

# Protein Molecular Solutions

Al Molecular Manufacturing Technology leverages bioinformatics and advanced Al data models to provide innovative molecular services. Synbio Technologies, as a professional Al molecular service manufacturer, has an advanced Al design and manufacturing integration platform, life science service platform, and industrial R&D production platform.

We provide end-to-end solutions for protein molecules, from Al-driven gene design to protein synthesis, expression, purification, and structural analysis. Our highly customizable services address challenges like long and difficult gene design, large molecular weight protein expression, and structural analysis, effectively shortening service cycles and saving costs.

We have successfully synthesized and expressed proteins with molecular weights up to 180 kDa.

# **One-stop Solution**

# Protein Molecular Solution

#### 1. Al Design & Optimization

- Protein Screening Al Design Codon Optimization Gene Analysis
  - 2. Gene Synthesis
  - Sequence Synthesis
     Vector Construction
     Sequence Verification
    - 3. Protein Expression & Purification
  - Expression System Construction Protein Purification
  - 4. Protein Structural Analysis
  - X-ray Crystallography Cryo-EM SPA MicroED
- 5. Large-Scale Production
- Provide Bacteria for Large-scale Fermentation
   Provide Expressed Proteins up to Gram Level

Application Area	Application Scenario	Proteins
CGT  Industrial Biotechnology	Target Protein Production  Therapeutic Protein Production  Viral Vector Production  Core Reagents for Cell Therapy  ADC Drug Development  Industrial Enzyme Customization  Biosensor Development	<ul> <li>GPCR proteins, ion channel proteins</li> <li>Insulin, growth factors, antibodies</li> <li>AAV capsid proteins, viral envelope proteins</li> <li>Cytokines, chemokines, CAR-T recognition domain proteins</li> <li>Antibodies, linkers, toxins</li> <li>Cellulase, lipase</li> <li>Antigen/antibody pairings, labeled proteins</li> </ul>
Research & Diagnostics	Artificial Protein Materials  Structural Biology Research  In Vitro Diagnostic (IVD) Reagents	<ul> <li>Collagen-like proteins, collagen</li> <li>Isotope-labeled proteins, membrane proteins</li> <li>COVID-19 antigens, antibodies, enzyme labels</li> </ul>

# **Service Advantages**

- An Al-driven design and manufacturing integration platform that ensures precision and efficiency (patented technology)
- Capable of expressing and analyzing proteins with molecular weights exceeding 180 kDa
- Advanced protein structural analysis techniques
- Stage-based service fees, offering high cost-effectiveness

## **Service Details**

Steps	Service Details	Deliverables	Turnaround Time
Al Design & Optimization	<ul><li> Protein Screening</li><li> AI Design</li><li> Codon Optimization</li><li> Gene Analysis</li></ul>	<ul> <li>2~5 µg lyophilized plasmid DNA</li> <li>Sequencing chromatogram</li> <li>1- 5 mg protein products</li> <li>COA report</li> <li>the report and the structure factor data and structure atomic coordinate files that comply with the PDB database reception standards</li> </ul>	3-5 Days
Gene Synthesis	Gene Synthesis     Vector Construction		Start from 1 Week
Protein Expression & Purification	Protein Expression Validation     Large-scale Cultivation &     Protein Purification		Start from 2 Weeks
Protein Structural Analysis	<ul><li>Protein Crystallization Optimization</li><li>X-ray Diffraction</li><li>MicroED</li><li>CryoEM-SPA</li></ul>		Start from 1 Month
Large-Scale Production	<ul> <li>Provide Bacteria for Large-scale Fermentation</li> <li>Provide Expressed Proteins up to Gram Level</li> </ul>		Quote



Why Choose Us



- Leading AI biological molecular technology services



- Mature protein service experience

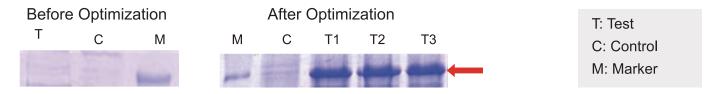


- One-stop solution

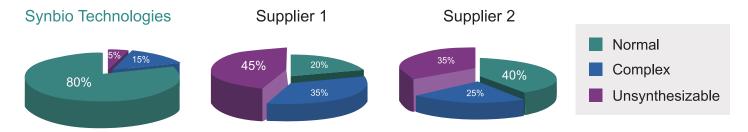
# 1. Al Design & Optimization

Synbio Technologies' AI sequence design & optimization platform integrates deep learning and advanced algorithms to refine the gene synthesis process. Through NG Codon technology, we improve the accuracy and efficiency of codon optimization, while the AI-TAT system forecasts production cycles with high precision. The Complexity Index (CI) system ensures 100% accuracy when synthesizing complex gene sequences, making the entire process faster, more reliable, and cost-effective.

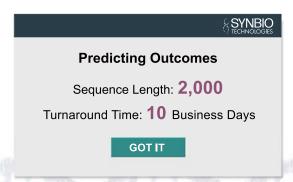
# **Advanced AI Sequence Design | NG Codon Optimization**



# Order Cycle Prediction | AI-TAT System

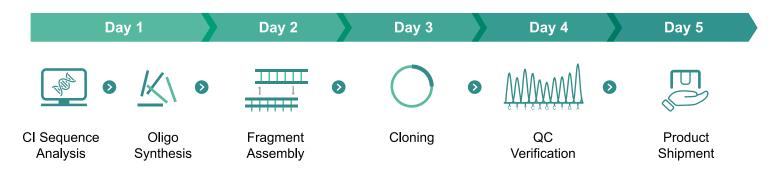


# **Unlimited Gene Synthesis | CI System**



# 2. Gene Synthesis

Our AI-powered gene synthesis platform delivers high-fidelity, fully optimized genes tailored to your unique research needs. By harnessing three advanced AI systems-NG Codon Optimization, Complexity Index (CI), and AI-TAT-we streamline the entire synthesis process, enhancing accuracy, speed, and cost-efficiency. With over 10 billion bases synthesized and millions of codons optimized, we tackle even the most complex genetic challenges, including high and low GC content, repetitive sequences, and hairpin structures. We guarantee 100% accuracy in DNA synthesis, offering competitive pricing and fast turnaround times. Additionally, we provide free codon optimization, vector storage, and vector design to ensure your complete satisfaction.



# Case Study

# Al-designed, ultra-complex collagen protein gene synthesis

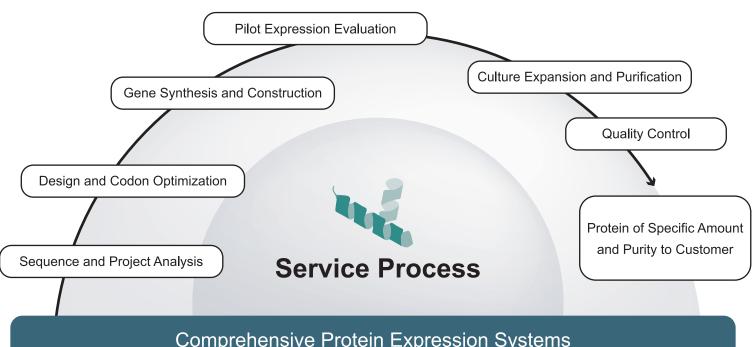
We successfully synthesized Al-designed recombinant collagen DNA sequence of 5121 bp in length containing





# 3. Protein Expression & Purification

Synbio Technologies offers protein expression services across four systems: bacterial, yeast, insect, and mammalian. Whether producing milligrams or grams, our synthetic biology platforms and NG Codon optimization technology generate high-purity, active proteins to accelerate your research. By carefully selecting from a variety of expression vectors, hosts, and fusion tags, we deliver optimal results tailored to the nature of your protein and your specific project requirements. Our customized services empower scientists worldwide to achieve their research goals efficiently.



## Comprehensive Protein Expression Systems



E.coli

Fast and cost-effective for high-yield protein production.



Yeast

Ideal for eukaryotic protein expression with post-translational modifications.



Mammalian

Best for producing complex proteins with accurate folding and glycosylation.



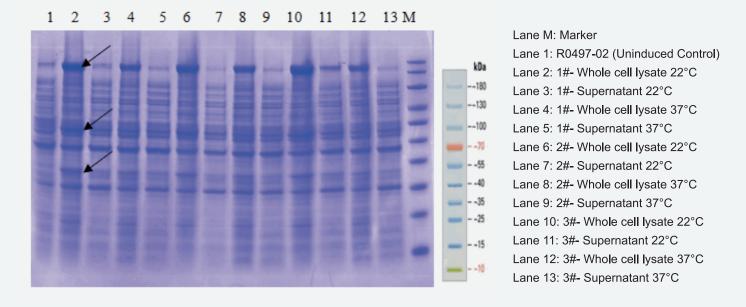
Insect

Efficient for expressing large, complex proteins with eukaryotic-like modifications.

# Case Study

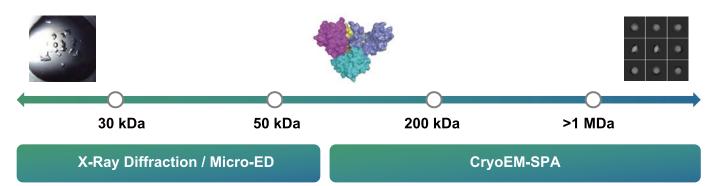
# Successful Expression of 180 kDa Multi-Enzyme Complex + Chaperone Proteins

Al-driven codon optimization significantly enhanced the co-expression efficiency of a large multi-enzyme complex (180 kDa) and two chaperone proteins in *E. coli*, overcoming the molecular weight limits of traditional expression systems. Compared to traditional methods, the soluble fraction was increased by more than 35%.



180kDa Multi-Enzyme Complex + Two Chaperone Proteins

# 4. Protein Structural Analysis



#### X-Ray Diffraction

- · Wide range of protein MW
- Highest resolution, typically <2Å</li>
- Suitable for soluble proteins, membrane proteins, protein complexes

#### Cryo-EM SPA

- No crystallization
- Closer-to-native state
- Minimal sample consumption
- Suitable for large MW proteins, such as membrane proteins and large protein complexes

#### MicroED

- Proteins with diverse MW
- Minimum crystal requirement
- · No need for crystal optimization
- High resolution, <3Å

## X Ray Diffraction Case Study

#### ► KRAS G12D

## **Technical difficulty:**

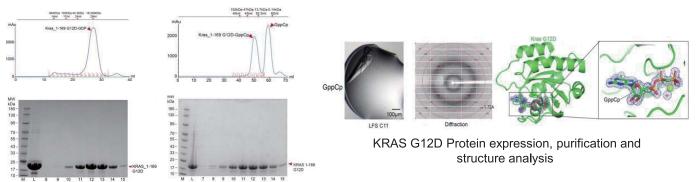
Kras 1-169 G12D-GDP

- KRAS binds to GTP and is in an activated state; when it binds to GDP, it switches to an off-state. The conversion of GTP to GDP is an extremely rapid process, making it difficult to obtain the activated state of KRAS bound to GTP.
- To study this target, it is necessary to obtain the protein in both the activated and off states.

Kras 1-169 G12D-GppCp

#### Solution:

We degraded the GDP bound to KRAS using an enzyme, then added GppCp and incubated it with the KRAS protein to
obtain the activated state of the KRAS protein. Finally, we solved the high-resolution structures of KRAS G12D-GDP/GppCp and KRAS G12D-GDP-1133 (~1.8 Å).



# **MicroED Case Study**

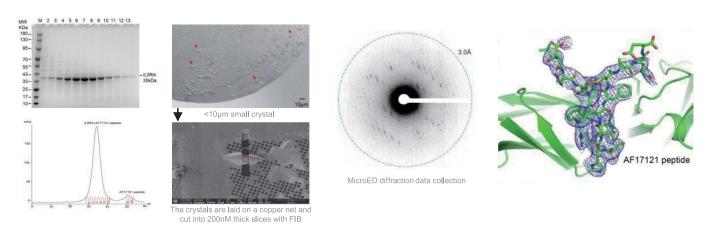
#### ► IL5RA

### **Technical difficulty:**

- IL5RA is highly prone to forming inclusion bodies during protein expression, typically appearing in an insoluble form.
- The cocrystals of IL5RA and peptides are extremely small (<10 μm), making it very difficult to optimize and obtain larger single crystals (>50 μm).

#### Solution:

- We purified IL5RA inclusion bodies and refolded the protein to obtain soluble IL5RA with >90% purity.
- We mounted the crystals on copper grids, prepared 200 nm-thick sections using Cryo-FIB, and collected diffraction data via MicroED. The structure was solved at 3 Å resolution, with clear electron density for the AF17121 peptide.



Cryo-FIB & MicroED

IL5RA structure determination

# **Cryo-EM SPA Case Study**

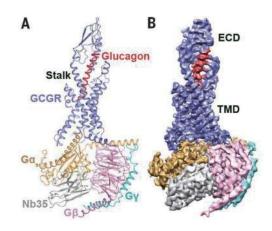
#### ▶ GPCR – GCGR

## **Technical difficulty:**

- The transmembrane regions of membrane proteins possess extensive hydrophobic surfaces, making them difficult to stabilize in polar aqueous solutions after dissociation from the membrane.
- The protein expression level of GCGR is extremely low, which prevents the completion of subsequent purification.
- In the protein sample preparation stage, the protein samples mostly adhere to the supporting film on the grid rather than being embedded in the ice layer within the holes of the supporting film.
- The protein samples exhibit significant preferential orientation.

#### Solution:

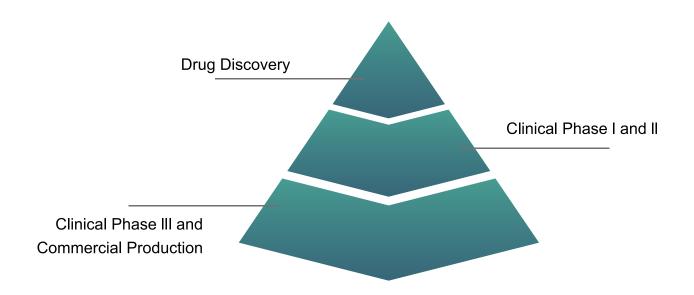
- We changed the expression vector and protein expression system to increase protein yield, and added monoclonal fragments during expression to enhance the stability of the complex.
- During the protein sample loading process, we adjusted the protein concentration and tried other types of grids, eventually achieving a higher number of protein particles embedded in the holes.
- Increased the thickness of the vitreous ice layer and applied a carbon film on the grid to average the number of particles in different orientations, thereby improving the preferential orientation of the protein particles.



GCGR CryoEM result

# 5. Large-Scale Production

Our CRDMO services offer comprehensive solution for your entire project lifecycle, from analysis and development to full CMC services and raw material production.



**One-stop Biomolecular Solutions CRDMO** 

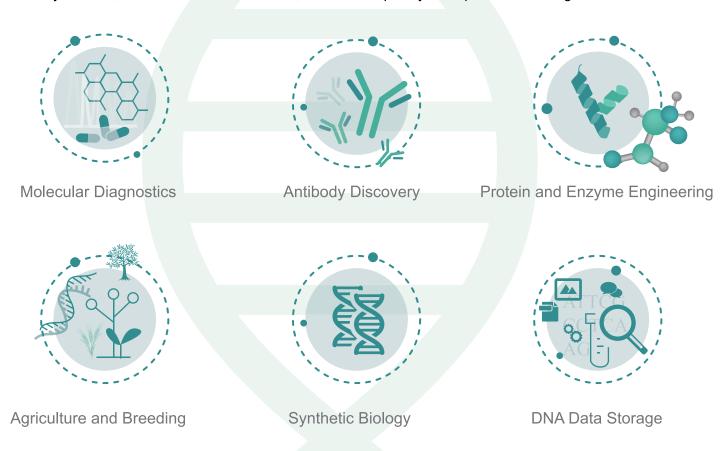


# **Synbio Technologies**

# Your One-Stop Partner for DNA, RNA, Protein & Al Molecule Solutions

# Turn Complexity into Speed and Savings

Founded in 2013, Synbio Technologies empowers innovation in life sciences through integrated solutions for DNA, RNA, protein, and AI-designed molecules. With cutting-edge synthesis platforms and AI-powered bio-design tools, we support researchers, biotechs, and pharma companies in accelerating discovery and development. Trusted by over 20,000 clients in 70+ countries, we turn complexity into speed and savings.









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