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omponent [1	ilename [2	dividualID [3	pecimenID [4	MultiSpecimen [5	ssay [6	leFormat [7	onsortium [8	tudy [9	ant [1	sourceType [1	taType [1	taSubtype [1	etadataType [1	alysisType [1	ModelSystem [1	romosome [1	ecies [18]

omponent [1	ilename [2	dividualID [3	pecimenID [4	MultiSpecimen [5	ssay [6	leFormat [7	onsortium [8	tudy [9	ant [1	isourceType [1	taType [1	taSubtype [1	etadataType [1	alysisType [1	ModelSystem [1	romosome [1	ecies [18]

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omponent [1	ilename [2	dividualID [3	pecimenID [4	MultiSpecimen [5	ssay [6	leFormat [7	onsortium [8	tudy [9	ant [1	sourceType [1	taType [1	taSubtype [1	etadataType [1	alysisType [1	ModelSystem [1	romosome [1	ecies [18]							

[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.