

Table S6 Over-represented GO terms among Europe-biased genes.

| Ontology | GO Term | Genes | Adj. P |
|--------------------|---|-------|---------|
| Molecular function | electron carrier activity | 32 | 0.0001 |
| Molecular function | oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxygen | 31 | 0.01 |
| Molecular function | cofactor binding | 31 | 0.01 |
| Molecular function | oxidoreductase activity, acting on the CH-OH group of donors, NAD or NADP as acceptor | 19 | 0.01 |
| Molecular function | ATP binding | 79 | 0.02 |
| Molecular function | iron ion binding | 28 | 0.02 |
| Molecular function | transporter activity | 99 | 0.03 |
| Molecular function | disulfide oxidoreductase activity | 10 | 0.03 |
| Cellular component | membrane part | 173 | 0.02 |
| Cellular component | mitochondrial envelope | 38 | 0.02 |
| Cellular component | cytoskeletal part | 69 | 0.03 |
| Cellular component | lipid particle | 33 | 0.04 |
| Biological process | oxidation-reduction process | 87 | 4.6e-08 |
| Biological process | organonitrogen compound metabolic process | 84 | 0.001 |
| Biological process | carboxylic acid metabolic process | 49 | 0.001 |
| Biological process | carbohydrate derivative metabolic process | 59 | 0.01 |
| Biological process | cellular response to stress | 56 | 0.02 |
| Biological process | single-organism transport | 138 | 0.02 |
| Biological process | cellular localization | 81 | 0.03 |
| Biological process | centrosome organization | 23 | 0.03 |
| Biological process | organic substance catabolic process | 61 | 0.04 |
| Biological process | cell redox homeostasis | 13 | 0.04 |