## **1** Supporting Information

- Evolutionary expansion of the amidohydrolase superfamily in bacteria in response to
  synthetic compounds: the molinate and diuron hydrolases.
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FIG. S1 Both MolA (A) and PuhB (B) have extensive salt bridge formation across the dimerdimer interface (orange, blue). The extended length of the MolA C-terminus allows for additional salt-bridge formation *via* R459 (A).



FIG. S2 (A) An exogenous supply of cobalt (•) and zinc (•) does not increase the activity of MolA. (B) Exogenous bicarbonate addition has no effect on Co-MolA (•) or Zn-MolA (•), in the presence of two equivalents of the respective metals. Error bars represent the standard deviation of two independent replicates. Equivalents refer to the relative concentration of exogenous metal or bicarbonate to protein concentration.

**TABLE S1** Effect of the metal cation on molinate hydrolase (MolA) thermostability

	21	
Enzyme	Melting point (°C)	
Zinc-MolA	46.0 ± 0.34 22	
Apo-MolA	50.8 ± 0.31 23	

FIGURE S3 (A) Substrates from which kinetic parameters could be calculated with molinate hydrolase. (B) Substrates with which there was detectable hydrolysis. (C) Substrates with which there was no detectable hydrolysis.



28





TABLE S2 The kinetic parameters for wild-type MolA and variants with molinate, ethiolate, paraoxon ethyl, *p*-nitrophenyl butyrate, L-alanine *p*-nitroanilide and diuron. The  $k_{cat}/K_m$  of diuron was estimated using a single fixed substrate concentration well below the  $K_m$ . Values shown are the mean and 95% confidence interval of the calculated kinetics.

Substrate	Variant	$K_{\rm m}$ ( $\mu$ M)	$k_{\rm cat}({\rm s}^{-1})$	$k_{\rm cat}/K_{\rm m}({\rm M}^{-1}{\rm s}^{-1})$
	Wild-type	6.88±1.85	8.24x10 <sup>-3</sup> ±6.54x10 <sup>-4</sup>	120 x10 <sup>1</sup>
	D327E	27.09±1.27	1.22x10 <sup>-2</sup> ±4.62x10 <sup>-4</sup>	450
Molinate	S319G	14.79±4.67	5.57x10 <sup>-2</sup> ±3.21x10 <sup>-3</sup>	376 x 10 <sup>1</sup>
N S	N69H	0	0	0
	H215N	0	0	0
	F318V	161.27±15.4	3.29x10 <sup>-3</sup> ±1.75x10 <sup>-5</sup>	20.4
	I73M	17.12±4.54	1.17x10 <sup>-2</sup> ±8.42x10 <sup>-4</sup>	153 x10 <sup>1</sup>
	V217C	59.42±15.56	1.17x10 <sup>-2</sup> ±2.54x10 <sup>-4</sup>	197
	F318M	62.17±5.40	1.16x10 <sup>-2</sup> ±3.16x10 <sup>-4</sup>	189
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	Wild-type	1976.37±322.46	6.66x10 <sup>-2</sup> ±1.79x10 <sup>-3</sup>	33.7
	D327E	1965.10±722.71	2.91x10 <sup>-2</sup> ±3.31x10 <sup>-3</sup>	14.8
Ethiolate	S319G	6266.73±1079.17	6.06x10 <sup>-2</sup> ±6.20x10 <sup>-3</sup>	9.67
	F318V	2914.30±2205.85	1.03x10 <sup>-2</sup> ±2.40x10 <sup>-3</sup>	3.54
	I73M	6099.57±2524.58	1.72x10 <sup>-2</sup> ±2.93x10 <sup>-3</sup>	2.81
	V217C	6234.20±286.81	$3.15 \times 10^{-2} \pm 9.23 \times 10^{-4}$	5.05
	F318M	7621.10±3023.30	2.43x10 <sup>-2</sup> ±3.95x10 <sup>-3</sup>	3.18
	•			
	Wild-type	1298.90±132.24	2.09x10 <sup>-2</sup> ±2.32x10 <sup>-3</sup>	16.1

	D327E	1756.13±138.53	$4.73 \times 10^{-2} \pm 5.52 \times 10^{-3}$	26.9
Paraoxon ethyl	S319G	1591.67±73.84	6.04x10 <sup>-2</sup> ±3.48x10 <sup>-3</sup>	38.0
	F318V	1965.10±605.32	2.03x10 <sup>-2</sup> ±5.16x10 <sup>-3</sup>	10.3
	I73M	1930.97±833.69	1.86x10 <sup>-2</sup> ±5.14x10 <sup>-3</sup>	9.61
0	V217C	539.47±88.89	1.04x10 <sup>-2</sup> ±5.43x10 <sup>-4</sup>	19.3
	F318M	580.65±433.36	3.72x10 <sup>-3</sup> ±2.38x10 <sup>-3</sup>	6.40
	Wild-type	407.20±18.21	1.37±5.47x10 <sup>-2</sup>	336 x10 <sup>1</sup>
	D327E	400.31±49.65	1.67±1.02x10 <sup>-1</sup>	418 x10 <sup>1</sup>
<i>p</i> -Nitrophenyl butyrate	S319G	464.57±28.55	8.97±2.42x10 <sup>-1</sup>	193 x10 <sup>2</sup>
	F318V	484.63±15.77	2.28x10 <sup>-1</sup> ±5.37x10 <sup>-3</sup>	470
	I73M	474.67±28.09	6.49x10 <sup>-1</sup> ±1.76x10 <sup>-2</sup>	137 x10 <sup>1</sup>
	V217C	345.06±48.44	8.28x10 <sup>-1</sup> ±5.60x10 <sup>-2</sup>	240 x10 <sup>1</sup>
	F318M	433.24±11.97	1.15±2.98x10 <sup>-2</sup>	266 x10 <sup>1</sup>
	Wild-type	1586.40±605.27	$1.65 \times 10^{-3} \pm 3.21 \times 10^{-4}$	1.04
	D327E	1583.23±331.47	1.83x10 <sup>-3</sup> ±2.22x10 <sup>-4</sup>	1.15
L-Alanine <i>p</i> -nitroanilide	S319G	1548.77±258.10	2.58x10 <sup>-3</sup> ±2.33x10 <sup>-4</sup>	1.67
	F318V	0	0	0
	I73M	816.55±46.60	1.78x10 <sup>-3</sup> ±3.44x10 <sup>-5</sup>	2.19
	V217C	696.18±18.09	1.72x10 <sup>-3</sup> ±5.69x10 <sup>-5</sup>	2.47
	F318M	0	0	0
	Wild-type			0.43

	D327E		1.67
Diuron	S319G		2.43
O CI	F318V	Could not be determined due to low solubility.	1.20
	I73M		0.28
	V217C		0
	F318M		0.09