

Unlock the Potential of Microbes

"bit-MAP®" Single-cell Genomic Analysis of Microbes

Single-cell level Analysis Service of Microbiome Sample

- Sequence uncultured bacterial genomes at once
- ✓ Obtain contamination-free genome sequences from rare bacteria
- ✓ Comparative genome analysis at the sub-strain resolution

bitBiome Inc.

bitBiome, a Japanese biotech Startup, provides a unique Single-cell, whole-genome analytical platform, bit-MAP®, specifically targeting microbes that can be applied to uncultured "raw" samples.



bit-MAP®

Whole genome sequencing provides valuable insights about types and functions of bacteria. Yet, accessible genomic information was limited to certain types of bacteria due to the difficulty in culturing most of bacteria.

We developed a new analytical platform "bit-MAP®" to sequence bacterial genomes from a single cell. bit-MAP® enables us to uncover new bacterial genomes nobody has ever discovered.

Encapsulation Cell Lysis & DNA Amplification Sorting & Sequencing Bioinformatic Analysis Single-cell isolation of microbial community Parallel lysis and DNA amplification within gel Gel capsules with genome are collected in well plates and functional information of







capsules



sequenced.

individual bacteria

Basic Services

Order No.	Service Menu		Lead Time	Deliverables	Minimum Order Quantity	Unit Price (per sample)
S-51	Single-cell Genomic Analysis (384 well)	Standard	8-10 weeks	Analysis Report	A CIV	USD 9,800
S-52		Express	3-4 weeks	Sequence Data (Details are at [O-51] on 3 rd page)	ASK -	USD 13,200

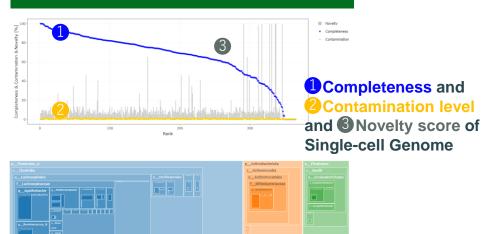
- Reduced sample size option for Single-cell Genomic Analysis is available to fit multiple samples in a 384-well plate (price may vary. Please see the price example).
- Lead Time above is estimated from the date of sample receipt.
- Transportation costs, such as freight and custom declaration costs, are not included in the price.

Sample Requirement

Sample	Amount			
Human Feces	≧300 mg			
Human Saliva	≧1 mL			
Suspension of Cells/Extracellular Particles	≧100 μL (Conc.: ≧14,000 cells/μL)			

- For other types of samples or low-volume samples, please contact us.

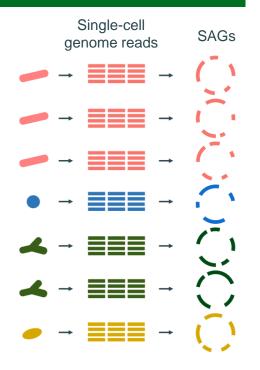
Analysis Report Example

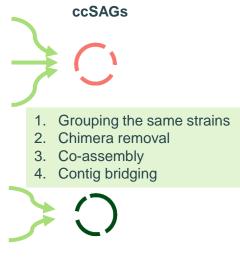


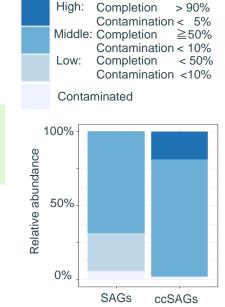
Taxonomy identification and statistical analysis

Basic: SAG Assembly

Option Service: ccSAG Assembly







Quality of draft genomes

SAGs are produced by *de novo* assembling of single-cell genomic reads, included in the Basic Analysis (O-51). For advanced single-cell genomics, bitBiome provides the original bioinformatics pipeline, ccSAG Assembly option (O-53). With the option, you can obtain non-redundant and high-quality draft genomes of microbial strains and perform strain-resolved genome analysis.

Optional Service

Order No.	Optional Service	Abstract	No. of Unit	Unit Price
O-51	Basic Analysis with S-51/S-52	 De novo assembly Evaluation of SAG (Single Amplified Genomes) Quality Taxonomy identification Gene prediction 	Included in Basic Menu "S-51/52"	
O-52	Option Additional Fee	• To be charged when one or more Option services (O-53, O-54, O-55, etc) is ordered.	1 order	USD 2,000
O-53	ccSAG Analysis	 Chimera removal and SAG composition Evaluation of ccSAG quality Taxonomy identification Gene prediction 	1 sample	USD 2,000

⁻ Additional analytical services are also available upon request (e.g., phylogenetic analysis, biosynthetic gene clusters identification, antibiotic-resistant gene identification, and SAG-guided MAG analysis). Please contact us.

Price Example

Case1: High-Quality Single-cell Genomic Analysis 2 sample with 2 plates

Case2: Single-cell Genomic Analysis 4 sample with 2 plates (192 well x 4)

	Service	Amount	Price (USD)	Service	Amount	Price (USD)
Basic	Single-cell Genomic Analysis (Standard) [S-51]	2 sample	19,600	Single-cell Genomic Anal (Standard) [S-51]	ysis 4 sample /2 plate	29,400
	Basic SAG Report [O-51]	-	Free of Charge	Basic SAG Report [O-51]	-	Free of Charge
Option	Option Additional Fee [O-52]	1 order	2,000		11 x 1.5 Charge of	
	ccSAG Analysis [O-53]	2 sample	4,000		allocation)	
	Total		USD 25.600	Total	ı	JSD 29.400

Other Service

Order No.	Other Service		Lead Time	Deliverables	Minimum Order Quantity	Unit Price (per sample)
M-53	Shotgun Metagenomic Analysis (Data: 5Gb)	Standard	8-10 weeks	Sequence Data	ASK	USD 1,580
M-54		Express	3-4 weeks			USD 2,360

- Additional analytical services are available upon request (e.g., Metagenomic Reads/Contig Analysis and MAG analysis). Please contact us.
- Lead Time above is estimated from the date of sample receipt.

Service Flow



Confirmation of REQ, QTN &Order, Shipment of Sample



Sequencing, and Data Analysis



Reporting