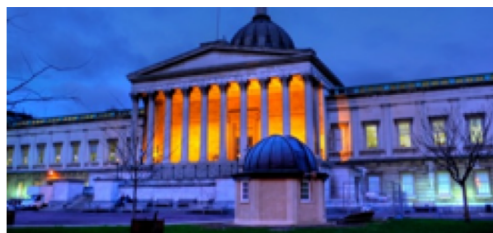


## Evidence For Bias Of Genetics Ancestry In Resting State Functional MRI



[@CombineLab](https://twitter.com/CombineLab)

**Andre Altmann**

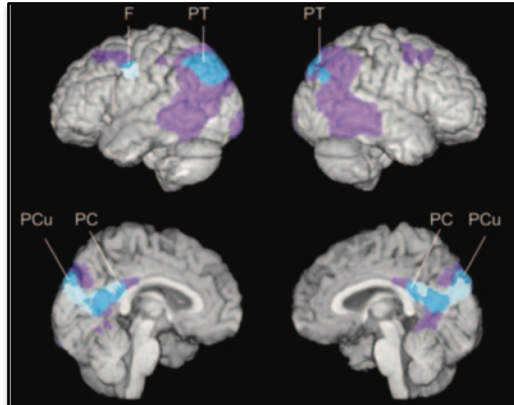
Department of Medical Physics and Biomedical Engineering

Centre for Medical Image Computing (CMIC)

[a.altmann@ucl.ac.uk](mailto:a.altmann@ucl.ac.uk)

Venice, April 8<sup>th</sup> 2019

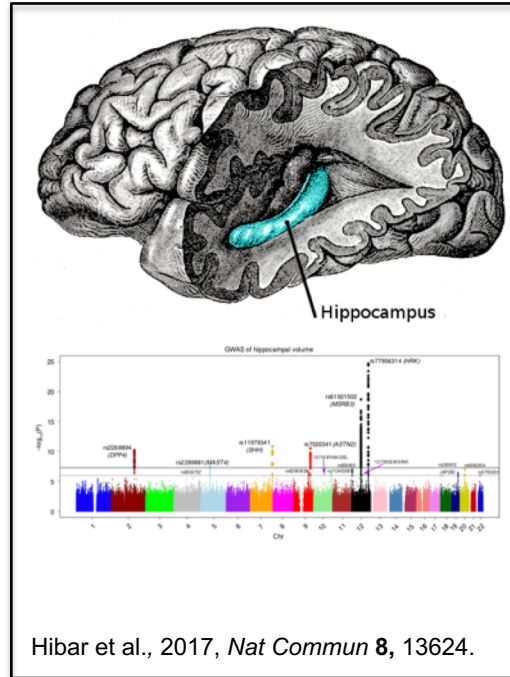




APOE- $\epsilon$ 4 effect  
on FDG-PET

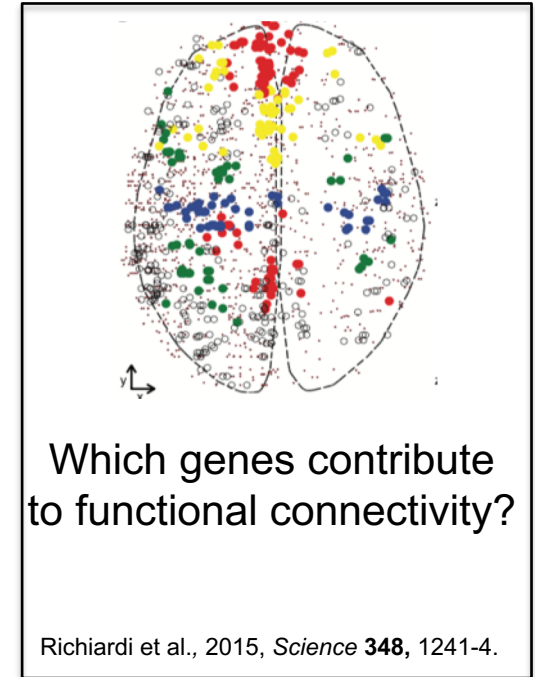
Reiman et al., 2005, *PNAS* **102**, 8299–8302.

Quantify genetic effects



Hibar et al., 2017, *Nat Commun* **8**, 13624.

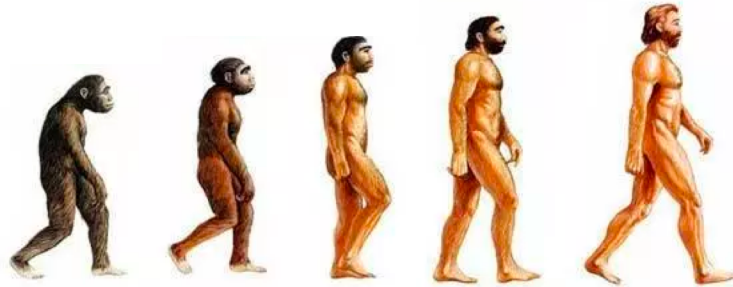
Discover genetic effects



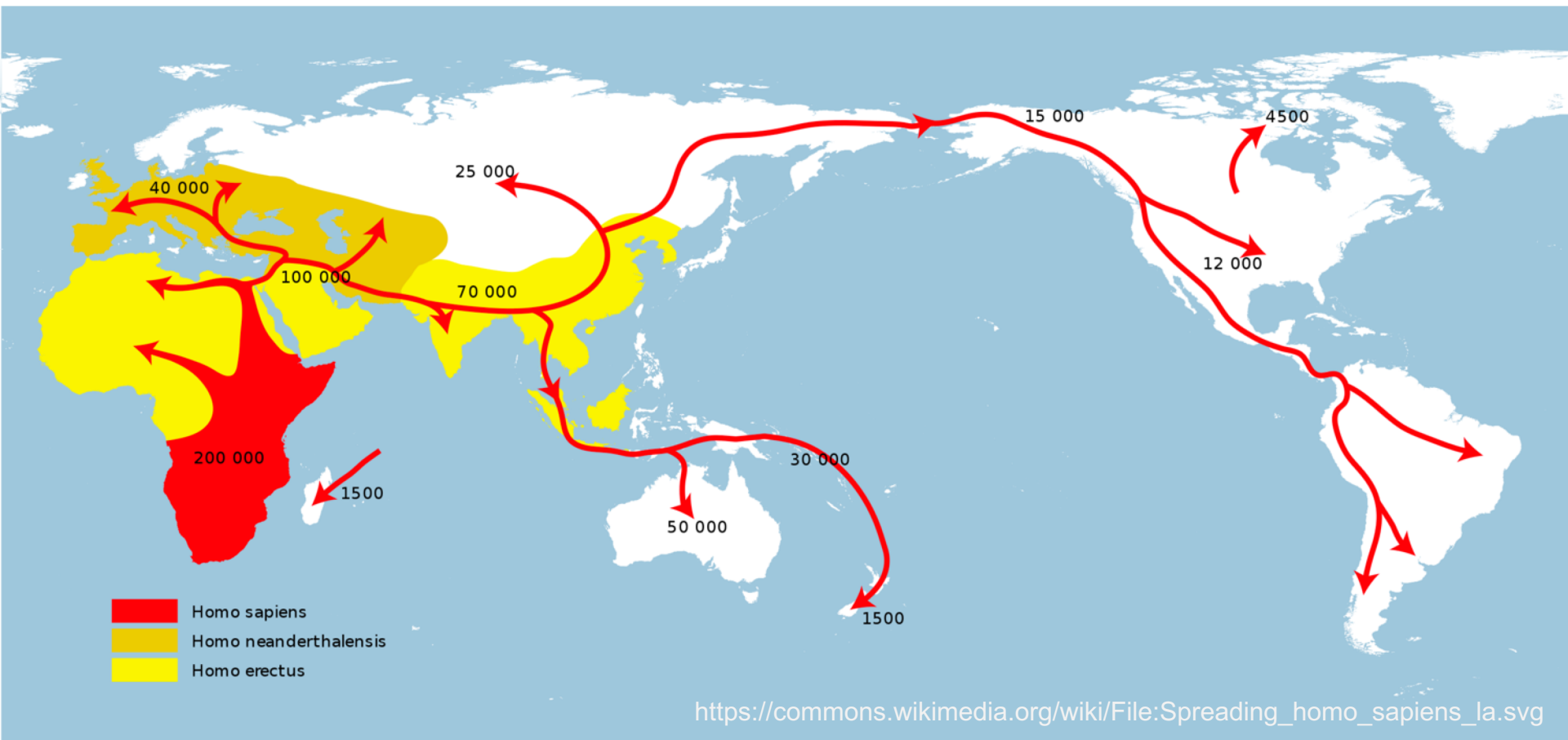
Which genes contribute  
to functional connectivity?

Richiardi et al., 2015, *Science* **348**, 1241–4.

Understand organization

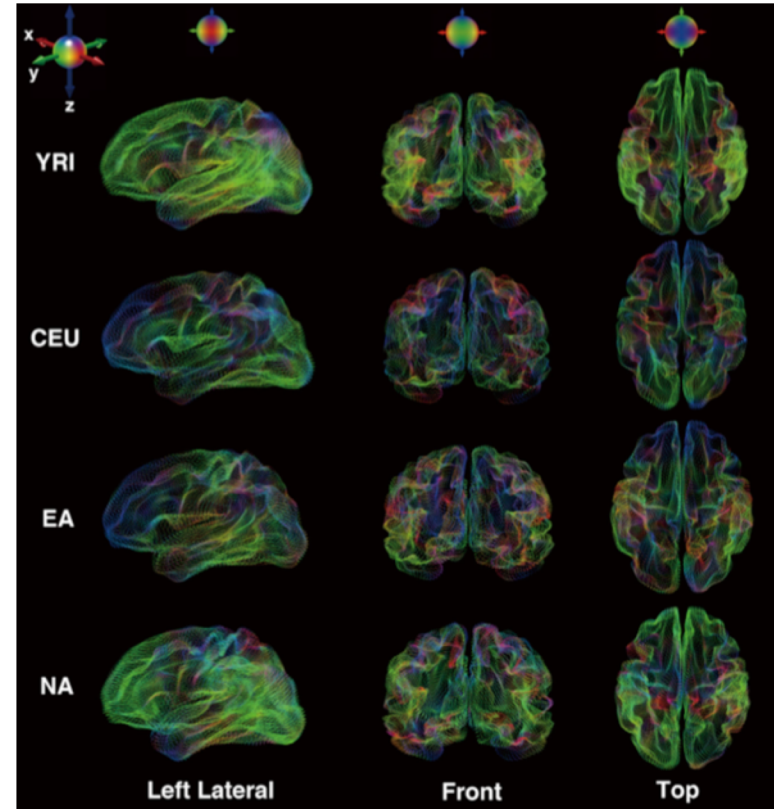
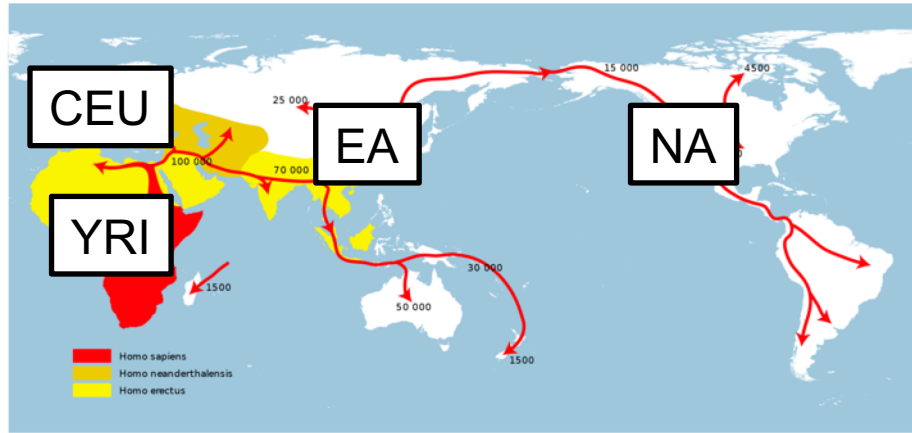


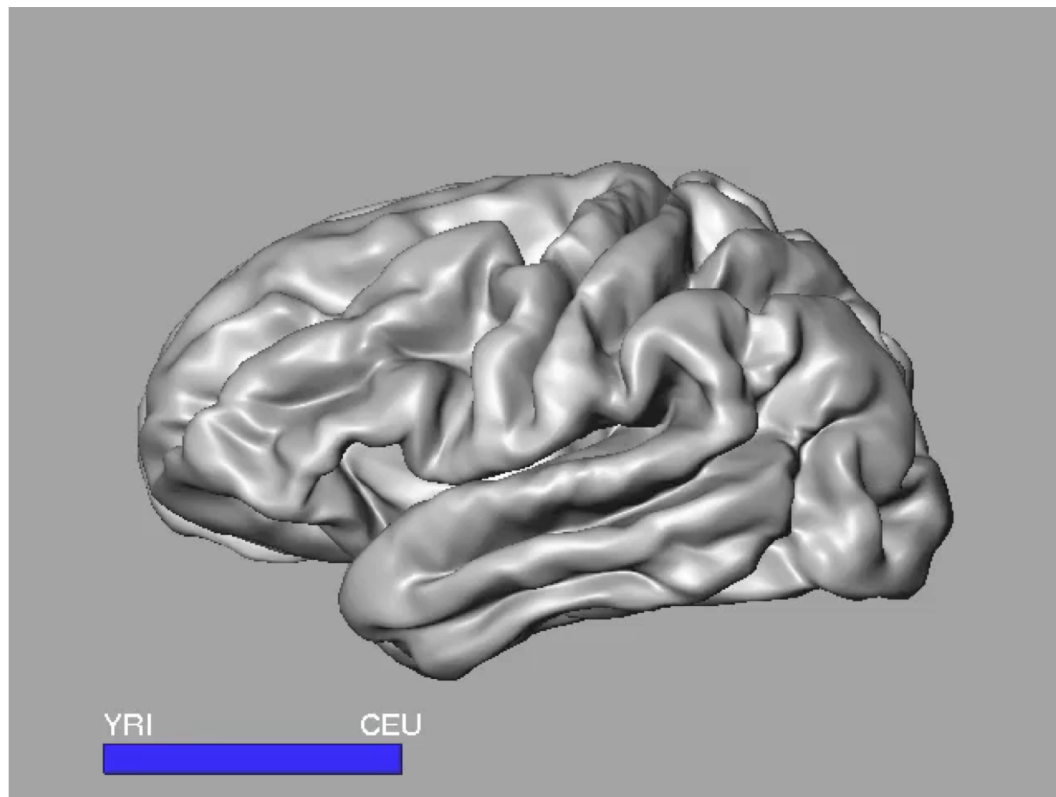
# Human migration



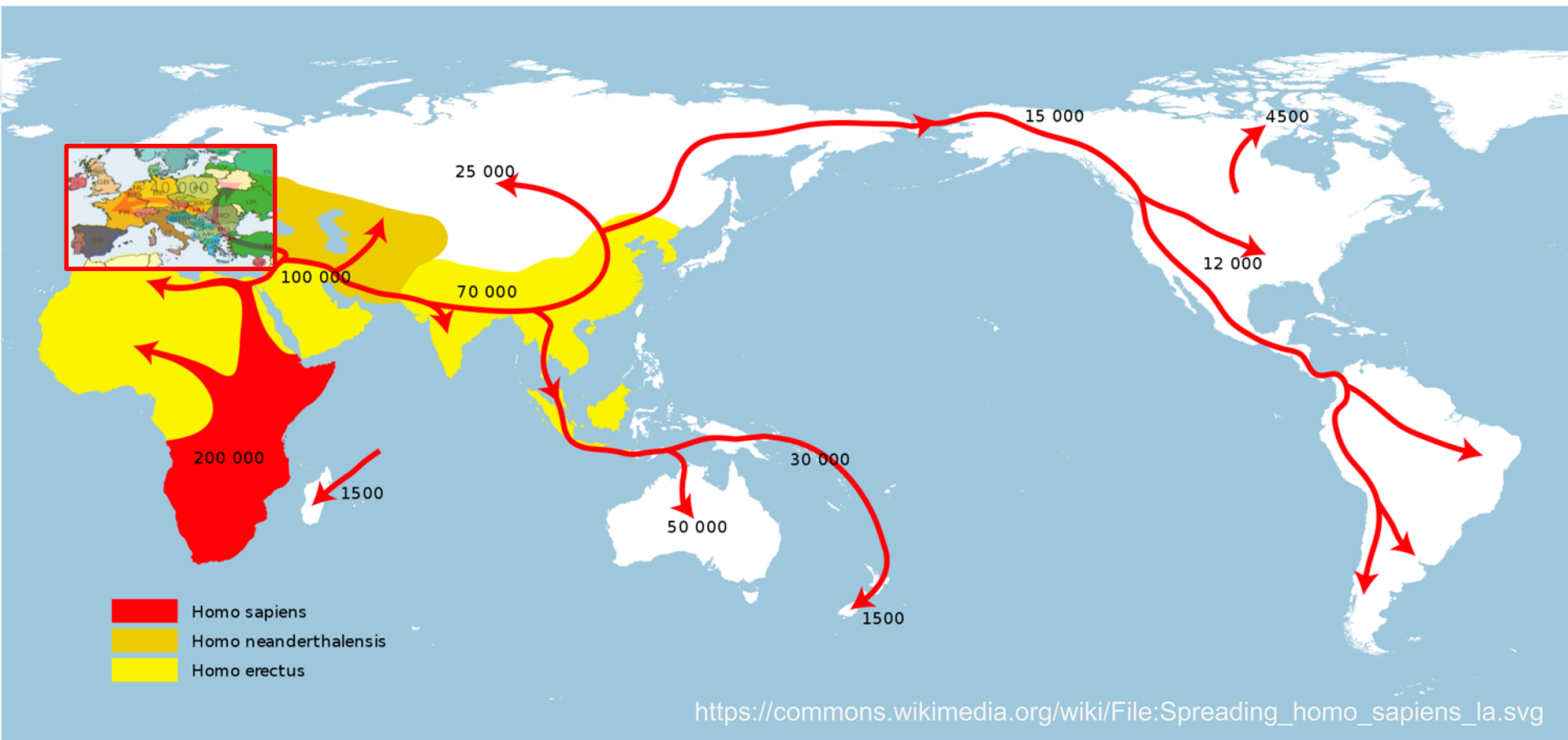


# Genetic ancestry and brain morphology





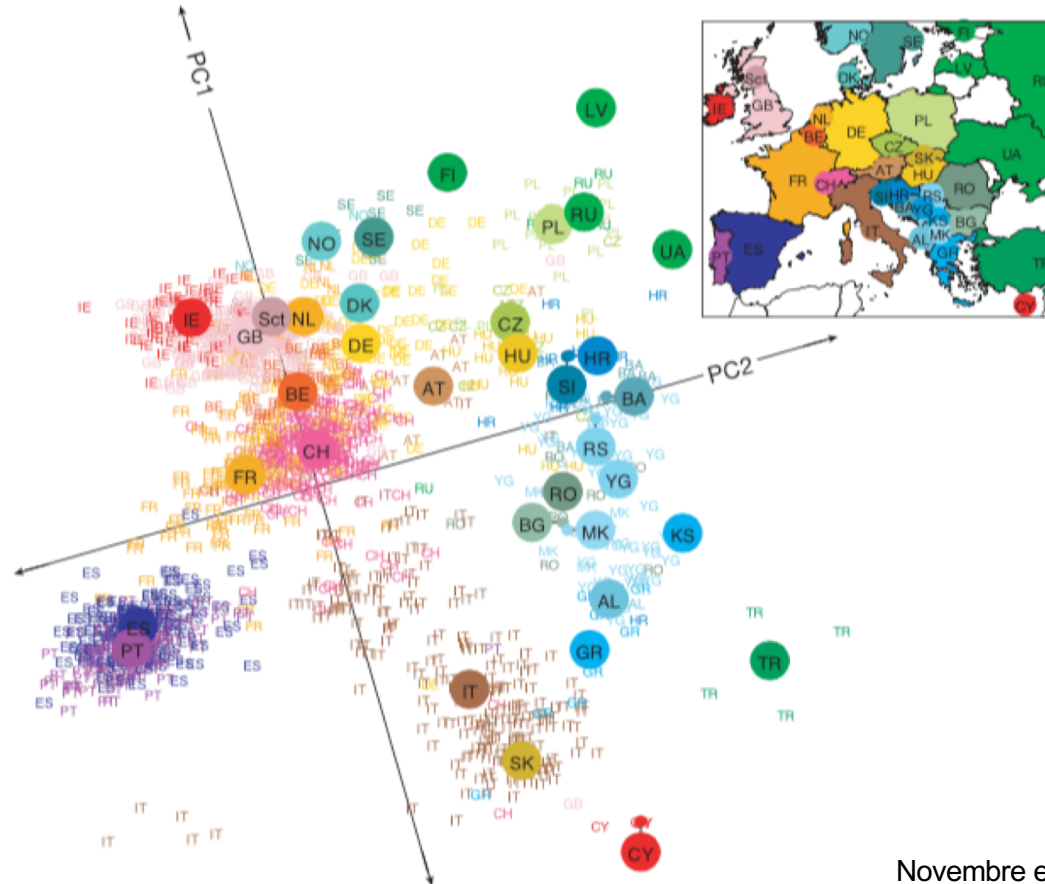
# Human migration



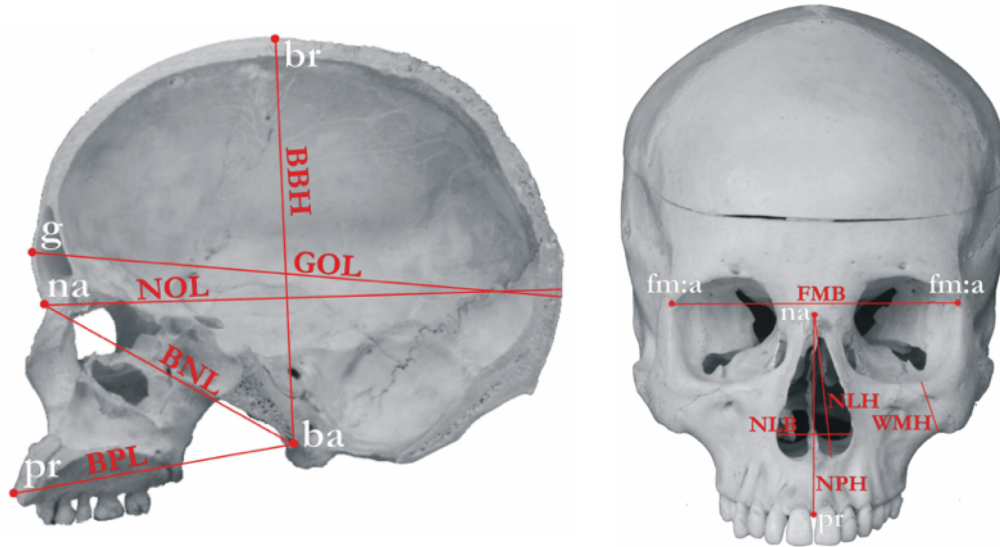
# Genes mirror geography within Europe



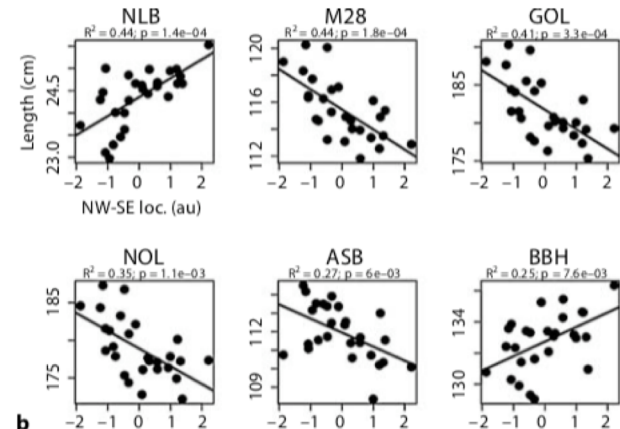
## Genes mirror geography within Europe



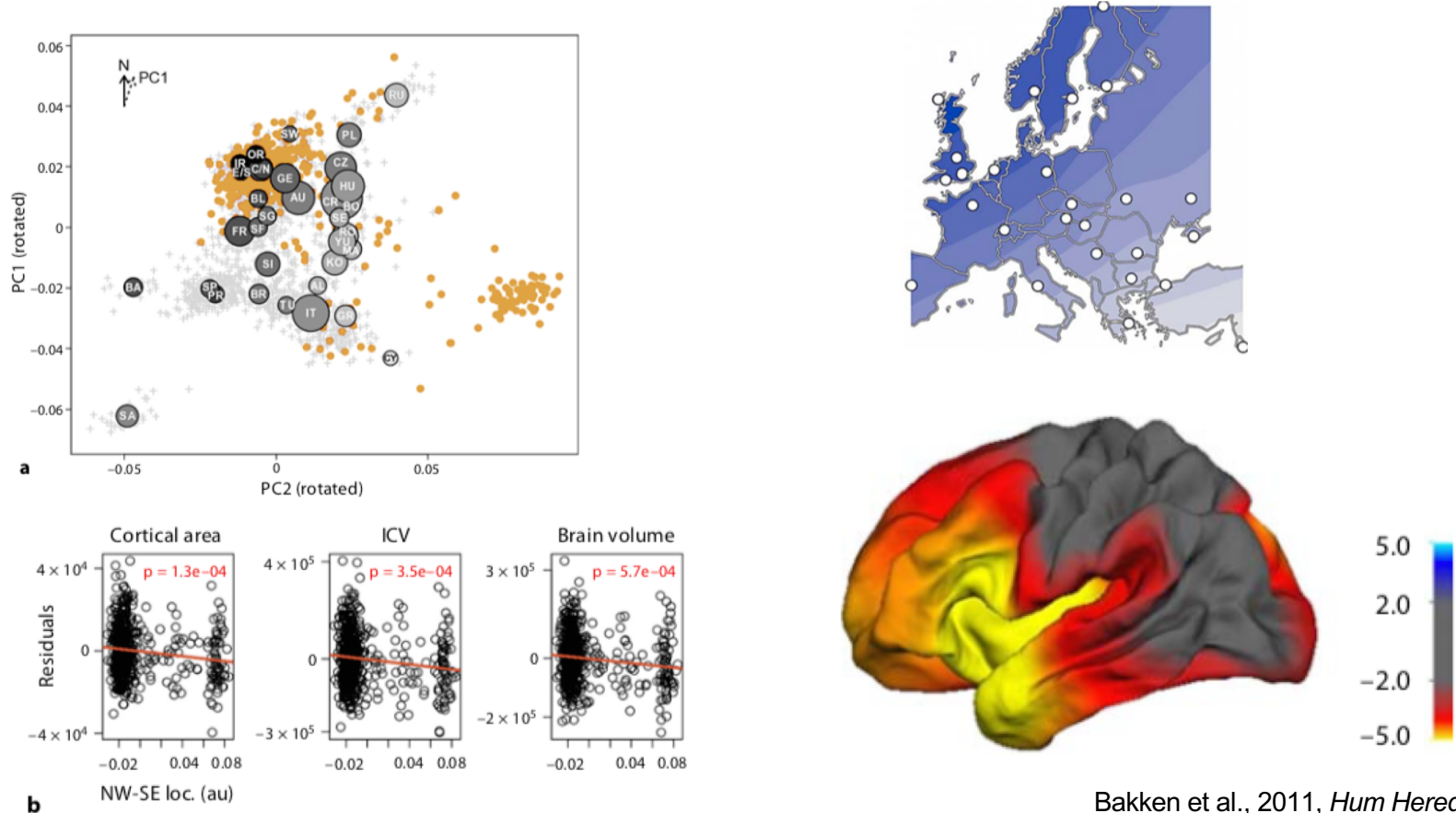
# Geography cline of skull and brain morphology in Europe



<http://osteomics.com/cranExplr/>



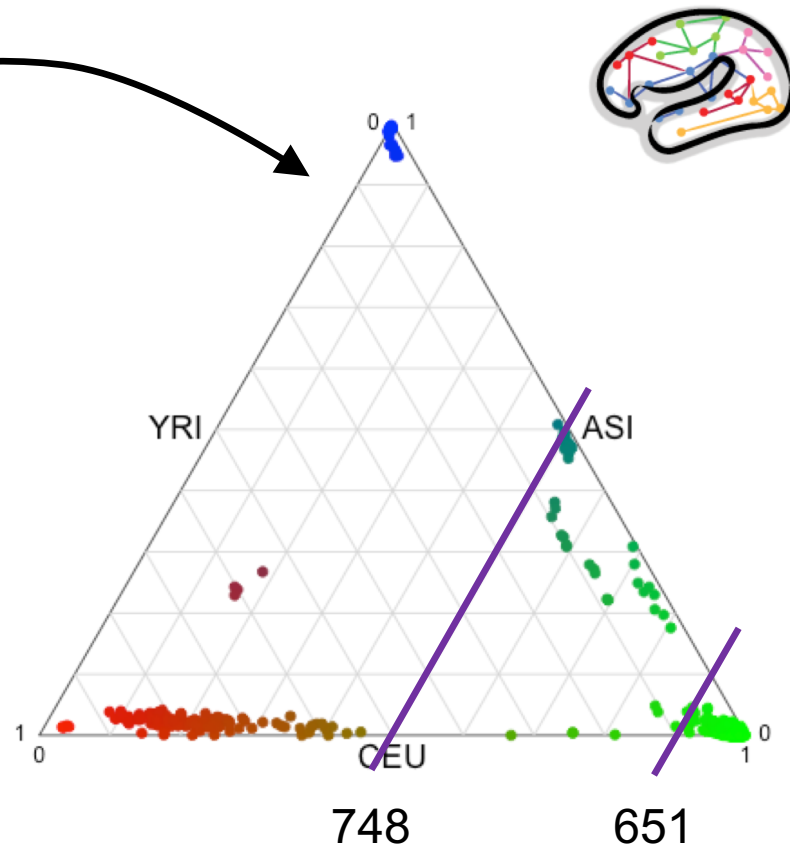
# Geography cline of skull and brain morphology in Europe



**Does this evolutionary bias extend to  
functional brain measures?**

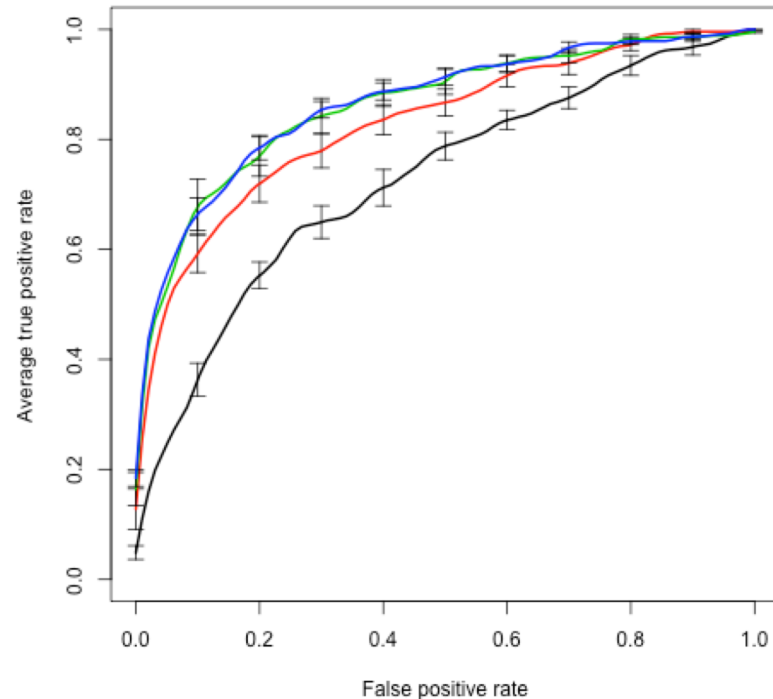


- 950 young adults with
  - Genome-wide SNP data
  - rs-fMRI (1h)
- Aim: classify CEU vs non-CEU
  - ICA: 15 to 300
  - Elastic net classifier
  - Nested CV
    - 10 outer folds (family aware)
    - 5 inner folds (optimize  $\lambda$ )
  - ROC curves



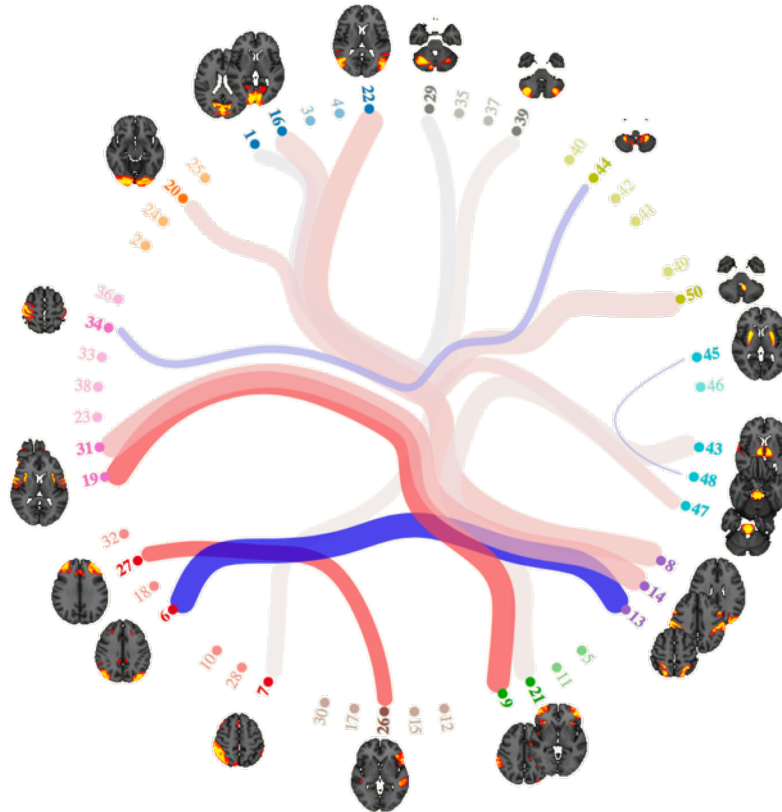
- Genetic ancestry can be predicted from rs-fMRI
- Works best with ICA  $\geq 100$
- Works better for 0.5 cutoff

ICA	dim	$p_{\text{CEU}} > 0.5$	$p_{\text{CEU}} > 0.9$
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)

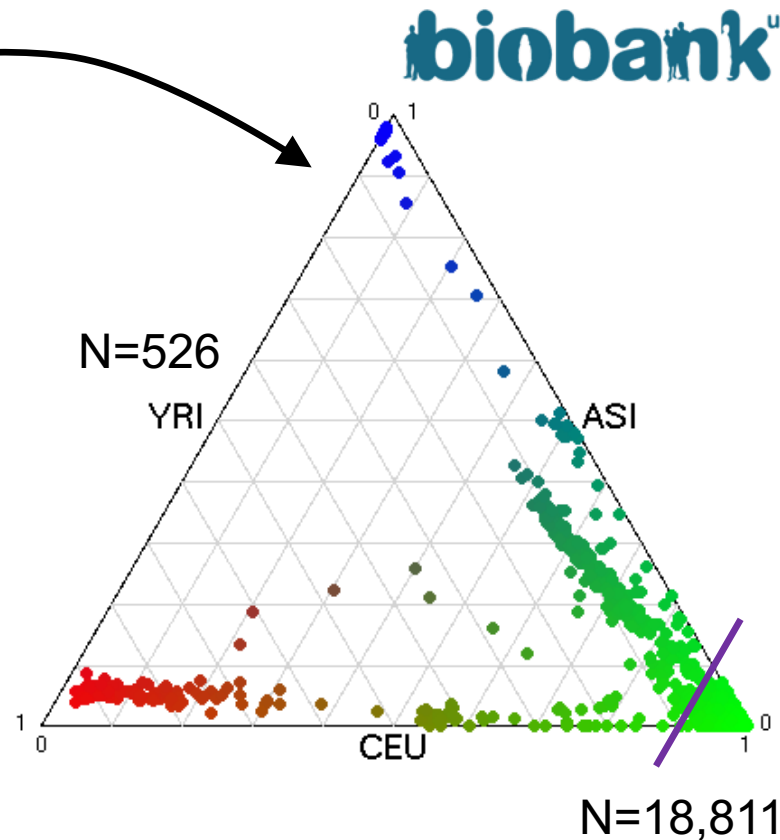


0.5 cutoff; ICA 15, 50, 100, 300

# Which connections are important?



- 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



- 19,337 adults with
  - Genome-wide SNP data
  - rs-fMRI (6 min)
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  - 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

ICA 25

- AUC 0.72

ICA100

- AUC 0.81

ICA	dim	$p_{\text{CEU}} > 0.5$	$p_{\text{CEU}} > 0.9$
15	105	0.78 (0.088)	0.72 (0.049)
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- 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