



Type	Exome			Targeted resequencing of genes panels
	Standard	Salive	Cancer	
Samples requirements	Quantity: 1 µg DIN: 8<	Quantity: 1 µg DIN: 8<	Quantity: 1 µg DIN: 8<	Quantity: 1 µg DIN: 8<
Protocol	TWIST Biosciences WES	TWIST Biosciences WES	TWIST Biosciences WES	Agilent XTHS2 DNA targeted sequencing
Sequencing technology	NovaSeq X	NovaSeq X	NovaSeq X	NovaSeq X
Sequencing	PE, 100	PE, 100	PE, 100	PE, 100
Recommended sequencing per sample	45 M	45 M	90 M	Variable
2025 price (subject to change in 2026)	313 €	408 €	408 €	From 173 €
Delay	≈ 1.5 months	≈ 1.5 months	≈ 1.5 months	≈ 1.5 months

For more information, please contact us on: genomique@users-imagine.fr



Amplicon Sequencing		WGS	
Type	Approach by ligation		
Samples requirements	Quantity: 250 ng w/o dimers primers (AMPure purified)	Quantity: 5 µg DIN: 8<	
Protocol	TruSeq DNA PCR free	TruSeq DNA PCR free	
Sequencing technology	NovaSeq X	NovaSeq X	
Sequencing	PE, 100 or 150	PE, 150	
Recommended sequencing per sample	2 M	400 M (Coverage ≈ 30X)	
2025 price (subject to change in 2026)	110 €	630 €	
Delay	≈ 1.5 months	≈ 1.5 months	

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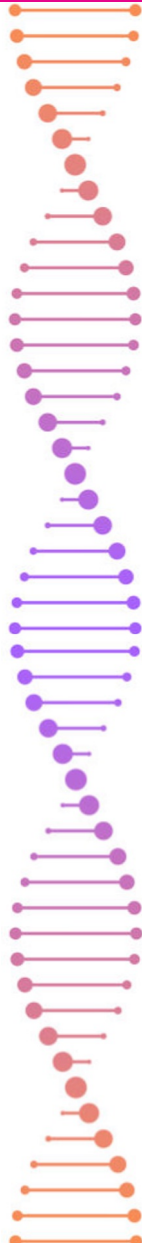


	mRNA	Total RNA
Type	Transcriptome standard	Transcriptome, low quantity
Samples requirements	Quantity: 2.5 µg RIN: 8<	Protocol input range: 5 ng → 200 ng – RIN: 8<
Protocol	Universal Plus mRNA-Seq UDI, Tecan	NEB Single Cell/Low Input
Sequencing technology	NovaSeq X	NovaSeq X
Sequencing	PE, 100	PE, 100
Recommended sequencing per sample	60 M	60 M
2025 price (subject to change in 2026)	300 €	300 €
Delay	≈ 1.5 month	≈ 1.5 month

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On request
mRNA capture
Ribodepletion
Very low RNA input (<5 ng)
Custom projects



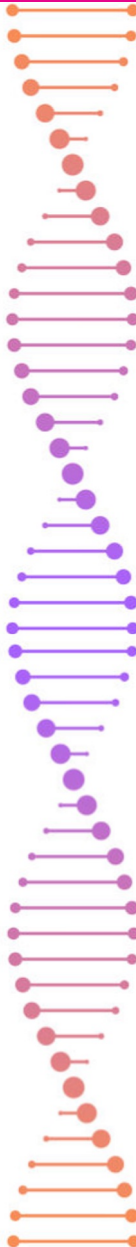
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**The NovaSeq X is available for all short read applications.
The AVITI24 will be progressively implemented for short read applications during 2026.**

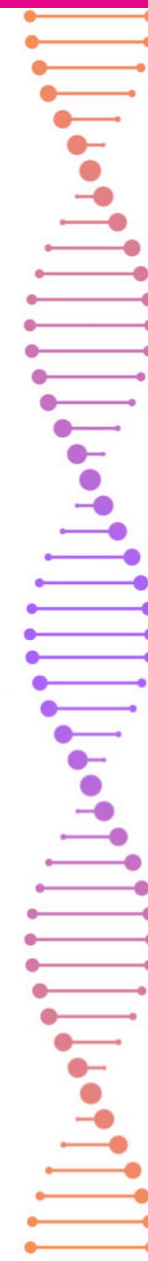


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	Plasmid	Amplicon	Transcriptomics
Samples requirements	100 ng high molecular weight plasmid DNA per sample q.s. 18 µl	250 ng for 1 Kb amplicons DNA per sample q.s. 25 µl	800 ng Minimum concentration > 40 ng/µl RIN > 10
Protocol	SQK-RBK114	SQK-NBD114	Kinnex
Sequencing technology	Oxford Nanopore Technology	Oxford Nanopore Technology	PacBio
Recommended sequencing per sample	7 000 reads	20 000 reads	10M reads
2025 price (subject to change in 2026)	38 €	47 €	720 €
Delay	1 week	≈ 1 month	≈ 1.5 months



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	WGS	WGS
Samples requirements	4 µg high molecular weight genomic DNA q.s. 100 µl DIN > 7	5 µg high molecular weight genomic DNA q.s. 50 µl GQN10Kb > 7
Protocol	SQK-LSK114	SMRTbell 3.0
Sequencing	Oxford Nanopore Technology	PacBio
Recommended sequencing per sample	≈ 25X coverage	~ 25X coverage
2025 price (subject to change in 2026)	1 090 €	798 €
Delay	≈ 1.5 months	≈ 1.5 months

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On request

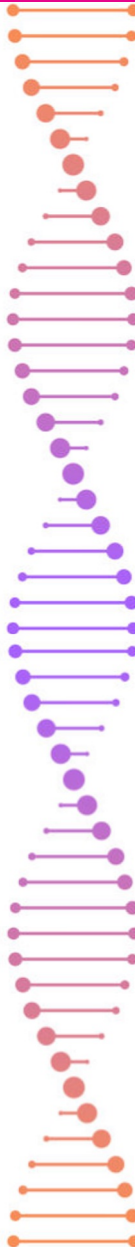
RNA on Oxford Nanopore Technologies

RNA Methylation on Oxford Nanopore Technologies

Capture Long-read Sequencing (> 3kb)

Methylation of specific regions

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Short read: flow cell prices (2025)			
Illumina NovaSeq X			
	1.5 B	10 B	25 B
100 cycles	2 590 €	9 435 €	N/A
200 cycles	3 200 €	11 720 €	N/A
300 cycles	3 440 €	12 445 €	21 160 €

Long read: flow cell prices (2025)	
PacBio Revio	Nanopore (DNA)
1 027 €	970 €

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