

Using ArcGIS Pro to Calculate Rarity Criteria for NatureServe Conservation Status Assessments

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This document describes the steps to be taken when using ArcGIS Pro to calculate rarity criteria for NatureServe conservation status assessments for species. More specifically, it describes how to gather species occurrence data, clip data to a boundary, calculate range extent, calculate number of occurrences, and calculate area of occupancy. This information can then be used directly to assign conservation status ranks using the NatureServe rank calculator (available at natureserve.org; NatureServe 2020b). These processes can be used to calculate criteria for an individual species or as a batch process to calculate criteria for multiple species simultaneously. Additionally, the processes can be applied at all geographic scales (global, national, and subnational) provided that the appropriate boundary is selected.

To follow these instructions, a basic understanding of ArcGIS software is required. Developed in 2025, these guidelines use ESRI ArcGIS Pro Version 3.0.0. Please note that as applications are continuously evolving, the applicability of these instructions may change slightly over time. Questions and further information related to the instructions below can be directed to Julia Prins (juliaprins19@gmail.com).

I would like to thank Dillon Blankenship (Tennessee Natural Heritage Program) and Nyssa Hunt (The University of Tennessee at Chattanooga Interdisciplinary Geospatial Technology Lab) for their GIS expertise and assistance with this manual.

Gathering Species Occurrence Data

Species occurrence records containing geographic coordinates are necessary to calculate the rarity criteria of range extent, number of occurrences, and area of occupancy. These records can be obtained from internal sources such as Biotics or external sources such as GBIF and the Symbiota portals.

Obtaining occurrences records from GBIF:

1. Visit gbif.org and select occurrences.
2. Select the All filters tab.
3. Under the Taxon category, apply a filter to only include records of the taxon of interest.
4. Under the Location category apply a filter to only include records from your area of interest by selecting the desired Country or area and State province. Under location select including coordinates.
5. Under the Event category apply a filter to exclude historic records using the desired cutoff year.
6. In the display window select Download and choose either the Simple or Darwin Core download option.

Obtaining occurrences records from a Symbiota portal:

Note: the instructions below are for the Symbiota portal of the SouthEast Regional Network of Expertise and Collections (SERNEC) but similar steps can be taken to obtain occurrence records from the other Symbiota portals (available at symbiota.org).

1. Visit <https://sernecportal.org/portal/>
2. Under the Specimen Search tab select Search Collections.
3. Under the Taxon category, apply a filter to only include records of the taxon of interest.
4. Under the Locality category, enter the Country or State of interest.
5. Under the Sample Properties category, check the Limit to Specimens with Geocoordinates box.
6. Under the Collecting Event category, enter the desired historic cutoff date.
7. Under Results Display Format, select Table and hit Search.
8. Click the Download Specimen Data button at the top of the page.
9. Select the settings of your choice and hit Download Data.

Prepping Occurrence Data for Upload:

If gathering occurrence data from multiple sources, all occurrences need to be merged into a single spreadsheet and several steps need to be taken to ensure information matches.

1. Ensure the same scientific names are being used across datasets. If any synonyms are being used in a dataset replace the synonym names with the correct scientific name. This can be done by obtaining a synonym list and cross checking it against the dataset.
2. Ensure the same coordinate system is being used across datasets. Coordinates should be in decimal degrees of latitude and longitude, with latitudes south of the equator

represented by negative values and longitudes west of the Prime Meridian also indicated by negative values.

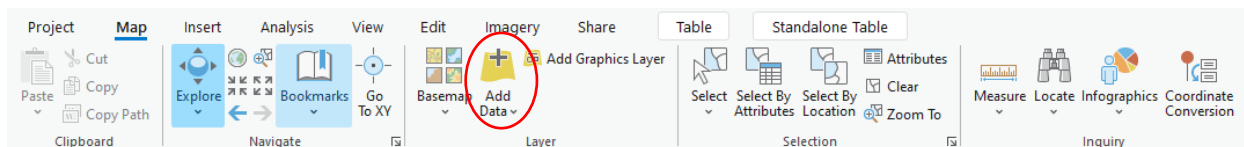
3. Next, the datasets can be combined into a single spreadsheet. One column with scientific name, one column with latitude, and one column with longitude must be present.
4. If ranking at the species level all infraspecific designations should be removed and simplified to species level (e.g. *Asplenium scolopendrium* var. *americanum* simplified to *Asplenium scolopendrium*). This ensures that all occurrences for a species are being accounted for in its calculations.
5. Lastly, ensure no spaces exist in column names or scientific names. Any spaces should be replaced with an underscore (“_”) using the find and replace tool. This is because ArcGIS Pro can have difficulty reading file names that contain a space.

Clipping the Data to Area of Interest

When using occurrence records from external sources, such as GBIF and Symbiota portals, it is crucial to verify that all records fall within the area of interest when mapped. Even if data is filtered during the initial collection, inaccuracies in locality information can lead to records being mapped outside the intended area. Therefore, to ensure the accuracy of data, it is important to clip the occurrence records to the boundary of the area of interest as described below.

Step 1: Upload a boundary file.

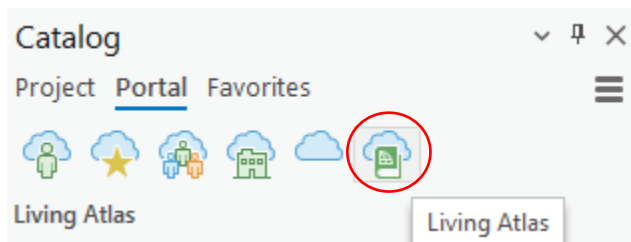
If you already have a layer defining the boundary of your area of interest add this to your map by navigating to the ribbon at the top of the page and under the Map tab in the Layer group, select Add Data. Select the boundary layer and hit OK.



If you do not have a layer defining the boundary of your area of interest, you can use a pre-existing layer from the ArcGIS Online Living Atlas (option 1 below) or define your own boundary (option 2 below).

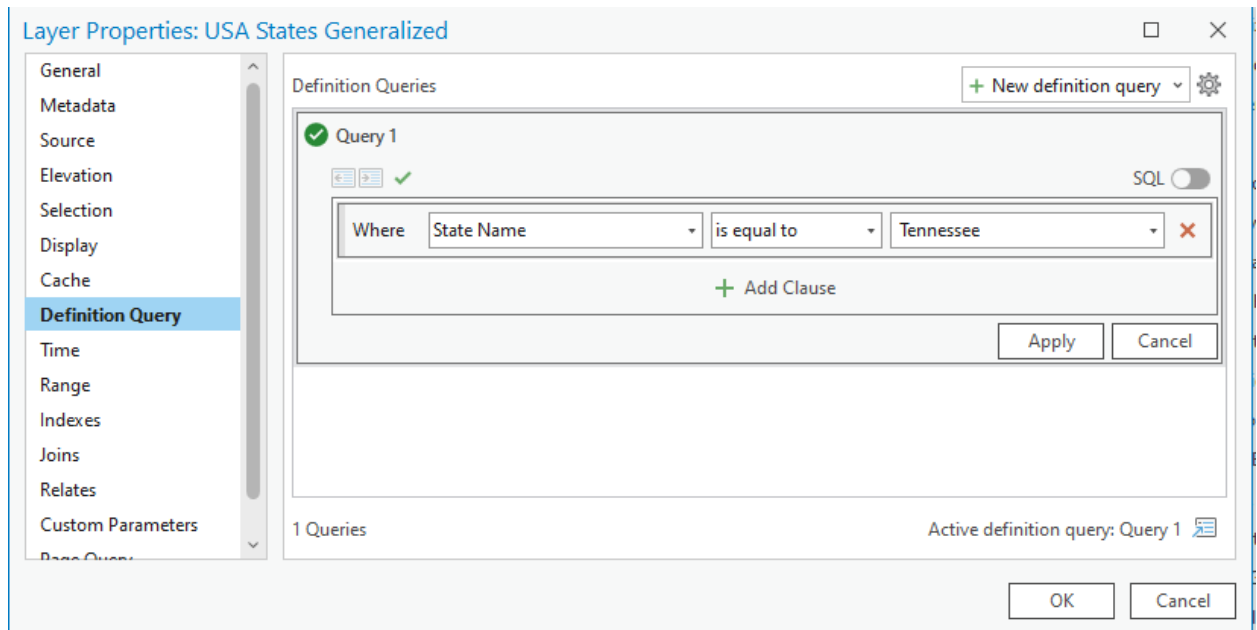
Option 1: Using a layer from the ArcGIS Online Living Atlas for U.S. States or Canadian Provinces/Territories

1. In the Catalog pane select the Portal tab. Select Living Atlas.

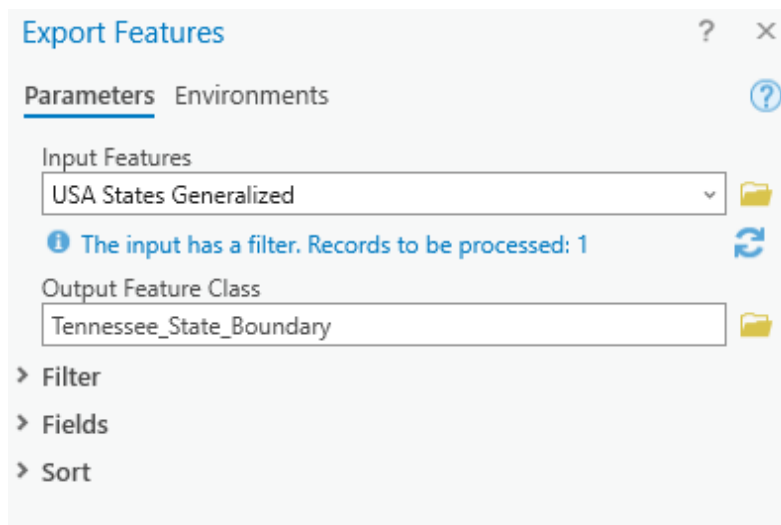


2. Search “USA State Boundaries” or “Provinces and Territories of Canada”. Multiple layers containing the boundaries of your search will appear. Right click the feature layer of your choice and select Add to Current Map.

3. To only include the boundary of your state or province/territory of interest a definition query can be applied to the layer. In the Contents pane right click the boundary layer and select Properties. In the Properties window select the Definition Query tab. Select New definition query and add a query where state or province/territory name is equal to state or province/territory of interest. Select Apply and hit OK.

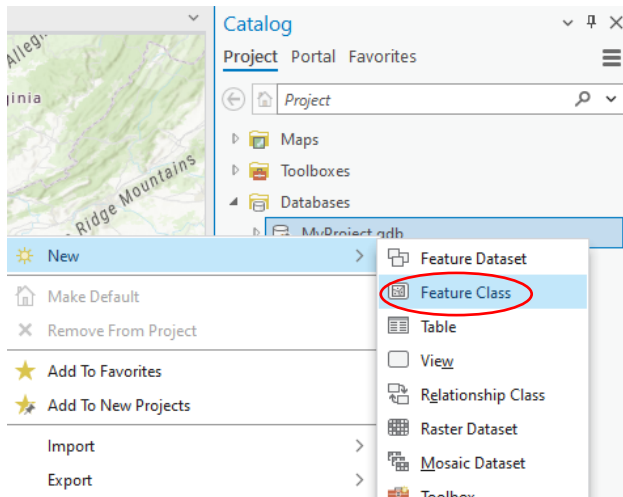


4. In the Contents pane right click the boundary layer and select Data and Export Features. Name the Output Feature Class as the state or province/territory boundary and select OK. This will add the living atlas layer to the local geodatabase so it can be used for geoprocessing tools.

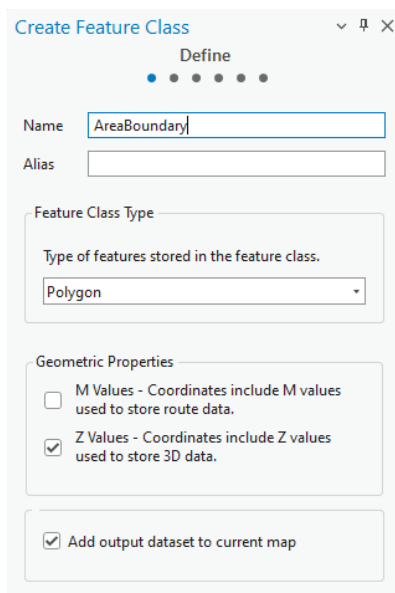


Option 2: Manually Defining a Boundary

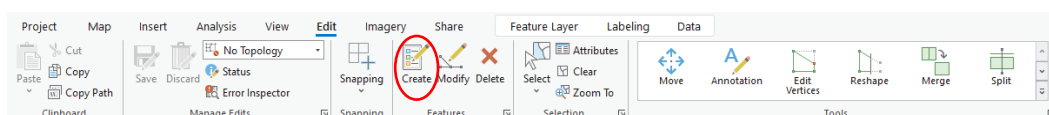
1. In the Catalog pane, under the Project tab expand Databases. Right click the current project geodatabase and select New and Feature Class.



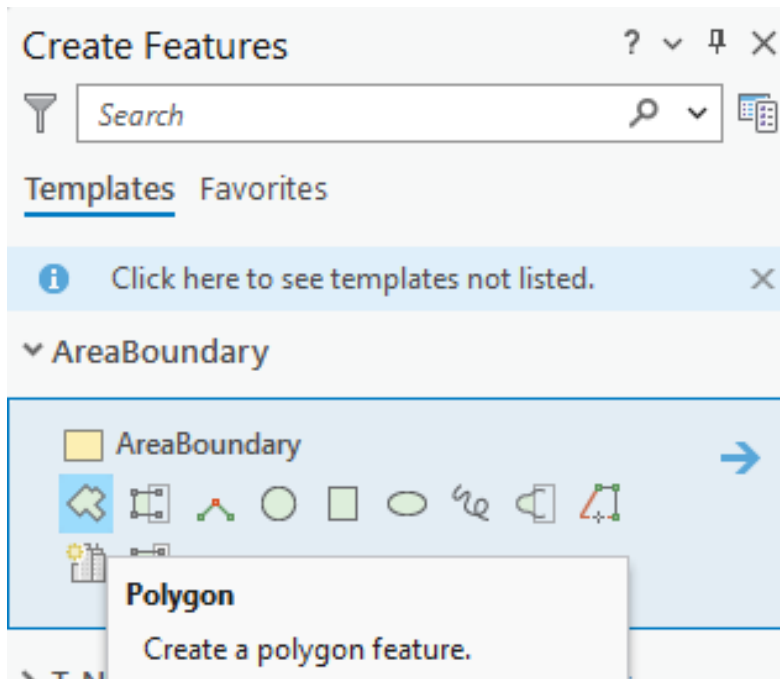
2. Name the Feature Class “AreaBoundary” and ensure Polygon is selected as the Feature Class Type. Select Finish.



3. Navigate to the ribbon at the top of the page. Under the Edit tab in the Features group select Create.



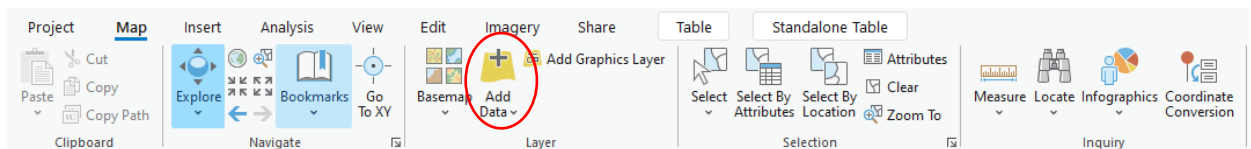
4. In the Create Features pane select the new AreaBoundary layer and use the Polygon tool to draw a boundary for your area of interest on the map. Double click to finish the polygon.



5. Navigate to the ribbon at the top of the page. Under the Edit tab in the Manage Edits group select Save.

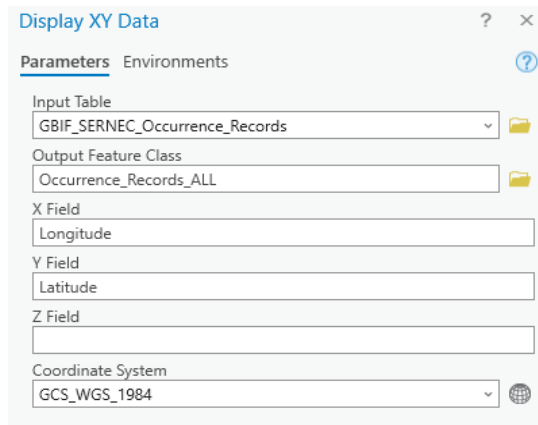
Step 2: Upload your occurrence records.

1. Ensure your spreadsheet is downloaded as a CSV file.
2. Navigate to the ribbon at the top of the page. Under the Map tab in the Layer group select Add Data.



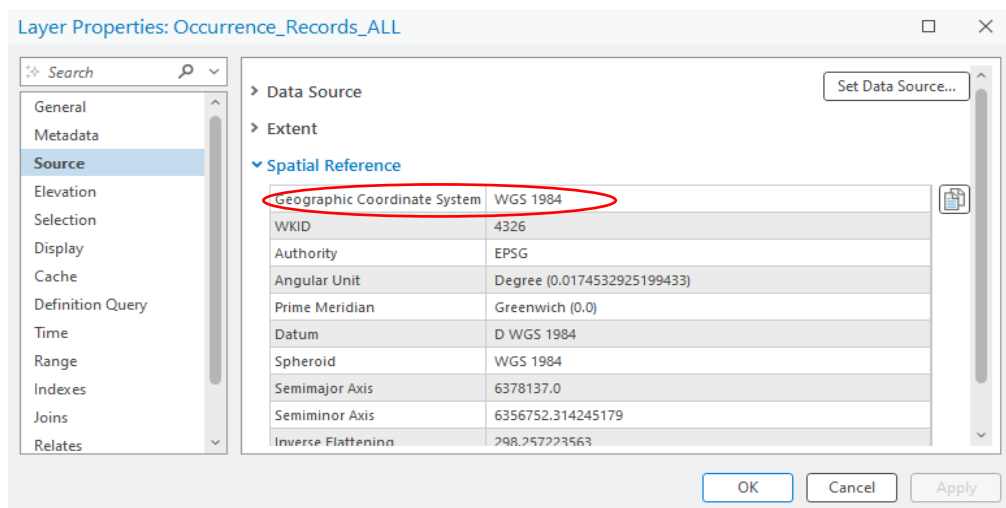
3. Select the file containing your occurrence records and hit OK.

4. In the Contents pane right click the newly added attribute table containing your occurrence records and select Display XY Data (Note: newer versions of ArcGIS Pro may require you to select Create Points From Table and then XY Table to Point instead).
 - a. Ensure your occurrence file is selected as the Input Table.
 - b. Name your Output Feature Class as “Occurrence_Records_ALL”.
 - c. For the X Field ensure the column containing longitude is selected.
 - d. For the Y Field ensure the column containing latitude is selected.
 - e. Hit OK.
 - f. This will add the occurrence points to your map.

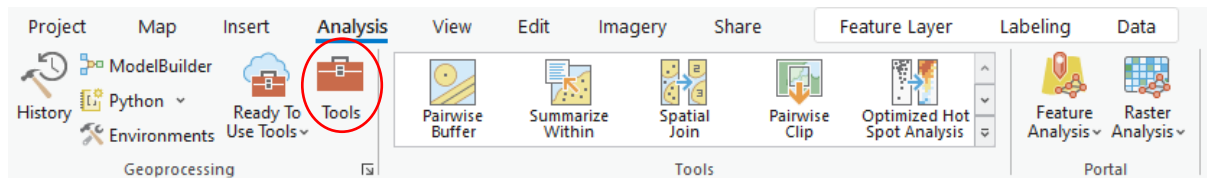


Step 3: Ensure the coordinate system of your dataset is UTM (Universal Transverse Mercator)

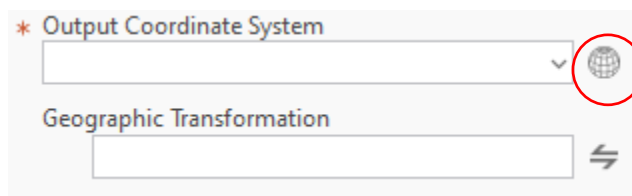
1. To check the coordinate system of your dataset, in the Contents pane right click the layer containing all occurrence records and select Properties. In the Properties window select the Source tab. Expand Spatial Reference and the Geographic Coordinate System will be listed. If the coordinate system is not UTM the occurrence record layer will need to be Projected into a UTM coordinate system using the following steps (Note: the image below shows a coordinate system *not* in UTM).



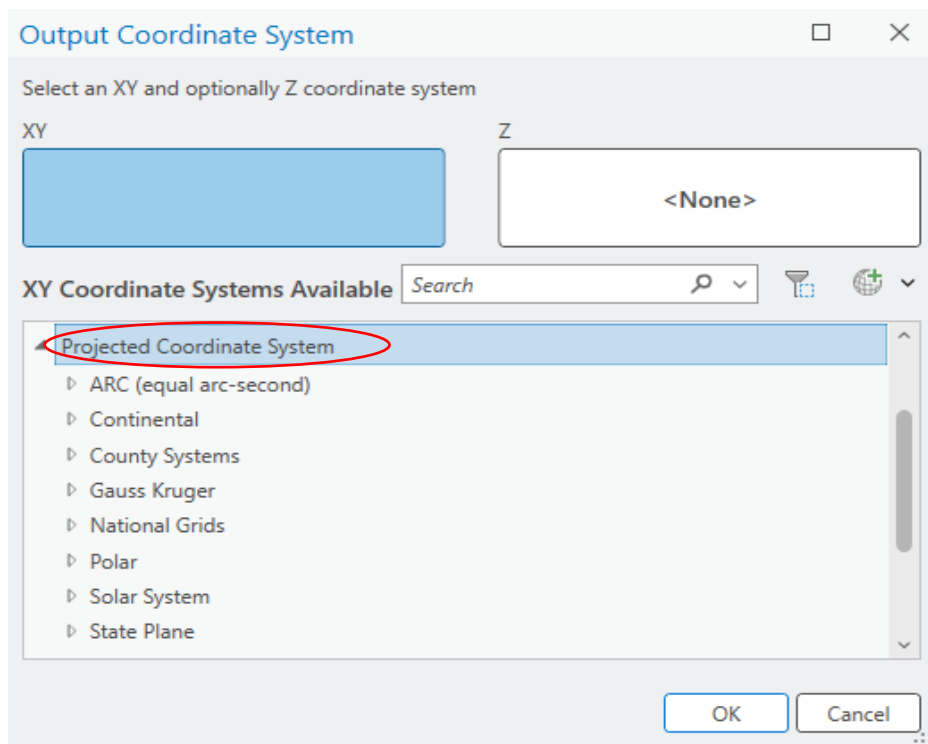
2. Open the geoprocessing pane by navigating to the ribbon at the top of the page. Under the Analysis tab in the Geoprocessing group select Tools.



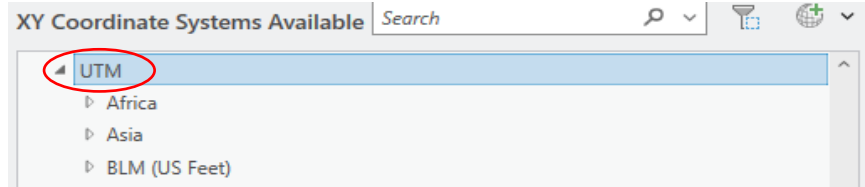
3. In the geoprocessing pane search for and open the *Project* tool.
4. For Input Dataset or Feature Class select the layer containing all occurrence records.
5. Name the Output Dataset or Feature Class “Occurrence_Records_UTM”.
6. Select a UTM Output Coordinate System.
 - a. Click the globe icon to select a coordinate system.



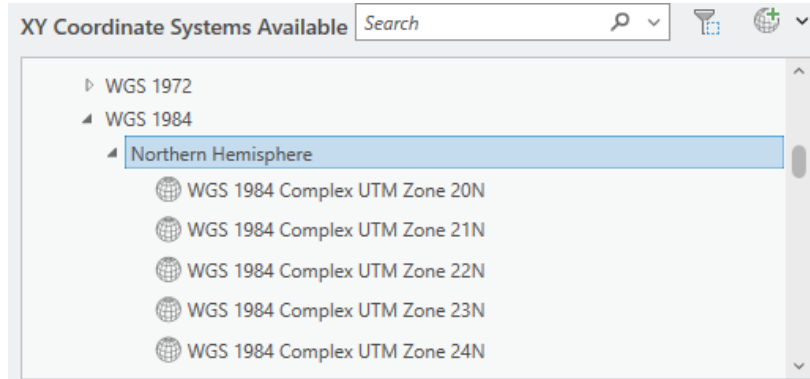
- b. In the Output Coordinate System window under XY Coordinate Systems Available expand Projected Coordinate System.



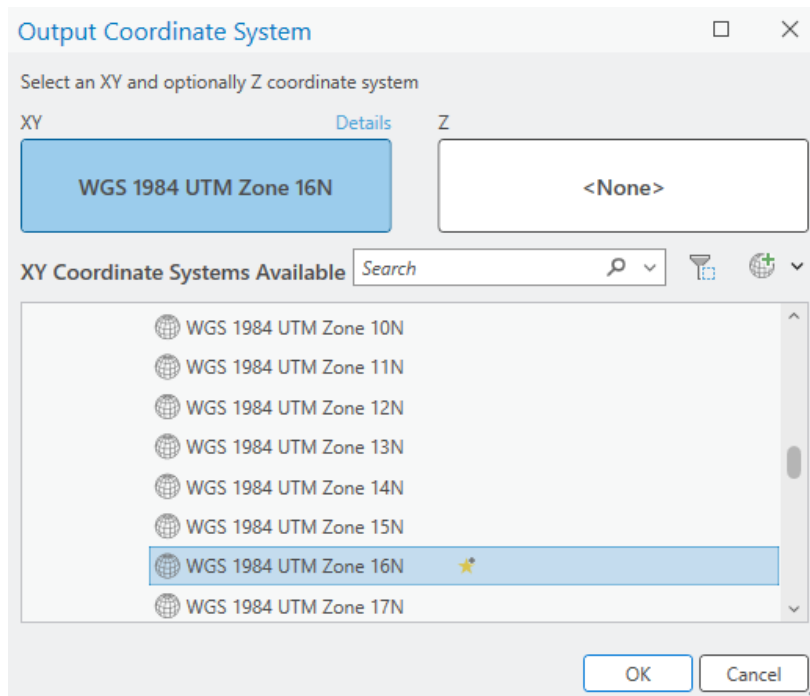
- c. Scroll down and expand UTM.



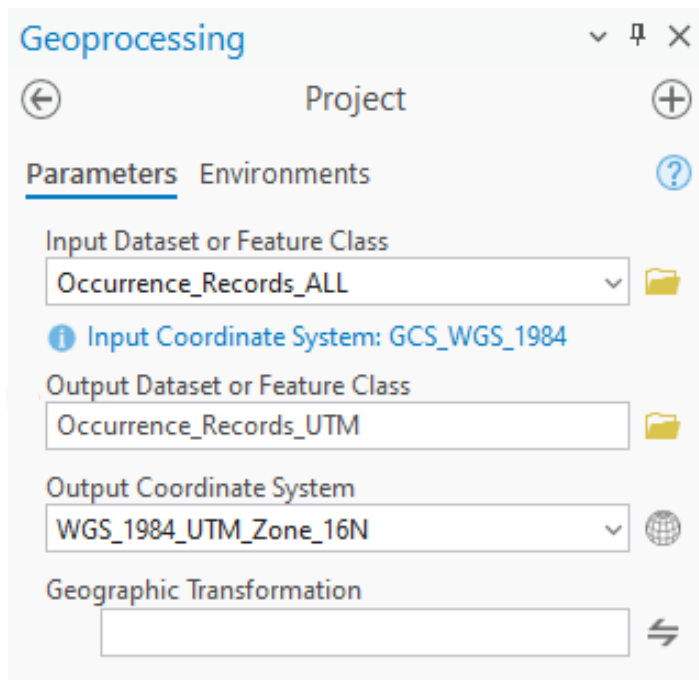
- d. Scroll down and expand WGS 1984 and expand Northern Hemisphere



- e. Find and select the UTM zone located at the center of your area of interest (this information is readily available on the internet).



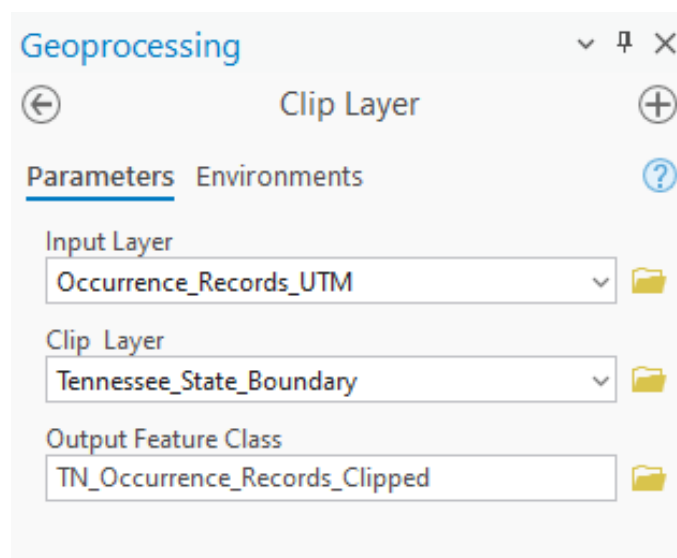
- f. Click OK.



7. Click Run.

Step 4: Use the *Clip Layer* tool to remove all occurrence points outside of the area of interest.

1. In the geoprocessing pane search for and open the *Clip Layer* tool.
 - a. For Input Layer select the layer containing all occurrence records in the UTM coordinate system created in step 3.
 - b. For Clip Layer select the layer containing the boundary of your area of interest.
 - c. Name the Output Feature Class "Occurrence_Records_Clipped".
 - d. Click Run.

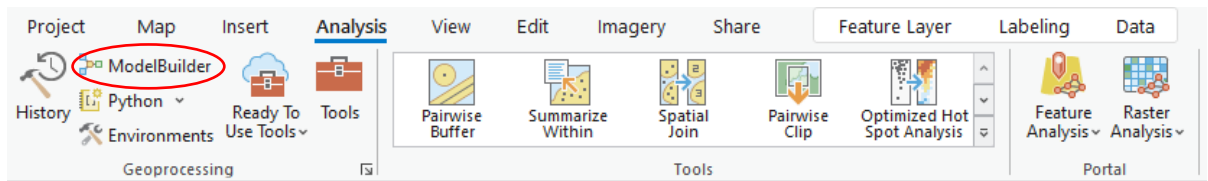


Creating An Individual Occurrence Layer for Each Species (Only Necessary for Batch Processing)

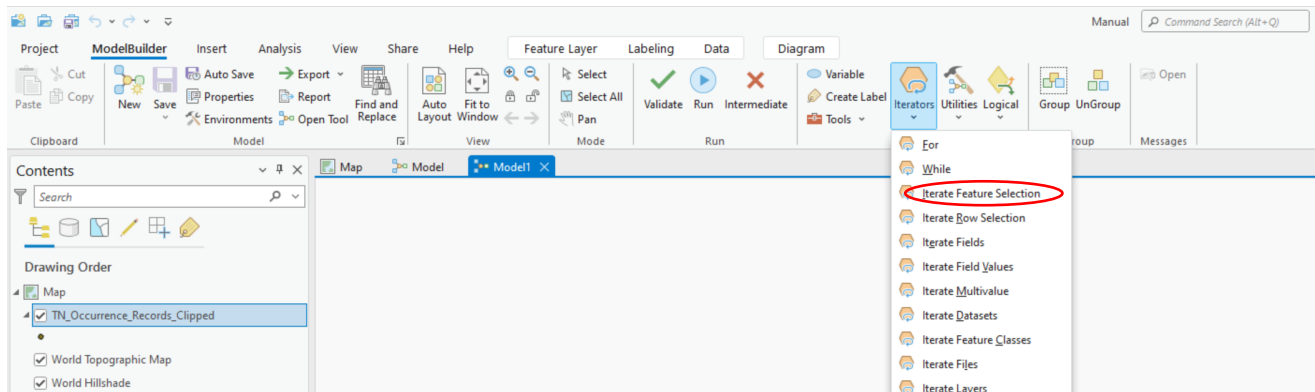
If calculating criteria for multiple species you will need to separate your occurrence data into individual layers for each species. This ensures that criteria calculations for each species are based solely on occurrences records of that species. This can be done by creating a ModelBuilder in ArcGIS Pro as outlined below. If calculating criteria for a single species this section can be skipped.

Create a ModelBuilder.

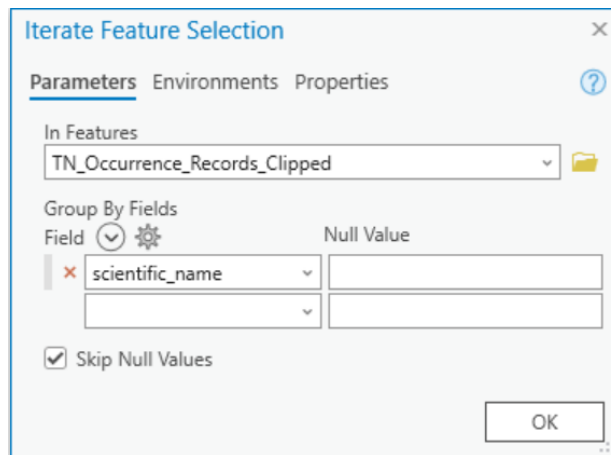
1. Open a ModelBuilder by navigating to the ArcGIS Pro ribbon at the top of the page. Under the Analysis tab in the Geoprocessing group select ModelBuilder. This will open a new blank model page.



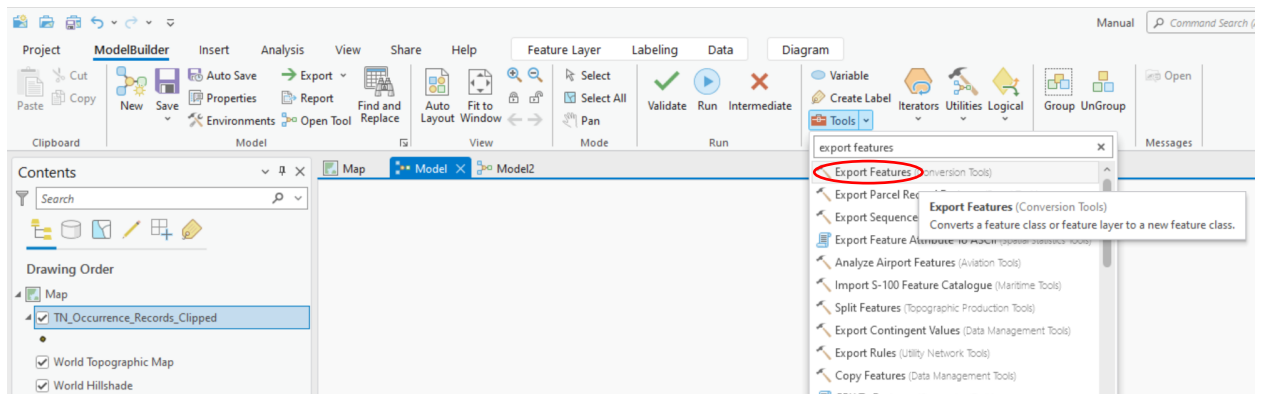
2. Under the ModelBuilder tab in the Insert group select Iterators and choose *Iterate Feature Selection*. This will add the iterator to your model.



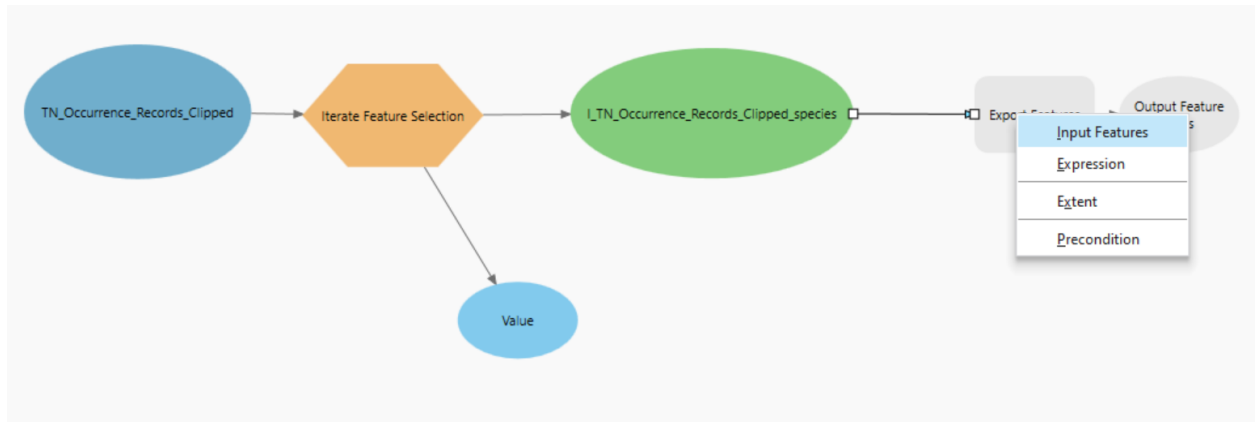
3. In your model open the Iterate Feature Selection element by right clicking and selecting open.
 - a. For In Features select the clipped occurrence layer created in the previous step.
 - b. Under Group By Fields from the dropdown select the field that contains the scientific name of your species. This will be whatever you named the column containing species names in your original occurrence file.
 - c. Ensure the Skip Null Values box is checked.
 - d. Select OK.



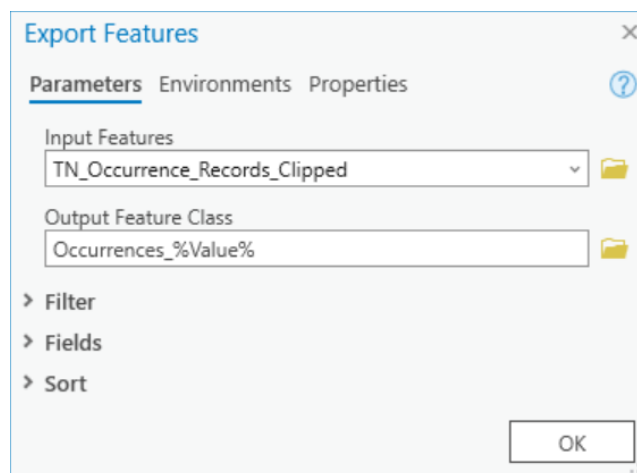
4. Under the ModelBuilder tab in the Insert group select Tools and search for the *Export Features* tool. Select this tool and drag it into your model.



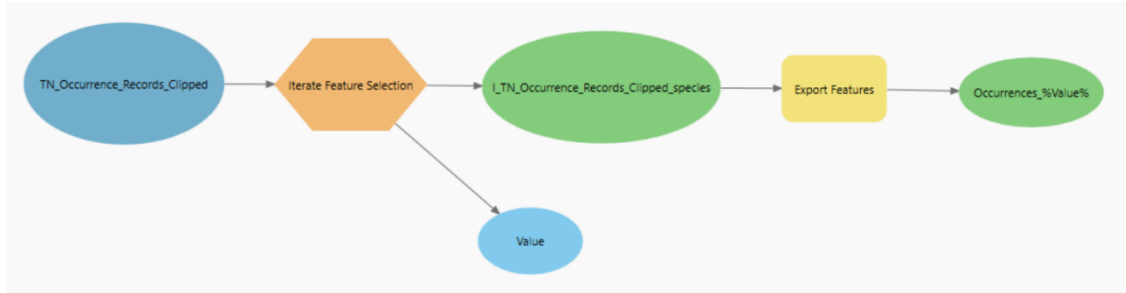
5. In your model drag an arrow from the selected features of your Iterate Feature Selection element to the Export Features element and select Input Features.



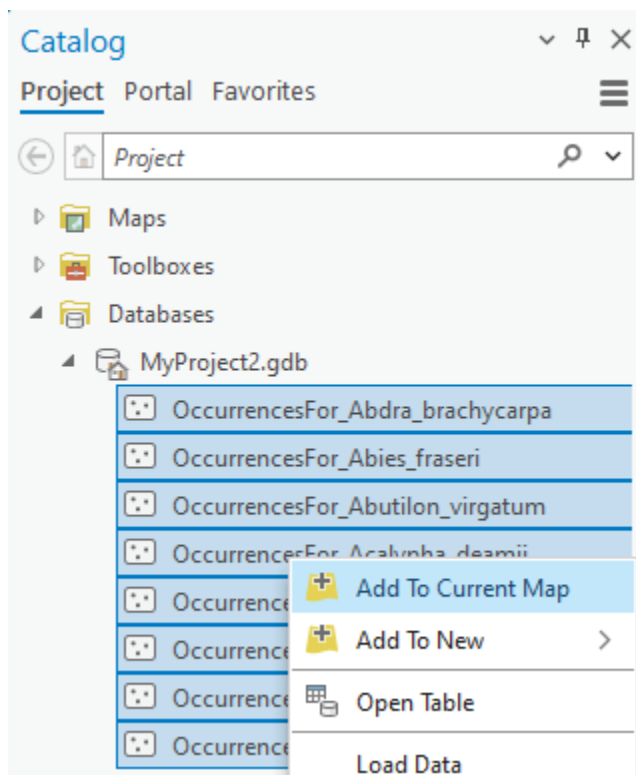
6. Open the Export Features element by right clicking and selecting open.
 - a. For Output Feature Class type “Occurrences_%%Value%”. The “%%Value%” command tells the model to pull in the name from the selected scientific name field from step 3 when creating and naming each unique layer. This ensures that each new layer will contain its scientific name (e.g. “Occurrences_Acer_rubrum”, “Occurrence_Abies_fraseri”, etc.).
 - b. Do not edit Input Features. This was determined when dragging the arrow to connect the Iterate Feature Selection element and the Export Features element.
 - c. Select OK.



7. Your model should now look something like this.



8. Navigate to the ribbon at the top of the page and under the ModelBuilder tab in the Run group select Run. This should start your model, and it will begin creating new occurrence layers for each species. Depending on how many occurrences records are in your file and the processing speed of your device this may take several hours (e.g. when running this model with 200,000 occurrence records this step took over 24 hours).
9. Once the model has finished running, in the Catalog pane, under the Project tab expand Databases and expand your project geodatabase. Select all newly created occurrences layers, right click, and select Add To Current Map (Note: if the new occurrence layers are not immediately showing up you may need to refresh the geodatabase by right clicking and selecting refresh).

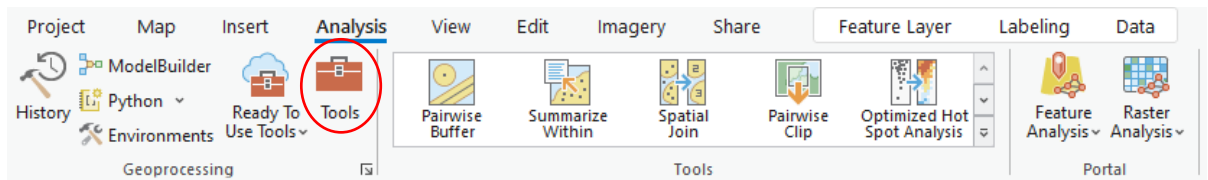


Calculating Range Extent

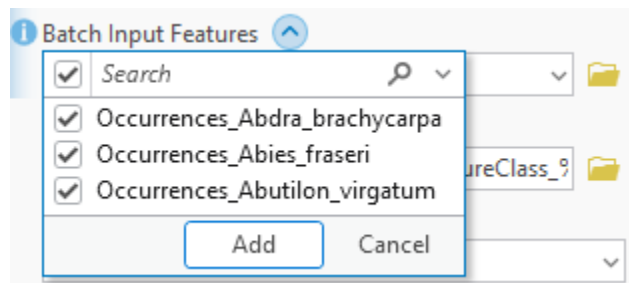
Range extent is defined as the smallest possible boundary that encompasses all known occurrence points of a taxon within the area of interest (Master et al. 2012). The NatureServe rank calculator uses square kilometers as the unit of measurement for this criterion.

Step 1: Use the *Minimum Bounding Geometry* tool to map species range.

1. Open the geoprocessing pane by navigating to the ribbon at the top of the page. Under the Analysis tab in the Geoprocessing group select Tools.



2. In the geoprocessing pane search for the *Minimum Bounding Geometry* tool.
 - a. For single species processing: Select and open the tool. For Input Feature select the species occurrence layer.
 - b. For batch processing: Right click the tool and select Batch. Ensure Input Features is selected as the batch parameter and hit Next. Next to Batch Input Features click the down arrow and select all individual species occurrence layers. Click Add.



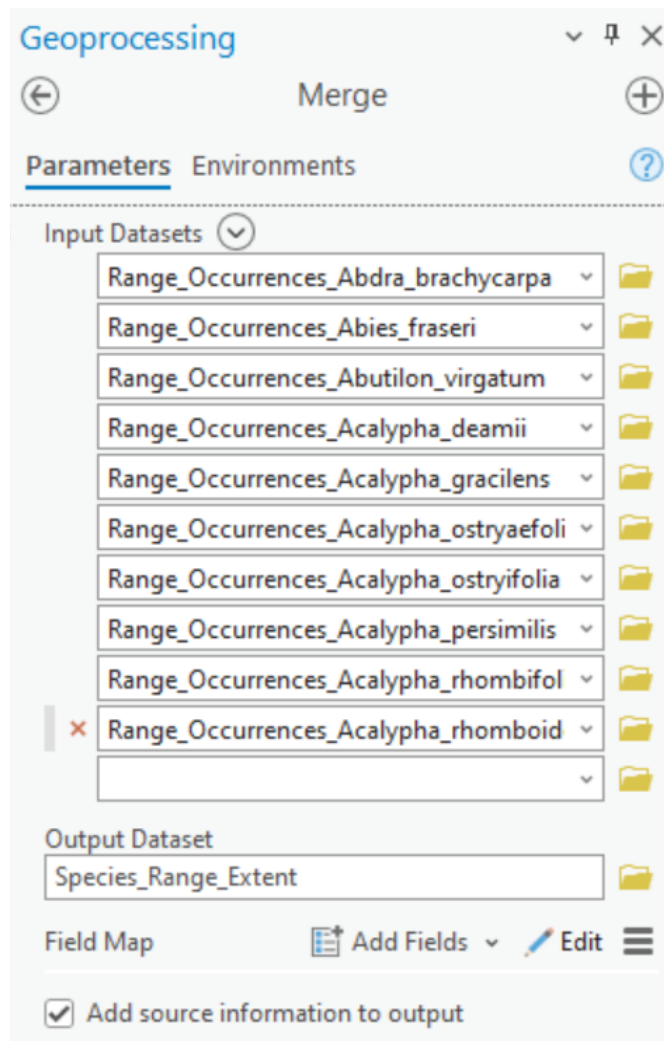
3. Name the output feature class “Range_ %Name%”. The “%Name%” command will be replaced by the name of each unique occurrence layer. This ensures that each output feature class contains the species name.
4. For Geometry Type select Convex hull.
5. For Group Option select All.
6. Click Run. This may take several hours depending on the number of layers being run.

The screenshot shows the 'Geoprocessing' window with the tool 'Batch Minimum Bounding Geometry' selected. The 'Parameters' tab is active. The 'Batch Input Features' list contains ten species occurrence layers: Occurrences_Abdra_brachycarpa, Occurrences_Abies_fraseri, Occurrences_Abutilon_virgatum, Occurrences_Acalypha_deamii, Occurrences_Acalypha_gracilens, Occurrences_Acalypha_ostryaefolia, Occurrences_Acalypha_ostryifolia, Occurrences_Acalypha_persimilis, Occurrences_Acalypha_rhombifolia, and Occurrences_Acalypha_rhomboides. The 'Output Feature Class' is set to 'Range_%Name%'. The 'Geometry Type' is set to 'Convex hull'. The 'Group Option' is set to 'All'. The 'Group Field(s)' field is empty. The checkbox 'Add geometry characteristics as attributes to output' is unchecked.

7. For single species processing: open the attribute table of the newly created species range layer and range extent in square meters will be listed under the “Shape_Area” column (Note: this will need to be converted into square kilometers before uploading to the NatureServe rank calculator).
8. For batch processing proceed to step 2 below.

Step 2: Use the *Merge* tool to create a single table listing range extent for all species (only necessary for batch processing).

1. In the geoprocessing pane search for and open the *Merge* tool.
2. Next to Input Datasets click the down arrow and select all individual range layers created in step 1. Click Add.
3. Name your Output Dataset.
4. Ensure the Add source information to output box **is checked**. This ensures that each file name, and therefore species name, is included in the output table along with its range.
5. Click Run.



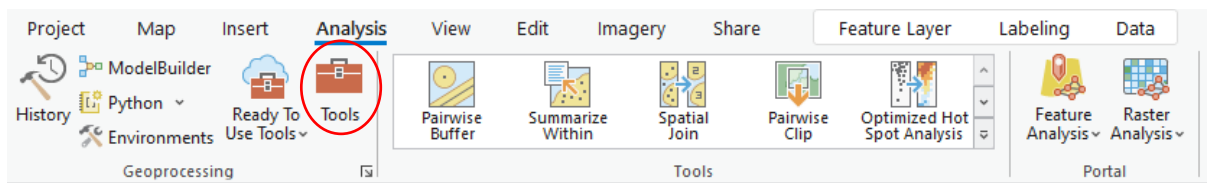
6. Open the attribute table of the new Output Dataset. Range extent in square meters for each species will be listed under the “Shape_Area” column and the input file name which contains species name will be listed under the “MERGE_SRC” column (Note: range will need to be converted into square kilometers before uploading to the NatureServe rank calculator).

Calculating Number of Occurrences

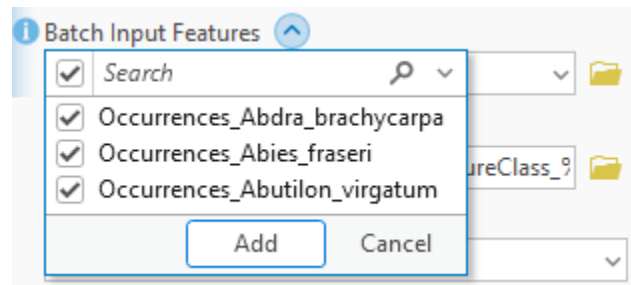
Number of occurrences (often referred to as Element Occurrences or EO) is the number of “on-the-ground” locations where a taxon is present (Master et al. 2012). To eliminate the effect of including multiple occurrence records of the same “on-the-ground” location, it is recommended that a minimum of 1 km separation distance be present between occurrences. If records are less than 1 km apart, they should be counted as a single occurrence (NatureServe 2020a).

Step 1: Use the *Buffer* tool to create a 500 m buffer around each point.

1. Open the geoprocessing pane by navigating to the ribbon at the top of the page. Under the Analysis tab in the Geoprocessing group select Tools.



2. In the geoprocessing pane search for the *Buffer* tool.
 - a. For single species processing: Select and open the tool. For Input Feature select the species occurrence layer.
 - b. For batch processing: Right click the tool and select Batch. Ensure Input Features is selected as the batch parameter and hit Next. Next to Batch Input Features click the down arrow and select all individual species occurrence layers. Click Add.



3. Name your output feature class “Buffer_ %Name%”. The “%Name%” command will be replaced by the name of each unique occurrence layer. This ensures the output feature class contains the species name.
4. For Distance enter 500 and as the unit select Meters.
5. Ensure the following parameters are selected.
 - a. Side Type: Full
 - b. End Type: Round
 - c. Dissolve Type: No Dissolve
 - d. Method: Planar
6. Click Run. This may take several hours depending on the number of occurrence layers being run.

Geoprocessing ⌵ 🔍 ✕

⬅ **Batch Buffer** ⊕²

Parameters Environments ?

Batch Input Features ⌵

Occurrences_Abdra_brachycarpa	⌵	📁	✎	⌵
Occurrences_Abies_fraseri	⌵	📁	✎	⌵
Occurrences_Acalypha_deamii	⌵	📁	✎	⌵
✕ Occurrences_Acalypha_gracilens	⌵	📁	✎	⌵
	⌵	📁	✎	⌵

Output Feature Class

Buffer_ %Name% 📁

Distance [value or field] Linear Unit ⌵

500 Meters ⌵

Side Type

Full ⌵

End Type

Round ⌵

Dissolve Type

No Dissolve ⌵

Dissolve Field(s) ⌵ ⚙

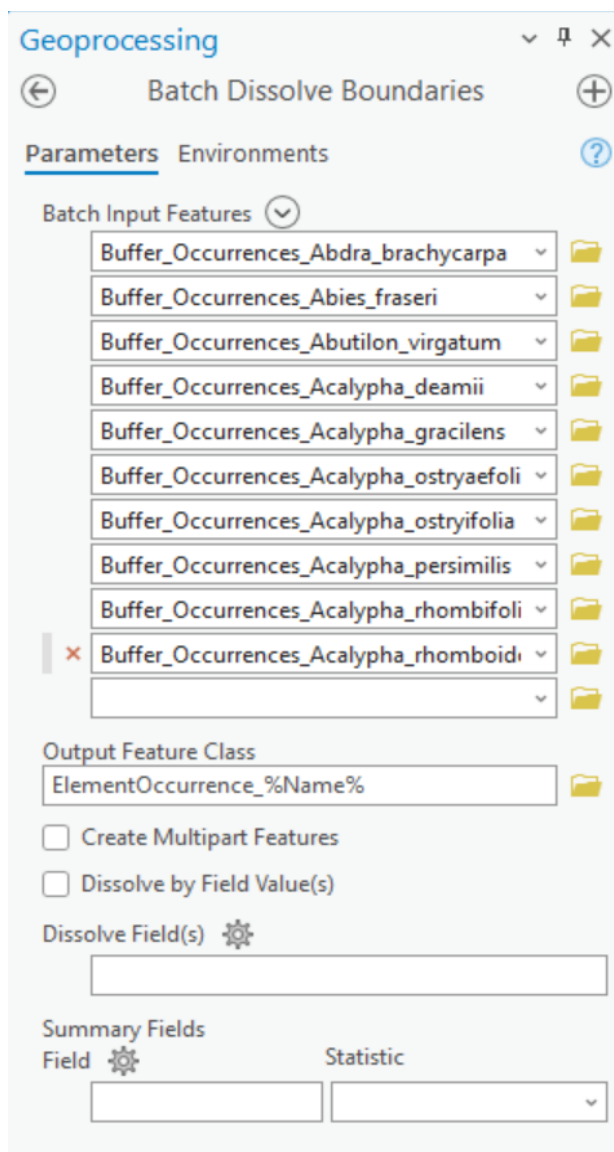
⌵

Method

Planar ⌵

Step 2: Use the *Dissolve Boundaries* tool to merge occurrence points with overlapping boundaries into a single point.

1. In the geoprocessing pane search for the *Dissolve Boundaries* tool.
 - a. For single species processing: Select and open the tool. For Input Feature select the buffer layer created in step 1.
 - b. For batch processing: Right click the tool and select Batch. Ensure Input Features is selected as the batch parameter and hit Next. Next to Batch Input Features click the down arrow and select all individual buffer layers created in step 1. Click Add.
2. Name the output feature class “ElementOccurence_ %Name%”.
3. Ensure all other parameters are left blank and boxes left unchecked.
4. Click Run. This may take several hours depending on the number of layers being run.



Step 3: Use the *Summary Statistics* tool to count the number of occurrences for each species.

1. In the geoprocessing pane search for the *Summary Statistics* tool.
 - a. For single species processing: Select and open the tool. For Input Table select the dissolved buffer layer created in step 2.
 - b. For batch processing: Right click the tool and select Batch. Ensure Input Table is selected as the batch parameter and hit Next. Next to Batch Table Features click the down arrow and select all individual dissolved buffer layers created in step 2. Click Add.
2. Name the output feature class “EO_Count_%Name%”.
3. Under Statistics Fields from the dropdown select the OBJECTID field and as the Statistic Type select Count.
4. Ensure all other parameters are left blank.
5. Click Run. This may take several hours depending on the number of layers being run.

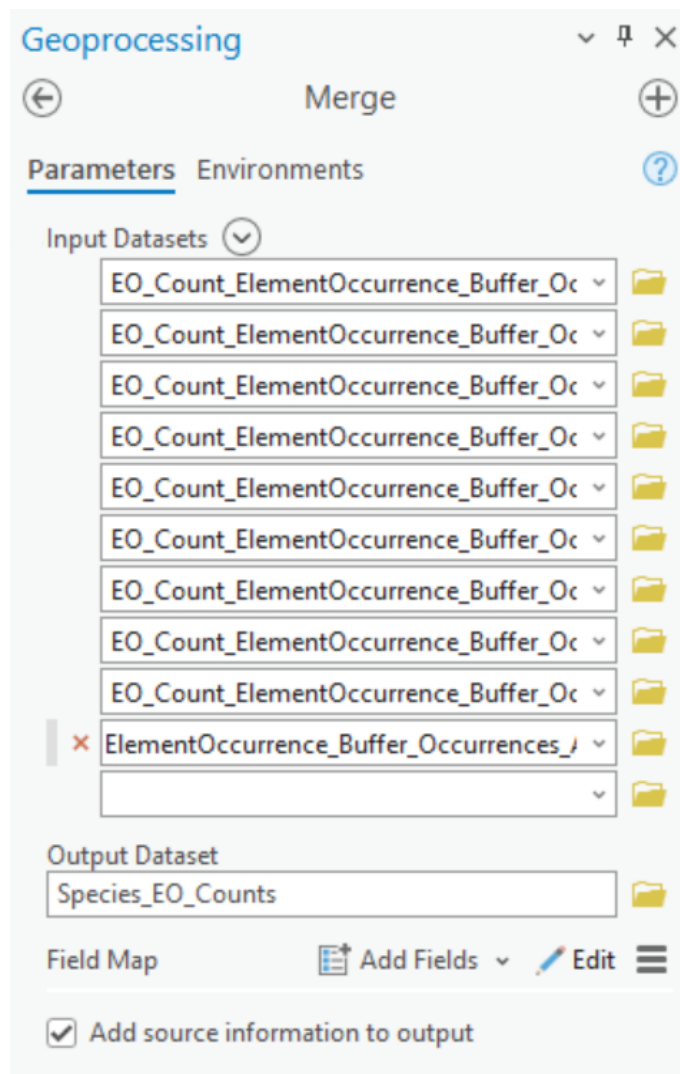
The screenshot shows the 'Geoprocessing' window with the 'Batch Summary Statistics' tool selected. The 'Parameters' tab is active. The 'Batch Input Table' section has a dropdown menu with four entries, all labeled 'ElementOccurrence_Buffer_Occurrences_'. The 'Output Table' section has a text box containing 'EO_Count_%Name%'. The 'Statistics Fields' section has a table with two columns: 'Field' and 'Statistic Type'. The first row has 'OBJECTID' in the 'Field' column and 'COUNT' in the 'Statistic Type' column. The 'Case Fields' section has a dropdown menu. The 'Concatenation Separator' section has a text box.

Field	Statistic Type
OBJECTID	COUNT

6. For single species processing: open the attribute table of the newly created occurrence count layer and the number of occurrences will be listed under the FREQUENCY column.
7. For batch processing proceed to step 4 below.

Step 4: Use the *Merge* tool to create a single table listing the number of occurrences for all species (only necessary for batch processing).

1. In the geoprocessing pane search for and open the *Merge* tool.
2. Next to Input Datasets click the down arrow and select all individual occurrence count layers created in step 3. Click Add.
3. Name the Output Dataset.
4. Ensure the Add source information to output box **is checked**. This ensures that each file name, and therefore species name, is included in the output table along with its number of occurrences.
5. Click Run.



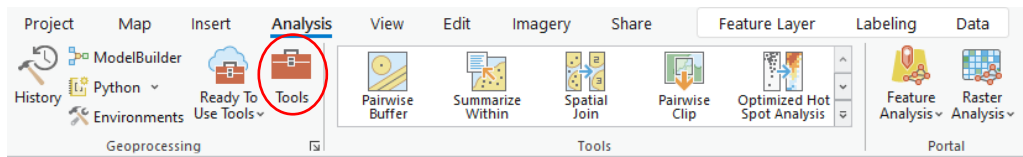
6. Open the new attribute table created. Number of occurrences for each species will be listed under the “FREQUENCY” column and the input file name which contains species name will be listed under the “MERGE_SRC” column.

Calculating Area of Occupancy

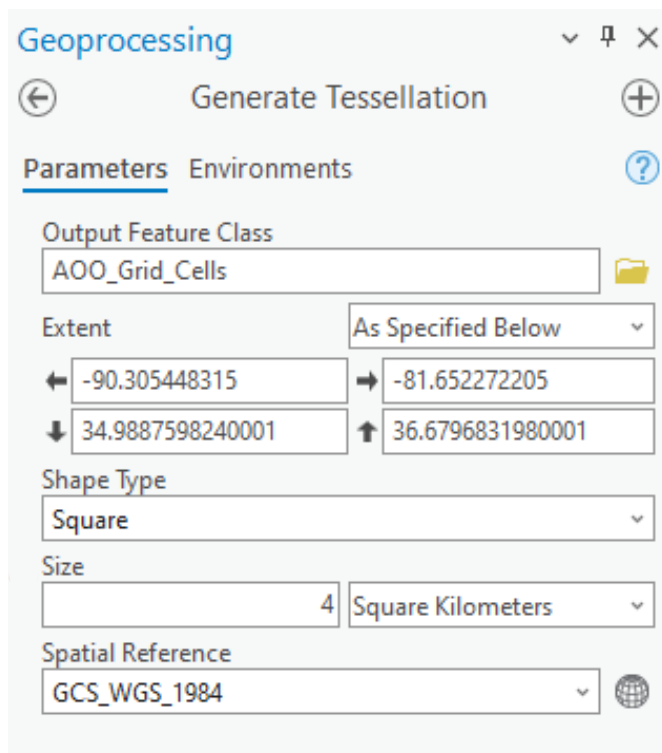
Area of occupancy (AOO) is the area within its range extent that is actually occupied by the taxon. It is calculated by counting the number of 2 x 2 km grid cells that contain an occurrence record. This reflects that a taxon may not occupy the entirety of its range extent due to unsuitable or unoccupied habitat (Master et al. 2012).

Step 1: Use the *Generate Tessellation* tool to create a layer of 2 x 2 km grid cells.

1. Open the geoprocessing pane by navigating to the ribbon at the top of the page. Under the Analysis tab in the Geoprocessing group select Tools.

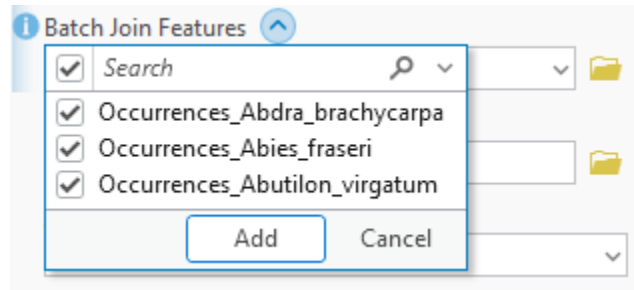


2. In the geoprocessing pane search for and open the *Generate Tessellation* tool.
3. Name the Output Feature Class “AOO_Grid_Cells”.
4. For Extent click the dropdown and select the layer containing the boundary for your area of interest. Extent will auto populate based on this layer.
5. For Shape Type select Square.
6. For Size enter 4 and select Square Kilometers as the unit.
7. Click Run.



Step 2: Use the *Spatial Join* tool to create a layer of occupied 2 x 2 km grid cells for each species.

1. In the geoprocessing pane search for the *Spatial Join* tool.
 - a. For single species processing: Select and open the tool. For Join Feature select the species occurrence layer.
 - b. For batch processing: Right click the tool and select Batch. For the batch parameter choose Join Features and hit Next. Next to Batch Join Features click the down arrow and select all individual species occurrence layers. Click Add.



2. For Target Features select the 2 x 2 km grid cells created in step 1.

3. Name your output feature class “SpatialJoin_ %Name%”. The “%Name%” command will be replaced by the name of each unique occurrence layer. This ensures the output feature class contains the species name.
4. For Join Operation select Join one to one.
5. Ensure the Keep All Target Features box is **unchecked**.
6. For Match Option select Intersect.
7. Ensure all other parameters are left blank.
8. Click Run. This may take several hours depending on the number of layers being run.

Geoprocessing Batch Spatial Join2

Parameters Environments

Target Features
AOO_Grid_Cells

Batch Join Features

- Occurrences_Abdra_brachycarpa
- Occurrences_Abies_fraseri
- Occurrences_Abutilon_virgatum
- Occurrences_Acalypha_deamii
- Occurrences_Acalypha_gracilens
- Occurrences_Acalypha_ostryaefolia
- Occurrences_Acalypha_ostryifolia
- Occurrences_Acalypha_persimilis
- Occurrences_Acalypha_rhombifolia
- Occurrences_Acalypha_rhomboidea

Output Feature Class
SpatialJoin_%Name%

Join Operation
Join one to one

☐ Keep All Target Features

Match Option
Intersect

Search Radius
Decimal Degrees

Distance Field Name

> Fields

> Attribute Matching

Step 3: Use the *Summary Statistics* tool to count the number of occupied 2 x 2 km grid cells for each species.

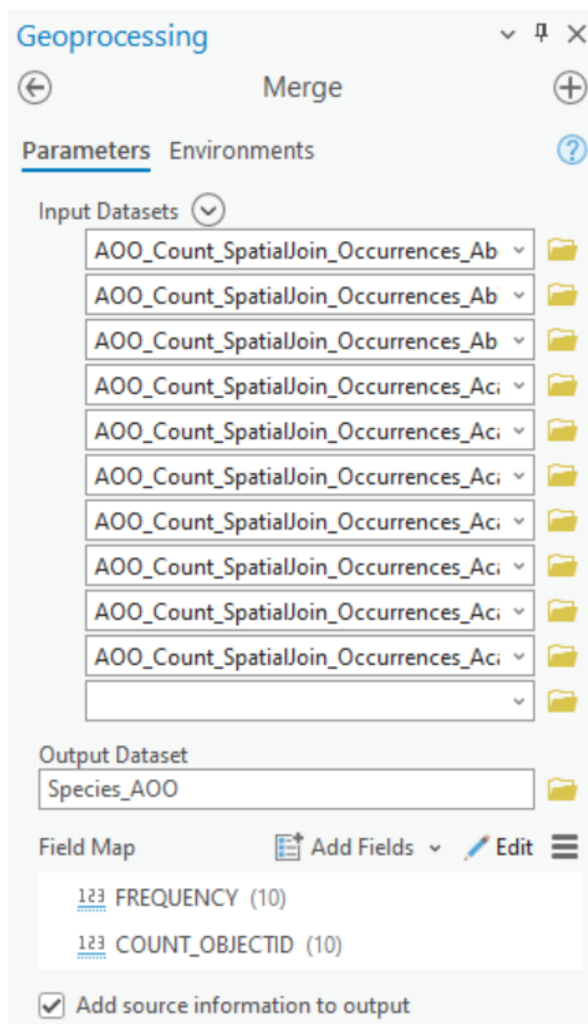
1. In the geoprocessing pane search for the *Summary Statistics* tool.
 - a. For single species processing: Select and open the tool. For Input Table select the spatial join layer created in step 2.
 - b. For batch processing: Right click the tool and select Batch. Ensure Input Table is selected as the batch parameter and hit Next. Next to Batch Input Table click the down arrow and select all the individual spatial join layers created in step 2. Click Add.
2. Name the output feature class “AOO_Count_%%Name%”.
3. Under Statistics Fields from the dropdown select the OBJECTID field and as the Statistic Type select Count.
4. Ensure all other parameters are left blank.
5. Click Run. This may take several hours depending on the number of layers being run.

The screenshot shows the 'Geoprocessing' pane with the tool 'Batch Summary Statistics1' selected. The 'Parameters' tab is active. The 'Batch Input Table' dropdown is expanded, showing a list of spatial join layers for various species: *Abdra brachyca*, *Abies fraseri*, *Abutilon virgatum*, *Acalypha deamii*, *Acalypha gracilis*, *Acalypha ostrya*, *Acalypha ostryifolia*, *Acalypha persinii*, *Acalypha rhomboides*, and *Acalypha rhomboides*. The 'Output Table' is set to 'AOO_Count_%%Name%'. Under 'Statistics Fields', the 'Field' is 'OBJECTID' and the 'Statistic Type' is 'Count'. The 'Case Field' is empty, and the 'Concatenation Separator' is also empty.

6. For single species processing: open the attribute table of the newly created area of occupancy count layer and area of occupancy will be listed under the “FREQUENCY” column.
7. For batch processing proceed to step 4 below.

Step 4: Use the *Merge* tool to create a single table listing area of occupancy for all species (only necessary for batch processing).

1. In the geoprocessing pane search for and open the *Merge* tool.
2. Next to Input Datasets click the down arrow and select all individual area of occupancy count layers created in step 3. Click Add.
3. Name the Output Dataset.
4. Ensure the Add source information to output box **is checked**. This ensures that each file name, and therefore species name, is included in the output table along with its area of occupancy.
5. Click Run.



6. Open the new attribute table created. Area of occupancy for each species will be listed under the “FREQUENCY” column and the input file name which contains species name will be listed under the “MERGE_SRC” column.

References

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<https://www.natureserve.org/conservation-tools/conservation-rank-calculator>
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