

ACRF Centre for Cancer Genomic Medicine

MHTP Medical Genomics Facility
Monash Health Translation Precinct
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<http://mhptmedicalgenomics.org.au>

Illumina Request Form

Name _____ Date ordered _____

Institute and group _____

Address _____

Email _____ Phone _____

Cost centre or P.O. _____ Authorising signature _____

Illumina Sequencing Required HiSeq3000 HiSeq1500 MiSeq NextSeq

Whole gDNA

Whole Exome Roche capture Agilent capture Illumina capture

Methylated Enriched

ChIP Enriched

pA RNA-Seq

Ribodepleted RNA-Seq

Small RNA

Other Please specify _____

Service required

Library Preparation Cluster Generation Sequencing

Barcoding No. of Samples per Lane _____

150 bp SR Sequencing 2 x 150 bp Paired End bp Mate Pair

100 bp SR Sequencing 2 x 100 bp Paired End

50 bp SR Sequencing

Special instructions _____

Sample No.	Sample Name	Conc	260:280 Ratio	260:230 Ratio	Office use only	
					Barcode	ULN
1						
2						
3						
4						
5						
6						
7						
8						
9						
10						
11						
12						
13						
14						
15						
16						
17						
18						
19						
20						

Office use only

Sample receipt checked

Library Preparation

Template Preparation

Sequencing