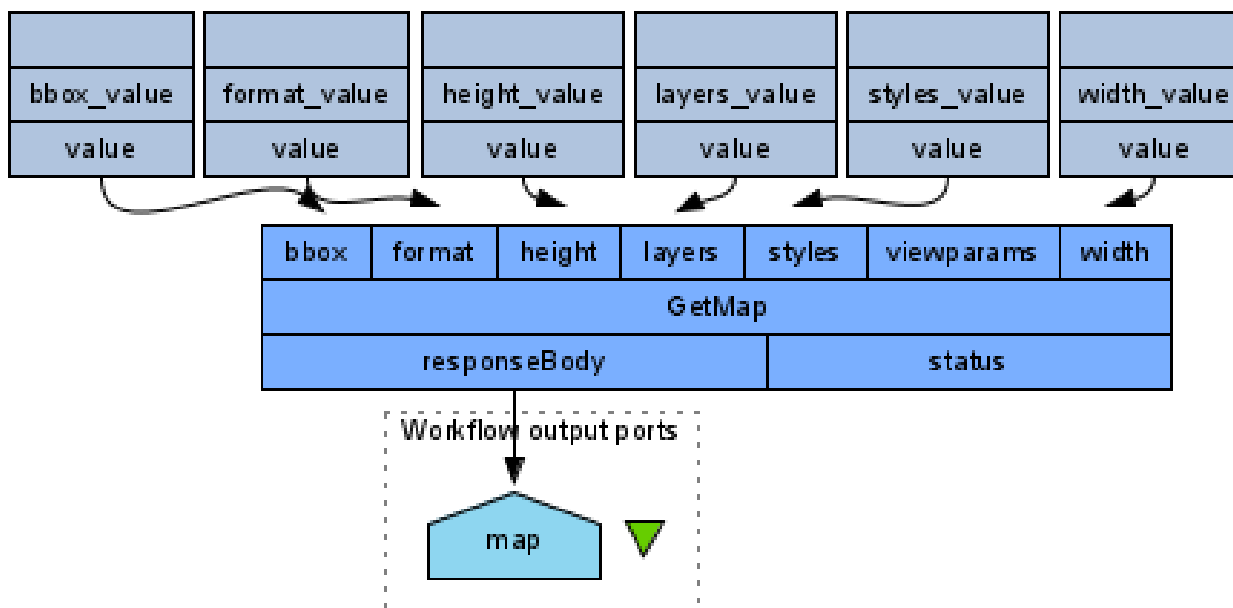


# Setting the return type

- Although we have set the format parameter to *image/jpeg*, Taverna does not know to expect *image/jpeg*.
- Right-click on *GetMap* and choose **Configure REST service**
- Change the 'Accept header' to *image/jpeg* and click **Apply** then **Close**
- Connect the *responseBody* port of *GetMap* to a new workflow output port



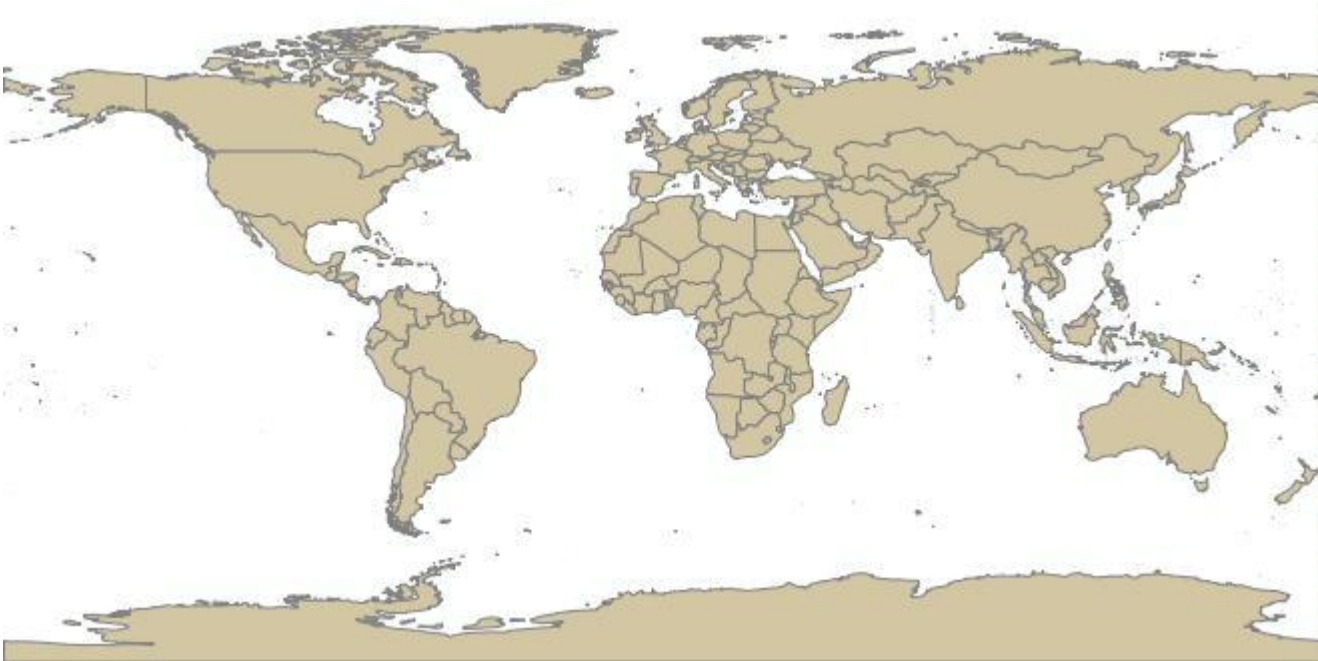
# GetMap workflow





# GetMap results

- When you run the workflow, you should see a world map:





# Save GetMap

- Save your workflow – *get\_map* is a good name
- We will use the workflow later on for more complicated maps
- You can now close the workflow – **Close workflow** under the **File** or **Taverna** menu



# GetFeature

- We are now going to get some species observations
- Create a new workflow
- Add the *GetFeature* service from the service
- In the workflow, merge in the Construct\_viewparams workflow from myExperiment – **Insert -> Merge workflow**
  - See: <http://www.myexperiment.org/workflows/4483>
- We covered merging workflows in the *Nested workflows* tutorial
- Connect the *viewparams* ports



# GetFeature parameters

- Set constant values for these ports of *GetFeature* and *construct\_viewparams*

Port name	Text value
maxfeatures	20
outputformat	csv
typename	OBIS:drs_with_woa

- Connect the *tname* port of *construct\_viewparams* to a new workflow input port



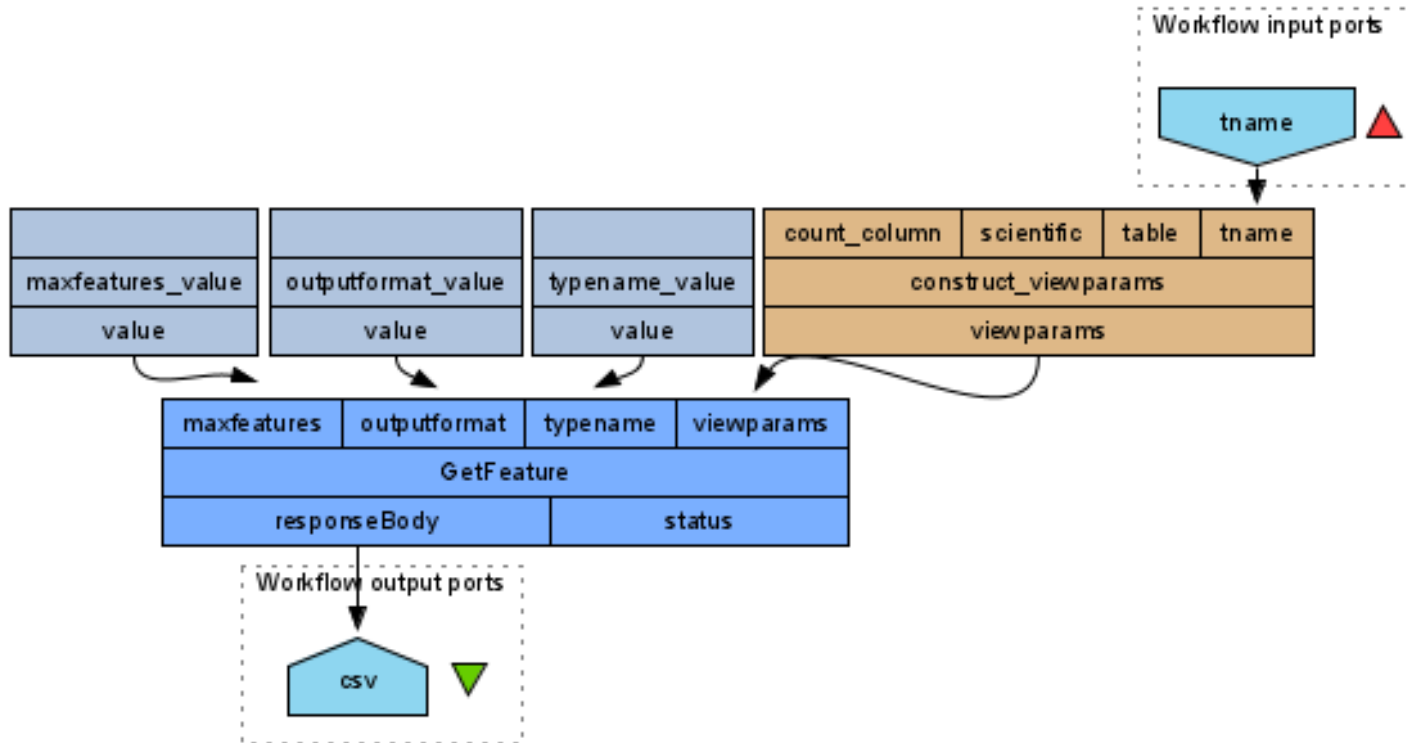


# What do the parameters mean?

- *maxfeatures* is the maximum number of features we want to return – so we will return at most 20
- *outputformat* says how we want the results returned – csv means as comma separated values
- *typename* is the layer we want to query – OBIS:drs\_with\_woa has occurrence information
- You can find the possible layers and their formats at <http://www.iobis.org/geoserver/web/> under Layer Preview
- *viewparams* is an SQL query into the table storing the layer



# GetFeature workflow





# GetFeature results

- Run the workflow with *tname* as Kogia breviceps
- The results of the GetFeature workflow are (in this case) csv data
- This data can be used, for example, to produce ecological niche models



# Gridded maps

- GetMap can be used to retrieve species distribution (gridded) maps
- Open the GetMap workflow you saved before
- We are now going to retrieve data from OBIS:dist\_sp – a species distribution layer - as well as OBIS:country
- We need to specify *viewparams* - an SQL query into the table storing the layer



# GetMap species distribution parameter changes

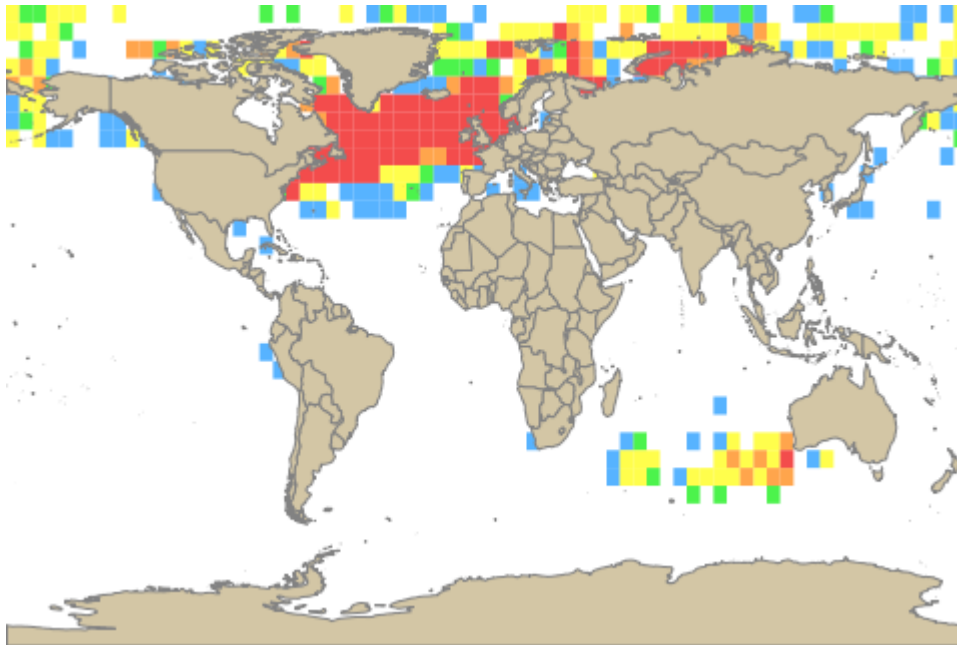
- We will need to edit the value for *layers* and also for *styles* – right-click on *layers\_value* and select **Edit value...**
- Also change *styles\_value*
- Add a constant value for the *viewparams* port

Port name	Text value
layers	OBIS:country,OBIS:dist_sp
style	country, grid_rainbow
viewparams	where:scientific='Calanus finmarchicus';table:dist_sp_5deg;



# GetMap species distribution results

- When you run the workflow, you should get a result like:





# Species distribution - issues

- The viewparams is difficult to understand
- What do the colours mean?



# Constructing viewparams

- To make it easier to construct the *viewparams*, merge in the Construct\_viewparams workflow from myExperiment – **Insert -> Merge workflow**
  - See: <http://www.myexperiment.org/workflows/4483>
- We covered merging workflows in the *Nested workflows* tutorial
- Construct\_viewparams is a helpful workflow that we have created for you
- Delete the *viewparams\_value* service





# Constructing viewparams - 2

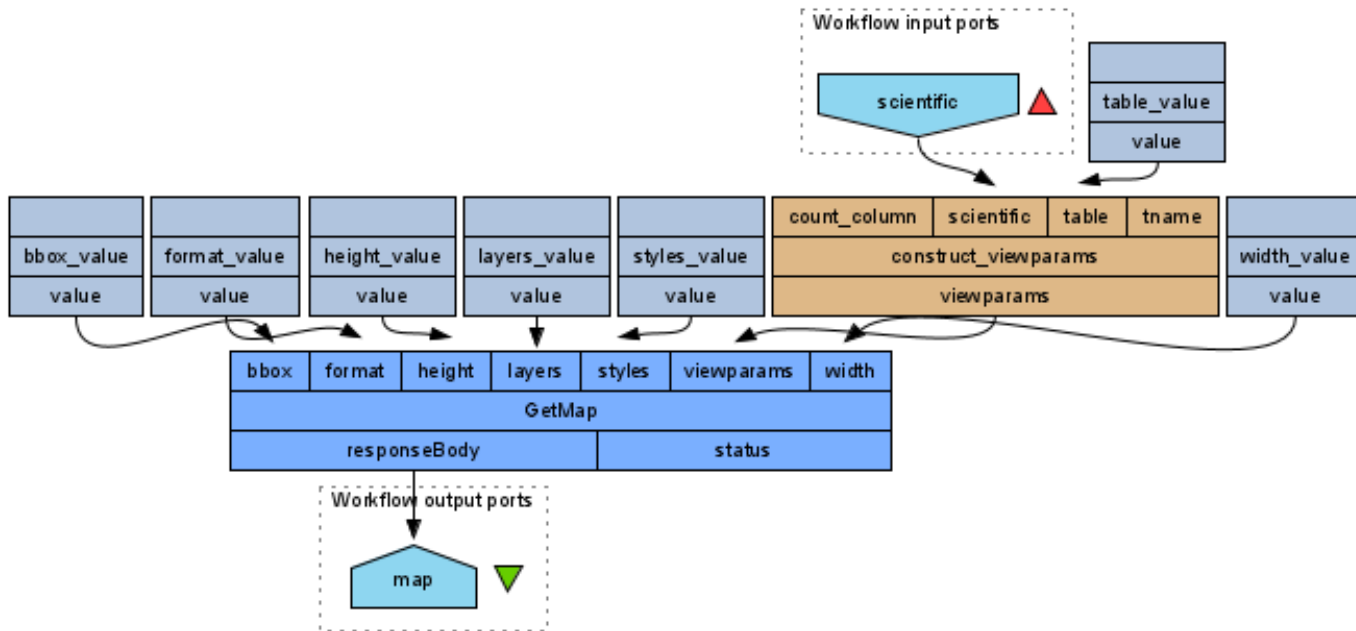
- Set a constant value for the table port of *construct\_viewparams* service to

Port name	Text value
table	dist_sp_5deg

- Connect the *scientificname* port of viewparams to a new workflow input port
- Connect the two *viewparams* ports



# Species distribution workflow





# Obtaining species distribution for a species

- You can now run the workflow and specify the species
- Try *Kogia breviceps* – you should get the same result as before
- Try *Limulus polyphemus* to see different results



# GetLegendGraphic

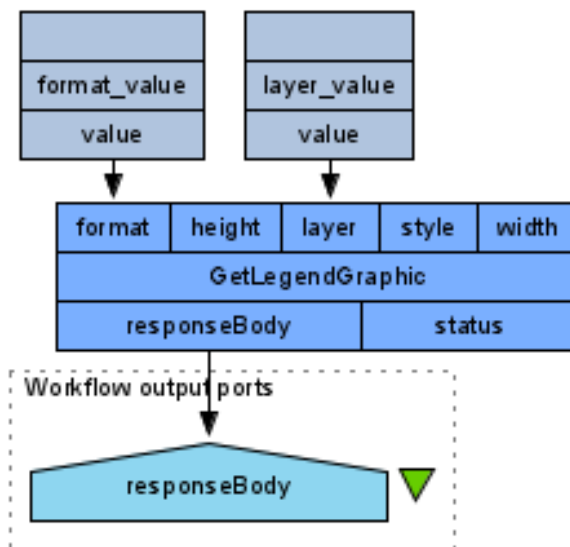
- We can find out what the colours mean with the *GetLegendGraphic* service
- Create a new workflow and add GetLegendGraphic from the Service Catalogue perspective
- Set constant values for its input ports

Port name	Text value
layer	OBIS:dist_sp
format	image/png

- Connect the responseBody port of GetLegendGraphic to a new workflow output port




# GetLegendGraphic workflow








# GetLegendGraphic - results

- Running the workflow will give:

  $> 101$

 51-100

 11-50

 6-10

 1-5



# OBIS services

- Congratulations – you are now able to
  - retrieve simple maps
  - retrieve species occurrence information
  - retrieve gridded maps for specific species
  - see what the colours on a map mean