

Resource Summary Report

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GraphRBF

RRID:SCR_025652

Type: Tool

Proper Citation

GraphRBF (RRID:SCR_025652)

Resource Information

URL: <https://github.com/Wssduer/GraphRBF>

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Description: Software tool as protein-protein/nucleic acid interaction site prediction model built by enhanced graph neural networks and prioritized radial basis function neural networks. Protein-protein and protein-nucleic acid binding site prediction via interpretable hierarchical geometric deep learning.

Resource Type: simulation software, software resource, source code, software application

Keywords: protein, nucleic acids, interactions, python, binding site prediction,

Availability: Free, Available for download, Freely available

Resource Name: GraphRBF

Resource ID: SCR_025652

License: GNU GPL v3.0

Record Creation Time: 20240827T053245+0000

Record Last Update: 20260819T044614+0000

Ratings and Alerts

No rating or validation information has been found for GraphRBF.

No alerts have been found for GraphRBF.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.