

Supplementary Table 1. SAXS parameters across the AtzF and AtzF₄₆₇ concentration series.

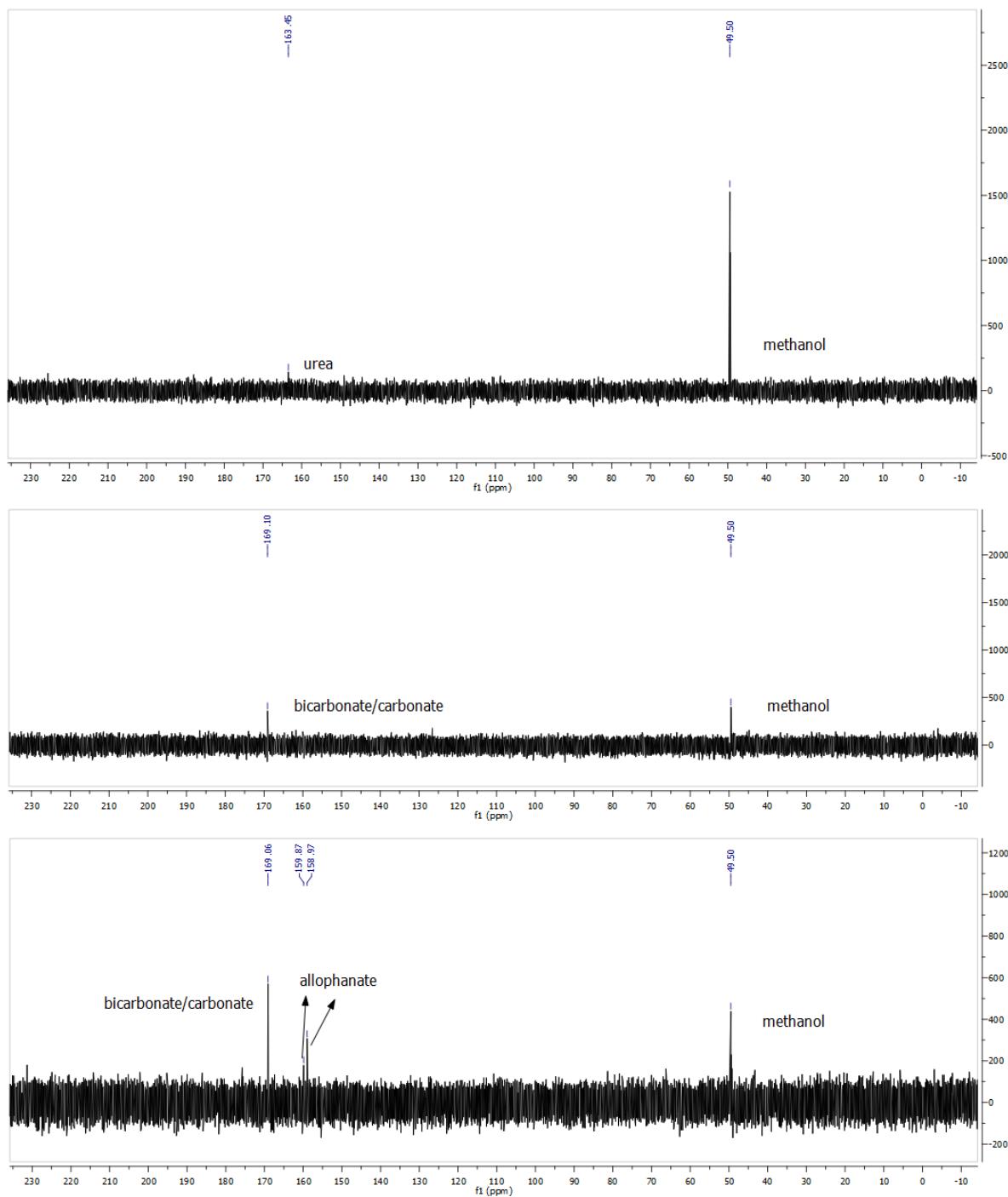
| AtzF | | | | |
|--------------|-------|----------|------|---------------|
| Conc (mg/ml) | I(0) | MW (Kda) | Rg | No. subunits* |
| 0.4 | 0.065 | 249.3 | 50.6 | 3.8 |
| 0.7 | 0.132 | 253.2 | 51.1 | 3.8 |
| 1.5 | 0.267 | 256.1 | 51.2 | 3.9 |
| 2.9 | 0.521 | 249.8 | 51.2 | 3.8 |
| 5.8 | 0.993 | 238.1 | 51.4 | 3.6 |

| AtzF ₄₆₇ | | | | |
|-------------------------|-------|----------------------|-----------|---------------|
| protein conc (mg/ml) | I(0) | Mol. Weight (Kda) | Rg (Å) | No. subunits* |
| 0.8 | 0.071 | 119.8 | 37.7 | 2.3 |
| 1.6 | 0.132 | 111.4 | 37.2 | 2.2 |
| 3.3 | 0.268 | 113.1 | 37.0 | 2.2 |
| 6.6 | 0.506 | 106.7 | 35.8 | 2.1 |
| 13.1 | 0.919 | 96.9 | 36.5 | 1.9 |

Supplementary Table 2. Effect of temperature and pH on the catalytic properties of AtzF,

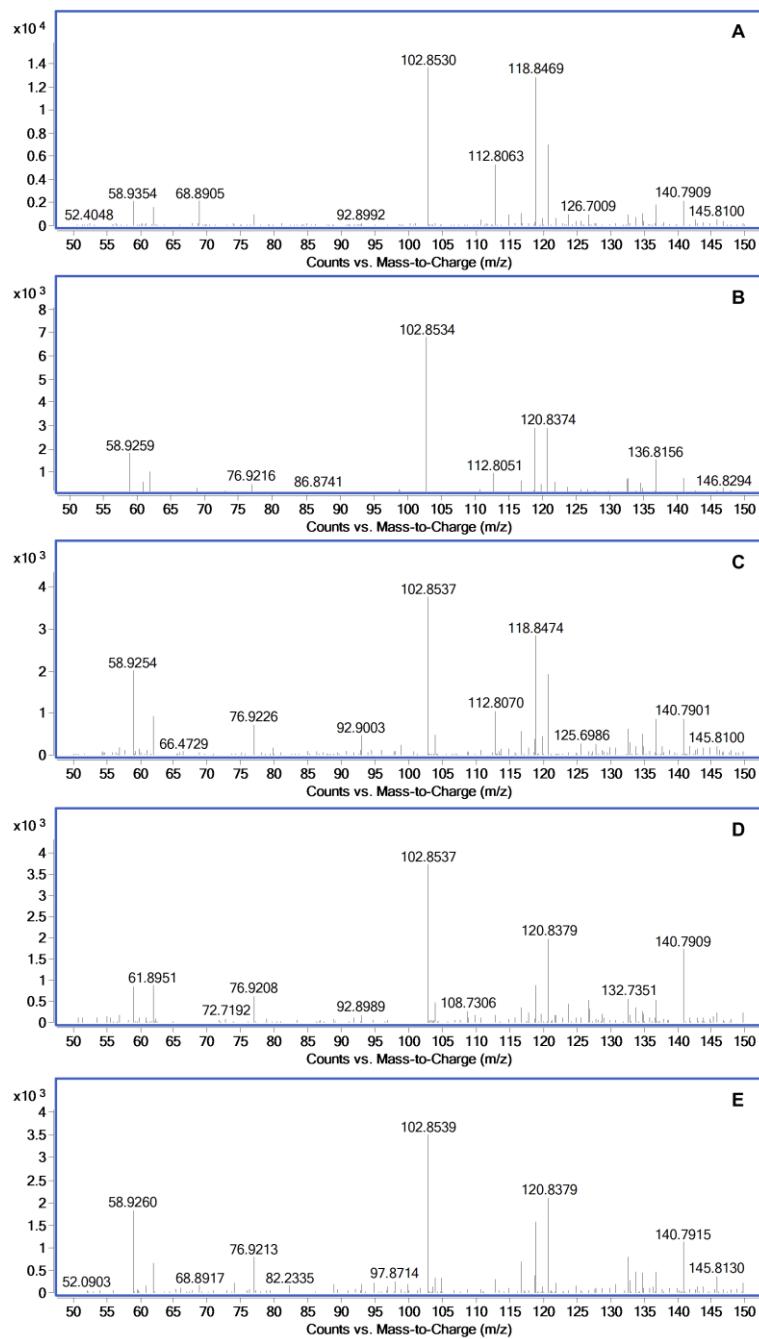
AtzF₄₆₇, and AtzFH488A.

| Enzyme | Temp (°C) | pH | K _M (μM) | k _{cat} (s ⁻¹) | k _{cat} /K _M (M ⁻¹ .s ⁻¹) × 10 ⁴ |
|---------------------|--------------|-----|------------------------|--|---|
| AtzF | 28 | 7.0 | 120.0 ± 21.4 | 8.6 ± 0.4 | 7.1 ± 1.3 |
| AtzF ₄₆₇ | | | 114.6 ± 17.2 | 7.4 ± 0.3 | 6.5 ± 0.1 |
| AtzF H488A | | | 127.6 ± 13.6 | 7.8 ± 0.2 | 6.1 ± 0.7 |
| AtzF | 28 | 7.5 | 149.5 ± 15.7 | 11.6 ± 0.3 | 7.7 ± 0.8 |
| AtzF ₄₆₇ | | | 168.6 ± 21.9 | 11.2 ± 0.4 | 6.7 ± 0.9 |
| AtzF H488A | | | 156.1 ± 11.2 | 11.6 ± 0.2 | 7.5 ± 0.5 |
| AtzF | 28 | 8.0 | 115.7 ± 8.8 | 13.3 ± 0.2 | 11.5 ± 0.9 |
| AtzF ₄₆₇ | | | 109.3 ± 4.2 | 11.8 ± 0.1 | 10.8 ± 0.4 |
| AtzF H488A | | | 113.9 ± 4.7 | 13.1 ± 0.1 | 11.5 ± 0.5 |
| AtzF | 28 | 8.5 | 78.6 ± 4.6 | 13.2 ± 0.2 | 16.8 ± 0.1 |
| AtzF ₄₆₇ | | | 71.0 ± 4.7 | 7.3 ± 0.1 | 10.2 ± 0.7 |
| AtzF H488A | | | 72.6 ± 5.0 | 12.0 ± 0.2 | 16.6 ± 1.0 |
| AtzF | 28 | 9.0 | 74.0 ± 4.6 | 10.7 ± 0.1 | 14.4 ± 0.9 |
| AtzF ₄₆₇ | | | 174.9 ± 26.0 | 3.9 ± 0.2 | 2.2 ± 0.3 |
| AtzF H488A | | | 82.2 ± 5.9 | 9.0 ± 0.1 | 10.9 ± 0.8 |
| AtzF | 4 | 7.0 | 380.2 ± 62.0 | 0.7 ± 0.04 | 0.19 ± 0.03 |
| AtzF ₄₆₇ | | | 363.3 ± 61.0 | 0.7 ± 0.04 | 0.19 ± 0.07 |
| AtzF H488A | | | 305.5 ± 46.9 | 0.7 ± 0.03 | 0.2 ± 0.01 |
| AtzF | 4 | 7.5 | 146.3 ± 5.4 | 2.6 ± 0.02 | 1.8 ± 0.1 |
| AtzF ₄₆₇ | | | 164.0 ± 12.7 | 2.6 ± 0.05 | 1.6 ± 0.1 |
| AtzF H488A | | | 124.1 ± 10.9 | 2.4 ± 0.06 | 1.9 ± 0.2 |
| AtzF | 4 | 8.0 | 44.6 ± 3.3 | 4.2 ± 0.05 | 9.4 ± 0.7 |
| AtzF ₄₆₇ | | | 57.4 ± 4.8 | 3.1 ± 0.05 | 5.4 ± 0.5 |
| AtzF H488A | | | 40.4 ± 2.5 | 4.0 ± 0.04 | 9.9 ± 0.6 |
| AtzF | 4 | 8.5 | 50.8 ± 3.9 | 4.8 ± 0.06 | 9.5 ± 0.7 |
| AtzF ₄₆₇ | | | 54.6 ± 4.3 | 3.7 ± 0.05 | 6.8 ± 0.5 |
| AtzF H488A | | | 47.4 ± 3.7 | 4.8 ± 0.06 | 10.1 ± 0.7 |
| AtzF | 4 | 9.0 | 59.6 ± 4.9 | 4.4 ± 0.07 | 7.3 ± 0.6 |
| AtzF ₄₆₇ | | | 66.0 ± 5.2 | 2.6 ± 0.04 | 3.9 ± 0.3 |
| AtzF H488A | | | 54.5 ± 3.2 | 4.3 ± 0.05 | 8.0 ± 0.5 |

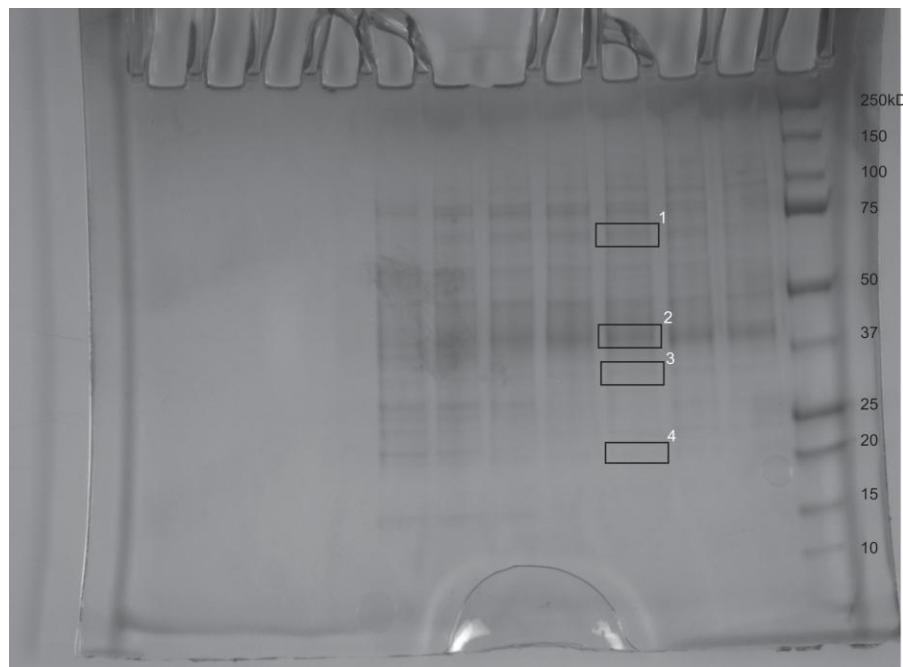


Supplementary figure 1. ^{13}C NMR of allophanate, urea and sodium bicarbonate.

^{13}C NMR shifts of urea, sodium bicarbonate and potassium allophanate in 1M potassium hydroxide prepared in D_2O . First two spectra are of pure urea and bicarbonate/carbonate, which were expected to be present as in allophanate sample as a result of autodecarboxylation, whereas, the third spectra correspond to allophanate sample.



Supplementary Figure 2. Allophanate and *N*-carboxycarbamate in AtzF catalyzed allophanate deamination reactions. Reactions were conducted at pH 9.0 and 4 °C to promote stabilization of *N*-carboxycarbamate (mass \approx 104). An ion consistent with the mass of allophanate (\approx 103) is observed in all reactions. A) Allophanate only, 0 minutes incubation, B) allophanate only at 45 minutes, C) allophanate with AtzF₄₆₈ (0.96 μM) after 45 minutes, D) allophanate with AtzF₄₆₈ (0.48 μM) after 45 minutes, and E) allophanate with AtzF (0.48 μM) after 45 minutes.



Supplementary figure 3. Gel fragments used in mass spectroscopy

Gel fragments used for mass spectroscopy of ~660kD complex that showed cyanuric acid hydrolase, biuret hydrolase and allophanate hydrolase activity and peptides.