

Supporting Information

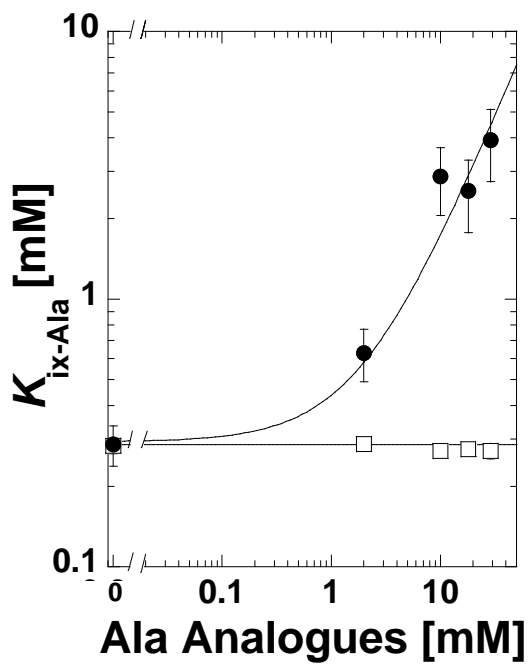


Figure S1. Examples of competitive binding, using the competitive binding of L-alanine methyl ester (●) to wild type hL-PYK. Not all *Ala* analogues that lack an allosteric response show competitive binding with *Ala* (e.g. butylamine, □).

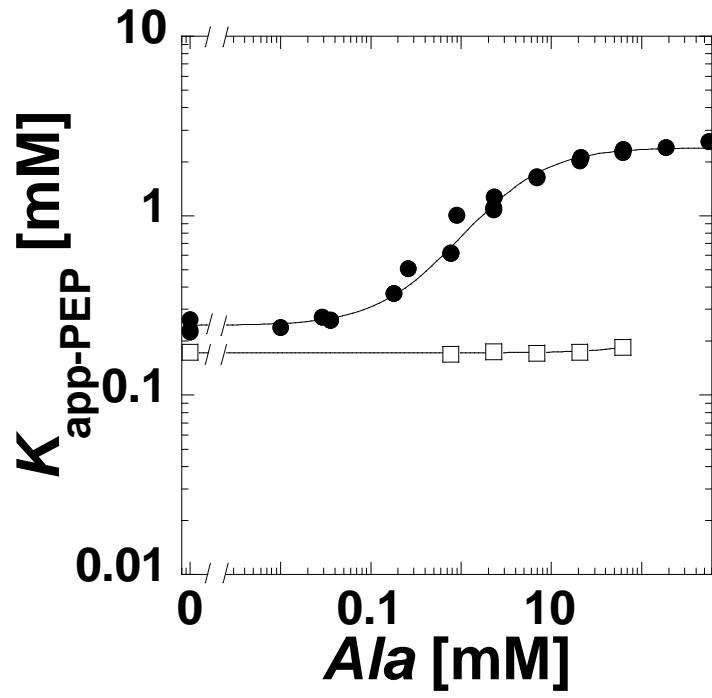
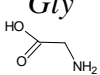
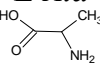
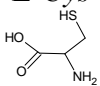
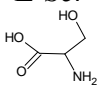
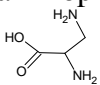
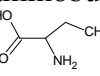
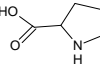
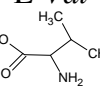
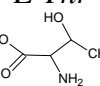
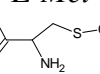
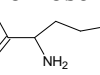
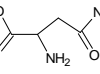
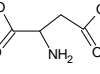
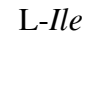
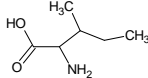
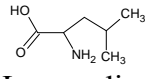
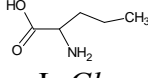
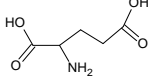
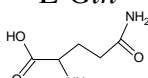
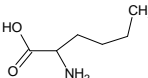
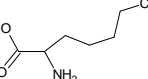
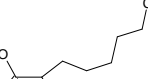
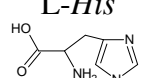
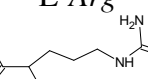
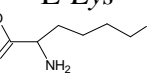


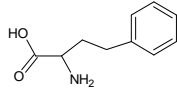
Figure S2. The response of hL-PYK wild type (●) and H476L (□) to 500mM *Ala*.

Table S1: Fit Parameters for Amino Acids

Amino Acid	K_{ix} (mM)	Q_{ax}
<i>Gly</i> 	23±2	0.26±0.01
<i>L-Ala</i> 	0.33±0.01	0.100±0.001
<i>L-Cys</i> 	0.101±0.003	0.124±0.002
<i>L-Ser</i> 	4.6±0.4	0.24±0.01
L-(+)-2,3-diaminopropionic acid 	8±3	0.15±0.05
L-(+)-2-aminobutyric acid 	0.43±0.02	0.05±0.01
<i>L-Pro</i> 	1.40±0.04	0.06±0.001
<i>L-Val</i> 	7.6±0.4	0.09±0.004
<i>L-Thr</i> 	19.6±0.7	0.160±0.003
<i>L-Met</i> 	21±2	0.35±0.01
L-homoserine 	10±4	No Upper Plateau ^b
<i>L-Asn</i> 	— ^a	— ^a
<i>L-Asp</i> 	— ^a	— ^a
<i>L-Ile</i> 	31±8	No Upper Plateau ^b

		
L-Leu	___ a	___ a
		
L-norvaline	29±3	No Upper Plateau ^b
		
L-Glu	___ a	___ a
		
L-Gln	___ a	___ a
		
L-norleucine	___ a	___ a
2-aminoheptanoic acid	___ a	___ a
		
2-aminocaprylic acid	___ a	___ a
		
L-His	___ a	___ a
		
L-Arg	___ a	___ a
		
L-Lys	___ a	___ a
		
L-Phe	3.9±0.3	0.39±0.01
(S)-(+)-2-phenylglycine	___ a	___ a
		
4-nitro-L-phenylalanine	___ a	___ a

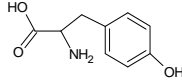
L-homophenylalanine



—^a

—^a

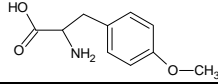
L-Tyr



—^a

—^a

O-methyl-L-tyr



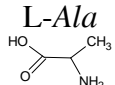
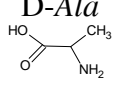
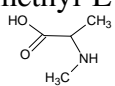
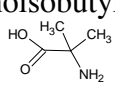
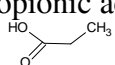
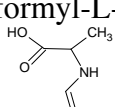
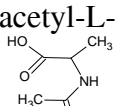
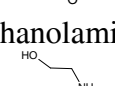
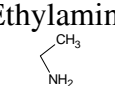
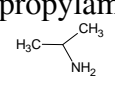
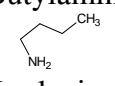
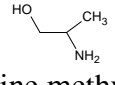
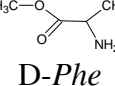
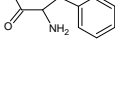
3.8±0.6

0.20±0.04

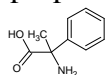
^a K_{a-PEP} was not responsive to the amino acid analogue within the working concentration defined in Materials and Methods.

^bAlthough the amino acid analogue caused a reduction in PEP affinity, within the working concentration range the upper plateau was not obtained.

Table S2: Fit Parameters for Ala and Phe Analogues^a

Ala Analogues	K_{ix} (mM)	Q_{ax}
<i>L-Ala</i> 	0.33±0.01	0.100±0.001
<i>D-Ala</i> 	10±2	0.37±0.02
<i>N-methyl-L-ala</i> 	1.1±0.2	0.16±0.01
2-aminoisobutyric acid 	11±2	0.16±0.01
Propionic acid 	— ^b	— ^b
<i>N-formyl-L-ala</i> 	— ^b	— ^b
<i>N-acetyl-L-ala</i> 	— ^b	— ^b
Ethanolamine 	50±20	No Upper Plateau ^c
Ethylamine 	40±10	No Upper Plateau ^c
Isopropylamine 	25±2	No Upper Plateau ^c
Butylamine 	—	—
L-alaninol 	25±1	No Upper Plateau ^c
L-alanine methyl ester 	1.90±0.05	0.120±0.002
<i>D-Phe</i> 	— ^b	— ^b

S(+)-2-amino-2-methyl-3-phenylpropionic acid



___^b

___^b

^a All responses corrected using the response of H476L as described in text.

^b K_{a-PEP} was not responsive to the amino acid analogue within the working concentration defined in Materials and Methods.

^c Although the amino acid analogue caused a reduction in PEP affinity, within the working concentration range the upper plateau was not obtained.