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Due to the molecular biology process, gene regulatory network inference can be understood as a group feature selection problem based on the dependencies between the expression data of TF complexes

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While discrepancies in noise variance are overlooked in routine molecular network inference, the results presented clearly elucidate their non-trivial impact on the conclusions that in turn can challenge the biological significance of the findings. The analytical treatment and arguments presented are generic and not restricted to molecular data sets. Introduction. Identifying associations and

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[Biological network inference - Wikipedia](#)

Biological network inference is the process of making inferences and predictions about biological networks. Biological networks. In a topological sense, a network is a set of nodes and a set of directed or undirected edges between the nodes. Many types of biological networks exist, including transcriptional, signalling and metabolic. Few such networks are known in anything approaching

their