

Validation analysis performed on 2025-03-13.

	Number of samples populated in example-table1.tsv	Number of samples populated in example-table2.tsv	Number of differences (exact match)	Validation Criteria	Number of samples failing the validation criteria
busco_results	5	5	4	SET,2	0
assembly_length	5	5	5	0.01,500	0
gambit_predicted_taxon	5	5	1	EXACT	1
amrfinderplus_amr_core_genes	4	5	5	SET	2
extra_column	2	2	2	IGNORE	0

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)
- RANGE
 - Tests if two values are more than +- the indicated number (must be in integer format)
- COMBINED
 - Tests both options and returns a pass for that row if at least one criteria is met