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component [filename] [2]individualID [3]specimenID [4]MultiSpecimen [5]assay [6]fileFormat [7]consortium [8]study [9]ant [10]sourceType [11]staType [12]itaSubtype [13]etadateType [14]analysisType [15]modeSystem [16]chromosome [17]ecies [18]

Table with 18 columns and multiple rows. Columns are labeled with codes and descriptions: component, filename, individualID, specimenID, MultiSpecimen, assay, fileFormat, consortium, study, ant, sourceType, staType, itaSubtype, etadateType, analysisType, modeSystem, chromosome, ecies.





- [1] TBD
- [2] Name of a file
- [3] Identifying string linked to the individual or animal being studied
- [4] Identifying string linked to a particular sample or specimen
- [5] Boolean flag indicating whether or not a file has data for multiple specimens
- [6] The technology used to generate the data in this file
- [7] Defined format of the data file, typically corresponding to extension, but sometimes indicating more general group of files produced by the same tool or software
- [8] The name of the consortium
- [9] study
- [10] Grant number including activity code, institute code, and serial number (e.g. U01MH103392)
- [11] The type of resource being stored and annotated
- [12] Types of input/output data in bioinformatics pipelines
- [13] Further qualification of dataType, which may be used to indicate the state of processing of the data, aggregation of the data, or presence of metadata.
- [14] metadata, a description of the type of metadata in the file.
- [15] Type of analysis
- [16] TBD
- [17] A structure composed of a very long molecule of DNA and associated proteins (e.g. histones) that carries hereditary information; number or designation of the particular chromosome that the data is associated with.
- [18] The name of a species (typically a taxonomic group) of organism.