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Molecular modeling the proteins from the *exo-xis* region of lambda and shigatoxigenic bacteriophages

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PDB file contents

File	Figure	Description
figure_3_lambda_ea22_alphafold.pdb	Figure 3	λ ea22 C-terminal domain AlphaFold prediction
figure_3_lambda_ea22_cterm_rosettafold.pdb	Figure 3	λ ea22 C-terminal domain RoseTTAFold prediction
figure_3_lambda_orf60_rosettafold.pdb	Figure 3	λ Orf60a AlphaFold prediction
figure_3_lambda_orf60a_alphafold.pdb	Figure 3	λ Orf60a RoseTTAFold prediction
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figure_3_lambda_orf63_rosettafold.pdb	Figure 3	λ Orf63 RoseTTAFold prediction
figure_3_phi24B_orf61_alphafold.pdb	Figure 3	ϕ 24B Orf61 AlphaFold prediction
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figure_9_phi24B_gp05_dimer.pdb	Figure 9	ϕ 24B Gp05 dimer AlphaFold prediction