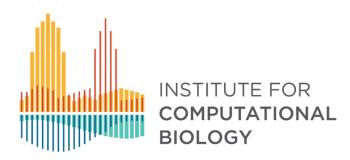
WHY BOTHER WITH DIVERSITY IN PRECISION MEDICINE RESEARCH?



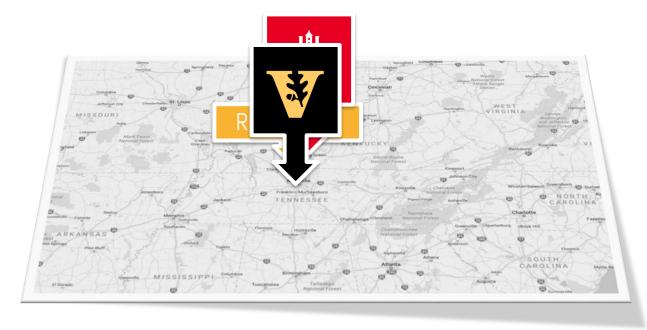


October 13, 2018

William S. Bush Associate Professor Institute for Computational Biology Population and Quantitative Health Sciences

My Journey to Genomics and Precision Medicine







WHAT IS PRECISION MEDICINE?

Before giving you a medical treatment, we ideally want to know:

- Will this treatment be effective for you?
- What dose do you need?
- Are you at increased risk for side effects from this treatment?





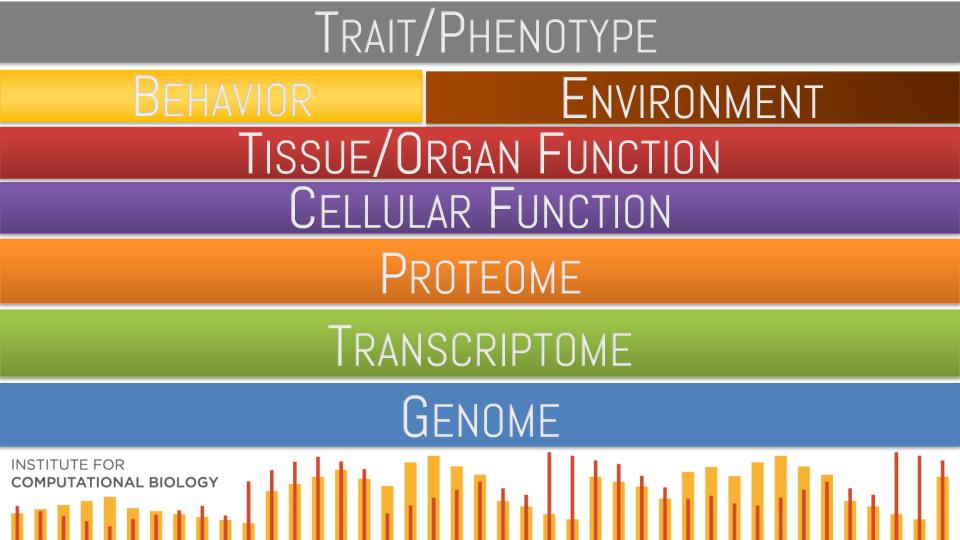
Infinite Diversity in Infinite Combinations



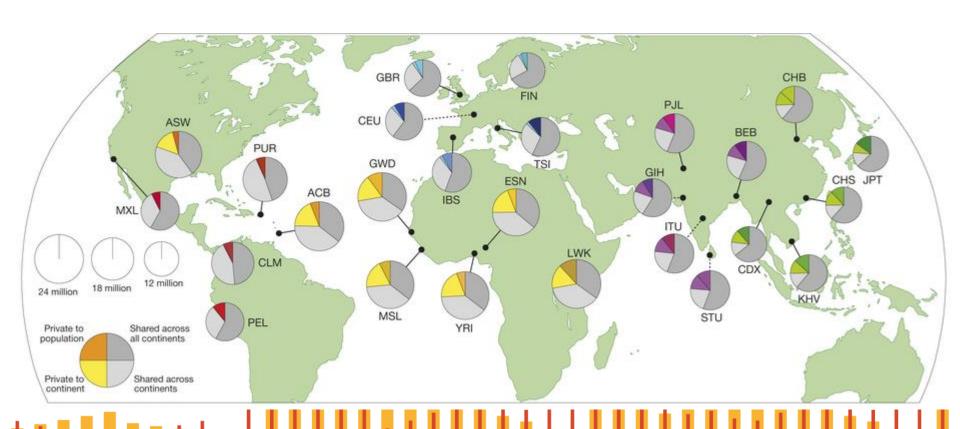




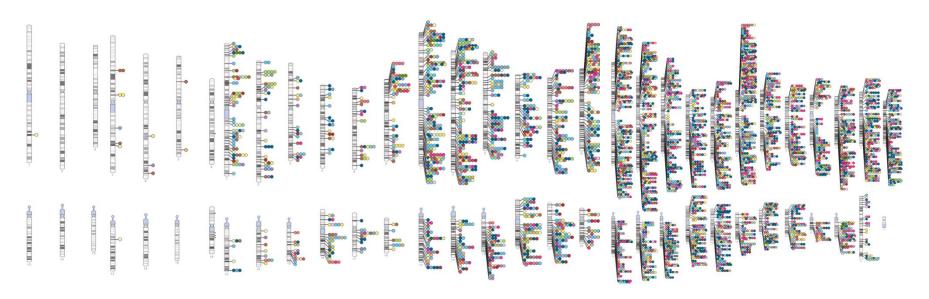




GENOMIC DIVERSITY OF HUMAN POPULATIONS



THE STATE OF GENOMIC DISCOVERY





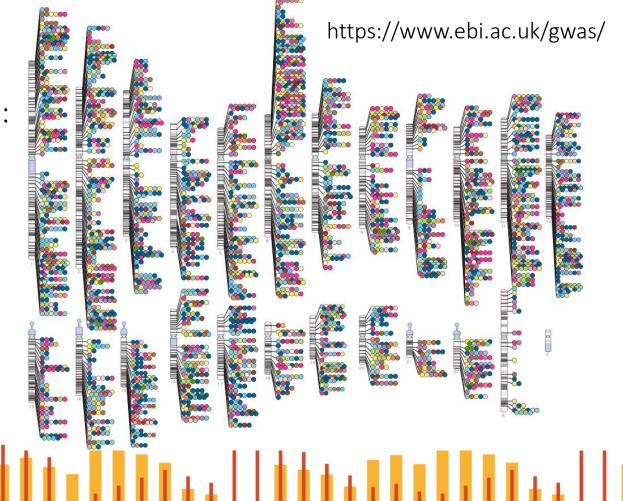
GWAS as of 9/30/2018:

3,596 publications

75,827 SNP-trait associations

INSTITUTE FOR

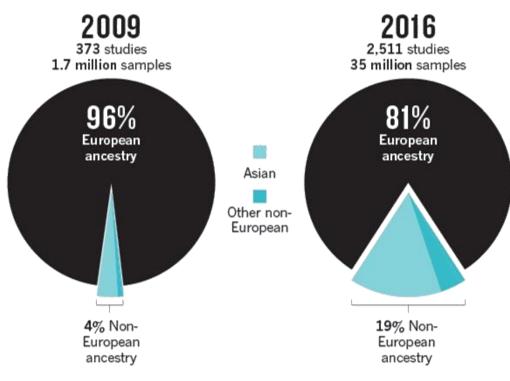
COMPUTATIONAL BIOLOGY



MOSTLY EUROPEAN-DESCENT

Bustamante, Burchard, de la Vega (2011) *Nature* 475:163-165

Popejoy and Fullerton (2016) *Nature* 538, 161–164



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DIVERSITY IN RESEARCH MATTERS

SPECIAL ARTICLE

Genetic Misdiagnoses and the Potential for Health Disparities

Arjun K. Manrai, Ph.D., Birgit H. Funke, Ph.D., Heidi L. Rehm, Ph.D., Morten S. Olesen, Ph.D., Bradley A. Maron, M.D., Peter Szolovits, Ph.D., David M. Margulies, M.D., Joseph Loscalzo, M.D., Ph.D., and Isaac S. Kohane, M.D., Ph.D.

N Engl J Med 2016; 375:655-665 August 18, 2016 DOI: 10.1056/NEJMsa1507092

REPORT

Association of Trypanolytic ApoL1 Variants with Kidney Disease in African Americans

Giulio Genovese^{1,2,*}, David J. Friedman^{1,3,*}, Michael D. Ross⁴, Laurence Lecordier⁵, Pierrick Uzureau⁵, Barry I. Freedman⁶, Donald W. Bowden^{7,8}, Carl D. Langefeld^{8,9}, Taras K. Oleksyk¹⁰, Andrea L. Uscinski Knob⁴, Andrea J. Bernhardy¹, Pamela J. Hicks^{7,8}, George W. Nelson¹¹, Benoit Vanhollebeke⁵, Cheryl A. Winkler¹², Jeffrey B. Kopp¹¹, Etienne Pays^{5,†}, Martin R. Pollak^{1,13,†}

+ Author Affiliations

 μ [†]To whom correspondence should be addressed. E-mail: mpollak@bidmc.harvard.edu (M.R.P.); epays@ulb.ac.be (E.P.) μ * These authors contributed equally to this work.

Science 13 Aug 2010: Vol. 329, Issue 5993, pp. 841-845 DOI: 10.1126/science.1193032



Precision Medicine Initiative Cohort Program

All of Us

Large

Linked to EHR

Oversample subgroups

Re-contact for return of results

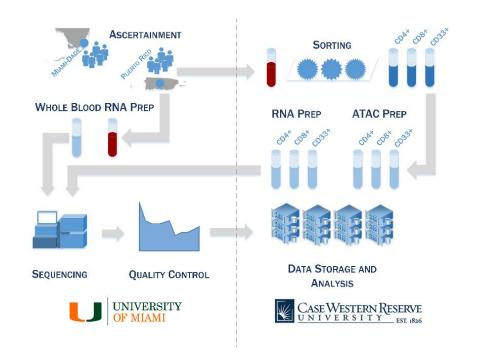
Wide age range Lifestyle data collection

Rare & common disease Extensive follow-up





MULTIPLE SCLEROSIS RISK IN PUERTO RICO



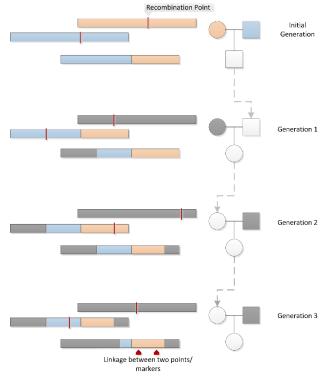


Alliance for Research in Hispanic Multiple Sclerosis





TRACING ANCESTRAL CHROMOSOMES



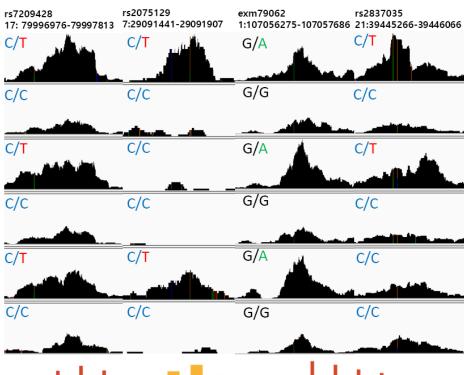
Initial Generation 100 Generations 1000 Generations

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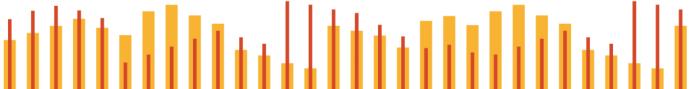
Population moves from Linkage Disequilibrium to Linkage
Equilibrium over time

Decay of Linkage over successive generations

ATAC-SEQUENCING



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ADDING DIVERSITY TO STUDIES OF ALZHEIMER'S DISEASE

- Alzheimer's Disease
 Sequencing Project
- Whole genome sequence
- Addition of Dominican,
 Puerto Rican, and African
 American participants





