## Supporting Information

## Enzymatic production of $\beta$ -glucose 1,6-bisphosphate through manipulation of catalytic magnesium coordination

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**Figure S1.** Initial rate measurements for the conversion of  $\beta$ G1P to G6P catalysed by  $\beta$ PGM<sub>D170N</sub> monitored using a glucose 6-phosphate dehydrogenase (G6PDH) coupled assay. (A) Reactions were conducted in 200 mM K<sup>+</sup> HEPES buffer (pH 7.2), 5 mM MgCl<sub>2</sub>, 1 mM NAD<sup>+</sup> and 5 U/mL G6PDH containing 10  $\mu$ M  $\beta$ PGM<sub>D170N</sub>, 1 mM  $\beta$ G1P and were initiated using increasing concentrations of  $\beta$ G16BP (10, 25, 50, 100, 150, 250, 350, 750, 1000, 1500, 2500  $\mu$ M). Initial rates of G6P production were obtained using an in-house Python non-linear least squares fitting program yielded an apparent  $K_m$  ( $\beta$ G16BP) = 150 ± 13  $\mu$ M. (B) Reactions were conducted in 200 mM K<sup>+</sup> HEPES buffer (pH 7.2), 5 mM MgCl<sub>2</sub>, 1 mM NAD<sup>+</sup> and 5 U/mL G6PDH containing 10  $\mu$ M  $\beta$ PGM<sub>D170N</sub> and increasing concentrations of  $\beta$ G1P (50, 100, 200, 300, 500, 700, 1000, 1500, 2000, 3000, 5000  $\mu$ M) and were initiated using 250  $\mu$ M  $\beta$ G16BP. Initial rates of G6P production were obtained using a linear least-squares fitting routine Subsequent fitting of these rates to Equation 2 using an in-house Python non-linear least squares fitting routine fitting of these rates to Equation  $\mu$ M and were initiated using 250  $\mu$ M  $\beta$ G16BP. Initial rates of G6P production were obtained using a linear least-squares fitting routine Subsequent fitting of these rates to Equation 2 using an in-house Python non-linear least squares fitting program yielded an apparent  $K_m$  ( $\beta$ G1P) = 6.9 ± 1.0  $\mu$ M and an apparent  $K_i$  ( $\beta$ G1P) = 1536 ± 170  $\mu$ M.



Figure S2. Activity of  $\beta PGM_{WT}$  and  $\beta PGM_{D170N}$  with increasing MgCl<sub>2</sub> concentration. Normalised initial rate measurements for the conversion of  $\beta$ G1P to G6P by either  $\beta$ PGM<sub>WT</sub> (black circles) or βPGM<sub>D170N</sub> (green circles) at different concentrations of MgCl<sub>2</sub> monitored using a G6PDH coupled assay. Reactions were conducted in 200 mM K<sup>+</sup> HEPES buffer (pH 7.2) containing different concentrations of MgCl<sub>2</sub> (0, 0.1, 0.3, 0.6, 1.0, 1.5, 2.5, 5, 10, 20, 50 and 100 mM), 1 mM NAD<sup>+</sup>, 5 U/mL G6PDH, 1 mM ßG1P and either 1 nM ßPGM<sub>WT</sub> with 100 μM βG16BP, or 10 μM βPGM<sub>D170N</sub> with 1250 μM βG16BP. Initial rates of G6P production were obtained using a linear least-squares fitting routine. Subsequent fitting of these rates to Equation 2 using an in-house Python non-linear least squares fitting program yielded an apparent  $K_{\rm m}$  (Mg<sup>2+</sup>) = 180 ± 40  $\mu$ M for  $\beta$ PGM<sub>WT</sub> and an apparent  $K_{\rm m}$  (Mg<sup>2+</sup>) = 690 ± 110  $\mu$ M for βPGM<sub>D170N</sub>. The standard error of the mean of three technical replicates falls within the radii of the data points. The discrepancy between the  $K_{\rm m}$  (Mg<sup>2+</sup>) value obtained using the G6PDH coupled assay and <sup>31</sup>P NMR time-course experiments (Fig. 3B-C) is likely due to the different conditions employed, although similar maximal initial rates of reaction are observed using each technique (maximal initial rate using G6PDH coupled assay =  $0.009 \text{ s}^{-1}$ ; maximal initial rate using <sup>31</sup>P NMR time-course experiments =  $0.012 \text{ s}^{-1}$ ). These observations indicate that a component of the reaction mixture used in the <sup>31</sup>P NMR experiments is competing with  $Mg^{2+}$  ions to bind to  $\beta PGM_{D170N}$ . One notable difference between the conditions of each technique is the 10-fold higher βG1P concentration used in the <sup>31</sup>P NMR experiments. Given that βPGM<sub>D170N</sub> experiences βG1P inhibition (Fig. S1B) at a comparable level to βPGM<sub>WT</sub>, this behaviour provides a likely source for the competitive inhibition observed in the <sup>31</sup>P NMR experiments. Although the mechanism for  $\beta$ G1P inhibition has not been structurally characterised, it is plausible that  $\beta$ G1P binds to Mg<sub>cat</sub>-free  $\beta$ PGM<sub>D170N</sub> to form a closed complex, thus preventing \u03b3G16BP production in Step 1 and G6P production in Step 2, until dissociation occurs.



**Figure S3.** Solution behaviour of substrate-free  $\beta$ PGM. Overlay of <sup>1</sup>H<sup>15</sup>N-TROSY spectra for substrate-free  $\beta$ PGM<sub>WT</sub> (black) and substrate-free  $\beta$ PGM<sub>D170N</sub> (green), recorded in 50 mM K<sup>+</sup> HEPES buffer (pH 7.2), 5 mM MgCl<sub>2</sub>, 2 mM NaN<sub>3</sub>, 10% <sup>2</sup>H<sub>2</sub>O (v/v) and 1 mM TSP. There is a broad correspondence between peaks of  $\beta$ PGM<sub>WT</sub> and  $\beta$ PGM<sub>D170N</sub>, indicating a similar solution behaviour and overall protein fold. Two conformers are present in slow exchange (~70% conformer A and ~30% conformer B) for both  $\beta$ PGM<sub>WT</sub> and  $\beta$ PGM<sub>D170N</sub>, which arise from *cis-trans* isomerisation at the K145-P146 peptide bond.<sup>23</sup> Additionally, ~15 peaks are present for  $\beta$ PGM<sub>D170N</sub>, which are absent in  $\beta$ PGM<sub>WT</sub> due to backbone conformational exchange on the millisecond timescale.<sup>23</sup> This observation indicates that residue N170 in  $\beta$ PGM<sub>D170N</sub> abolishes the intermediate exchange dynamic that residue D170 propagates in  $\beta$ PGM<sub>WT</sub>.



**Figure S4.** Comparative overlays of a section of  ${}^{1}H^{15}N$ -TROSY spectra for substrate-free  $\beta$ PGM recorded in 50 mM K<sup>+</sup> HEPES buffer (pH 7.2), 5 mM MgCl<sub>2</sub>, 2 mM NaN<sub>3</sub>, 10%  ${}^{2}H_{2}O$  (v/v) and 1 mM TSP. (A) Comparison of substrate-free  $\beta$ PGM<sub>D170N</sub> that had been preincubated at 25 °C for 0 h (light green) and 48 h (dark green). Near-identical spectra indicate that the incubation process has a negligible effect on the stability of substrate-free  $\beta$ PGM<sub>D170N</sub>. (B) Comparison of substrate-free  $\beta$ PGM<sub>D170N</sub> preincubated at 25 °C for 0 h (light green) and substrate-free  $\beta$ PGM<sub>D170N</sub> preincubated at 25 °C for 0 h (light green) and substrate-free  $\beta$ PGM<sub>WT</sub> (black). (C) Comparison of substrate-free  $\beta$ PGM<sub>D170N</sub> preincubated at 25 °C for 48 h (dark green) and substrate-free  $\beta$ PGM<sub>WT</sub> (black). The absence of observable  $\beta$ PGM<sub>WT</sub> peaks in the  $\beta$ PGM<sub>D170N</sub> spectrum indicates that reversion of  $\beta$ PGM<sub>D170N</sub> to  $\beta$ PGM<sub>WT</sub> through deamidation is not a process that occurs readily under these sample conditions.



**Figure S5.** Activity of  $\beta$ PGM<sub>D170N</sub> in 200 mM K<sup>+</sup> HEPES buffer (pH 7.2) and 100 mM MgCl<sub>2</sub> monitored using <sup>31</sup>P NMR time-course experiments. (A) Reaction kinetics for the equilibration of 10 mM  $\beta$ G1P with G6P catalysed by 200  $\mu$ M  $\beta$ PGM<sub>D170N</sub> that had been preincubated at 25 °C for 0 h (light green symbols), 24 h (medium green symbols) and 48 h (dark green symbols). The reactions were initiated by and timed from the addition of 20 mM AcP. Normalised integral values of the <sup>31</sup>P resonances of  $\beta$ G16BP and G6P have been converted to concentrations and are plotted as a function of 10 mM  $\beta$ G1P with G6P catalysed by 200 nM  $\beta$ PGM<sub>D170N</sub> to  $\beta$ PGM<sub>D170N</sub> containing 200 nM  $\beta$ PGM<sub>WT</sub> (representative of 0.1% reversion of  $\beta$ PGM<sub>D170N</sub> to  $\beta$ PGM<sub>WT</sub> through deamidation). The reaction was initiated by and timed from the addition of 20 mM AcP. Normalised integral values of the <sup>31</sup>P resonances of the <sup>31</sup>P resonances of  $\beta$ G16BP (open circles) and G6P (closed circles). (B) Reaction kinetics for the equilibration of 10 mM  $\beta$ G1P with G6P catalysed by 200  $\mu$ M  $\beta$ PGM<sub>D170N</sub> to  $\beta$ PGM<sub>WT</sub> through deamidation). The reaction was initiated by and timed from the addition of 20 mM AcP. Normalised integral values of the <sup>31</sup>P resonances of  $\beta$ G16BP and G6P have been converted to concentrations and are plotted as a function of time for  $\beta$ G16BP and G6P have been converted to concentrations and are plotted as a function of the <sup>31</sup>P resonances of  $\beta$ G16BP and G6P have been converted to concentrations and are plotted as a function of time for  $\beta$ G16BP (open circles) and G6P (closed circles).