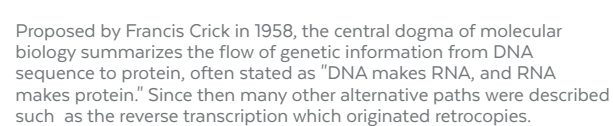




First classified as pseudogenes lacking introns, retrocopies are fascinating genomic elements that defy the central dogma of molecular biology. These elements are genomic copies of a gene (defined as parental gene) but have no introns and have a polyA tract, indicating a RNA-dependent mechanism of gene duplication. Retrocopies are often classified as “dead on arrival” since the lack regulatory elements in the majority of regions where these copies are inserted doesn’t favor transcriptional activity. But, when retrocopies are inserted inside a gene (a host gene), they can use its elements to be transcribed! These hard to believe elements are rare to find among Eukaryotes, but in mammals, including us, retrocopies are way more common!



The most retrocopied gene in the human genome are the RPL21 with 105 retrocopies. RPLs genes are genes which encode ribosomal proteins and are ubiquity expressed through most human tissues. This expression profile could explain why they are the most retrocopied genes and the importance of these genes could provide a possible evolutionary advantage to have many copies.

Mammals, including us, are by far the species with more retrocopies. This interesting aspect can be most the presence of long interspersed nuclear elements (LINEs) present in mammals, in special LINE1 elements, that provide the needed machinery to the retroposition of processed RNAs with polyA tails.