|  |  |  |  |
| --- | --- | --- | --- |
|  | | **Read 1** | **Read 2** |
| **Raw reads** | | **2,296,478** | |
| **Sequence length (pb)** | | **150** | |
| **% of trimmed sequences** | **Vector and Illumina adaptors trimming** | **95.51** | **87.26** |
| **Circularization adaptor trimming** | **27.03** | **38.10** |
| **Average sequence length after trimming** | | **101** | **103** |
| **Useful paired end sequences after trimming** | | **1,438,662** | |

Sequencing metrics after in-house quality control on a BES library.