TEMPLATE_DataFields_Vouchers_BioAssets **Form Submitted** Data Uploaded to Symbiota Data Uploaded to Research DB with occurrenceID & onlineInformation added **Specimens Processed and Accessioned** catalogNumber and image info **EXPORTED from Symbiota and UPDATED in Research DB**

This workflow illustrates the process to produce complete data records (specimen and tissue) that can be exported from the Research Database and Uploaded to GBIF.

based on occurrenceID OR recordedBY and recordNumber