

# Formation of human long intergenic non-coding RNA genes and pseudogenes: ancestral sequences are key players

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**Supplementary Figure S4.** **a.** Color highlighted sections represent the *FAM230B*--*LOC105372935*--*GGT2* sequences that are found in *FAM230C*--*LOC101060145*--*GGT4P* with the respective percent identities. xxx represents sequences from the clincRNA region (*LOC105372935*) of *FAM230B*--*LOC105372935*--*GGT2* that are missing in *FAM230C*--*LOC101060145*--*GGT4P*. The unhighlighted section, |----| represents the 5' half sequence of *FAM230C* that does not form part of *FAM230B*. **b.** Schematic of *FAM230B*--*LOC105372935*--*spacer*--*GGT2* for comparisons. The % identities shown in Figure S3a are relative to the *FAM230B*--*LOC105372935*--*GGT2* sequence.

