



*Individuals from Family datasets have Pedigree role assigned among their attributes

A Variant can be found in may Run-Analysis from same or different Biosamples/Individuals/ Datasets

**Interactors are Individual/ entities which are not sequenced in study. Only its metadata/ phenotypic data was collected in relation to the main (sequenced) individual. Interactor is characterized by relation type with the main (sequenced) Individual such as Pathogen, Host, etc

m samples can be in 1 "Run-Analysis" when 1 analysis includes many Runs (Pedigree multi-VCF). Otherwise, 1 biosample can be in n run-analysis as many times as runs where run for sample x analysis performed on same run, leading possibly different variants called

