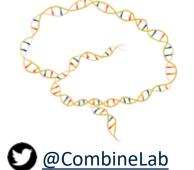


Evidence For Bias Of Genetics Ancestry In Resting State Functional MRI



Andre Altmann

Department of Medical Physics and Biomedical Engineering



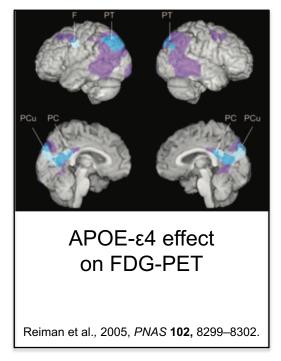
Centre for Medical Image Computing (CMIC)

a.altmann@ucl.ac.uk

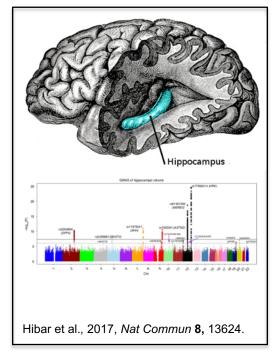
Venice, April 8th 2019



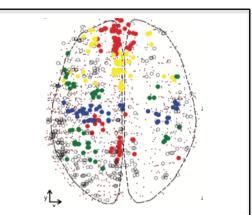
Topics in imaging genetics



Quantify genetic effects



Discover genetic effects

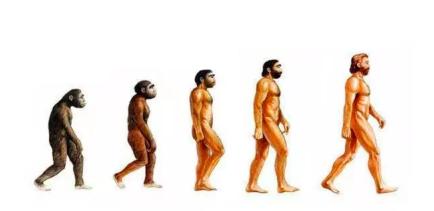


Which genes contribute to functional connectivity?

Richiardi et al., 2015, Science 348, 1241-4.

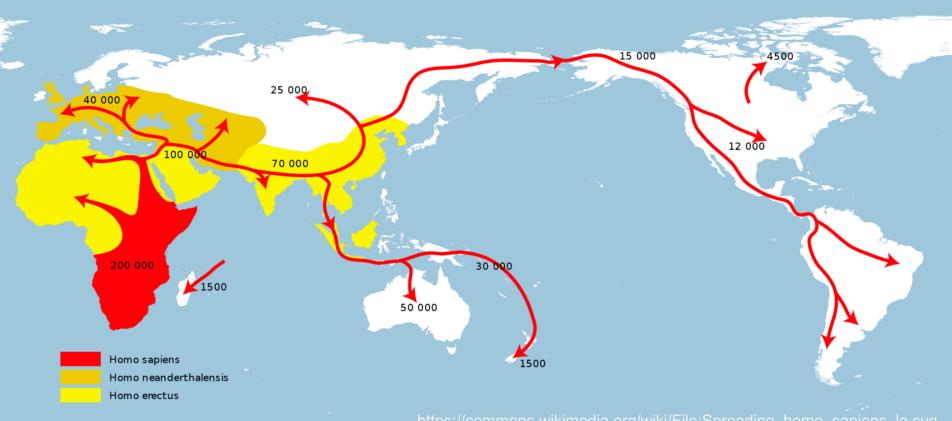
Understand organization

Genetics as 'historic baggage' in precision medicine



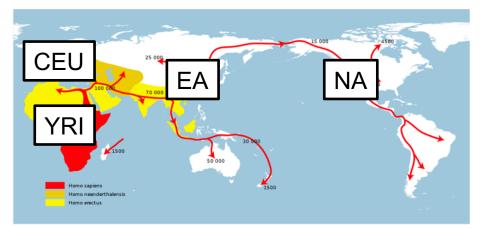
https://www.quora.com/

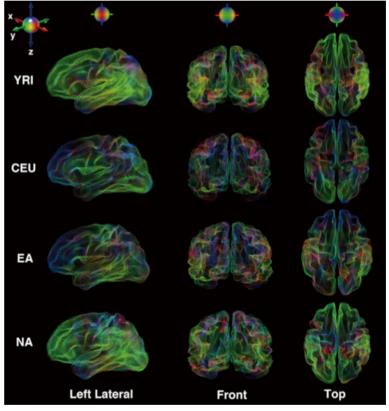
Human migration



ittps://commons.wikimedia.org/wiki/File:Spreading_homo_sapiens_la.svg

Genetic ancestry and brain morphology

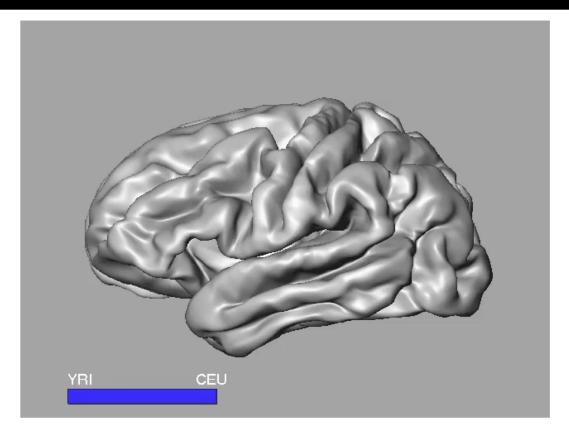




Fan et al., 2015, Current Biology 25, 1988–1992

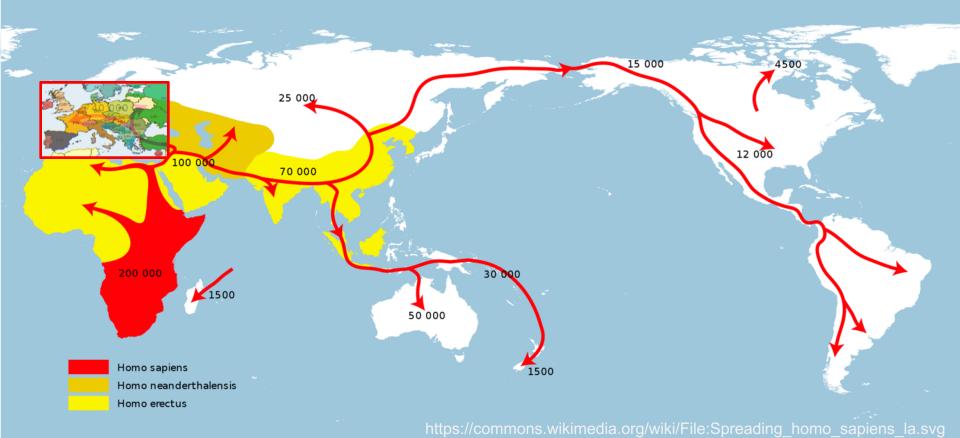
Genetic ancestry and brain morphology





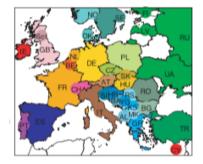
Fan et al., 2015, *Current Biology* 25, 1988–1992

Human migration



Genes mirror geography within Europe

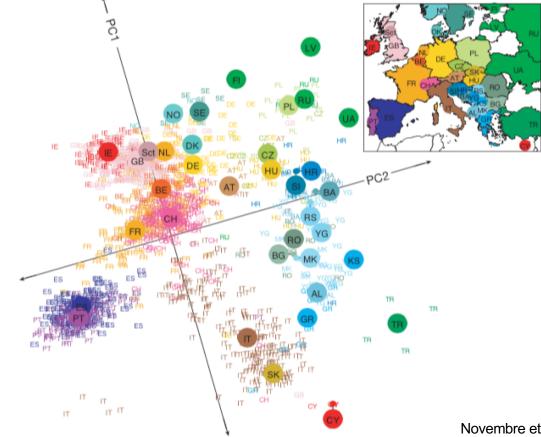




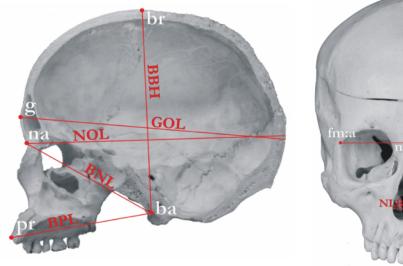
Novembre et al., 2008, Nature 456, 98–101

Genes mirror geography within Europe



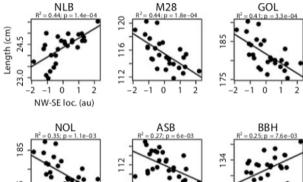


Geography cline of skull and brain morphology in Europe



http://osteomics.com/cranExplr/





 $\mathbf{b} = \begin{bmatrix} \mathbf{b} \\ -\mathbf{c} \\ -\mathbf{c$

Bakken et al., 2011, Hum Hered 72, 35-44

Geography cline of skull and brain morphology in Europe

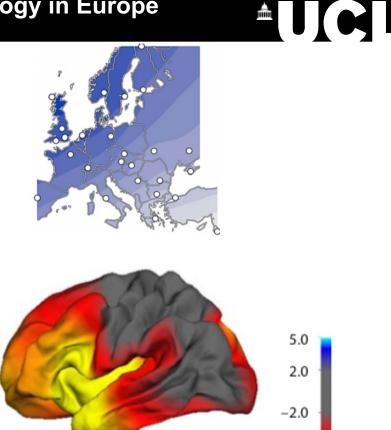
0.06 PC1 0.04 0.02 PC1 (rotated) 0 0 -0.04 -0.06 а -0.05 Ó 0.05 PC2 (rotated) Cortical area ICV Brain volume 4×10^{5} 00 4×10^{4} orght p = 3.5e-04 3×10^5 p = 5.7e - 04p = 1.3e - 04Residuals -2 -4×10^{4} -3×10^{5}

-0.02

0.04 0.08

0.04 0.08

-0.02



Bakken et al., 2011, Hum Hered 72, 35-44

-5.0

b

0.04 0.08

NW-SE loc. (au)

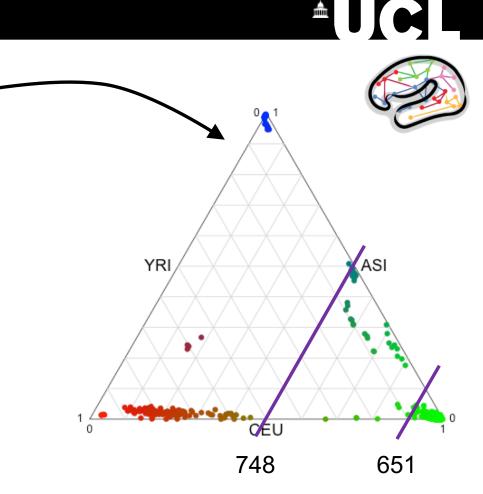
-0.02



Does this evolutionary bias extend to functional brain measures?

Genetic ancestry in HCP

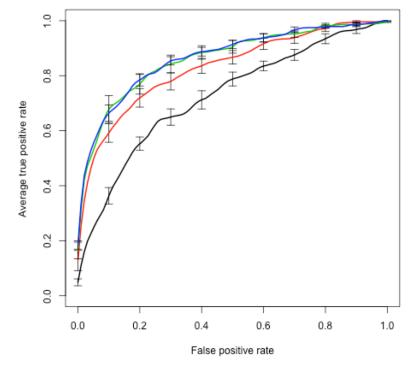
- 950 young adults with
 - Genome-wide SNP data
 - rs-fMRI (1h)
- Aim: classify CEU vs non-CEU
 - ICA: 15 to 300
 - Elastic net classifier
 - Nested CV
 - 10 outer folds (family aware)
 - 5 inner folds (optimize λ)
 - ROC curves

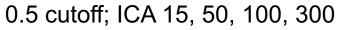


Genetic ancestry in HCP

- Genetic ancestry can be predicted from rs-fMRI
- Works best with ICA \ge 100
- Works better for 0.5 cutoff

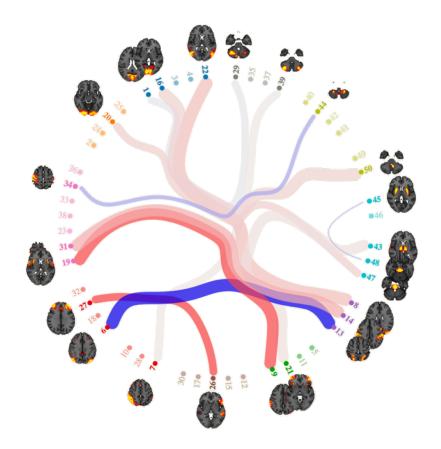
ICA	dim	$p_{\scriptscriptstyle ext{CEU}} > 0.5$	$p_{ m CEU} > 0.9$
15	105	0.78 (0.088)	0.72 (0.049)
25	300	0.81 (0.060)	0.76 (0.061)
50	1225	0.86 (0.055)	0.83 (0.063)
100	4950	0.91 (0.046)	0.86 (0.061)
200	19900	0.92 (0.039)	0.88 (0.055)
300	44850	0.93 (0.036)	0.87 (0.032)





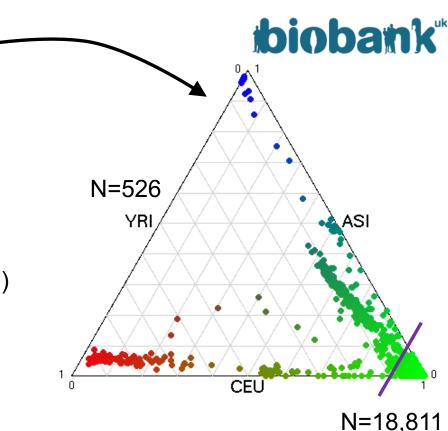
Which connections are important?





Genetic ancestry in UK Biobank

- 19,337 adults with
 - Genome-wide SNP data
 - rs-fMRI (6 min)
- Aim: classify CEU vs non-CEU
 - ICA: "25" and "100" (21 and 55)
 - Elastic net classifier
 - 100 bootstraps, subsample CEU (3:1)
 - Nested CV
 - 10 outer folds
 - 5 inner folds (optimize λ)
 - AUC



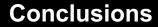
Genetic ancestry in UK Biobank

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ICA 25 • AUC 0.72 ICA100 • AUC 0.81					
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	44850	0.93 (0.036)	0.87 (0.032)		

thinhank







- Genetic ancestry influences skull and brain morphology
 - Interpretation in precision medicine
- Genetic ancestry can be predicted from rs-fMRI
 Presumably effects from brain normalization
- Consider to adjust brain imaging studies for genetic ancestry