AQ1: Give the project name or acronym to be used

AQ2 what kind of data do you handle

Selection: image, RNAseq, DNASeq, other NGS, Metabolomics, Proteomics, targeted assays (e.g. glucose and fructose content), models, code, excel, cloned DNA

AQ3: how much data do you think you will generate

AQ4: are there any standards that are relevant for you [some are preselected]

MIAPPE, MinSeqE,

AQ5: Are you going to use DataPLANT ARC

If yes 🡺 end

AQ5: Where will you submit your data as endpoints [some answers are preselected]

ENA, GEO, NCBI, genbank, pride, proteomexchange, zenodo, dataDyrad

AQ6: Will you adhere to any high level metadata submission standards

Dublin Core

AQ6: When will you make your data public?

AQ6b If late🡺 are there intellectual property (IP) issues to adhere to (e.g. patents you might want to file) or private partners involved whose IP has to be taken into account

AQ7: