**PINOT update version 1.1 (implemented 04/01/2021)**

**Update:** HGNC complete list of human genes with multiple ID types, used in the ID conversion step of the pipeline, updated (downloaded on 29.09.2020). Note, approved names for genes/proteins may have changed in accordance with this release from the HGNC.

**Update:** *C. elegans* protein interaction data updated, derived from the Alliance of Genome Resources release 3.1.1. This updated dataset contains 36,768 *C. elegans* PPIs, 6994 more interactions (over 23% more) than available in PINOT version 1.0. The updated difference between querying this dataset using lenient vs stringent filters is 575 interactions.

**Update:** Two new detection method IDs have been added to the method code reassignment step. MI:0314 (complex) and MI:0403 (colocalization) are both reassigned to “General” when using the lenient filter or “UNSPM” when using the stringent filter.

**Bug fixed:** We noticed that in some rare cases where only 1 PPI was available for a single queried protein, the corresponding interactor IDs (SwissProt & Entrez) were in the incorrect orientation in relation to interactor A and interactor B. This has been fixed with a modification to the “FirstPosition.R” file.

**Bug fixed:** Previously within the method code reassignment step, detection method ID MI:0004 (affinity chromatography technology) was reassigned to “Chromatography”, this is now reassigned to “CoIP” instead.