Bibliografía:

Andrews, R. M., Kubacka, I., Chinnery, P. F., Lightowlers, R. N., Turnbull, D. M., y Howell, N. (1999). Rea-nalysis and revision of the Cambridge reference sequence for human mitochondrial DNA. Nature Genetics, 23 (2) 147.

Bandelt, H. J., Forster, P., Röhl, A. (1999). Median-joining networks for inferring intraspecific phylogenies. Molecular Biology and Evolution, 16(1), 37–48.

González, A.M., Larruga, J.M., Abu-Amero, K.K. et al. (2007). Mitochondrial lineage M1 traces an early human backflow to Africa. BMC Genomics 8, 223

Kloss-Brandstätter, A., Pacher, D., Schönherr, S., Weissensteiner, H., Binna, R., Specht, G. and Kronenberg, F. (2011), HaploGrep: a fast and reliable algorithm for automatic classification of mitochondrial DNA haplogroups. Hum. Mutat., 32: 25-32.

Malhi Ripan S., Kemp Brian M., Eshleman Jason A., Cybulski Jerome, Glenn Smith David, Harold Harry Scott Cousins. (2007). Mitochondrial haplogroup M discovered in prehistoric North Americans. Journal of Archaeological Science,Volume 34, Issue 4,2007, Pages 642-648.

Nicholas, K.B. y Nicholas, H.B. (1997). GenDoc: a tool for editing and annotating multiple sequence align-ment (http://www.psc.edu/biomed/genedoc).

Schwab, M (2019). "Afinidades filogeográficas y estructuración geográfica de los linajes maternos y paternos presentes en poblaciones humanas del noroeste argentino". Tesis de doctorado. Universidad Nacional de la Plata (Argentina). Facultad de Ciencias Naturales y Museo.

Weissensteiner, H., Pacher, D., Kloss-Brandstätter, A., Forer, L., Specht, G., Bandelt, H.J. Kronenberg F, Salas A, y Schönherr, S. (2016). HaploGrep 2: mitochondrial haplogroup classification in the era of high-throughput sequencing. Nucleic Acids Research, 44(W1), W58-63.