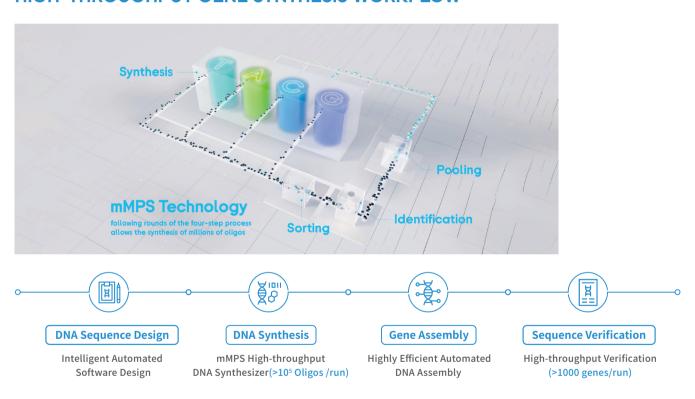


High-throughput Gene Synthesis

GCATbio is revolutionizing high-throughput gene synthesis by leveraging novel oligo synthesis and cutting edge assembly tools to deliver high quality, low cost genes. The underlying DNA synthesizer, called mMPS (microchip-based Massive in Parallel Synthesis) produces up to 100,000 oligos per run with unmatched yield, precision, and flexibility. Coupled with 100% NGS sequence verification, this platform empowers laboratories to tackle large-scale synthetic biology projects at a cost that's within reach of nearly all budgets.

In addition to gene synthesis, the mMPS synthesizer also supports diverse applications, including ultra pure oligo synthesis, chromosome synthesis, variant library construction, DNA storage, and beyond.

HIGH-THROUGHPUT GENE SYNTHESIS WORKFLOW



KEY BENEFITS



High throughput:

Annual synthesis capacity exceeding 1 billion base pairs; 100% verified by high-throughput sequencing



Highly automated:

An industry-leading modular automation platform, ensuring high efficiency and stable quality



Extensive genome synthesis experience:

Supports projects involving the synthesis of chromosomes and genomes at Mbp level

CASE

01

Synthesis of Eukaryotic Chromosomes (Sc2.0)



Several core members of GCATbio have made significant contributions to the Sc2.0 project. GCATbio's proprietary high-throughput gene synthesis platform supported the synthesis of three million bases, one quarter of the whole genome. The team also developed SynGenomeDesigner, a software platform for efficient and automated hierarchical genome design. This platform is suitable to efficiently assemble genomes for various model organisms including *E. coli, C.elegans etc.*

02 Applications in DNA Storage



The DNA data storage research is greatly supported by the high-throughput gene synthesis platform, which synthesized and assembled more than 54 Kb ultra-long fragments. Their sequences are verified to 100% accurate by NGS sequencing.

References

Richardson S M, Mitchell L A, Stracquadanio G, et al. Design of a synthetic yeast genome[J]. Science, 2017, 355(6329): 1040-1044. Shen Y, Wang Y, Chen T, et al.

Deep functional analysis of synll, a 770-kilobase synthetic yeast chromosome[J]. Science, 2017, 355(6329): eaaf4791.

Ping Z, Chen S, Zhou G, et al. Towards practical and robust DNA-based data archiving using the yin-yang codec system[J]. Nature Computational Science, 2022, 2(4): 234-242.

Boost your scientific endeavors with GCATbio High-throughput Gene Synthesis

