



AmpValue **NEW**

\$4.99 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.

 **\$4.99** 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

Sample Input

- Unpurified PCR product (10µL, 2 ng/µL per 100bp)
- Purified PCR product (10µL, 2 ng/µL per 100bp)

Deliverables

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

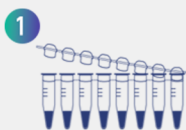
5 Free Samples

Valid Through April 30 2026

The promotional pricing applies to all customers ordering AmpValue for the first time. The discount will be applied automatically when you place your order through the QuinGo online portal. Quintara Biosciences reserves all rights to this promotion.

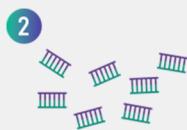
Nanopore Amplicon Sequencing

Various solutions to meet your needs



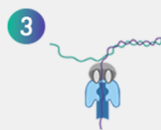
Sample Submission

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



Library Preparation

- ONT library prep workflows
- Quintara's proprietary workflows



Sequencing

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



Data Analysis

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

Choose the Services According to Your Needs

	AmpValue Overnight \$4.99 New	AmpStandard Overnight \$15 Upgrade	AmpPro Overnight \$30
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 500 reads	Up to 1,000 reads	Up to 4,000 reads
Variants/Sample	1-3	1	10-200+
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*	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	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

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

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