

Validation analysis performed on 2025-03-27.

| | 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