

# **Molecular docking and dynamics simulation studies uncover the host–pathogen protein–protein interactions in soybean (*Glycine max* (L.) Merr.) and Groundnut bud necrosis virus: first report**

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## **Abstract**

<https://www.frontiersin.org/journals/bioinformatics/articles/10.3389/fbinf.2026.1821527/full>

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