

## **Sample Submission Guidelines**

Service	Sample Type	Sample Source	Optimum Input Amount	Volume	Concentration	Purity and Quality
hWGS (PCR-free)	gDNA	Fresh frozen tissue, cell line, saliva, blood	≥ 1.5 µg	≥ 20 µl	≥ 20 ng/µl	OD260/280 = 1.8 - 2.0 OD 260/230 ≥ 1.7 DIN ≥ 7
hWGS (PCR-based)	gDNA	Fresh frozen tissue, cell line, saliva, blood	≥ 500 ng	≥ 20 µl	≥ 20 ng/µl	OD260/280 = 1.8 - 2.0 OD 260/230 ≥ 1.7 DIN ≥ 7
	gDNA	FFPE	≥ 700 ng	≥ 20 µl	≥ 20 ng/µl	Main band of > 1,500 bp (Q Score > 0.5) DIN ≥ 4

### Human Whole Genome Sequencing

Recommended suspension buffer: TE buffer or Nuclease-free water

#### Human Whole Exome Sequencing

Service	Sample Type	Sample Source	Optimum Input Amount	Volume	Concentration	Purity and Quality
hWES	gDNA	Fresh frozen tissue, cell line, saliva, blood	≥ 300 ng	≥ 20 µl	≥ 15 ng/µl	OD260/280 = 1.8 - 2.0 OD 260/230 ≥ 1.7 DIN ≥ 7
	gDNA	FFPE	≥ 500 ng	≥ 20 µl	≥ 20 ng/µl	Main band of > 1,500 bp (Q Score > 0.5) DIN ≥ 4

Recommended suspension buffer: TE buffer or Nuclease-free water

#### **RNA Sequencing**

Service	Sample Type	Sample Source	Optimum Input Amount	Volume	Concentration	Purity and Quality
RNA-Seq	Total RNA	Fresh frozen tissue, cell line, saliva, blood	≥ 400 ng	≥ 20 µl	≥ 20 ng/µl	OD260/280 = 1.8 - 2.0 OD 260/230 ≥ 2.0 RIN ≥ 6
	Total RNA	FFPE	≥ 500 ng	≥ 20 µl	≥ 20 ng/µl	OD260/280 = 1.8 - 2.0 OD 260/230 ≥ 2.0 DV200 > 50% RIN ≥ 2
RNA-Seq (Directional)	Total RNA	Fresh frozen tissue, cell line, saliva, blood	≥ 800 ng	≥ 20 µl	≥ 20 ng/µl	OD260/280 = 1.8 - 2.0 OD 260/230 ≥ 2.0 RIN ≥ 6
	Total RNA	FFPE	≥ 1 µg	≥ 20 µl	≥ 20 ng/µl	OD260/280 = 1.8 - 2.0 OD 260/230 ≥ 2.0 DV200 > 50% RIN ≥ 2

Recommended suspension buffer: TE buffer or Nuclease-free water



# **Sample Submission Guidelines**

### Human Whole Genome Methylation Sequencing

Service	Sample Type	Sample Source	Optimum Input Amount	Volume	Concentration	Purity and Quality
hWGMS	gDNA	Fresh frozen, cell line, saliva, blood	≥ 1.2 µg	≥ 20 µl	≥ 20 ng/µl	OD260/280 = 1.8 - 2.0 OD 260/230 ≥ 1.7 DIN ≥ 7
	gDNA	FFPE	≥ 1.2 µg	≥ 20 µl	≥ 20 ng/µl	Main band of > 1,500 bp (Q Score > 0.5) DIN ≥ 4

Recommended suspension buffer: TE buffer or Nuclease-free water

#### **DNA Extraction Service**

Service	Sample Type	Sample Source	Optimum Input Amount
	gDNA	Blood	3-5ml whole blood
	gDNA	Fresh frozen	ТВА
gDNA Extraction	gDNA	Cell line	TBA
	gDNA	Saliva	ТВА
	gDNA	FFPE	10 mm <sup>2</sup> , 5-10 slides

### Pe-made Library Sequencing Service

Service	Library Concentration (measured by Qubit)	Total Amount	Total Volume
Pre-made library sequencing	≥ 2 ng/µl	≥ 200 ng	≥ 20 µl

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### Shipping Information FAQ

#### 1. How do I prepare and pack my samples for shipping?

Your MiRXES representative will provide you a Service Order Form and please fill in the required information, and then return it to us by both emailing a soft copy and attaching a printed copy in your sample package.

- a) Ensure all tubes or plates are clearly labelled per your descriptions in the Service Order Form
- b) Ensure all tubes or plates are tightly sealed to prevent leakage of samples
- c) Attach a copy of the printed Service Order Form
- d) Ensure adequate dry ice is used for the specimen storage & transport
- e) For Singapore customers: Inform your MiRXES representative to come collect the package For non-Singapore customers: Ship your sealed package with your preferred courier to the address below

For Japanese Customer