Import of sample data to Corema - Avian samples

Basic sample data

Data common to all items originating from the same "sampling event" should be entered in the yellow (taxon), green ("collection event", individual data) and orange (locality) coloumns. Try to fill in as much information as possible, but at least the shaded coloumns should be filled in.

For many of the coloumns (fields) hints and comments on how to fill them in are included as comments to the header cells (indicated by the small red triangle in the upper right corner of the cells). You should also read the information in the *Info* sheet before you start filling in your data.

Actual sample data

The actual, physical samples should be registered as items belonging to a specific accession. Given that an accession typically corresponds to one sampling event of one individual, e.g. a blood sample, a sperm sample and a sperm motility recording sampled from the same bird during one sampling event would all be registered as separate items under the same accession. If, on the other hand, the same bird was recaptured later the same or some other day, that would typically represent a new sampling event and thus also a new accession with new items.

The import template is by default set up for three items. To register more items, just select all the blue coloumns for one of the items (e.g. all coloumns ending with "…_1") and add as many copies of these as needed before the *Associated documents* coloumn. Make sure to change the trailing number (e.g. "…_1") to "…_4", "…_5" etc. in BOTH header rows (row 1 and 2) for the newly added coloumns.
You don't have to have the same number of items for all accessions, just make sure to fill in the coloumns from the left so that you use e.g. item columns 1-3 and not 1, 3 and 5 to register 3 items.

Some comments on data entry:
- Remember to fill in the status of the samples (hopefully mostly "OK" but still…)
- Fill in the date and collector (person) coloumns also for the items
- Remember to indicate type of tube/container and medium in which the sample is stored
- Remember to fill in any relevant restrictions on use of the samples – if none apply use the "None" option!

The following item types can currently be registered (contact the Corema administrator if you need to register other types of samples):
- Blood
- Tissue
- Feather
- Egg
- Sperm
- Sperm motility recording
- Testicle
- DNA extract
- Sperm morphology slide
- Preservation (skins etc.)