Illumina Sample Drop Off Form	Date:
PI Name on account:	Name:
Account to be billed:	Phone Number:
	Email:

	Sample #1	Sample #2	Sample #3	Sample #4	Sample #5	Sample #6	Sample #7
Sample type?							
Are you submitting a library?							
Are we making a library? Bar-coded?							
Fragment size?*							
Tube type and sample name							
Sample Volume and Concentration (ng/ul or nM)							
If bar-coded samples, indicate Illumina or 5' bar-coding.							
Method of quantifying**							
Type of Run (SR or PE), # of cycles, and # of lanes							
Organism?							
Kind of sample (mRNA-seq, genomic, ChIP, etc)							
Special Instructions (eg, analysis prefs, priority, pooling required)							

<sup>\*</sup>For PE genomic DNA libraries, we recommend at least 350 bp. For SR libraries, we recommend 200 bp. For mRNA-seq, 200 bp is recommended, which may preclude long PE runs.

<sup>\*\*</sup>We require bioanalyzer traces and qPCR quantitation on all samples to ensure quality data. If you cannot provide, we will do for a fee.