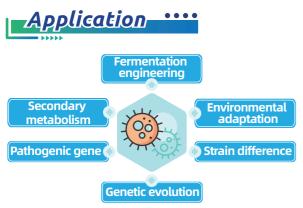


# **Bacterial Genome Sequencing**

Tailored to meet various research goals, BMKGENE offers genome sequencing and assembly services for bacteria, including whole genome, draft genome, and complete genome. By combining next-generation sequencing with third -generation sequencing data, to achieve high-quality genome assembly and accurate functional annotation. For complete bacterial genome assembly, BMKGENE guarantees 0 gap, providing researchers with a comprehensive understanding of their target bacteria.



## Bacterial Genome Solutions

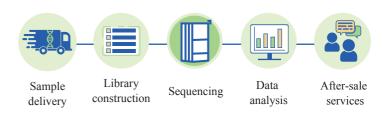
#### **Option 1 Genome re-sequencing:**

Bacterial whole genome sequencing (WGS):
 SNP calling.

#### **Option 2 Genome assembly:**

- Bacterial Draft Genome (NGS): Genome com
   -ponent analysis and function annotation.
- Bacterial Complete Genome (TGS+NGS):
   Genome component analysis, function annota
   -tion, and circos of genome.

### Service Workflow ....



#### Service Advantages · · · ·

- Multiple sequencing strategies are available for different bacterial genome research goals.
- The species identification and classification of bacteria are conducted to study their evolutionary relationships and more.
- Bacteria complete genome sequencing by PacBio or ONT platform with **0** Gap assembly guaranteed.
- Highly experienced in bacterial genome assembly with over 10,000 microbial genomes assembled.
- Professional after-sale technical support team fulfilling more specific research needs.



Library	Read Length	Recommended Strategy	Assembly	
Illumina	PE150	100X NGS	Draft map	
PacBio	HiFi-15 Kb	30X HiFi	Complete map 0-Gap	
Nanopore	10-20 Kb	100X ONT + 100X NGS	Complete map 0-Gap	

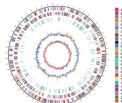
### Sample Requirements ....

Platform	Qubit Conc. (ng/μL)	Amount (μg)	Volume (µL)	Purity
Illumina	1	0.06	20	OD260/280: 1.7-2.2;
PacBio	20	1.2	20	OD260/230: ≥ 1.0;
Nanopore	40	2	20	Nanodrop/Qubit: 0.8-2.5

#### Demo Results ....







A A May processing and modification(2):

L Deepy production and construction(2):

C L Deepy production and construction(2):

C L Construction and construction(2):

C L Construction and construction(2):

P A Reducedist response and metabolism(12):

E C Construction (2):

E C Construction (2):

E C Construction (2):

E C Construction (2):

E C Transcription (2):

L Construction (2):

L C

1.SNP classification and annotation

2. Coding gene annotation

3. Genome visualization

### Featured Publications ....

Year	Journal	Article	Applications	DOI
2023	Science Advances	Ampicillin-controlled glucose metabolism manipulates the transition from tolerance to resistance in bacteria	Bacterial resistance	10.1126/sciadv.ade8582
2023	International Journal of Molecular Sciences	Discovery of Bacteroides uniformis F18-22 as a Safe and Novel Probiotic Bacterium for the Treatment of Ulcerative Colitis from the Healthy Human Colon	Disease treatment	10.3390/ijms241914669
2022	Science of The Total Environment	Integrative chemical and omics analysis of the ammonia nitrogen removal characteristics and mechanism of a novel oligotrophic heterotrophicnitrification-aerobic denitrification bacterium	Pollution treatment	10.1016/j.scitotenv. 2022.158519



Biomarker Technologies (BMK) GmbH	
② BioZ, Johann-Krane Weg       ☑ tech@bmkcloud.com       42, 48149 Münster, Germany      ⑪ www.bmkgene.com	
Copyright©2009-2023 Biomarker Technologies (BMK) GmbH . All Rights Reserved. Information and specifications are subject	
to change at any time without notice.	