

Validation analysis performed on 2025-04-02.

	Number of populated cells (table1/table2)	Number of differences (exact match)	Validation Criteria	Number of samples requiring further investigation	Notes
busco_results	5/5	4	SET,2	1	
assembly_length	Discordant: 5/4	5	0.01,500	2	
gambit_predicted_taxon	5/5	1	EXACT	1	
amrfinderplus_amr_core_genes	Discordant: 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