	RNA Illumina NovaSeq Submis	
Please indicate special instructions (pooling schemes, combination of projects in one NOTE: lane, etc.) in the Comments section below. Email chains with Facility personnel will NOT qualify as evidence of instruction in case of disputes.		
	Date (mm/dd/yyyy)	
Contact Information	Principal Investigator	Principal Investigator Email / Phone
	Department	Cancer Center Member? Yes No
	Experiment Contact	Experiment Contact Email / Phone
	Billing Administrator	Billing Administrator Email / Phone
	Sample Prepara	tion and Delivery
	AB-1	ATCGA
Use 1.5 Eppendorf tubes Total RNA: 13 ul at 100ng/ul "Initials-number" Email Us with and Sample Details genomics@bsd.uchicago.edu 9:00am-4:00pm N Use dry ice for maile		
Project Information	Sample Species: Human Mouse Rat	Other:
	Number of Tubes Submitted: (minimum 12 samples)	Have the samples been treated with DNase? Yes No (Note: DNAse treatment mandatory for Ribo-Zero)
	Please Submit Excel Sheet Listing Sample Labels	
	Library Type & Sequencing: NovaSeq Special	
	PE100 – S1 flowcell PE100 – S ~30M clusters/sample ~60M clusters	mRNA directional S1 flowcell Sters/sample E reads/sample Ribo-Zero depletion PE100 – S1 flowcell ~60M clusters/sample ~120M PE read/smpl
	NOTE: additional charges will be applied for any dilutions, cleanups or DNAse treatments performed by the Core.	
**Please describe the source of the RNA (There is a big difference in processing RNA from quiescent cells versus tissue culture tumor cell lines in terms of mRNA content that affects the specifics of processing the samples on our end):		

