## UNDERREPRESENTATION OF MINORITIES IN **GENOMIC DATABASES** Sara Martín García, Simon Alkema, Esmeé Haenen and Niels Boersbroek

Genomic research should be for everyone. However, our modern society is very diverse, and genomic databases are still not representative for everyone in society.

The current Genome of Europe project showcases this lack of equal representation of different ethnic groups.

## HELP: what is a genome again?

A genome is the complete set of DNA (genetic material) of a person. While everyone's genome is unique, differences between people's genome are very small. Scientists combine the genomes of many people to create a reference genome, which is a representive average for those people.

**BE CAREFUL!** There is no genetic basis for race, it is a social construct. However, genetic differences do exist and they can affect diseases and drug effectiveness.



## **Consequences of lack of diversity**



**Misrepresentation of** cultural identities can have legal and economic consequences, negative



The underrepresentation of diverse populations limits genetic



There's a participation barrier due to historial mistrust and systemic racism and discrimination



**Structural barriers like lack** of culturally sensitive strategies, research focuses more on data rather than



The lack of representation leads to an inequitable benefit distribution as health disparities widen and the cycle of exclusion

stereotypes can be reinforced

understanding and therefore leads to a less accurate diagnosis

this furthers polarisation

people

and inequity is reinforced

## Website of The Genome of Europe



How do we make everyone benefit from genomic research?

- Advocating for **policies that promote inclusivity**, such as including information about genetic variation inside of countries in guidelines in the Genome of Europe project.
- Ensuring **representation** of marginalised communities.
- **Building trust** between scientists and underrepresented groups.
- **Raising awareness** in society and the scientific community.

CBS. (2022). StatLine—Bevolking; kerncijfers. Bevolking; kerncijfers. https://opendata.cbs.nl/statline/#/CBS/nl/dataset/85496NED/table?ts=1719306614591

Lemke AA, Esplin ED, Goldenberg AJ, Gonzaga-Jauregui C, Hanchard NA, Harris-Wai J, Ideozu JE, Isasi R, Landstrom AP, Prince AER, Turbitt E, Sabatello M, Schrier Vergano SA, Taylor MRG, Yu JH, Brothers KB, Garrison NA. Addressing underrepresentation in genomics research through community engagement. Am J Hum Genet. 2022 Sep 1;109(9):1563-1571. doi: 10.1016/j.ajhg.2022.08.005. PMID: 36055208; PMCID: PMC9502069.

Images from Canva