QUINTARA

AmpValue NEW

$2.99 Amplicon Sequencing

Introducing Quintara's new **AmpValue** Nanopore sequencing service, a streamlined, affordable solution for your PCR product sequencing needs. It offers rapid turnaround at a competitive price, making high-quality long-read sequencing more accessible than ever.

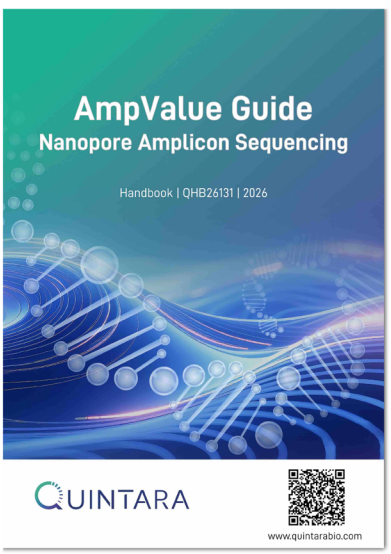
- $2.99 — Flat price, cost-effective, and no hidden fees
- >1,000 drop boxes — Easy submission with >1,000 drop boxes
- <1 day — Overnight data delivery

Service Name	Size	Turnaround Time	Listing Price
AmpValue New	200bp – 25kb	Overnight	~~$4.99~~ $2.99

Sample Input	
PCR Product	Unpurified PCR Product; Purified PCR Product
Bacteria & Plasmid	Colony Suspension; Overnight Culture; Glycerol Stock; Agar Plate; Plasmid
Yeast	Colony in Water; Culture; Agar Plate
Genomic DNA	Genomic DNA

Deliverables

- FASTA File
- AB1 File (chromatogram)
- Genbank File
- Per-Base Data File
- Raw Data FASTQ File
- QC Report
- Alignment File
- Summary File and Report
- Interactive QC Tool



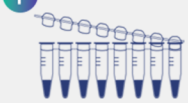
Scan the QR code to download for free.



Nanopore Amplicon Sequencing

Various solutions to meet your needs

1



Sample Submission

- No primers required
- Accepts purified and unpurifled samples
- 1,000+ drop boxes

2



Library Preparation

- ONT library prep workflows
- Quintara's proprietary workflows

3



Sequencing

- ONT long-read technology
- Ultra-high coverage depth
- Real-time data acquisition

4



Data Analysis

- Data processing and alignment
- Variant analysis and quality assessment
- Ready-to-use deliverables

Choose the Services According to Your Needs

	AmpValue Overnight $2.99 (New)	AmpStandard Overnight $15	AmpPro Overnight $30 (Upgrade)
Designed For	Budget-friendly option for sequencing short to medium-length amplicons. Ideal for high-throughput projects.	Classic long-read sequencing for clonal amplicons of any length. Ideal for routine amplicon verification.	Deep sequencing with full-length reads for resolving amplicon mixtures. Ideal for amplicon library verification.
Sample Size	200bp - 25kb	500bp - 25kb	100bp - 25kb
Sample Input	Unpurified PCR product (10µL, 2 ng/µL per 100bp) Purified PCR product (10µL, 2 ng/µL per 100bp)	Unpurified PCR product (10µL, 20-100ng/µL) Purified PCR product (10µL, 20-100ng/µL)	Unpurified PCR product (20µL, 1 ng/µl per 100bp) Purified PCR product (20µL, 1 ng/µl per 100bp)
Coverage	Up to 1,000 reads	Up to 1,000 reads	Up to 16,000 reads
Variants/Sample	1 - 2 +	1	1 – 10 +
Deliverables	1. FASTA File 2. AB1 File (chromatogram) 3. Genbank File* 4. Per-Base Data File* 5. Raw Data FASTQ File* 6. QC Report * 7. Alignment File* 8. Summary File and Report* 9. Interactive QC Tool	1. FASTA File 2. AB1 File (chromatogram) 3. Genbank File 4. Per-Base Data File 5. Raw Data FASTQ File 6. QC Report 7. Alignment File 8. Summary File and Report 9. Interactive QC Tool	1. FASTA File 2. AB1 File (chromatogram) 3. Genbank File 4. Per-Base Data File 5. Raw Data FASTQ File 6. QC Report 7. Alignment File 8. Summary File 9. Variant Analysis Results 10. Interactive QC Tool

* Improvements made to data package based on direct feedback from our beta testing customers.

\+ Variant consensus sequences require at least ~5% difference in sequence or length; closer variants may collapse into a single consensus.

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