AR Lab Network Form for Phylogenetic Tree-level Mycotics Reporting (C. auris Whole Genome Sequencing)

Data element Name	Data element Definition
tree_id	Tree ID or REDCap IDUnique ID for this record. Use a tree ID if this has been established by your lab or is available. Please include your lab's two- or three-letter jurisdiction abbreviation as a prefix to the id (e.g., CDC-Tree1).
performingorgstate	ARLN PHL state or JurisdictionThe testing lab's state, territory, or jurisdiction.
performing_facility	Public health laboratory nameName of your public health laboratory.
redcap_reporting_date	REDCap reporting dateThe date this form was created and the data was initially reported in REDCap. Format: YYYY-MM-DD
last_update_date	Date this form was last updated. Please update this field every time the form is resubmitted. Format: YYYY-MM-DD
mycosnp_performed	Did your lab use MycoSNP for QC and SNP analysisIndicate whether MycoSNP was used for QC and SNP identification.
bioinformatic_tools_others	If other bioinformatics tools were used for QC and SNP analysis, specifyFor C. auris WGS only. If MycoSNP was not used, specify which other bioinformatics tools were used.
strain_reference	Reference sequence usedThe reference used (strain and genbank accession) in the bioinformatics analysis for whole genome sequencing data.
other_reference	Other referenceIf available, please provide the NCBI Accession (e.g., GenBank Accession) of the reference used.
tree_file	Phylogenetic tree file attachmentThe attachments files used as input for phylogenetic tree visualization tool. (e.g. .nwk, .tre)
tree_metadata_file	Phylogenetic tree metadata file attachmentThe metadata file used as input for the phylogenetic tree visualization tool (e.g., .csv)
tree_visualized_file	Visualized phylogenetic tree file attachmentA file attachment of the visualized tree, if a URL is unavailable.
tree_visualization_url	Visualized phylogenetic tree linkThe URL to the phylogenetic tree visualization (e.g., Microreact link).
tree_interpretations	Phylogenetic tree interpretationsPlease provide any tree interpretations.
tree_comments	Additional comments on the phylogenetic treePlease share any additional comments related to the phylogenetic tree.