

Release Notes for BAO version 2.8.12

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About BAO

The BioAssay Ontology (BAO) has been developed to formally describe biological screening assays and their results including high-throughput screening (HTS) data; specifically in the domain of small molecule drug and probe development. BAO enables categorization of assays and results based on several concepts that are important to interpret and analyze screening data with the goal to infer the mechanism of action of small molecules based on the known aggregate screening results from many assays.

Changes in BAO v 2.8.12

[Github Milestone BAO v 2.8.12](#)

<https://github.com/BioAssayOntology/BAO/issues/120>

doubled: "cell per microliter" #120

<https://github.com/BioAssayOntology/BAO/issues/124>

update UBERON module #124

<https://github.com/BioAssayOntology/BAO/issues/125>

doubled up Cells per well #125

Public Location:

<http://www.bioassayontology.org/bao>

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