

Table 1: List of resource name discrepancies.

Summary of data file identification discrepancies found between the Galaxy instance set up at BGI and the original assemblathon/GAGE data used in manuscript by [1].

Galaxy Server file names	original GAGE dataset file names
saureus_B.L4.37bp3500IS45X.2.fq	http://gage.cbc.umd.edu/data/Staphylococcus_aureus/Data.original/frag.1.fastq.gz
saureus_B.L3.37bp3500IS45X.1.fq	http://gage.cbc.umd.edu/data/Staphylococcus_aureus/Data.original/shortjump.1.fastq.gz
saureus_A.L2.101bp180IS45X.2.fq	http://gage.cbc.umd.edu/data/Staphylococcus_aureus/Data.original/frag.2.fastq.gz
saureus_A.L1.101bp180IS45X.1.fq	http://gage.cbc.umd.edu/data/Staphylococcus_aureus/Data.original/shortjump.2.fastq.gz

References

- [1] Luo, *et al.* SOAPdenovo2: an empirically improved memory-efficient short-read de novo assembler. GigaScience. 2012 Dec;1(1):18. Available from: <http://dx.doi.org/10.1186/2047-217X-1-18>.