

## **Evangelos Papadopoulos, Ph.D.**

Newton, MA | 617-480-9485 | USA citizenship

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### **Professional Summary**

Accomplished research scientist with extensive experience in oncology, structural biology, and computational drug discovery. Proven expertise in high-throughput screening, gene editing technologies, and the design of novel therapeutic molecules. Passionate about advancing genomic medicine and translating complex biological insights into impactful healthcare solutions.

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### **Professional Experience**

#### **Research Scientist**

**Dana Farber Cancer Institute** | Boston, MA | 2020 – Present

- Led research on disease-relevant mutations in multiple myeloma and breast cancer using CRISPR/Cas9 and bioinformatics.
- Conducted high throughput docking of millions of compounds for novel drug discovery and repurposing.
- Managed a team to perform DNA cloning, cell culture, and drug assays, ensuring experimental precision and reproducibility.
- Visualized protein structures to assess mutation impacts and align findings with FDA-approved therapies.

#### **Senior Scientist**

**Stablix Inc.** | New York, NY | 2021 – 2023

- Designed bifunctional chemical molecules for targeted de-ubiquitination therapies.
- Implemented high-throughput virtual screens, docking 5 million compounds to identify novel binding pockets.
- Collaborated with cross-functional teams to optimize workflows for protein purification and drug assays.
- Curated extensive ELN databases, ensuring traceability for thousands of experimental results.

## **Research Scientist**

### **PIC Therapeutics** | Boston, MA | 2017 – 2020

- Developed protein translation inhibitors as cancer therapeutics, including testing on diverse cell lines.
- Docked over 100 million compounds to crystal structures, advancing small molecule drug development.
- Partnered with CROs to optimize assay pipelines, accelerating discovery timelines.

## **Research Scholar**

### **Harvard Medical School** | Boston, MA | 2009 – 2017

- Conducted structural biology research on protein translation initiation inhibitors using NMR and X-ray crystallography.
- Published groundbreaking findings, including the first structure of eIF4E with a novel drug (PDB Molecule of the Month).
- Designed and tested PROTAC analogs to inhibit oncogenic proteins.

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## **Education**

### **Post Doctoral in Molecular Biology**

Beth Israel Deaconess Medical Center | Boston, MA | 2008-2009

### **PhD in Biophysics**

Stockholm University | Stockholm, Sweden | 2008

### **MSc in Biophysics**

Stockholm University | Stockholm, Sweden

### **MSc in Applied Physics**

National Technical University of Athens | Athens, Greece

### **Bachelor diploma in Physics**

Kapodistrian University of Athens | Athens, Greece

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## **Key Skills**

- Computational drug design and virtual screening (ZINC, ENAMINE, MCULE)
- High-throughput assay development and execution (SPR, MST, ITC, ASMS, DEL)
- Structural biology techniques (NMR, X-ray crystallography, AlphaFold)
- Gene editing technologies (CRISPR/Cas9) and bioinformatics pipelines
- Cross-disciplinary project leadership and team management

## Publications and Awards

- Multiple peer-reviewed publications on oncology and drug discovery (e.g., *PNAS*, *Nature Communications*). <https://orcid.org/0000-0003-4050-4521>
  - Recognized with the PDB Molecule of the Month for contributions to structural biology. <http://pdb101.rcsb.org/motm/230>
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## Nonprint Materials

- NMR constraints [Script Library]. Stockholm (SE); 2006  
[https://pymolwiki.org/index.php/Nmr\\_cnstr](https://pymolwiki.org/index.php/Nmr_cnstr)
  - NMR solution structure of peptide from Doppel protein, in DHPC micelles.  
<https://evanspap.w3spaces.com/>
  - GitHub scripts for Computational Structural Biology:  
<https://github.com/evanspap/>
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## Professional References

- Gerhard Wagner, Ph.D., Harvard Medical School.,  
Elkan Blout Professor of Biological Chemistry and Molecular Pharmacology.  
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