

Validation analysis performed on 2023-12-12.

	Number of samples populated in table1_with_files.tsv	Number of samples populated in table2_with_files.tsv	Number of differences (exact match)	Validation Criteria	Number of samples failing the validation criteria
amrfinderplus_amr_core_genes	4	5	5	SET	1
assembly_length	5	5	5	PERCENT_DIFF: 1.00%	1
extra_column	2	2	2	IGNORE	0
file_column	4	3	3	EXACT	3
gambit_predicted_taxon	5	5	1	EXACT	1
sort_file_column	4	3	3	SET	2

Validation Criteria:

- EXACT
 - Performs an exact string match
- IGNORE
 - Ignores the values; indicates 0 failures
- SET
 - Compares items in a list without regard to order
- PERCENT_DIFF
 - Tests if two values are more than the indicated percent difference (must be in decimal format)