

# Gene expression under human self-domestication: an *in silico* exploration of modern human high-frequency variants

Univerza *v Ljubljani* 



Thomas O'Rourke<sup>1</sup>, Pedro Tiago Martins<sup>1,2</sup>, Alejandro Andirkó<sup>1\*</sup>

<sup>1</sup>University of Barcelona, <sup>2</sup>University of Ljubljana

#### Introduction



Homo sapiens



Canis lupus familiaris



Homo neanderthalensis

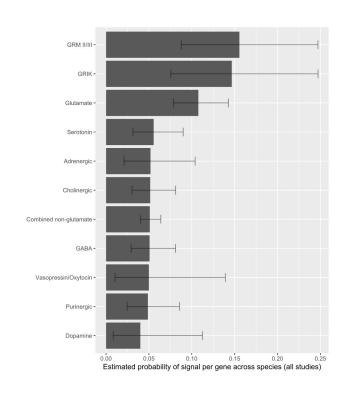


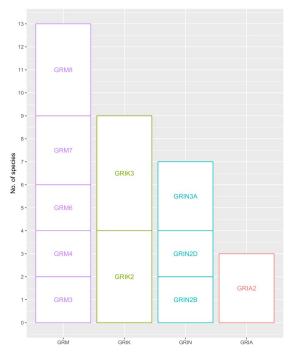
(adapted from Theofanopoulou et al. 2017, PLoS ONE)

Canis lupus lupus

## Differences:

- Braincase shape & size
- Browridge
- Nasal bone
- Tooth size
- Jaw progression





(O'Rourke & Boeckx 2020, Neurosci. Biobehav. Rev.)

# Hypothesis

## **Methods**

- Gene expression values per highfreq/fixed variant (ExPecto)
- Genes in glut. signaling pathway (based on GO categories)
- Tissues: blabla out of 228 in ExPecto

### **Discussion**

- Main interpretation of results
- Main conclusion

## Results

**GRAPH WITH RESULTS** 

References & further info: [maybe a github link with code, refs, more text, extra figs.]