**Table S2. Scripts used to estimate quality metrics and downsample WGS data files.**

|  |  |  |  |
| --- | --- | --- | --- |
| Script | Programmatic Dependencies | Source | Parameters |
| run\_assembly\_readMetrics.pl*for calculating coverage and PHRED scores* | perl-5.16.1-MT | github.com/lskatz/CG-Pipeline  | Coverage: -e 5700000 for *E. coli* O26, -e 4591637 for *S. enterica* ser. Reading, -e 4490000 for *S. enterica* ser. Pomona, and -e 4839734 for *Shigella sonnei* |
| countReadsWithAmbig.py | perl/5.22.1, python/3.4, and prinseq v0.20.3 | github.com/darlenewagner/NGS\_Multi\_Heal | --format integer |
| run\_assembly\_removeDuplicateReads.pl | perl-5.16.1-MT | github.com/lskatz/CG-Pipeline  | --downsample calculated based upon calculation: (desired coverage) ÷ (calculated coverage) |