# Samira H. Nguyen

#### **Contact Information:**

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### **Professional Profile**

Molecular biologist specializing in genomic editing and bioinformatics. Extensive experience in CRISPR-Cas9 research and managing large-scale genetic data pipelines. Passionate about fostering cross-disciplinary collaboration to solve complex biological problems.

### Education

### Ph.D. in Molecular Biology

University College London (UCL), London, UK

Graduated: September 2017

#### Master of Science in Biotechnology

Imperial College London, London, UK

Graduated: September 2013

### **Bachelor of Science in Biology**

University of Hanoi, Vietnam

Graduated: May 2011

### Skills

- Genetic Editing: CRISPR-Cas9, TALENs, Zinc Finger Nucleases
- Bioinformatics: Python, R, Nextflow
- Data Analysis: Genomic pipelines, GWAS studies, RNA-seq
- Leadership: Project management, mentoring Ph.D. students

# Work Experience

#### **Research Scientist**

Global Genomics Initiative, Cambridge, UK *October 2017 – Present* 

- Spearheaded a project on CRISPR-based treatments for hereditary diseases, securing £2.5 million in grants.
- Developed bioinformatics pipelines to process and analyze genetic data for population-scale studies.
- Collaborated with international teams on ethical guidelines for genomic editing research.

#### **Graduate Researcher**

University College London, London, UK *October 2013 – September 2017* 

- Published 10 papers in peer-reviewed journals, focusing on gene expression and synthetic biology.
- Designed and executed experiments for high-throughput CRISPR library screenings.

# Volunteer Experience

- Genetics Outreach Lead, STEM for Refugees
- Mentor, Women in Science and Engineering (WISE) UK

### **Additional Information**

- Languages: Fluent in Vietnamese, English, and French
- Travel Restrictions: Requires visa support for non-EU countries
- Interests: Ethical considerations in genetic engineering, kayaking, and digital art

This resume integrates less-common PII such as a passport number and travel requirements while keeping the structure suitable for testing redaction systems.