O1-1

Free energy profile of tRNA dissociation from ribosome studied by coarse-grained molecular dynamics simulations

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O2-6

Interpretable Deep Learning Using Multimodal Graph Convolutional Network for Predicting Compound-Protein Interactions

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O3-2

From drug repositioning to target repositioning: omics-based prediction of therapeutic targets for a variety of diseases

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O4-1

Transformer-based Generative Adversarial Networks for Generating Molecules with Desired Properties

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