

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

<u>Data element Name</u>	<u>Data element Definition</u>
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 referencelf 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.

Public reporting burden of this collection of information is estimated to average 6 minutes per response, including the time for reviewing instructions, searching existing data sources, gathering and maintaining the data needed, and completing and reviewing the collection of information. An agency may not conduct or sponsor, and a person is not required to respond to a collection of information unless it displays a currently valid OMB Control Number. Send comments regarding this burden estimate or any other aspect of this collection of information, including suggestions for reducing this burden to CDC/ATSDR Reports Clearance Officer, 1600 Clifton Road NE, MS H21-8, Atlanta, Georgia 30333; ATTN: PRA 0920-1310