

# **ChEMBL** Deposition quick checklist

This checklist has been designed to catch the most common errors in deposited ChEMBL data. If you check your data against this list and <u>the example dataset here</u>, it will substantially reduce the risk that your data will require corrections.

## **Depositor-Defined Identifiers**:

- □ Do your RIDX, AIDX and CIDXs all give a meaningful unique identifier to each reference compound and assay in their datasets?
- □ Is every RIDX a unique, non-redundant reference to a single dataset or publication?
- Does every CIDX refer to a unique compound?
- Does every AIDX refer to a unique assay?

### ASSAY:

- Does every Assay have an RIDX that links to a REFERENCE in this dataset?
- Does every assay have a description, assay\_organism and assay\_tax\_ID?
- □ Are variants, isoforms and other mutations captured within the assay description?
- ☐ Are cell-lines, tissues and subcellular fractions captured in the corresponding assay\_cell\_type, assay\_tissue and assay\_subcellular\_fraction columns?
- □ For functional cell-based assays, has the target protein been recorded as well as the cell-line?

#### ASSAY PARAMETERS:

☐ Are there assay parameters that should be captured as part of the assays (e.g. cell-lines, concentrations, tissues can be captured as part of the assays table)?.

### ACTIVITY:

- Does every CRIDX match a valid RIDX in this dataset?
- Does every ACTIVITY have a VALUE or TEXT VALUE?
- Does every VALUE have a RELATION and UNIT field?
- Does every TEXT\_VALUE have a blank RELATION field?
- Are any TEXT\_VALUES actually a VALUE plus a RELATION, and therefore should be in these fields?

### **ACTIVITY\_PROPERTIES**:

- □ Do the ACT\_IDs in ACTIVITY match to the ACT\_IDs for the relevant ACTIVITY\_PROPERTIES and vice versa?
- Does every ACTIVITY have a VALUE or TEXT VALUE?
- Does every VALUE have a RELATION and UNIT field?
- Does every TEXT\_VALUE have a blank RELATION field?
- Are any TEXT\_VALUES actually a VALUE plus a RELATION, and therefore should be in these fields?



## **REFERENCE**:

- Does every Reference have a TITLE, AUTHORS and ABSTRACT?
- □ Do all DATASETS have a title, authors and an abstract that provides an easy to understand explanation for database users?

## COMPOUND\_CTAB:

- □ Is every CTAB valid? You can check using the 'check' endpoint of the <u>ChEMBL APIs</u> <u>here</u>.
- □ Is every CIDX in the SDFile a CIDX that is in the COMPOUND\_RECORDS file?

## COMPOUND\_RECORDS:

- □ Have both a compound\_name and a compound\_key (this is often a depositor identifier) been supplied?
- □ Is there a valid RIDX for each compound, linking them to an RIDX in this dataset?