

FRED HUTCHINSON CANCER RESEARCH CENTER

A LIFE OF SCIENCE

Illumina Expression Arrays - **FHCRC Internal Request**

Contact/Billing Information

PI Name: _____ Project ID: _____
 Contact Name: _____ Signature: _____
 Contact Phone: _____ Date: _____
 FRED Account ID/E-mail: _____

Sample Information

Organism: _____ RNA Source: _____
 Array Name: _____ Extraction Method*: _____

*If a phenol-based extraction method is used, steps must be taken to ensure that there is no residual phenol contamination in the sample. Phenol strongly absorbs in the UV range and will distort RNA quantitation. Test for phenol: If the A₂₃₀ is approximately half of the A₂₆₀ reading, the sample is ok.

	Sample ID	RNA (ug/ul)**	A230/A260*	A260/A280
Chip 1	A:			
	B:			
	C:			
	D:			
	E:			
	F:			
	G:			
	H:			
Chip 2	A:			
	B:			
	C:			
	D:			
	E:			
	F:			
	G:			
	H:			

**Preferred RNA Quantity: at least 400ng of total RNA at 50ng/ul or higher for each sample. Please note that RNA is kept only until hybridizations are complete and satisfactory; we cannot store your leftover RNA long-term nor can we return it to you.

<i>Internal Use Only</i>		PI Name: <input type="text"/>	Project ID: <input type="text"/>
# of Samples _____ _____	Quantity	Charge ID	
	\$ _____	Illumina Expression Labeling GRI10003	
	\$ _____	Illumina Expression Processing GRI10003	