

DOI

Progress in Biochemistry and Biophysics ISSN 1000-3282,CN 11-2161/Q

BLUF TePixD Y8F	()
10.16476/j.pibb.2023.0095 2023-05-29	
	BLUF TePixD Y8F (
)[J/OL]	. https://doi.org/10.16476/j.pibb.2023.0095



ISSN 2096-4188 CN 11-6037/Z

Published online : 2023-05-29 19:51:33 URL : https://kns.cnki.net/kcms2/detail/11.2161.q.20230526.1541.001.html

PIBB

* This work was supported by the National Key Research and Development Program (2020YFA0509700) to LU Qing.

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Received: March 23, 2023 Accepted: May 23, 2023

DOI: 10.16476/j.pibb.2023.0095



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摘要



2020YFA0509700

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2023-05-23

Y8F* Wavelength **Resolution range** Space group Unit cell Unique reflections Multiplicity Completeness (%) Mean I/sigma(I) Wilson B-factor **R-merge R**-meas R-pim CC1/2 **Reflections used in refinement Reflections used for R-free R-work R-free** Number of non-hydrogen atoms macromolecules ligands solvent **Protein residues** RMS (bonds) RMS (angles) Ramachandran favored (%)

 Table 1
 Data collection and refinement statistics (PDB ID: 8198)

Ramachandran allowed (%) Ramachandran outliers (%) Rotamer outliers (%) Clashscore Average B



