

Preprints

P. García-Galindo*, Paula, S. E. Ahnert, **N S. Martin**. The non-deterministic genotype-phenotype map of RNA secondary structure. *bioRxiv* (2023): 2023.02.27.530309.

<https://doi.org/10.1101/2023.02.27.530309> *corresponding author

N.S. Martin*, S.E. Ahnert. The Boltzmann distributions of folded molecular structures predict likely changes through random mutations. *bioRxiv* (2023): 2023.02.22.529545.

<https://doi.org/10.1101/2023.02.22.529545> *corresponding author

Publications

N.S. Martin*, S.E. Ahnert. Thermodynamics and neutral sets in the RNA sequence-structure map. *EPL*, 139: 37001 (2022). <https://doi.org/10.1209/0295-5075/ac7c34> *corresponding author

N.S. Martin*, S.E. Ahnert. Fast free-energy-based neutral set size estimates for the RNA genotype-phenotype map. *J. R. Soc. Interface* 19: 20220072 (2022).

<https://doi.org/10.1098/rsif.2022.0072> *corresponding author

N. S. Martin*, S.E. Ahnert. Insertions and deletions in the RNA sequence–structure map. *J. R. Soc. Interface* 18: 20210380 (2021) <https://doi.org/10.1098/rsif.2021.0380> *corresponding author

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A reply to comments was published in 2022 with the same co-authors.