

Information Sheet Library Sequencing at Xpress Genomics

Dear Customer,

Please fill and return this information sheet with your library.

Customer Name	
Date	
Sequencer / Flow Cell Type	☐ G99 (80-100M reads) ☐ G400 (1600M reads) ☐ T7 (4-5B reads)
Library DNA	☐ dsDNA (requires conversion) ☐ circular ssDNA
Type of Library	☐ Amplicon ☐ RNA-seq/DNA-seq ☐ Other: _____
Concentration (Qubit)	ng/uL
Average Fragment Size	bp
Volume of submitted library	uL
Sequencing Primer (tick all applicable)	☐ MGI ☐ Illumina TruSeq ☐ Illumina Nextera ☐ Custom ¹
Read length layout	Read1: bp Read2: bp Barcode (i7): bp Dual Barcode (i5): bp
Demultiplexing of fastq	☐ No ☐ Yes ²

¹Please contact us to discuss feasibility of your custom sequencing primer setup.

²Please submit the barcode annotation for demultiplexing together with this information sheet.