

Comprehensive and Informative

Genomic DNA is fully sequenced by long reads (ONT) with high consensus accuracy. You will receive the **fully reconstructed and annotated genome sequence** of your bacterial species.

Speedy and Cost-effective

Results are delivered within 3 to 7 working days after sample receipt. The complete sequence can now be unraveled at a fraction of the cost compared to previous sequencing technologies.

Easy Handling

All you need is to send as little as 1 µg of bacterial genomic DNA via **Microsynth's drop box system**, which **includes free and fast shipping**. Do you need to outsource your DNA isolation? Microsynth has you covered.



Overview of BacterialSeq

Microsynth's new bacterial genome sequencing service is based on the latest long-read sequencing technology from Oxford Nanopore Technologies (ONT). It is designed for de novo whole genome sequencing and assembly of genomic DNA from a clonal population of bacteria (single species, genome sizes up to 12 Mb). The sequencing yields on average a 30x genome coverage.

Microsynth has developed a sophisticated bioinformatics pipeline to accurately reconstruct and thoroughly annotate microbial chromosomes.

Outstanding Features & Benefits

This new service is **fast**, **cost-effective**, **complete**, **and hypothesis-free**. Provide a clean, good quality DNA sample and receive **annotated contigs** of your bacterial genome, with a **wealth of supporting information** including complementary

species identification methods, and quality metrics such as coverage analysis. Compared to Illumina sequencing, both costs and turnaround times are significantly reduced. Long sequencing reads greatly enhance the accuracy of the assembly, even

in the presence of repetitive elements. Useful add-on services are available for an additional fee: DNA isolation, assembly polishing using supplemental Illumina sequencing, and variant calling based on a reference genome.

ONT vs. Illumina Sequencing

Research Question or Application	ONT	Illumina
Fast and cost-effective <i>de novo</i> sequencing of an entire genome sequence	+++	+
Sequencing of GC-rich and repetitive regions without gaps	++	+
Verifying number of repeated regions	++	-
Direct sequencing of native DNA, minimally fragmented and without the need for PCR amplification	+++	F
Identification of structural variations	++	-
Nucleotide level accuracy	++	+++

Incredibly Straightforward

- 1. Just submit 1 µg of a high quality DNA preparation.
- 2. Put your samples in one of our sample drop boxes.
- 3. Sit back, your sequencing results will be delivered in 3 to 7 working days following sample receipt. If you choose DNA isolation service in addition, another 4 days will be needed.

Products

3272	BacterialSeq	3273	BacterialSeq - Polishing
3271	BacterialSeq - DNA Isolation	3274	BacterialSeq - Large Genome (7-12 Mb)q
		3276	RacterialSeg - Variant Analysis

Never Tried?		
Write an email to your sales manager or info@microsynth.ch and ask for trial set .*		
How to Order?		
Enter our webshop via www.microsynth.com		
Click on "ONT Sequencing" in the green "Analysis Services" area		
Click on "BacterialSeq" service and follow the further instructions		
Need More Information?		
Call us at +41 71 726 10 04 or		
E-mail us at sanger.support@microsynth.ch		

^{*} Trial Set: 50% discount on up to 3 bacteria, per research group