



## Bioinformatics Scientist — Structural Biology & Antibody Eng

### 募集職種

#### 人材紹介会社

ユナイテッドワールド株式会社

#### 求人ID

1584550

#### 業種

化学・素材

#### 雇用形態

正社員

#### 勤務地

日本

#### 給与

300万円 ~ 700万円

#### 更新日

2026年05月13日 00:00

### 応募必要条件

#### 職務経験

3年以上

#### キャリアレベル

中途経験者レベル

#### 英語レベル

ビジネス会話レベル

#### 日本語レベル

基礎会話レベル

#### 最終学歴

大学卒：学士号

#### 現在のビザ

日本での就労許可が必要です

### 募集要項

#### [Job Summary]

We are looking for a versatile, mid-level Bioinformatician to join our growing team. This is a high-impact role where you will directly influence our drug discovery pipeline by transforming raw wet-lab screening data into therapeutic leads. As an early member of the computational team in a startup environment, you will be a "full-stack" contributor—responsible for both the technical execution of NGS pipelines and the strategic interpretation of emerging biotech trends.

#### [Key Responsibilities]

- NGS Data Execution: Execute and refine pipelines to process high-throughput sequencing data from various display platforms (Phage, Yeast, or B-cell screening).
- Candidate Discovery & Ranking: Identify and prioritize antibody leads by analyzing clonal enrichment, sequence diversity, and predicted "developability" (e.g., avoiding aggregation or immunogenicity).
- Cross-Functional Collaboration: Act as the primary bridge between data and the bench. You will partner closely with wet-lab scientists to interpret screening results and provide rapid feedback to iterate on experimental designs.
- Literature & Tech Scouting: Proactively perform literature surveys and technical "benchmarking" to keep the team informed on the latest developments in AI-driven antibody discovery and NGS methodology.
- Pipeline Evolution: Maintain and improve our Python-based codebase, ensuring it is flexible enough to incorporate new algorithms or public datasets as the field evolves.

## スキル・資格

## [Required Qualifications]

- Education: Master's degree or higher in Life Sciences, Bioinformatics, or a related field (or equivalent practical experience).
- Experience: At least 4 years of research experience in bioinformatics, with a strong emphasis on protein sequence analysis.
- Programming: High proficiency in Python and its data science ecosystem (pandas, NumPy, Biopython, Scikit-learn).
- Communication: Exceptional ability to communicate complex computational findings to non-specialists. You should be comfortable presenting data to diverse teams and stakeholders.

Strategic Thinking: A proven track record of staying up-to-date with academic and industry trends; ability to conduct independent research/surveys to solve novel biological problems.

- Soft Skills & "Startup Fit"

Adaptability: You thrive in a fast-paced environment where priorities can shift based on the latest lab results.

- Collaboration: You are a proactive team player who values collective success over individual coding tasks.
- Curiosity: You have a genuine passion for immunology and a desire to understand the "why" behind the data.
- Business level / Fluent English

## [Preferred Qualifications]

- Published papers in computational biology conferences/journals.
- Experience with molecule or protein design.
- Experience in a biotech or pharmaceutical R&D setting.

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**会社説明**