

AI-Powered Tools for Ancestry and Population Genomics in Healthcare

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Abstract

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The convergence of Artificial Intelligence (AI) and genomics is rapidly transforming precision medicine. The application of AI-powered tools to **ancestry and population genomics** is unlocking new capabilities in disease risk prediction, drug development, and personalized healthcare strategies. For digital health and AI professionals, understanding this synergy is crucial, as it represents a fundamental shift toward more equitable and accurate genomic medicine [1].

The Role of AI in Genomic Data Analysis

Genomic data is characterized by its immense scale and complexity. Population-level studies aggregate and analyze data from thousands to millions of individuals. Traditional statistical methods often struggle to discern the subtle, non-linear patterns within this data that are indicative of disease risk or drug response.

This is where AI, particularly **Machine Learning (ML)** and **Deep Learning (DL)**, excels. AI algorithms can process vast, high-dimensional genomic datasets—including Single Nucleotide Polymorphisms (SNPs), transcriptomic data, and clinical records—to identify complex genotype-phenotype relationships [2] [3]. Key applications include:

Variant Interpretation: Identifying and classifying genetic variants of uncertain significance (VUS) by predicting their pathogenicity [4]. **Disease Risk Prediction:** Developing sophisticated models, such as **Polygenic Risk Scores (PRS)**, that integrate thousands of genetic markers to estimate an individual's lifetime risk for complex diseases like cancer, diabetes, and cardiovascular conditions [5].

Ancestry and the Challenge of Bias in Genomic Data

A critical challenge in population genomics is the historical

also guide preventative interventions and personalize treatment pathways with unprecedented accuracy, ultimately transforming healthcare from reactive to proactive.

References

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