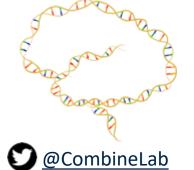
### 

#### **Evidence For Bias Of Genetics Ancestry In Resting State Functional MRI**



#### Andre Altmann

Department of Medical Physics and Biomedical Engineering



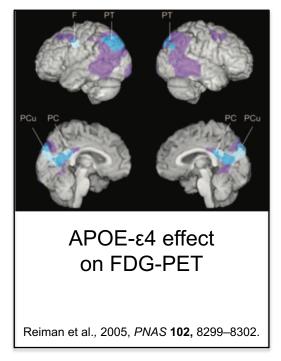
Centre for Medical Image Computing (CMIC)

a.altmann@ucl.ac.uk

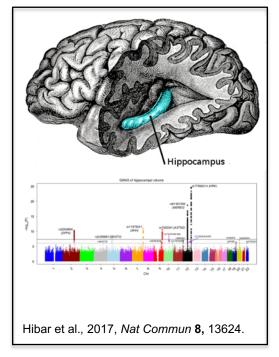
Venice, April 8th 2019



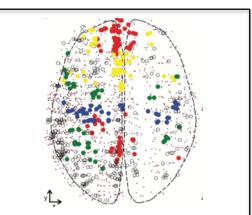
#### **Topics in imaging genetics**



Quantify genetic effects



Discover genetic effects

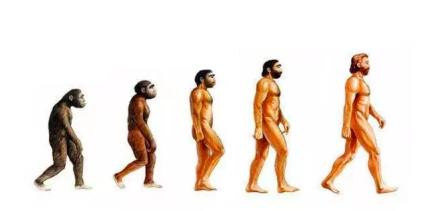


Which genes contribute to functional connectivity?

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

Understand organization

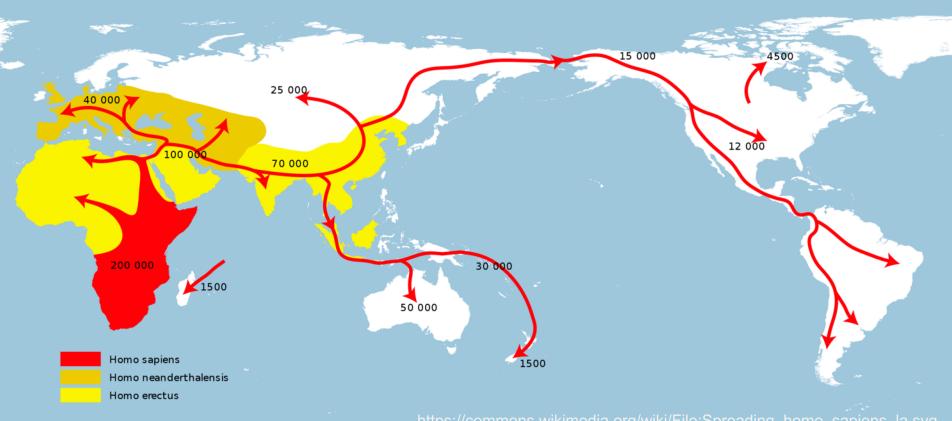
#### Genetics as 'historic baggage' in precision medicine



https://www.quora.com/

#### Human migration

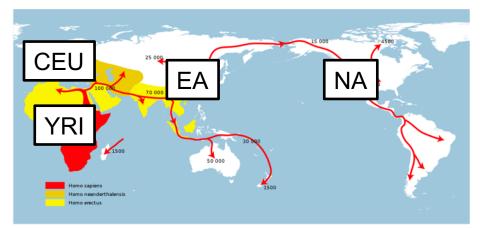
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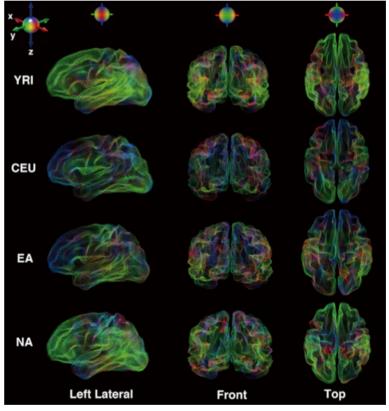


ittps://commons.wikimedia.org/wiki/File:Spreading\_homo\_sapiens\_la.svg

#### Genetic ancestry and brain morphology

# 

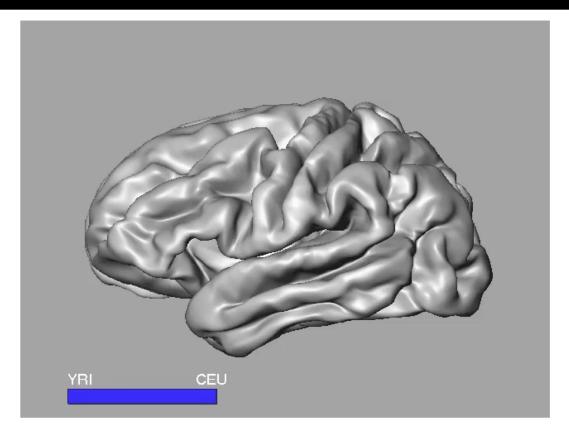




Fan et al., 2015, Current Biology 25, 1988–1992

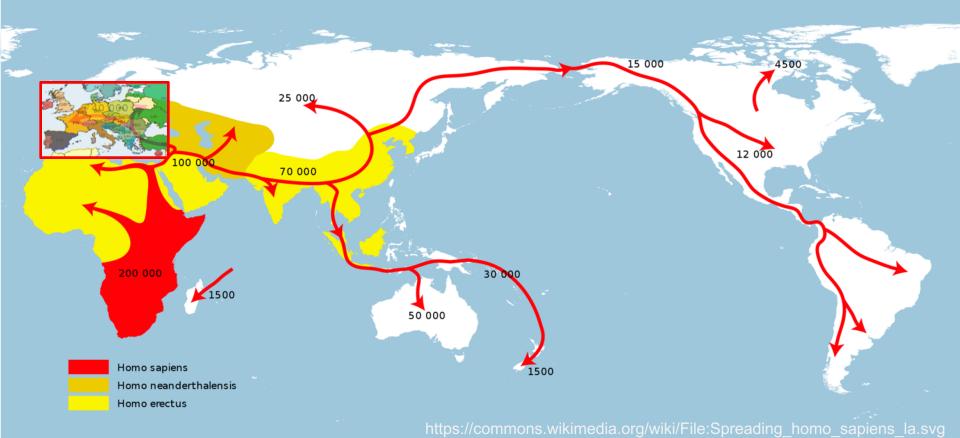
#### Genetic ancestry and brain morphology





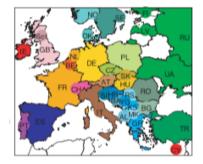
Fan et al., 2015, *Current Biology* 25, 1988–1992

#### Human migration



#### Genes mirror geography within Europe

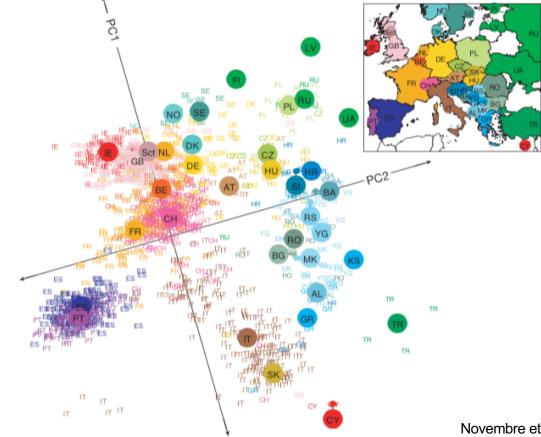




Novembre et al., 2008, Nature 456, 98–101

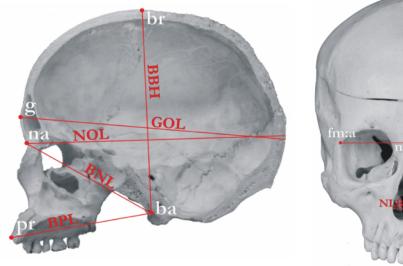
#### Genes mirror geography within Europe





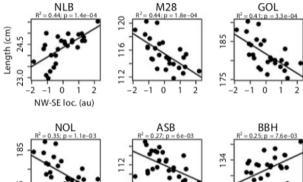
#### Geography cline of skull and brain morphology in Europe

## 



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

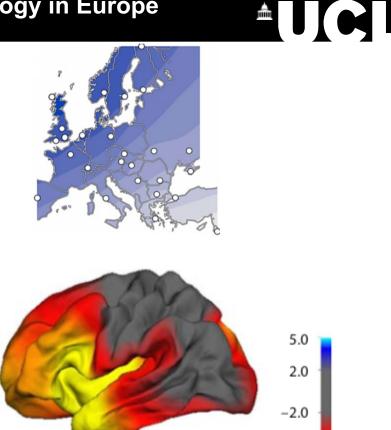
#### 0.06 PC1 0.04 0.02 PC1 (rotated) 0 0 -0.04 -0.06 а -0.05 Ó 0.05 PC2 (rotated) Cortical area ICV Brain volume $4 \times 10^{5}$ 00 $4 \times 10^{4}$ orght p = 3.5e-04 $3 \times 10^5$ p = 5.7e - 04p = 1.3e - 04Residuals -2 $-4 \times 10^{4}$ $-3 \times 10^{5}$

-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)

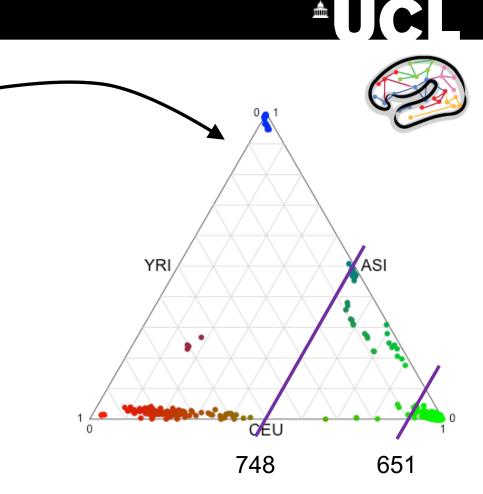
-0.02



# Does this evolutionary bias extend to functional brain measures?

#### **Genetic ancestry in HCP**

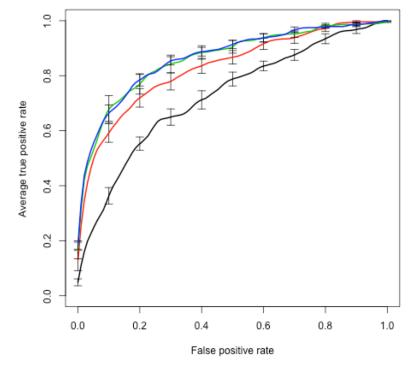
- 950 young adults with
  - Genome-wide SNP data
  - rs-fMRI (1h)
- Aim: classify CEU vs non-CEU
  - ICA: 15 to 300
  - Elastic net classifier
  - Nested CV
    - 10 outer folds (family aware)
    - 5 inner folds (optimize  $\lambda$ )
  - ROC curves

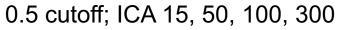


#### **Genetic ancestry in HCP**

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

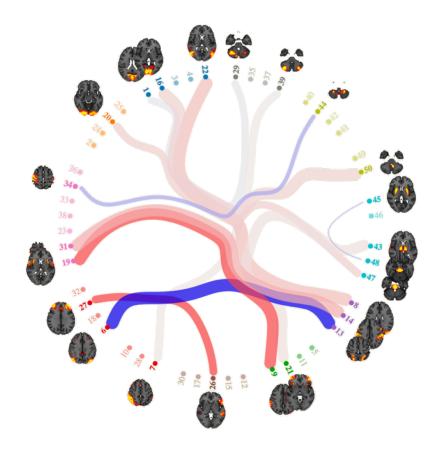
| ICA | dim   | $p_{\scriptscriptstyle 	ext{CEU}} > 0.5$ | $p_{ m CEU} > 0.9$ |
|-----|-------|--|--------------------|
| 15  | 105   | 0.78 (0.088)                             | 0.72 (0.049)       |
| 25  | 300   | 0.81 (0.060)                             | 0.76 (0.061)       |
| 50  | 1225  | 0.86 (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)       |





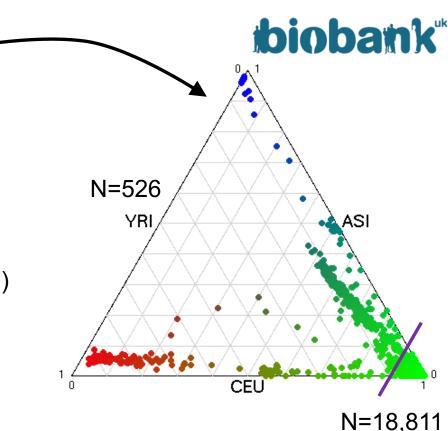
#### Which connections are important?





#### 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  $\lambda$ )
  - AUC



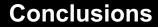
#### Genetic ancestry in UK Biobank

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    - 10 outer folds
    - 5 inner folds (optimize  $\lambda$ )
  - AUC

| ICA 25<br>• AUC 0.72<br>ICA100<br>• AUC 0.81 |       |                       |                    |  |  |
|--|-------|-----------------------|--------------------|--|--|
| ICA  | dim   | $p_{	ext{CEU}} > 0.5$ | $p_{ m CEU} > 0.9$ |  |  |
| 15   | 105   | 0.78 (0.088)          | 0.22 (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)       |  |  |
|  | 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