



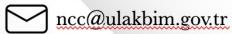


# EURO<sup>4SEE</sup>

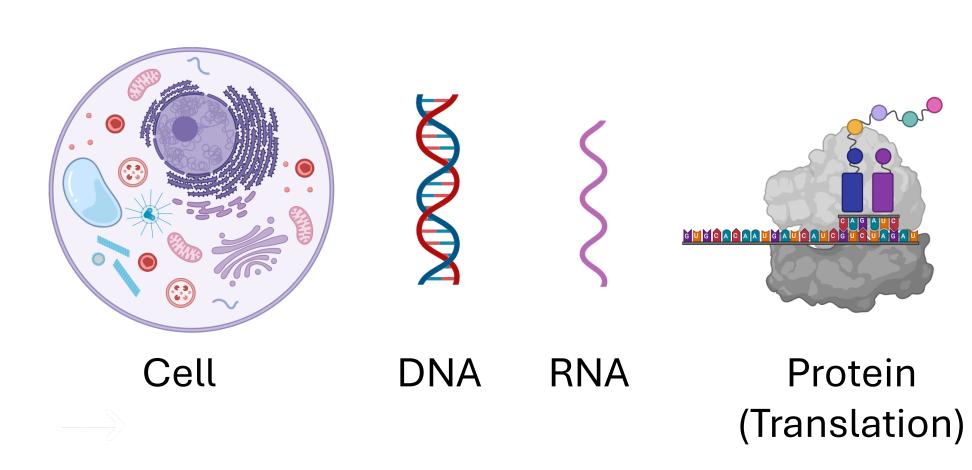
Generating Mutant Protein Structures Using Advanced Techniques

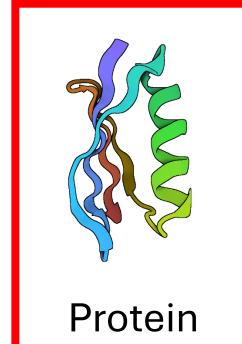
Tandac Furkan Guclu

Sabancı University



## Proteins?



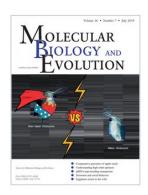


(Structure)

### Antibotic Resistance

#### Molecular Biology and Evolution

Issues More content ▼ Submit ▼ Alerts About ▼ Search



Volume 36, Issue 7 July 2019

**Article Contents** 

Abstract

JOURNAL ARTICLE

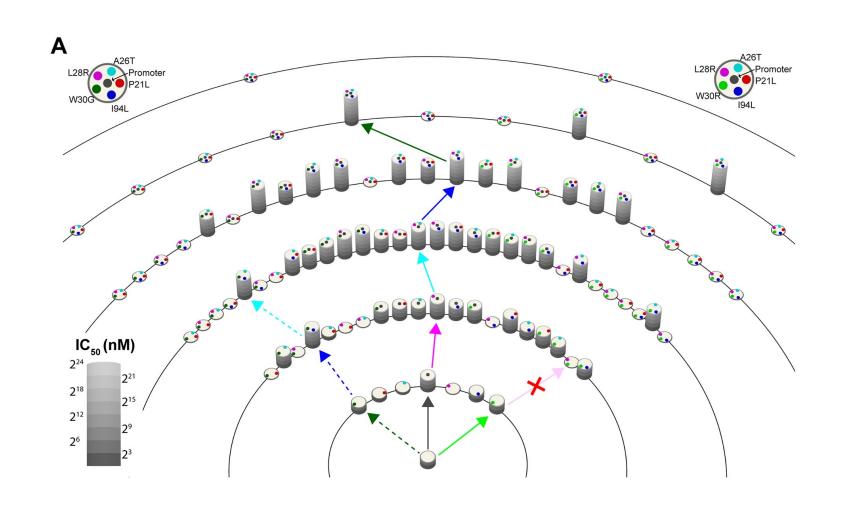
High-Order Epistasis in Catalytic Power of Dihydrofolate Reductase Gives Rise to a Rugged Fitness Landscape in the Presence of Trimethoprim Selection 8

Yusuf Talha Tamer, Ilona K Gaszek, Haleh Abdizadeh, Tugce Altinusak Batur, Kimberly A Reynolds, Ali Rana Atilgan, Canan Atilgan, Erdal Toprak ☒

Molecular Biology and Evolution, Volume 36, Issue 7, July 2019, Pages 1533–1550, https://doi.org/10.1093/molbev/msz086

Published: 15 April 2019

## Antibotic Resistance



## Fitness Landscape

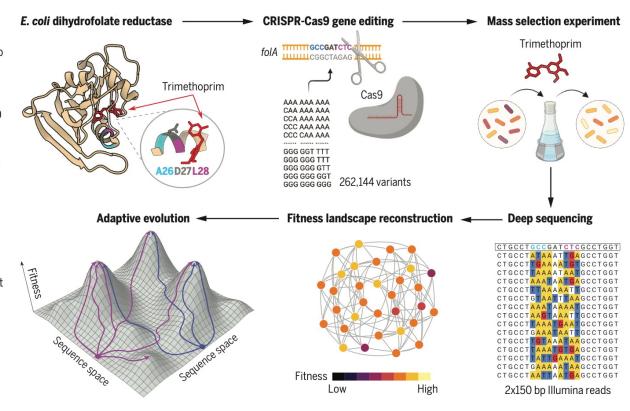
#### **EVOLUTIONARY BIOLOGY**

#### A rugged yet easily navigable fitness landscape

Andrei Papkou, Lucia Garcia-Pastor, José Antonio Escudero, Andreas Wagner\*

#### Empirical fitness landscape of dihydrofolate reductase.

We edited the E. coli genome to create a fitness landscape of all 64<sup>3</sup> codons encoding three consecutive amino acids (A, Ala; D, Asp; L, Leu) of the protein dihydrofolate reductase. We measured the fitness of each genotype in this landscape in the presence of the antibiotic trimethoprim using a mass selection experiment and deep sequencing. Even though the landscape is highly rugged, adaptive evolution can find the highest peaks from most starting locations via short and abundant fitness-increasing paths. [Created with BioRender.com1



## Deep Mutational Scanning

#### nature

Explore content > About the journal > Publish with us >

nature > articles > article

Article Published: 06 April 2022

## Mapping the energetic and allosteric landscapes of protein binding domains

Andre J. Faure, Júlia Domingo, Jörn M. Schmiedel, Cristina Hidalgo-Carcedo, Guillaume Diss & Ben Lehner 

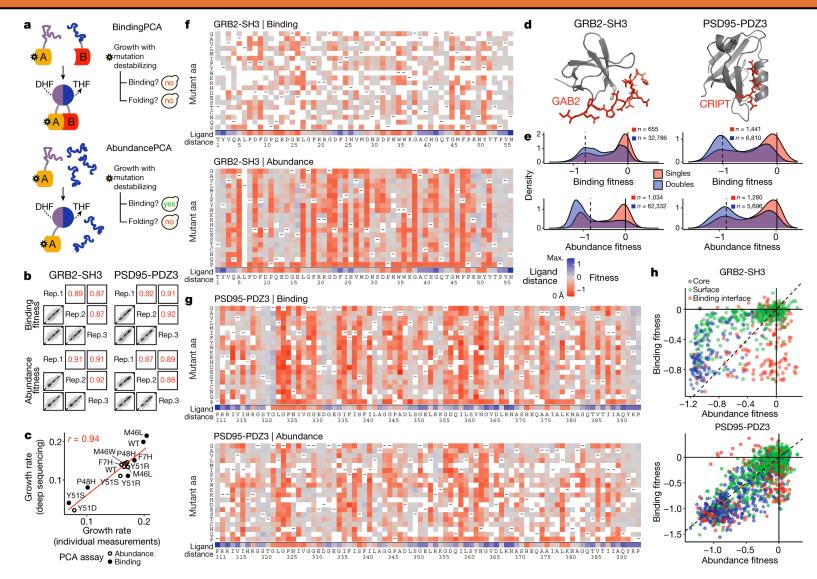
☐

Nature 604, 175–183 (2022) | Cite this article

44k Accesses | 226 Citations | 290 Altmetric | Metrics



## Deep Mutational Scanning





## Thanks!





This project has received funding from the European High-Performance Computing Joint Undertaking (JU) under grant agreement No 101191697. The JU receives support from the Digital Europe Programme and Germany, Türkiye, Republic of North Macedonia, Montenegro, Serbia, Bosnia and Herzegovina.