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The software *Group Contribution Method* is designed to estimate the standard Gibbs free energy of formation ($\Delta_f G'^{\circ}$) and reaction ($\Delta_r G'^{\circ}$) for the majority of the biochemical compounds and reactions found in the *i*JR904 and *i*AF 1260 genome-scale metabolic models of Escherichia coli and in the Kyoto Encyclopedia of Genes and Genomes and University of Minnesota Biocatalysis and Biodegradation Database.

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