## Potential Biases in AI Treatment Recommendations

The Cancer Genome Atlas (TCGA) represents a landmark achievement in cancer genomics, yet its application in Al-driven personalized medicine raises critical ethical concerns regarding representation and fairness. \*\*Racial and ethnic underrepresentation\*\* constitutes the most significant bias: approximately 77% of TCGA participants are of European ancestry, while African Americans comprise only 10%, Hispanic/Latino patients 5%, and Asian populations 3%. This demographic skew creates Al models that perform optimally for white patients while potentially recommending suboptimal or even harmful treatments for underrepresented minorities.

\*\*Genomic diversity disparities\*\* compound this issue. Tumors from different ethnic groups exhibit distinct mutational signatures and drug response profiles—for instance, EGFR mutations in lung cancer occur in 40% of Asian patients versus 15% in Caucasians. An Al trained predominantly on European data may fail to recognize these population-specific biomarkers, leading to missed therapeutic opportunities or inappropriate treatment selection for minority patients. \*\*Socioeconomic and geographic biases\*\* further stratify access: TCGA datasets overrepresent patients from well-resourced urban academic medical centers who could afford genomic sequencing, systematically excluding rural, uninsured, and economically disadvantaged populations. This creates algorithms that inadvertently optimize treatments for affluent patients while neglecting feasibility constraints facing underserved communities.

\*\*Survival outcome biases\*\* pose additional concerns. Patients with access to cutting-edge therapies and multidisciplinary care—disproportionately those from advantaged backgrounds—demonstrate longer survival, which the AI interprets as treatment efficacy rather than systemic healthcare inequality. The model then perpetuates these disparities by recommending aggressive interventions that assume similar resource availability for all patients. Finally, \*\*temporal and technological biases\*\* mean that older sequencing technologies overrepresent certain cancer types studied earlier, while emerging immunotherapies remain underrepresented, potentially causing AI systems to overlook breakthrough treatments for rare malignancies.

Addressing these biases requires comprehensive technical and policy interventions. \*\*Diversifying training datasets\*\* must be the foundation: actively recruiting participants from underrepresented ethnicities through community partnerships, providing free genomic testing to low-income patients, and collaborating internationally to include global genomic diversity. Organizations should mandate that AI training sets reflect population demographics, with minimum representation thresholds (e.g.,  $\geq 15\%$  for each major ethnic group).

\*\*Algorithmic fairness techniques\*\* can mitigate learned biases. Implementing fairness constraints during model training—such as demographic parity or equalized odds—ensures that prediction accuracy and treatment recommendations remain consistent across ethnic groups. Regular algorithmic audits disaggregated by race, ethnicity, socioeconomic status, and geography can identify disparate impacts before deployment. \*\*Federated learning approaches\*\* allow collaborative model training across institutions without centralizing sensitive patient data, enabling contribution from safety-net hospitals and international centers serving diverse populations.

\*\*Transparency and interpretability\*\* are essential for identifying bias. Explainable AI frameworks (SHAP values, attention mechanisms) should clearly articulate which genomic features drive each recommendation, allowing clinicians to detect when decisions rely on ethnicity-correlated features rather than causal biomarkers. Establishing independent ethics review boards with diverse stakeholder representation—including patient advocates from minority communities—provides oversight and accountability. Finally, \*\*clinical trial equity\*\* must become mandatory: requiring AI-recommended therapies to demonstrate efficacy across diverse subgroups before regulatory approval, with adequately powered minority cohorts, prevents replicating historical clinical research disparities where new treatments primarily benefited white patients.

Ultimately, ethical AI in personalized medicine demands recognizing that technical solutions alone cannot overcome structural healthcare inequities. Parallel investments in improving healthcare access, addressing social determinants of health, and dismantling institutional racism remain essential for ensuring AI advances medicine equitably for all patients, regardless of ancestry, geography, or economic status.