Generating Ography Files

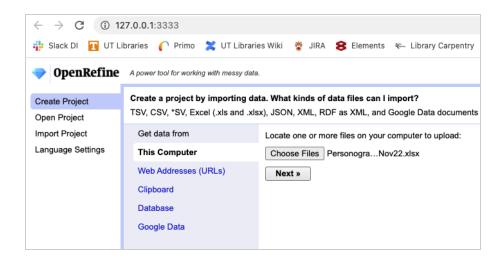
- 1. Go to the TaxonomyTerms spreadsheet in the Edgeworth Drive
- Assess the completeness of the "ography" file you are going to generate
 - a. Does every row have a xml:id?
- 3. Download the TaxonomyTerms spreadsheet and delete all of the tabs except for the one you plan to create an "ography" file for. These instructions will create a personography file using the "People" tab of the spreadsheet.
- 4. Delete the second row of the spreadsheet (which contains unnecessary instructional information).



becomes



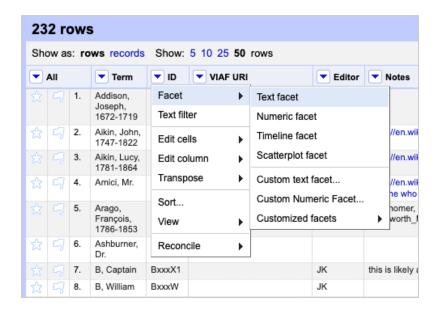
5. Upload edited spreadsheet to OpenRefine by going to "Create Project" and then "Choose Files." Once the spreadsheet is uploaded, click "Next."



6. On the next screen, ensure that the data looks correct. Rename the project if desired and be sure to uncheck "Store blank rows" and "Store blank rows as nulls" in the lower-right-hand corner. Click "Create Project" in the upper-right-hand corner when satisfied.



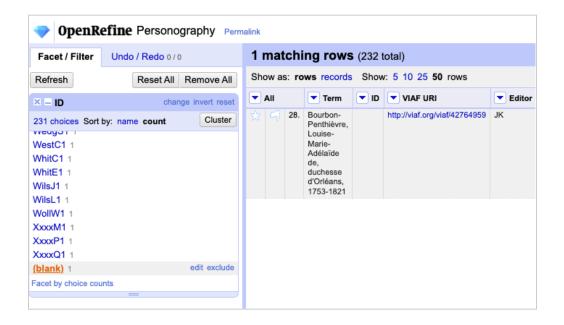
- 7. Once the project is created, check the "ID" column for duplicate and blank values.
 - a. To do this you can complete a "Text facet." Click the drop-down arrow on the "ID" column and then select "Facet" > "Text facet."



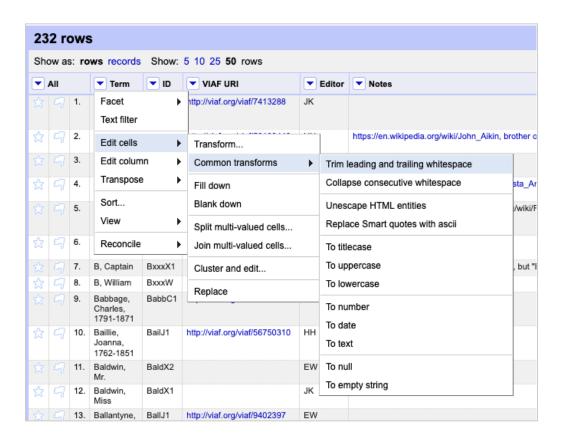
b. Then on the side facet, click "count" (rather than "name") and see if any ID values appear twice or more. If so, the duplicate ids will need to be distinguished from one another on both the OpenRefine project and back on the TaxonomyTerms spreadsheet. You will get an error in Oxygen later if duplicates are not resolved. The screenshot below shows a facet that requires no changes (if there are any IDs that appear twice, the "count" sort by would make them appear at the top).



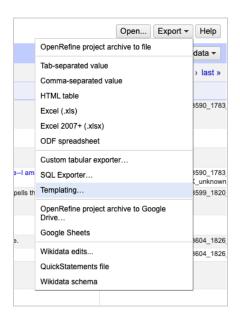
c. You need to also check the facet to see if there are any blank values (or names without established xml:id values). Scroll to the bottom of the facet to see if any appear. If present, click on "blank" to see what names need IDs. Then create an ID based on the <u>ID Conventions doc</u> and fill in this value in the OpenRefine project and back on the <u>TaxonomyTerms spreadsheet</u>.



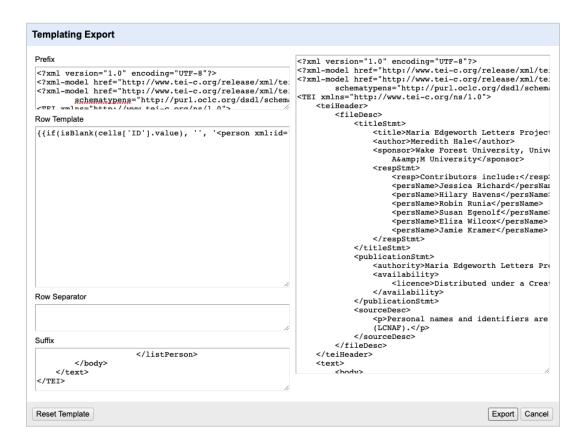
8. Next clean up the "Term" column values by trimming off any whitespace. To do this, click the drop-down arrow on the "Term" column, then select "Edit cells", "Commons transformations", and finally "Trim leading and trailing whitespace." This ensures that there are no extra spaces within persName> in the resulting XML. Repeat this action on the "VIAF URI" column.



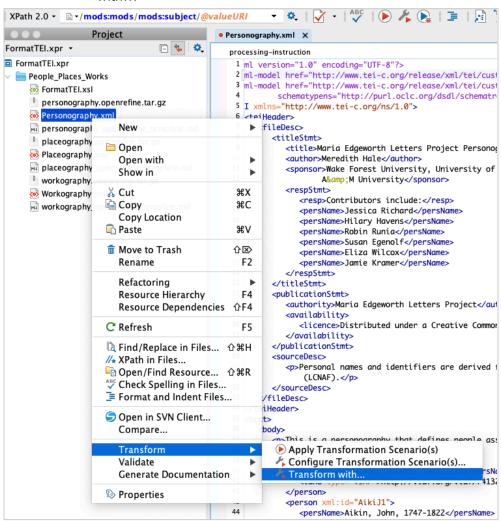
- 9. Next, use "Templating" and existing export templates on GitHub to create the XML.
 - a. First find the associated export template on the Maria Edgeworth me-tei GitHub repository by going to the People_Places_Works folder. Each export template is named by the type of "ography" it is and then "open_refine_template." For instance, the file you need for people is named
 - "<u>personography open refine template</u>." Either clone the repository code to GitHub desktop or download the single export file you need.
 - b. Open the file in Oxygen or MacDown. You'll notice that the template is composed of three separate sections. Those sections need to be copied and pasted into OpenRefine one by one.
 - c. In OpenRefine, go to "Export" in the upper-right-hand corner and then click on "Templating..."



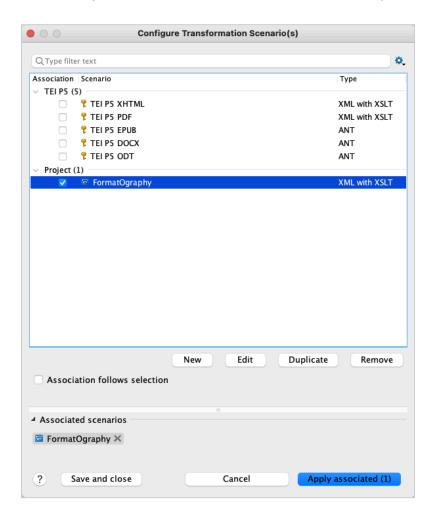
d. Next, fill out the "Templating Export" menu that pops up with the GREL code already present in the open refine export template you've opened from GitHub. Copy all the text (minus the back ticks ```) in ####Prefix to the "Prefix" section in the OpenRefine templating interface. Everything in the ####Body section goes to "Row Template". Delete the comma in the "Row Separator" box and leave it blank. Click "Export."



- 10. The exported file will be in your Downloads as a text file. Find and move the downloaded file to your Desktop and then change the extension (.txt) to .xml. Then open the file in Oxygen.
- 11. In Oxygen, ensure that you get a green box in the upper-right-hand corner and the message "Validation successful" at the bottom of the interface. If not, read and troubleshoot error messages. Save the file to your local GitHub repository and replace any older version of the "ography" you are working on with this new version (named the same thing as the previous file to track changes).
- 12. Transform the XML file using XSLT to streamline the formatting.
 - a. Open the <u>FormatTEI.xpr</u> file in Oxygen from the People_Places_Works folder within the GitHub repository to achieve this. Then right click on the appropriate "ography" file that you just saved/created and go to "Transform" > "Transform with..."



b. In the resulting pop up menu, select "" and then click "Apply selected scenarios".



13. Save the resulting file to your local GitHub repository for Edgeworth and then commit the new "ography" file to GitHub.