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- [1] Type of metadata template; provide the same one for all items/rows.
- [2] The name of the file.
- [3] 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
- [4] The type of resource being stored and annotated
- [5] Links an entity to data types that the entity represents/contains. This is closely tied to the assay property. For example, a file of dataType `genomicVariants` might have an assay value of `whole genome sequencing`.
- [6] 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.
- [7] The technology used to generate the data in this file.
- [8] A sequencing platform, microscope, spectroscope/plate reader, or other platform for collecting data.
- [9] A unique identifier (non-PII) that represents the subspecimen (subsample) from which the data came, e. g. an ID that distinguishes between different parts of the same parent tumor specimen.
- [10] Type of material in the characterization
- [11] Numeric value for concentration of the material
- [12] Unit used for the material concentration, e.g. mg/mL
- [13] Numeric value for NaCl concentration
- [14] Unit used for the NaCl concentration, e.g. mM
- [15] Numeric value for pH (range 0-14)