Identity-by-descent in HPRCv2

Franco Marsico, PhD

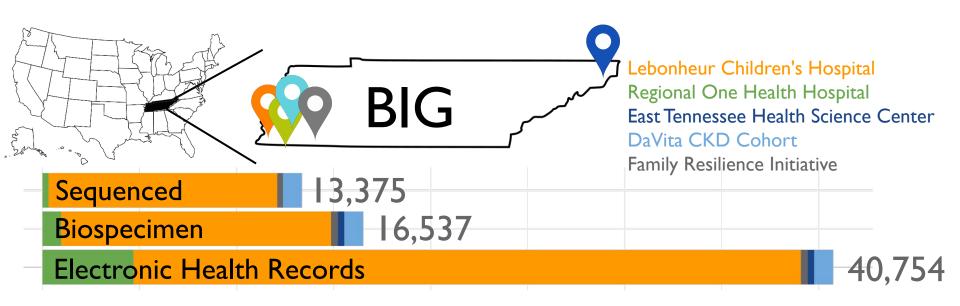
franco.lmarsico@gmail.com

University of Tennessee Health Science Center

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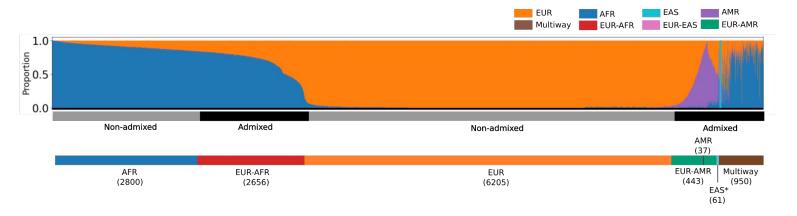


The Biorepository for Integrative Genomics initiative





Ancestry inference in BIG



- Four continental level ancestries
- Relevant proportion of admixed individuals
- High genetic diversity geographically concentrated

nature communications

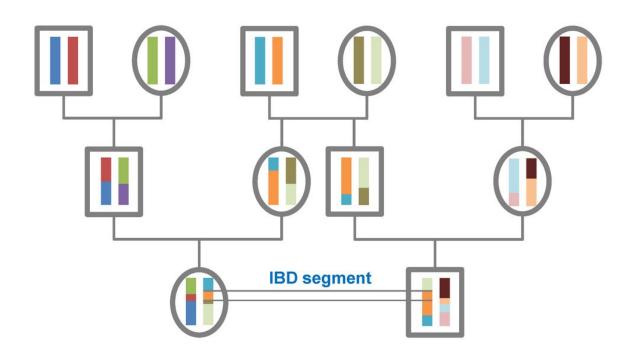
Insights from the Biorepository and Integrative Genomics pediatric resource

Received: 6 December 2024	Silvia Buonaiuto @ 1.10°, Franco Marsico 1.2.10°, Akram Mohammed @ 3, Lokesh K. Chinthala @ 3, Ernestine K. Arnos-Abanyie @ 3, Regeneron Genetics Center*, Pjotr Prins*, Khyobeni Mozhui @ 1.4°, Robert J. Rooney 1.5°, Robert W. Williams 1.6°, Robert L. Davis @ 3, Terri H. Finkel 7, Chester W. Brown 1.4° & Vincenza Colonna @ 1.2.5° \(\)
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Check for updates	

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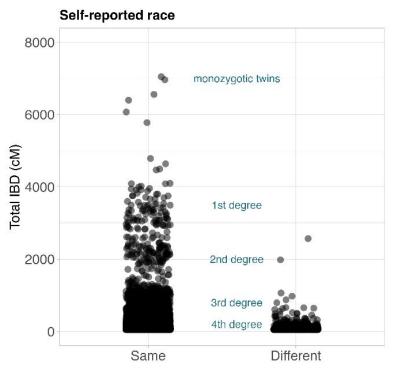


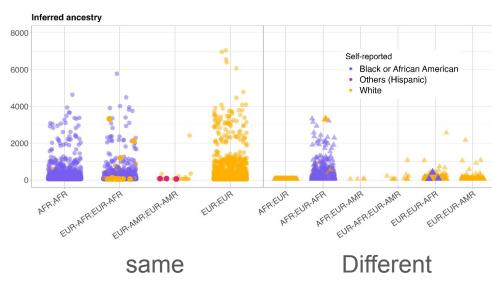
Identity by Descent (IBD)





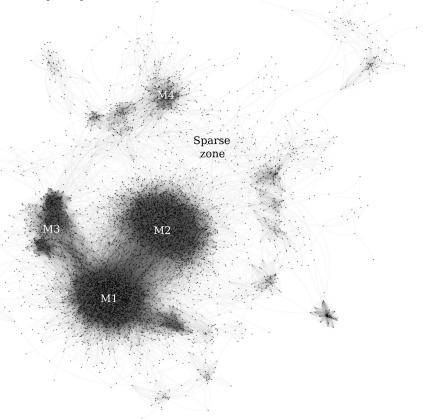
Self-reported race and genetic ancestry through the lens of relatedness



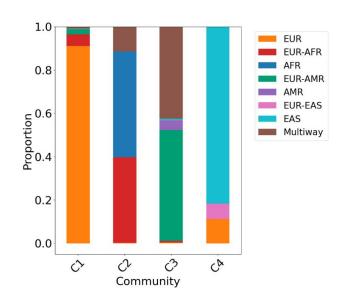




From pairwise IBD to population structure



Leiden algorithm for community detection



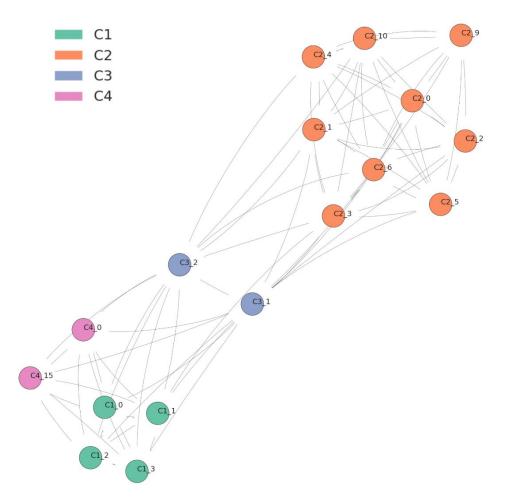


Beyond ancestry: grouping individuals based on how much genome they actually share

Identity-by-descent captures Shared Environmental
Factors at Biobank Scale

Franco Marsico^{1, 2, †}, Silvia Buonaiuto¹, Ernestine K Amos—Abanyie¹, Lokesh K Chinthala³, Akram Mohammed³, Regeneron Genetics Center⁴, Robert J Rooney^{1,5}, Robert W Williams^{1,6}, Robert L Davis³, Terri H Finkel⁷, Chester W Brown^{1, 8}, Pjotr Prins¹, and Vincenza Colonna^{1, 2, 5, †}

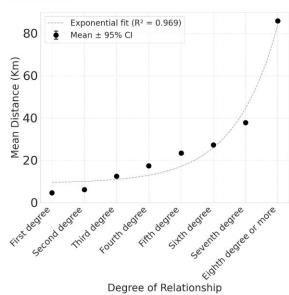
https://www.biorxiv.org/content/10.1101/2025.05.03.652048v1.full

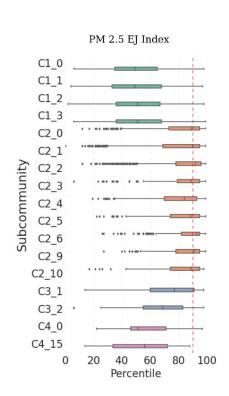


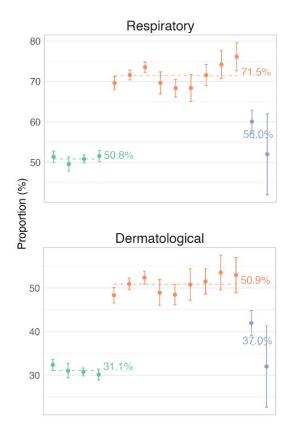


Identity-by-descent captures shared environment

Genealogical and geographic distance







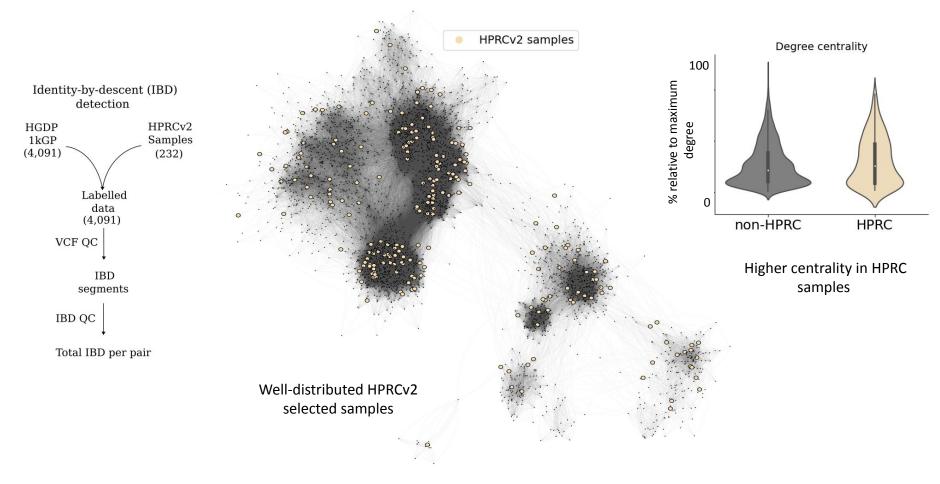


Can a similar approach be

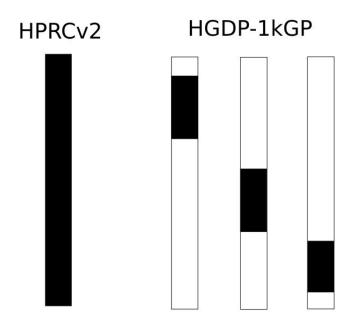
useful on HPRCv2?

Preliminary/pilot results

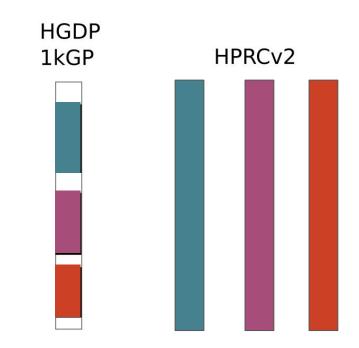








Haplotype informativeness: Compute the amount of cM explained by IBD sharing between the selected samples and public resources (HGDP-1kGP).



Proportion already explained: Compute the percentage (amount of cM) already explored by IBD sharing of a potential HGDP-1kGP sample to be incorporated to HPRCv2.



Thank you

https://github.com/MarsicoFL/HPRCv2-IBD

