SEQUENCING SERVICES ENQUIRY FORM

KENEPURU SCIENCE CENTRE (KSC) CHRISTCHURCH SCIENCE CENTRE (CSC)

NOTE: ESR DOES NOT OFFER Human Genomic Sequencing

E/S/R
Science for Communities
He Pūtaiao, He Tāngata

This form is to provide information to assess if your sequencing needs can be met by ESR. Please do not send any samples before receiving confirmation from our team that your request can be fulfilled. Send completed forms to ngs.requests@esr.cri.nz.

INSTRUCTIONS FOR USING FILLABLE FORMS: Please complete this form in Acrobat Reader DC and "SAVE AS PDF" to your computer. Email completed form to: ngs.requests@esr.cri.nz

SAMPLING/SOURCE INFORMATION		REQUESTOR CONTACT DETAILS	
Is the sample related to a clinical case or human derived sample?		Name:	
Sample origin:		Org:	
Human Animal		Phone:	
Food	Environmental	Email (1):	
Other: (please specify):		Email (2):	
		Address:	
SAMPLE INFORMATION			
Sample type: DNA RNA Bacterial susp./isolate Viral swab (VTM)			
PCR product: single-plex or multi-plex			
Other: (please specify)			
PROJECT DETAILS			
Total number of samples to be sequenced:			
Purpose of sequencing: (please provide details and attach any additional paperwork or laboratory results as required).			
SEQUENCING REQUIREMENTS			
Genome size of organism (if known):			
Type of sequencing: Viral Bacterial Metagenomic Other:			
Sequencing method: Illumina (short read) Nanopore (long read) Amplicon sequencing Undecided			
Sequencing method. Italiana (short read) Aniphreon sequencing ondecided			
DATA AND ANALYSIS REQUIREMENTS			
Would you like ESR to perform the data analysis? YES NO			
IF YES: Do you require the data to be compared with any other samples? (e.g. previous isolates/public data)			
Details:			
What should the report contain, and which files should be sent with the report? (select all that apply). Results of specific analysis Interpretation Assembly/Alignment Data Fastg files All raw data			
Results of specific analysis Interpretation Assembly/Alignment Data Fastq files All raw data (as requested above or under project details)			
If large amounts of data need to be returned, how will it be transferred?			
Server Web service TBD Other (specify)			
Would you be interested in sharing your data on a public database such as NCBI or ENA? YES NO Open to discussion			
Using Acrobat Reader DC, SAVE AS pdf with a ne	w name and email this request form to: ngs.requests@esr.cri.	nz	RESET FORM
Disclaimer: If samples are accepted for sequencing, the ESR sequencing team will be perform quality checks to ensure the samples are suitable for the analysis. Failed quality checks will be reported to the requestor. The requestor can agree to			
proceed with failed QC samples, but in this event ESR will not be responsible for sequencing failures. Repeated attempts at sequencing failed QC samples will incur additional charges.			