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Unraveling the function of the two Entner–Doudoroff branches in the thermoacidophilic Crenarchaeon Sulfolobus solfataricus P2

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Supplementary Material

Derivation of GK activity at constant ATP concentration (Eqn.1)

From Fig. 6 the rate equation for forming product P is given as:

$$\mathbf{v} = 2\mathbf{k}_{cat}[\mathbf{ES}_2] + 3\mathbf{k}_2[\mathbf{ES}_3] \tag{S1}$$

Affinity constant (K_s) and inhibitor constant (K_i) are defined as (assuming rapid equilibria):

$$K_{S} = \frac{[E][S]^{2}}{[ES_{2}]}; \qquad K_{I} = \frac{[ES_{2}][S]}{[ES_{3}]}$$
 (S2)

The total enzyme concentration [E]₀ is given as:

$$[E]_{0} = [E] + [ES_{2}] + [ES_{3}]$$
(S3)

Solving for [E] and [ES₃] from the equations defining K_S and K_i , and inserting these expressions into Eq. S3, leads to:

$$[E]_{0} = \frac{K_{S}[ES_{2}]}{[S]^{2}} + [ES_{2}] + \frac{[ES_{2}][S]}{K_{i}} = [ES_{2}] \left(\frac{K_{S}}{[S]^{2}} + 1 + \frac{[S]}{K_{i}}\right)$$
(S4)

Solving for [ES₂] and [ES₃] from Eqs. S4 and S2, respectively, gives:

$$[ES_{2}] = \frac{[E]_{0}}{\frac{K_{s}}{[S]^{2}} + 1 + \frac{[S]}{K_{i}}}; \quad [ES_{3}] = \left(\frac{[S]}{K_{i}}\right)[ES_{2}]$$
(S5)

Inserting the expressions for $[ES_2]$ and $[ES_3]$ from Eq. S5 into Eq. S1 gives Eq. 1:

$$v = 2k_{cat}[ES_{2}] + 3k_{2}\left(\frac{[S]}{K_{i}}\right)[ES_{2}] = 2k_{cat}[ES_{2}]\left(1 + \frac{3k_{2}}{2k_{cat}} \cdot \frac{[S]}{K_{i}}\right)$$

$$= \frac{2k_{cat}[E]_{0}}{\left(\frac{K_{s}}{[S]^{2}} + 1 + \frac{[S]}{K_{i}}\right)}\left(1 + \frac{3k_{2}}{2k_{cat}} \cdot \frac{[S]}{K_{i}}\right) = \frac{V_{max}[S]^{2}}{\left(K_{s} + [S]^{2} + \frac{[S]^{3}}{K_{i}}\right)}\left(1 + \alpha \cdot \frac{[S]}{K_{i}}\right)$$
(S6)

where $V_{max} = 2k_{cat}[E]_0$, and $\alpha = 3k_2 / 2k_{cat}$

$$E + ATP \underset{K_{M}^{ATP}}{\rightleftharpoons} E(ATP) \xrightarrow{k_{cat}} E + P$$

$$+$$

$$ATP$$

$$\downarrow \uparrow \kappa_{i}$$

$$E(ATP)_{2}$$

Figure S1. Possible catalytic mechanism at 80°C for GK (represented as E) at constant glycerate concentration leading to the rate equation

 $v = \frac{V_{\text{max}}[ATP]}{K_s^{ATP} + [ATP] + \frac{[ATP]^2}{K_i}}$ by binding a second ATP to the enzyme-substrate

complex E(ATP), and leading to an inactive (dead-end) complex E(ATP)₂. The following (rapid) equilibria are assumed with the constants $K_{M}^{ATP} = \frac{[E][ATP]}{[E \times ATP]}$

(Michaelis constant) and $K_i = \frac{[E \cdot (ATP)][ATP]}{[E \cdot (ATP)_2]}$. The reaction rate is defined as

$$v = k_{cat} [E \times ATP].$$



Figure S2. Construction of the PBL2025 Δ 3195 mutant. Exchange of the KDGK encoding gene SSO3195 with the marker gene *lacS* was used for blue/white screening with X-gal. The correct mutant was confirmed by PCR and sequencing with the primers 1557 and 1558. The mutant product runs slower on agarose gel, since the *lacS* gene is larger than SSO3195.



Figure S3. qRT-PCR analysis of PBL2025 \triangle 3195. The C_t values of SSO3194 and SSO3197 were standardized to the C_t values of SSO0685 (*secY*) for each biological replica.