



STRUCTURAL
BIOLOGY

Volume 78 (2022)

Supporting information for article:

**Molecular insight into 2-phosphoglycolate activation of
bisphosphoglycerate mutase phosphatase activity**

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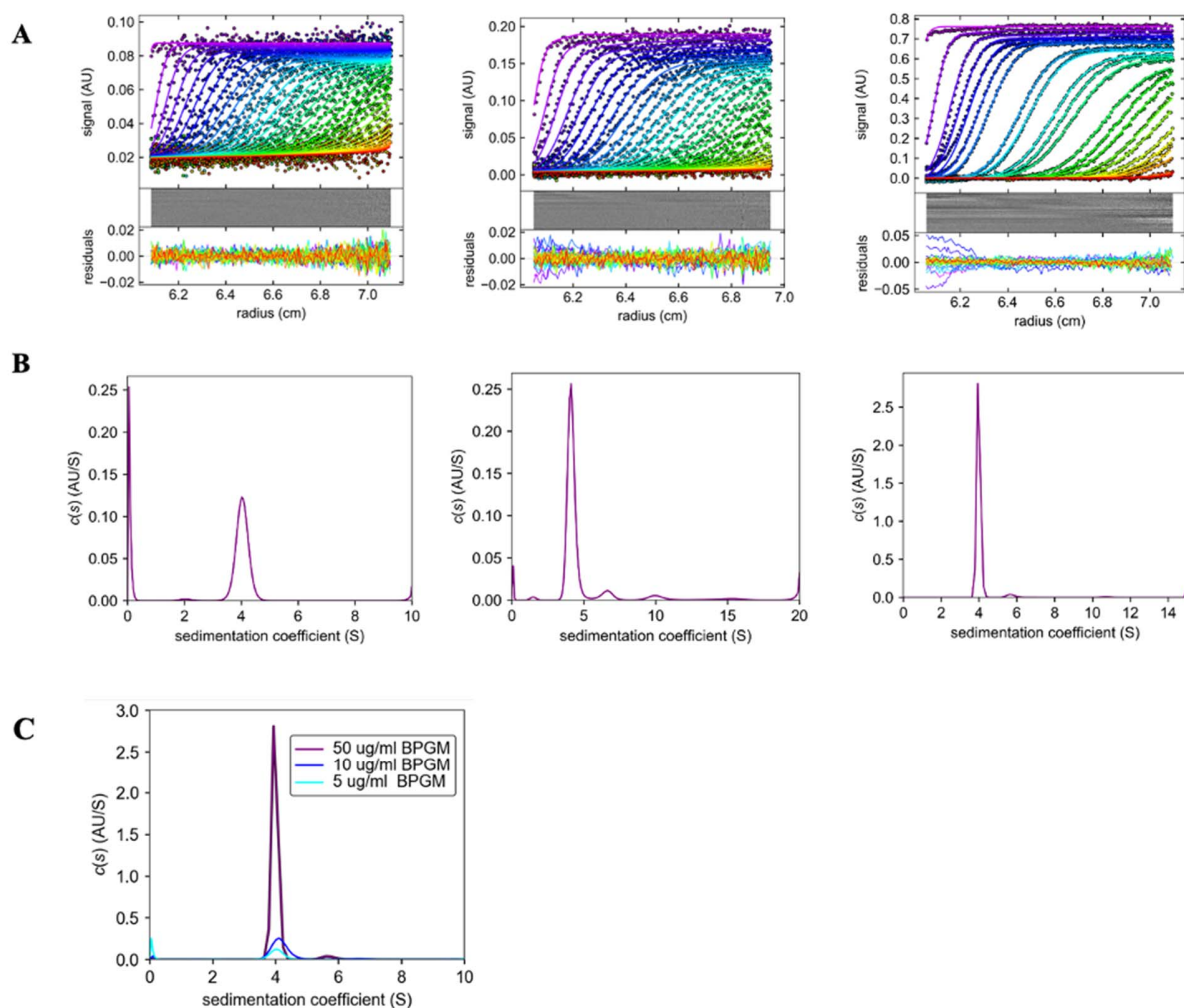


Figure S1 Sedimentation velocity analytical ultracentrifugation analysis of BPGM (A) raw sedimentation scan at 230 nm of BPGM at different concentrations 5, 10, 50 µg/ml BPGM (left to right). (B) Sedimentation coefficient distribution of BPGM at different concentrations 5, 10, 50 µg/ml BPGM (left to right) (C) Sedimentation coefficient distribution of BPGM calculated by SEDFIT for the three SV experiments

Table S1 . Characteristics of the BPGM corrected to pure water

Sample name	$S_{20,W}$, S	Molar Mass	Concentration	Friction ratio
5 ug/ml	4.12S	63.6Kda	0.064 A230	1.309
10 ug/ml	4.23S	62.2KDa	0.159 A230	1.374
50 ug/ml	4.05S	64.0KDa	9.72 A230	1.374
Dimer ^a	4.27S	64.0KDa	0 A 230	N.D.
Monomer	2.91S	32.0KDa	0 A 230	N.D.

^aDimer. Values calculated using the program hydropro10 and the X-ray structure factors from the PDB files for dimer and monomer.

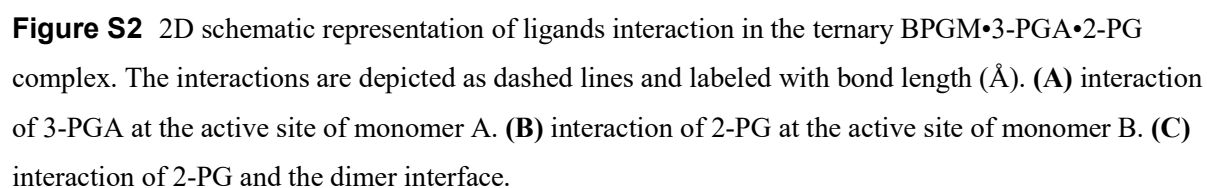


Figure S2 2D schematic representation of ligands interaction in the ternary BPGM•3-PGA•2-PG complex. The interactions are depicted as dashed lines and labeled with bond length (Å). **(A)** interaction of 3-PGA at the active site of monomer A. **(B)** interaction of 2-PG at the active site of monomer B. **(C)** interaction of 2-PG and the dimer interface.

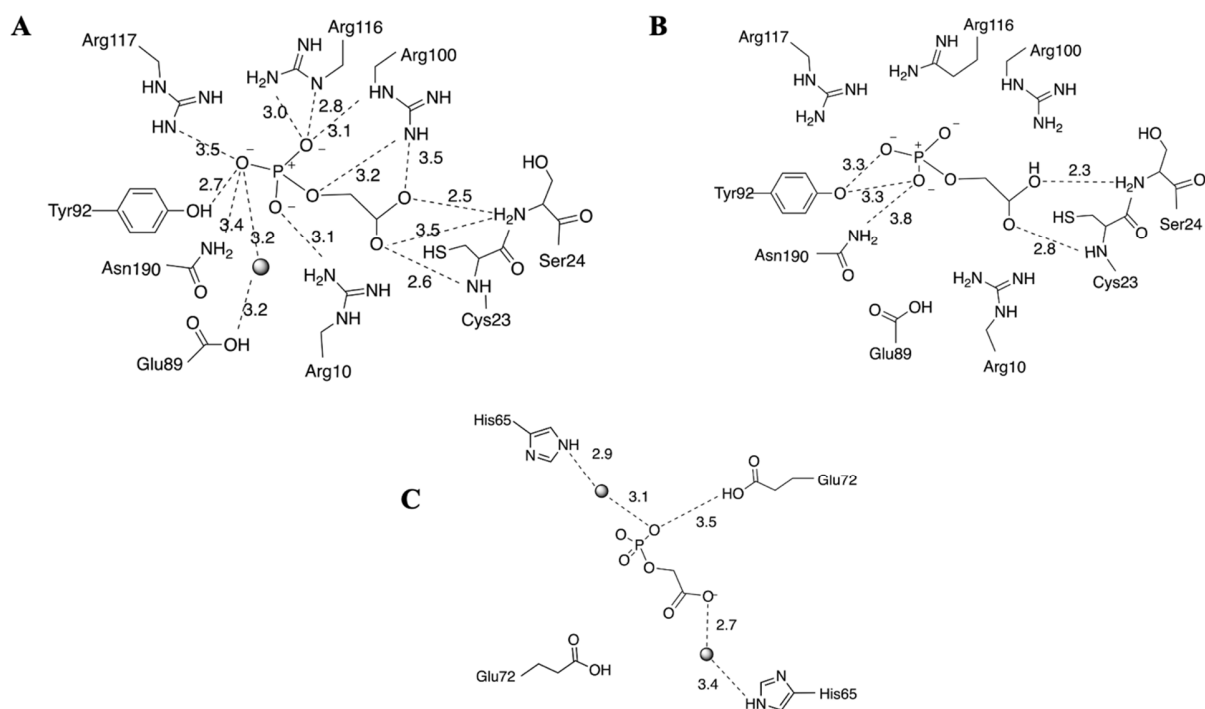


Figure S3 2D schematic representation of ligands interaction in the binary BPGM•2-PG complex. The interactions are depicted as dashed lines and labeled with bond length (Å). **(A)** interaction of 2-PG at the active site of monomer A **(B)** interaction of 2-PG at the active site of monomer B **(C)** interaction of 2-PG at the dimer interface.