Ardell Lab UC Merced

Computational Biology and Evolution of Gene Expression

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genes systems identity translation extend tRNAs quantitative genome also diversification accuracy expressed environments Ardell aim cells bacterial DNA future models translational study stable Metagenomics proteins history RNA prokaryotic Gene signals currently done example function living code comparative evolution life expression folding plan research genomics ics co conversion Group predict organisms genomes gene evolutionary better inversion role flow novel infer genomic operons Research RNAs use first matter massive important genetic tRNA stable data Translational characterizing ribosomal processes energy conditions discover model transcriptional annotated diversity mechanisms algorithm protein protein-coding work Systems codes

A Wordle of our Research Interests.

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