Bibliography

Attfield, K.E. *et al.* (2022) 'The Immunology of Multiple Sclerosis', *Nature Reviews Immunology*, 22(12), pp. 734–750. doi:10.1038/s41577-022-00718-z.

Barrie, W. et al. (2024) 'Elevated genetic risk for multiple sclerosis emerged in steppe pastoralist populations', *Nature*, 625(7994), pp. 321–328. doi:10.1038/s41586-023-06618-z.

Koller, D. *et al.* (2022) 'Denisovan and Neanderthal archaic introgression differentially impacted the genetics of complex traits in modern populations', *BMC Biology*, 20(1). doi:10.1186/s12915-022-01449-2.

Llamas, B. *et al.* (2016) 'Ancient mitochondrial DNA provides high-resolution time scale of the peopling of the Americas', *Science Advances*, 2(4). doi:10.1126/sciadv.1501385.

López, S., Van Dorp, L. and Hellenthal, G. (2015) 'Human dispersal out of Africa: A lasting debate', *Evolutionary Bioinformatics*, 11s2. doi:10.4137/ebo.s33489.

Pedro, N. et al. (2020) 'Papuan mitochondrial genomes and the settlement of Sahul', *Journal of Human Genetics*, 65(10), pp. 875–887. doi:10.1038/s10038-020-0781-3.

Skoglund, P. *et al.* (2015) 'Genetic evidence for two founding populations of the Americas', *Nature*, 525(7567), pp. 104–108. doi:10.1038/nature14895.

Stone, R. (2021). A Land Bridge to Nowhere? Science.org. [online] doi:https://doi.org/10.1126/article.35167

Zeberg, H. and Pääbo, S. (2020) 'The major genetic risk factor for severe COVID-19 is inherited from neanderthals', *Nature*, 587(7835), pp. 610–612. doi:10.1038/s41586-020-2818-3.