

CMMT / BCRI 2100 BioAnalysis Core Facility

Sample Submission Form

Please Choose a Service	
Next Scheduled Run (default service)	X
Don't require until (may cost less)	d / m / y
IMMEDIATELY (50% surcharge)	

PI: _____
User: _____
Grant: _____
Phone #: _____
Date: _____

Please submit or send samples to:
 Xiaohua Han
 CMMT
 950 West 28th Ave., Room 3028
 Vancouver BC, V5Z 4H4
 Ph. 604-875-3814
 Fax. 604-875-3819
 Email: 2100@cmmmt.ubc.ca

LabChip (wells/chip)	Assay	Check One
RNA 6000 Pico (11)	Total RNA (0.2 - 5 ng/ul)	<input type="checkbox"/>
RNA 6000 Pico (11)	mRNA (0.5 - 5 ng/ul)	<input type="checkbox"/>
RNA 6000 Nano (12)	Total RNA (5 - 500 ng/ul)	<input type="checkbox"/>
RNA 6000 Nano (12)	mRNA (25 - 250 ng/ul)	<input type="checkbox"/>
For Total RNA, please indicate if your sample is prokaryotic		<input type="checkbox"/>
DNA 1000 (12)	25 to 1000 bp (0.5 - 50 ng/ul)	<input type="checkbox"/>
DNA 7500 (12)	100 to 1000 bp (0.5 - 50 ng/ul)	<input type="checkbox"/>
Protein 200 (10)	14 to 200 kDa	<input type="checkbox"/>
Other		<input type="checkbox"/>

Sample #	Sample Name	Est. Conc. (ng/ul)	Comments
1			
2			
3			
4			
5			
6			
7			
8			
9			
10			
11			
12			
13			
14			
15			
16			
17			
18			
19			
20			
21			
22			
23			
24			
25			

Sample Submission Guidelines

Please:

- ⇒ Submit samples for different assays on separate forms.
- ⇒ Submit 2.5 ul of sample for RNA and DNA assays, and 10 ul for protein assays.
 - These volumes are enough for a repeat assays if required. If sample volumes are limited, please contact the Facility for instructions.
- ⇒ Email expected arrival date and shipping details when sending samples (particularly if frozen)
- ⇒ RNA samples should be resuspended in RNase free water. Salt (TRIS) 10mM or higher can interfere with analysis.
- ⇒ Please visit our web site for further sample preparation suggestions
www.cmmmt.ubc.ca/2100
- ⇒ Note the number of wells on each chip. Charges for empty wells can be minimized by submitting samples in multiples of this number.
 - Empty well costs will be reduced if your samples are run with other user's samples though may lead to your samples being run on multiple chips (multiple data files). If you wish to keep your samples on the same chip (at potentially higher cost) please indicate this below.

Submission Notes:

Submission ID:	
Run ID:	