

Supporting Information

Development of mechanism-based antibacterial synergy of Fmoc-phenylalanine hydrogel with aztreonam

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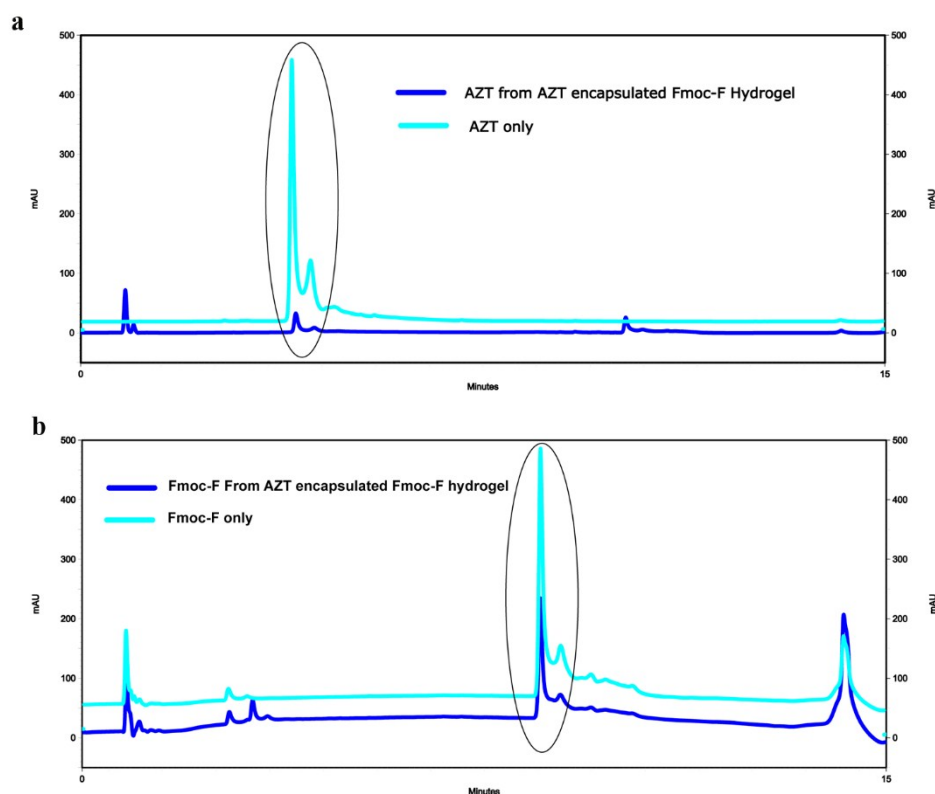


Figure S1: a) RP-HPLC profile of AZT from AZT encapsulated Fmoc-F hydrogel. b) HPLC profile of Fmoc-F from AZT encapsulated Fmoc-F hydrogel.

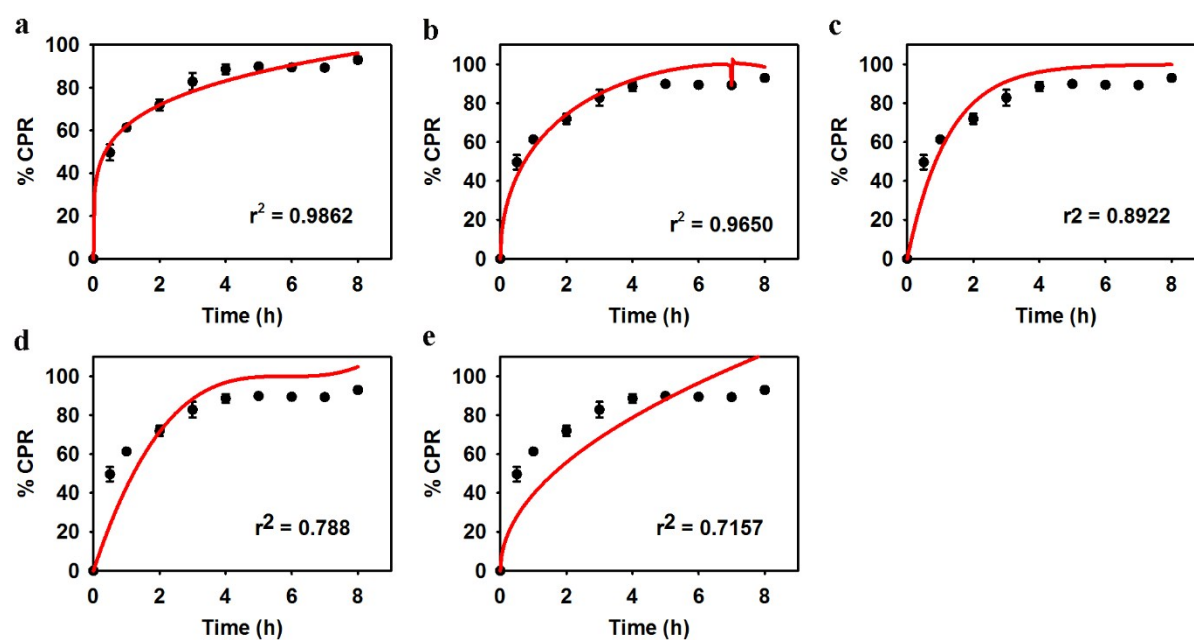


Figure S2: Different release kinetic models for simultaneous release profile of AZT from AZT encapsulated Fmoc-F hydrogel a) Kormeryer-Peppas model b) Baker and Lonsdale model c) First order kinetics model d) Hixon and Crowel model e) Higuchi Square Root Time model.

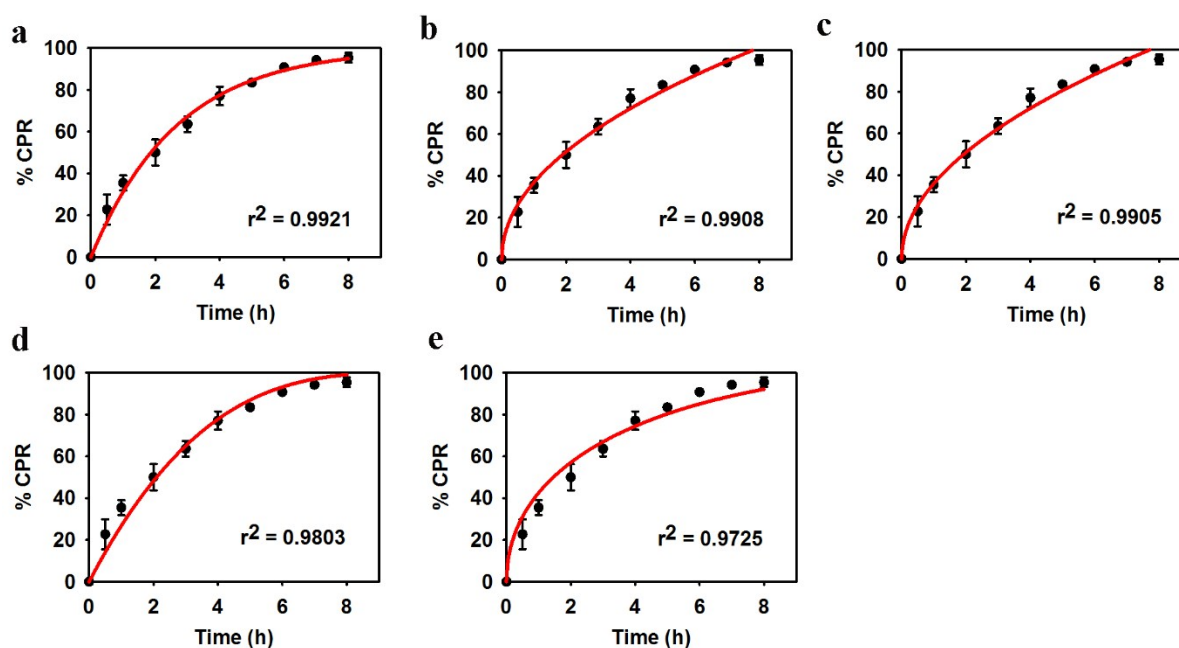


Figure S3: Simultaneous release profile of Fmoc-F from AZT encapsulated Fmoc-F hydrogel and different release kinetic models. a) First order kinetics model b) Kormeryer-Peppas model c) Higuchi Square Root Time model d) Hixon and Crowel model e) Baker and Lonsdale model.

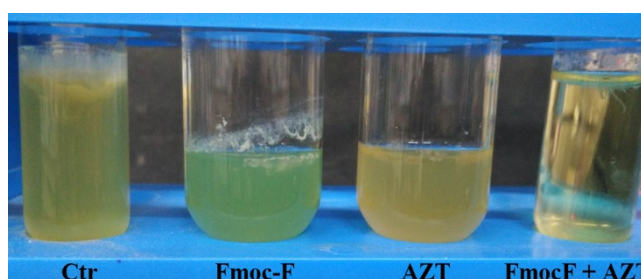


Figure S4: Effect of AZT encapsulated Fmoc-F hydrogel against *S. aureus* and *P. aeruginosa* co-culture.

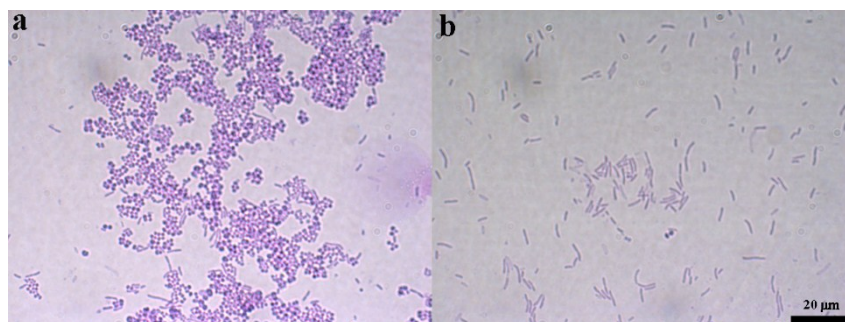


Figure S5: Gram-staining of *S. aureus* and *P. aeruginosa* co-culture treated with AZT (a) or Fmoc-F hydrogel (b) alone.

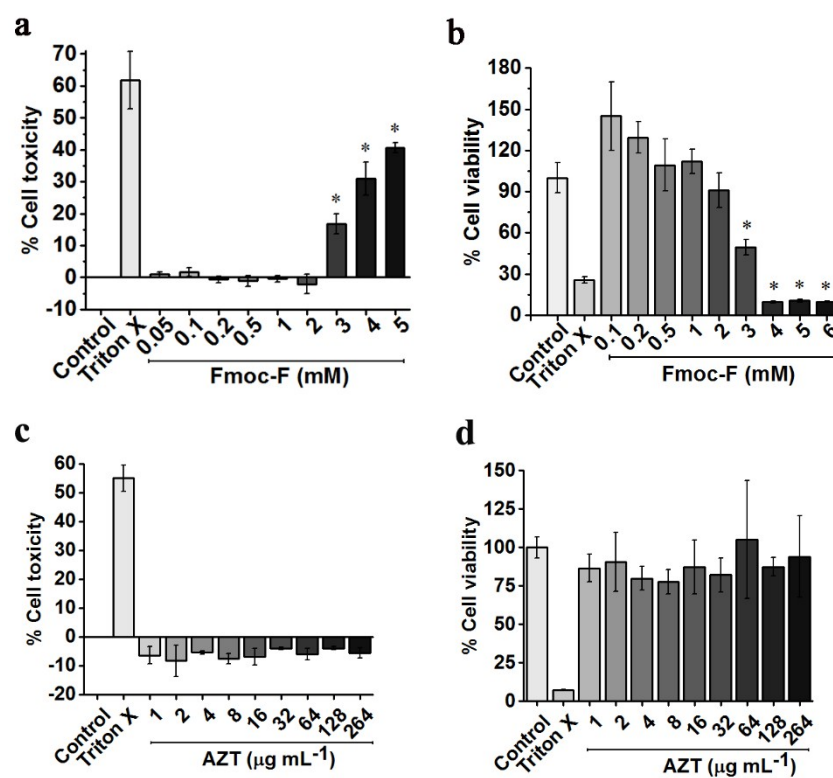


Figure S6: Cytotoxic effect of various concentrations of Fmoc-F (0.05 – 6 mM) on MDCK cells by LDH assay (a) or MTT assay (b). Cytotoxic effect of various concentrations of AZT (1 – 264 µg mL⁻¹) on MDCK cells by LDH assay (c) or MTT assay (d). Triton X (1%) was used as positive

control. * $p < 0.05$ vs control, One-way ANOVA followed by Dunnett's multiple comparison test.

Table S1: Result parameters obtained after data analysis by Gompertz equation to determine MIC of AZT or AZT in Fmoc-F hydrogel or AZT + Fmoc-F solution.

	AZT	AZT in Fmoc-F hydrogel	AZT + Fmoc-F solution
Best-fit values			
logMIC	1.767	1.031	-0.2489
Slope	3.279	7.514	2.219
Bottom	1.644	0.05348	1.091
Span	97.66	50.79	250.9
MIC	58.53	10.74	0.5638
Std. Error			
logMIC	0.09443	0.003513	0.2938
Slope	0.8979	0.1299	1.803
Bottom	4.464	0.05348	0.5696
Span	10.08	0.1753	431.2
95% Confidence Intervals			
logMIC	1.467 to 2.068	0.9864 to 1.076	-1.513 to 1.015
Slope	0.4216 to 6.136	5.863 to 9.164	-5.540 to 9.979
Bottom	-12.56 to 15.85	-0.6260 to 0.7329	-1.360 to 3.542
Span	65.59 to 129.7	48.57 to 53.02	-1605 to 2106
MIC	29.30 to 116.9	9.692 to 11.90	0.03068 to 10.36
Goodness of Fit			
Degrees of Freedom	3	1	2
R square	0.9891	1.000	0.9992
Absolute Sum of Squares	111.2	0.005719	1.924
Sy.x	6.088	0.07563	0.9809
Number of points			
Analyzed	7	5	6

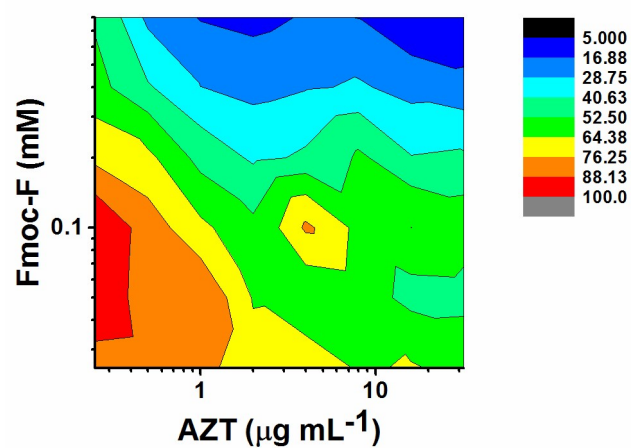


Figure S7: Heat plot showing the effect of AZT and Fmoc-F solution combination on *S. aureus*.

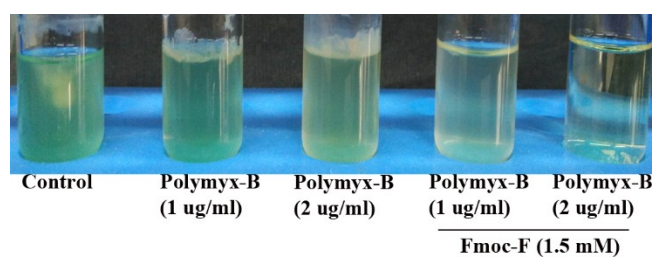


Figure S8: Effect of polymyxin B (1 or 2 $\mu\text{g mL}^{-1}$) + Fmoc-F (1.5 mM) combination or polymyxin B alone against *P. aeruginosa* using broth micro-dilution assay.

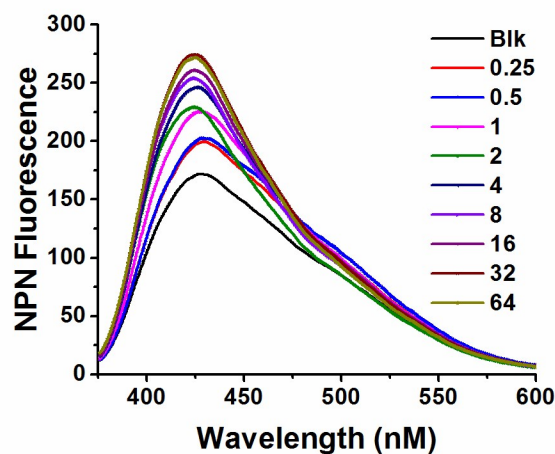


Figure S9: Effect of different concentrations (0.25 – 64 $\mu\text{g mL}^{-1}$) of polymyxin B on NPN fluorescence.

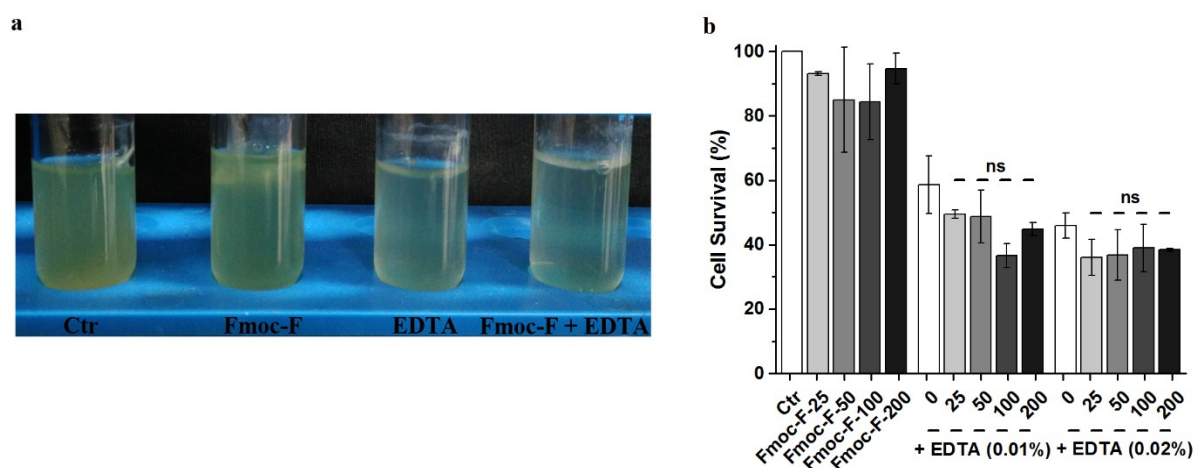


Figure S10: a) Effect of EDTA (0.02 %) + Fmoc-F (1.5 mM) combination or EDTA (0.02 %) alone against *P. aeruginosa* using broth micro-dilution assay. b) Effect of various concentrations of EDTA (0.01 or 0.02 %) + Fmoc-F (25 – 200 μM) combination or EDTA (0.01 or 0.02 %) alone against *P. aeruginosa* using broth micro-dilution assay. ns – nonsignificant, One-way ANOVA followed by Newman-Keuls multiple comparison test.