Revisiting the paper on "Applications of graph theory to enzyme kinetics and protein folding kinetics: steady and non-steady state systems" (Short Communication)

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Short Communication

About 30 years ago a very important paper on "Applications of graph theory to enzyme kinetics and protein folding kinetics: steady and non-steady state systems" was published [1].

Ever since then, a series of papers for graph theory to study enzyme kinetics [2-8] and protein folding kinetics [9-15].

The graph approach can provide an intuitive feeling for analyzing very complicated biological systems, as clearly demonstrated in the eight masterpieces papers [16-23] by the then Chairman of Nobel Prize Committee.

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