
R1: FastQC module status (11 modules)
R2: FastQC module status (11 modules)
Basic Statistics
Per base sequence quality
Per tile sequence quality
Per sequence quality scores
Per base sequence content
Per sequence GC content
Per base N content
Sequence Length Distribution
Sequence Duplication Levels
Overrepresented sequences
Adapter Content
PASS
WARN
FAIL
Raw
Adapter-only (cutadapt)
Q20 cut_right only (fastp)
fastp default