# Sequence modeling and design from molecular to genome scale with Evo

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#### Introduction

- 1. DNA: The Blueprint of Life
- 2. Using Machine Learning with DNA
- 3. Current Limits of Machine Learning in Biology

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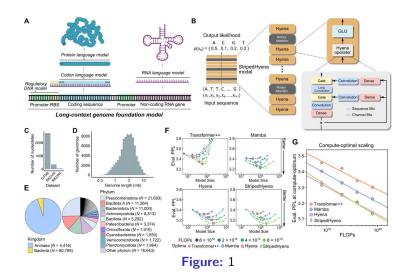
### Pretraining a genomic foundation model across prokaryotic life

- A: Long-Context Genome Foundation Model
- B: StripedHyena Model Processing Genomic Sequences
- C and D: Dataset Overview
- E: Dataset Composition by Kingdom and Phylum
- F: Model Performance Comparison (Perplexity vs. Model Size)

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G: Compute-Optimal Scaling

## Pretraining a genomic foundation model across prokaryotic life



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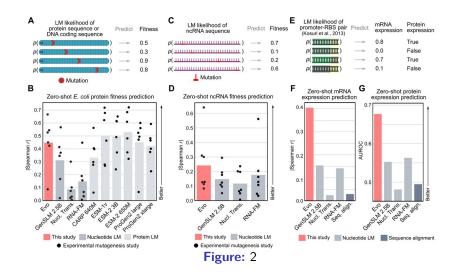
# Evo performs zero-shot function prediction for proteins, non-coding RNAs, and regulatory DNA

- Panel A: Language Model-Likelihood and Fitness Prediction
- Panel B: Zero-Shot E. coli Protein Fitness Prediction
- Panel C: Language Model Likelihood for Non-Coding RNA
- Panel D: Zero-Shot ncRNA Fitness Prediction
- Panel E: Promoter-RBS Pair Predictions (Gene Regulation)

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- Panel F: Zero-Shot mRNA Expression Prediction
- Panel G: Zero-Shot Protein Expression Prediction

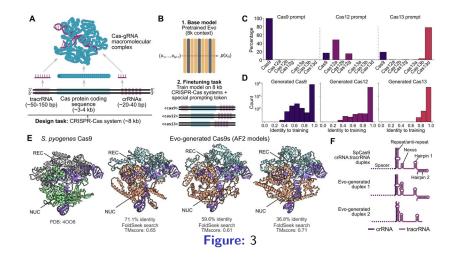
## Evo performs zero-shot function prediction for proteins, non-coding RNAs, and regulatory DNA



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- Panel A: CRISPR-Cas System Components
- Panel B: Training Evo for CRISPR-Cas Design
- Panel C: CRISPR Subtype Generation Preferences
- Panel D: Sequence Identity Between Evo-Generated and Training Data
- Panel E: Structural Comparisons of Natural vs. Evo-Generated Cas9 Proteins
- Panel F: Evolution of crRNA-tracrRNA Duplex Structures

## Finetuning on CRISPR-Cas sequences enables generative design of protein-RNA complexes



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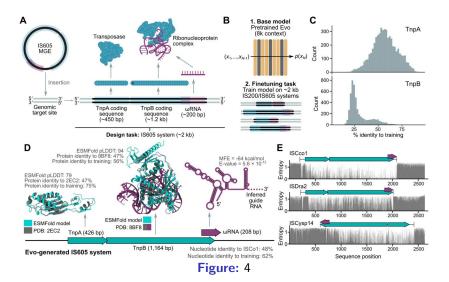
# Finetuning on IS200/IS605 sequences enables generative design of transposable biological systems

- Panel A: IS605 System Overview and Design Task
- Panel B: Training Evo for IS605 Generation
- Panel C: Sequence Identity Distribution of Generated IS605 Components
- Panel D: Structural Comparisons of Evo-Generated IS605 Proteins

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Panel E: Sequence Conservation and Entropy Analysis

# Finetuning on IS200/IS605 sequences enables generative design of transposable biological systems

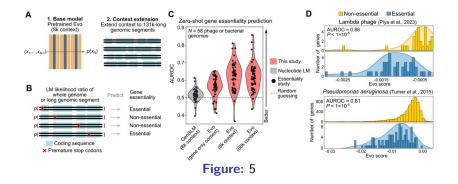


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## Evo performs zero-shot gene essentiality prediction across diverse bacterial and phage genomes

- Panel A: Expanding Context for Gene Essentiality Prediction
- Panel B: How Evo Predicts Gene Essentiality
- Panel C: Zero-Shot Gene Essentiality Prediction Performance
- Panel D: Gene Essentiality Scores Across Different Organisms

### Evo performs zero-shot gene essentiality prediction across diverse bacterial and phage genomes

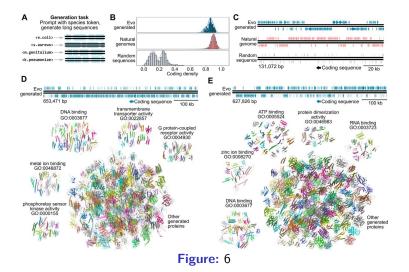


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### Evo generates genome-scale sequences with dense coding architecture

- 1. Panel A: Sequence Generation Task
- 2. Panel B: Coding Density Comparison
- 3. Panel C: Coding Sequence Comparison
- 4. Panel D: Functional Classification of Generated Proteins
- 5. Panel E: Functional Classification of Generated Genome

## Evo generates genome-scale sequences with dense coding architecture



#### 1. Evo: A Breakthrough in Genomic Modeling



Evo: A Breakthrough in Genomic Modeling
Ethical and Biosafety Considerations

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1. Evo: A Breakthrough in Genomic Modeling

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- 2. Ethical and Biosafety Considerations
- 3. Technical Challenges and Limitations

1. Evo: A Breakthrough in Genomic Modeling

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- 2. Ethical and Biosafety Considerations
- 3. Technical Challenges and Limitations
- 4. Future Research Directions

#### Thank You

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