



Wan Care Scientific Sdn Bhd (1214556-D)

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Illumina® offers the Illumina Genome Network (IGN) for whole-genome sequencing, and the Illumina CSpRo® (CSpRo) Program, supporting a range of genetic applications.

Next Generation Sequencing Request Form

Date: _____

Customer Name: _____

Name of University/ Institute/ Dept: _____

Address: _____

Mobile Number: _____

Email Address: _____

Supervisor Name: _____

Numbers of Sample: _____

Sample Type: gDNA / Total RNA / Plasmid / PCR Product /
 Small RNA, please specify: _____ /
 Others, please specify: _____

Sample Origin: Human / Animal / Microbes / Plant / Fungi /
 Others, please specify: _____

Type of Sequencing Service:

<input type="checkbox"/>	Whole Genome <i>de novo</i> Sequencing
<input type="checkbox"/>	Whole Genome Re-sequencing
<input type="checkbox"/>	Whole Genome Bisulfate Sequencing
<input type="checkbox"/>	Exome Sequencing
<input type="checkbox"/>	Target Region Sequencing
<input type="checkbox"/>	Target Region Bisulfate Sequencing
<input type="checkbox"/>	Human MHC Capture Sequencing
<input type="checkbox"/>	RAD-Seq Population Evolution
<input type="checkbox"/>	RAD-Seq Genetic Map Construction
<input type="checkbox"/>	Ultra Low Input Single Cell mRNA-Sequencing
<input type="checkbox"/>	Low Input, Degraded/ PFTE Total RNA-Sequencing
<input type="checkbox"/>	Epigenetic Profiling
<input type="checkbox"/>	Immune Repertoire Sequencing
<input type="checkbox"/>	Full Length Plasmid Sequencing
<input type="checkbox"/>	Transcriptomic <i>de novo</i> Sequencing
<input type="checkbox"/>	Transcriptomic Re-sequencing
<input type="checkbox"/>	miRNA Sequencing
<input type="checkbox"/>	Circular RNA Sequencing
<input type="checkbox"/>	shRNA/ sgRNA CRISPR Denovolution Screening NGS
<input type="checkbox"/>	NGS CRISPR Validation
<input type="checkbox"/>	Meta rRNA Amplicon Sequencing
<input type="checkbox"/>	Genotyping by Sequencing
<input type="checkbox"/>	ChIP Sequencing
<input type="checkbox"/>	Lentivirus & Retrovirus Integration Site Sequencing

Type of Platform:

<input type="checkbox"/>	Illumina HiSeq 2500/ HiSeq4000
<input type="checkbox"/>	Illumina MiSeq
<input type="checkbox"/>	PacBio
<input type="checkbox"/>	Affymetrix
<input type="checkbox"/>	Wafergen
<input type="checkbox"/>	BGISeq-500
<input type="checkbox"/>	Others, Please specify: _____

Data Output/ Sample Required:

_____ Mb/ Gb

No. of Reads/ Sample Required:

_____ Million

Bioinformatics Analysis Required:

Budget for this NGS Project:

Estimated Samples Submission Date:

Estimated NGS Completion Date:

Objective/ Goal for NGS project:

Other Requirements/ Remarks:

Thank you for your interest of our NGS service!
We will get in touch with you soon.