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REVISITING THE PAPER ON “APPLICATIONS OF GRAPH THEORY TO ENZYME KINETICS AND PROTEIN FOLDING KINETICS: STEADY AND NON-STEADY STATE SYSTEMS”

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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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