

## Modulation of Acetylcholinesterase Activity Using Molecularly Imprinted Polymer Nanoparticles

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**Fig. S11.** Direct measure of substrate conversion by enzyme alone and in the presence of FRF-MIP.

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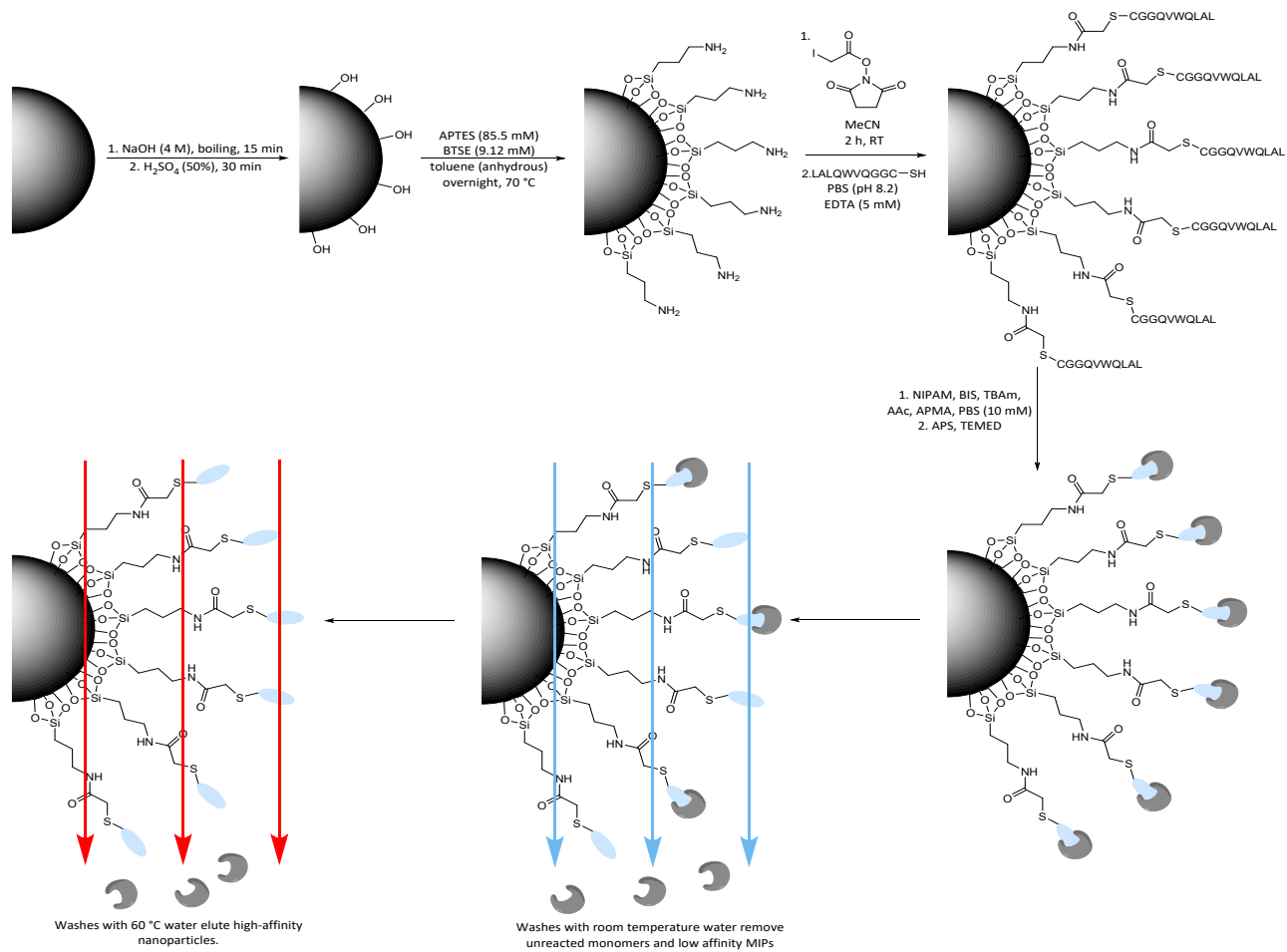
**Fig. S13.** Michaelis-Menten plot obtained for FRF-MIP.

#### Supplementary Data Tables

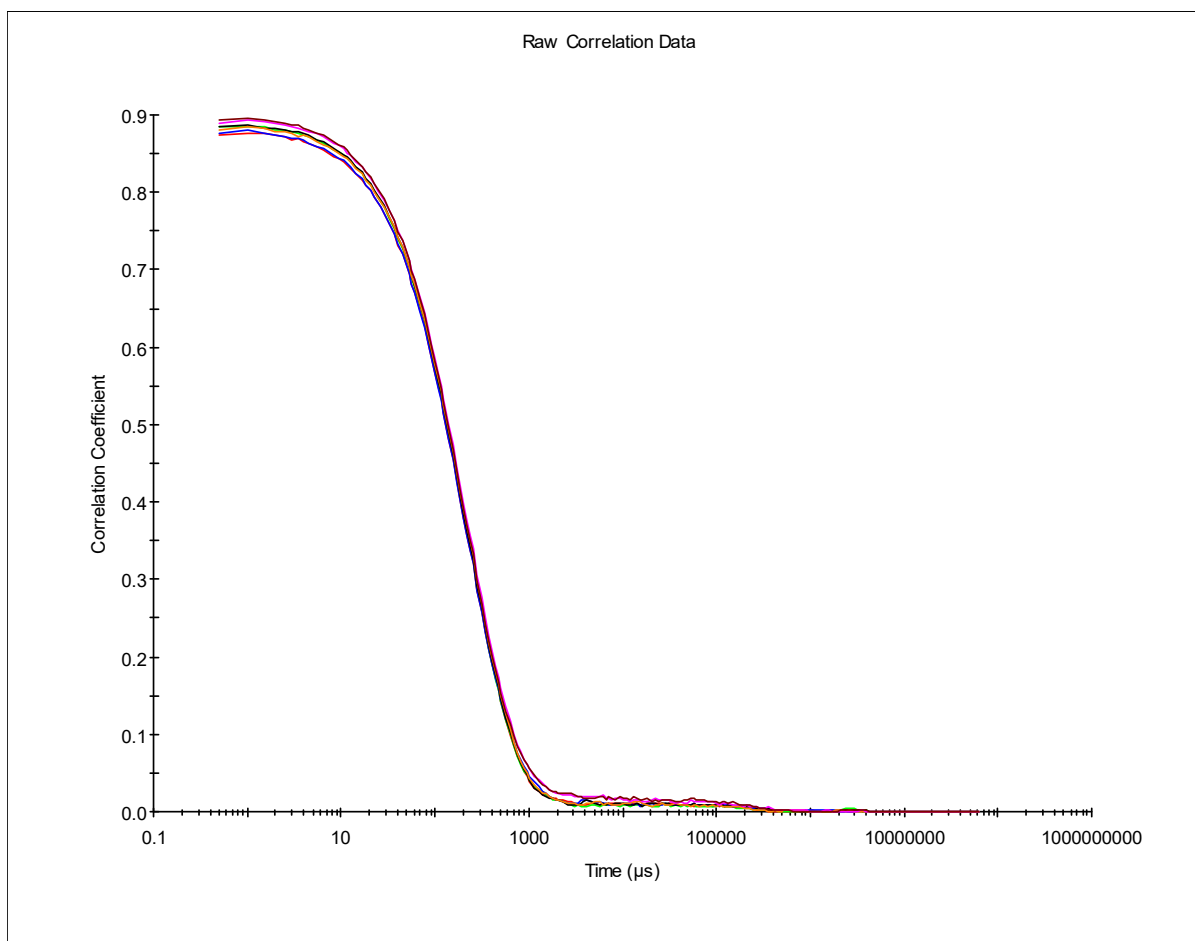
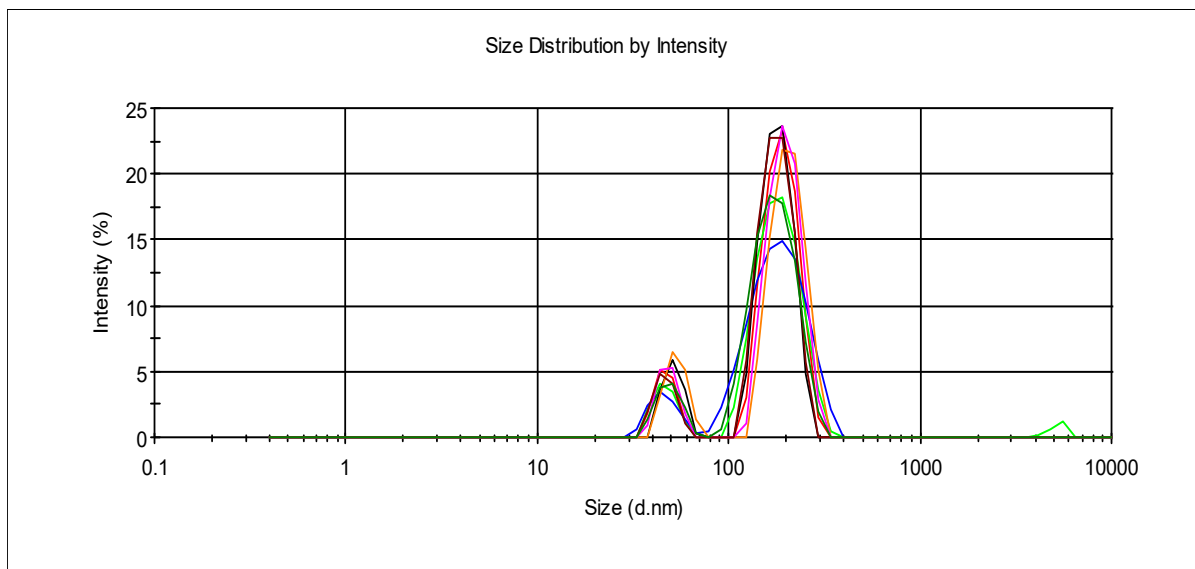
**Table S1:** Size values and distribution for the nanoMIPs measured using DLS.

**Table S2.** Affinity coefficient ( $K_D$ ) of the synthesized MIP nanoparticles upon interaction with EeAChE enzyme measured using SPR.

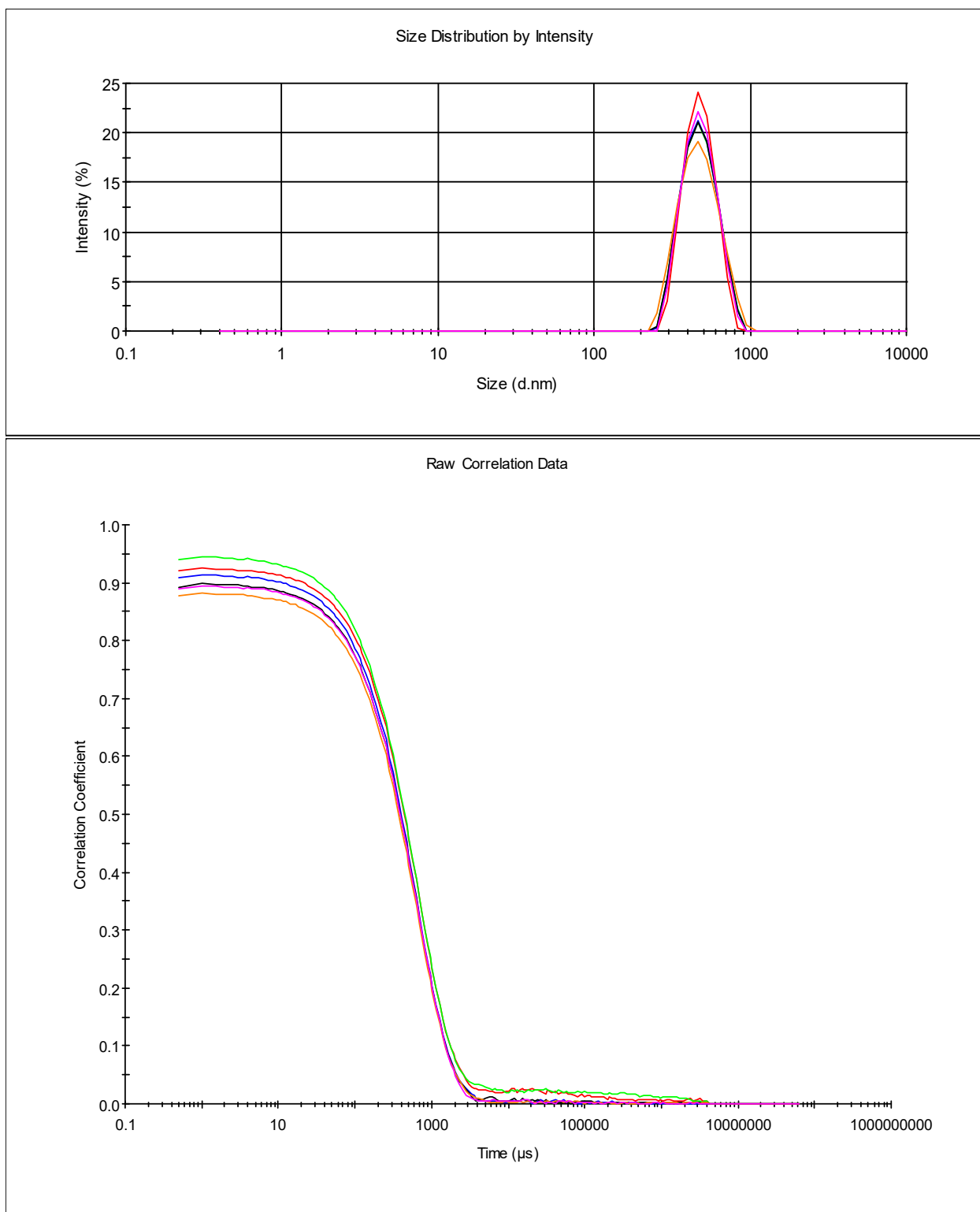
**Table S3.** Values obtained from Michaelis-Menten plot for FRF-MIP.



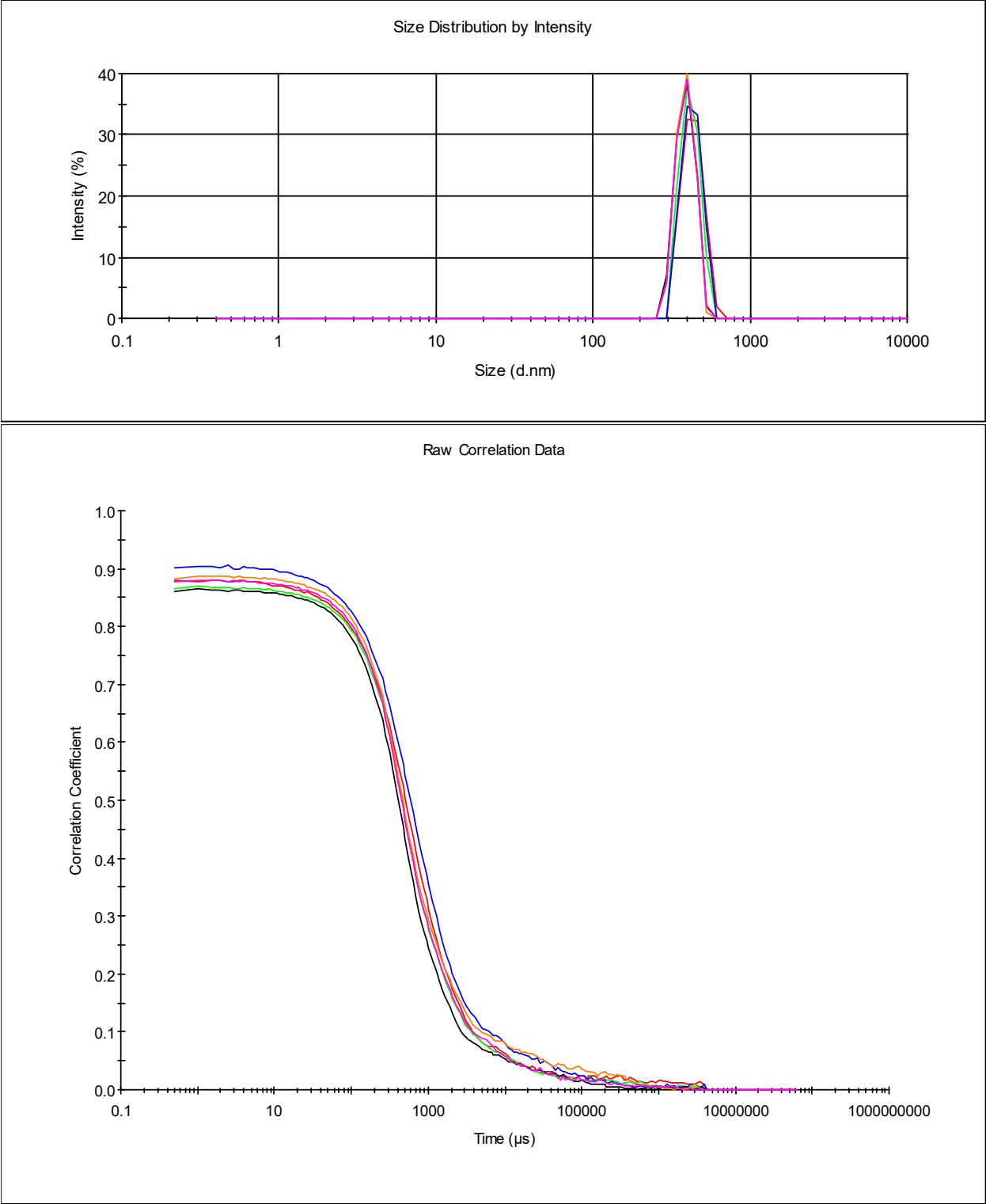
**Fig. S1.** Solid phase synthesis of MIP nanoparticles.



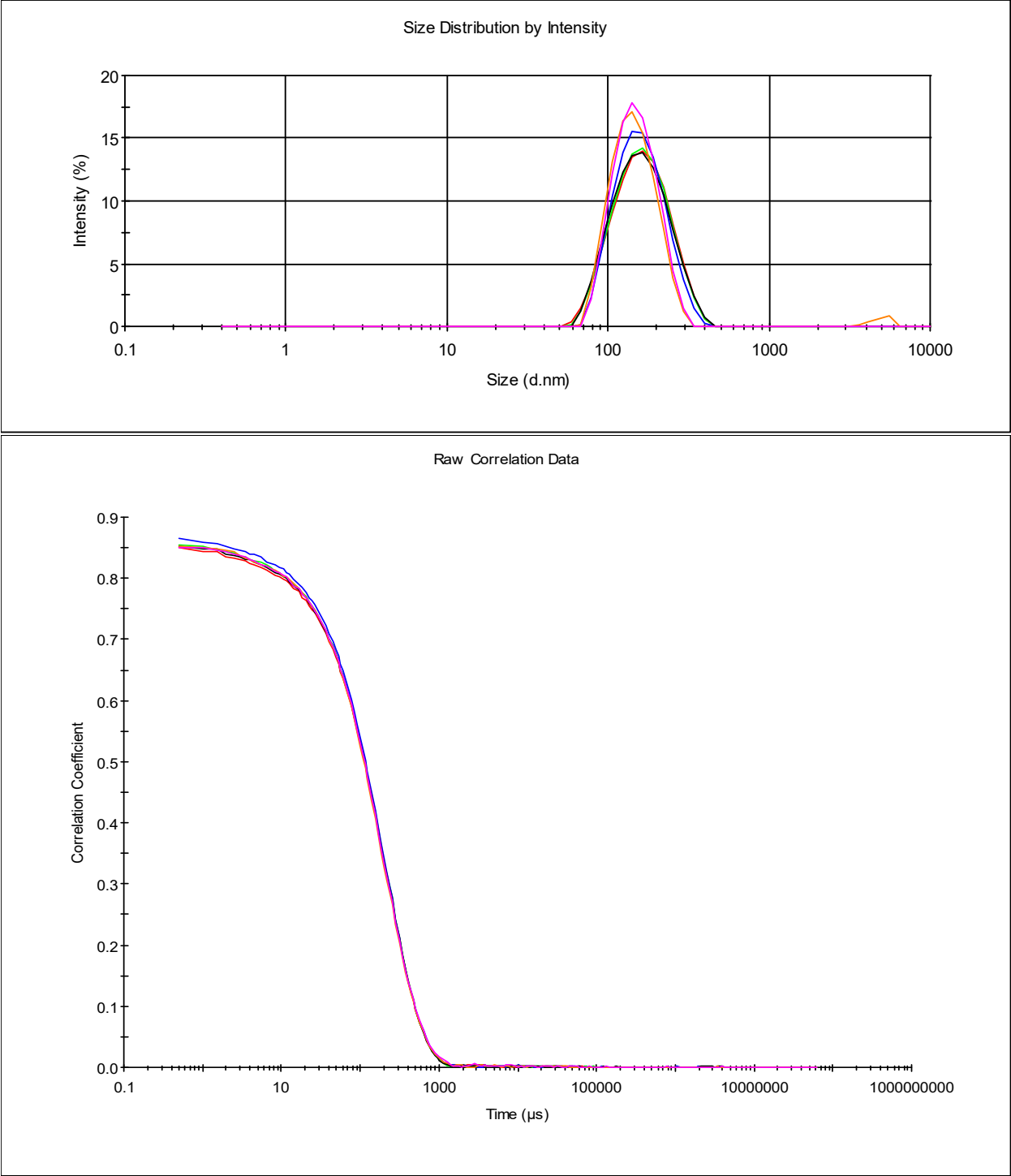
**Fig. S2.** Size distribution by intensity (top) and correlograms (bottom) for YWA-MIPs synthesized in this work (for the size and distribution values, see Table S1).



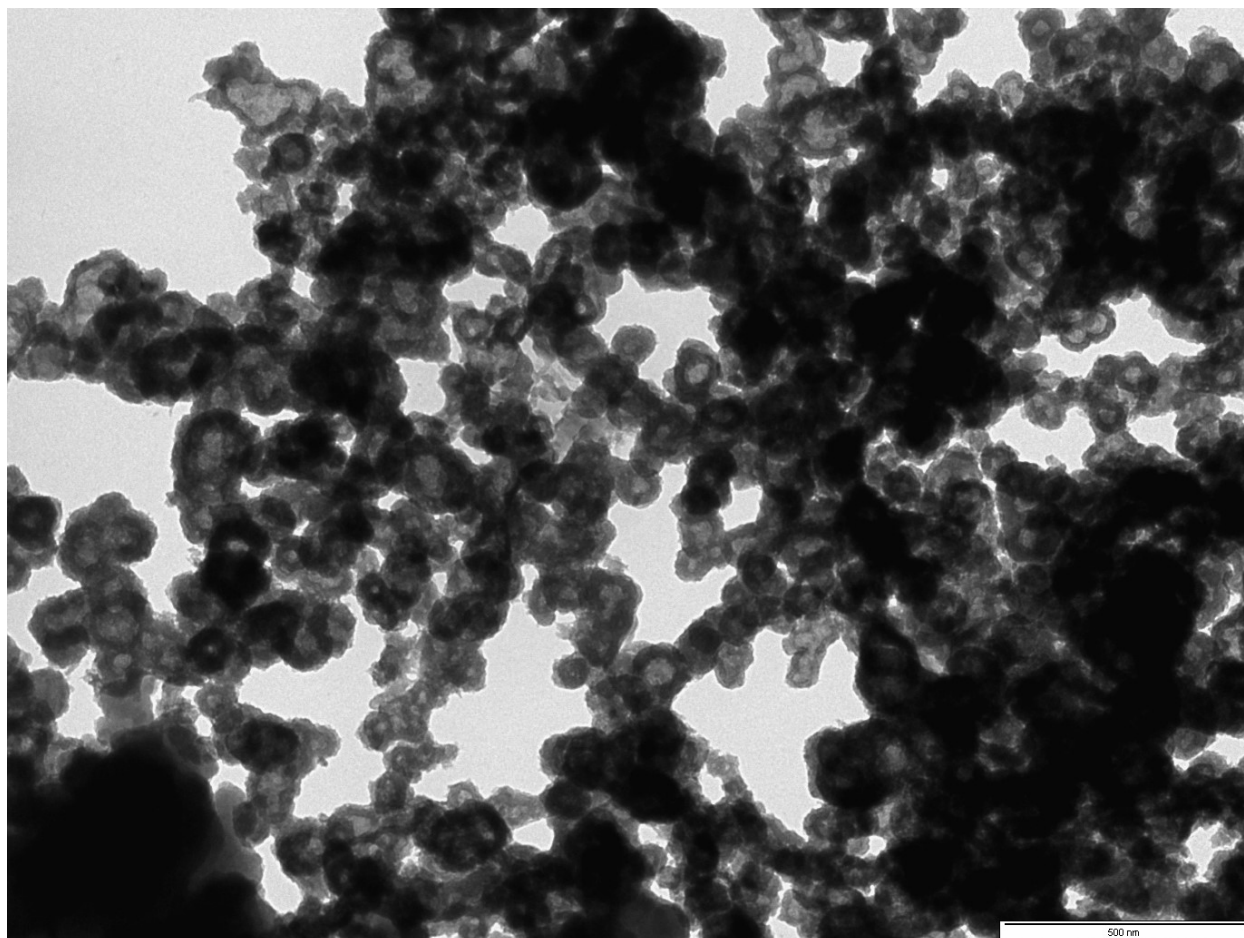
**Fig. S3.** Size distribution by intensity (top) and correlograms (bottom) for FRF-MIPs synthesized in this work (for the size and distribution values, see Table S1).



**Fig. S4.** Size distribution by intensity (top) and correlograms (bottom) for LAL-MIPs synthesized in this work (for the size and distribution values, see Table S1).



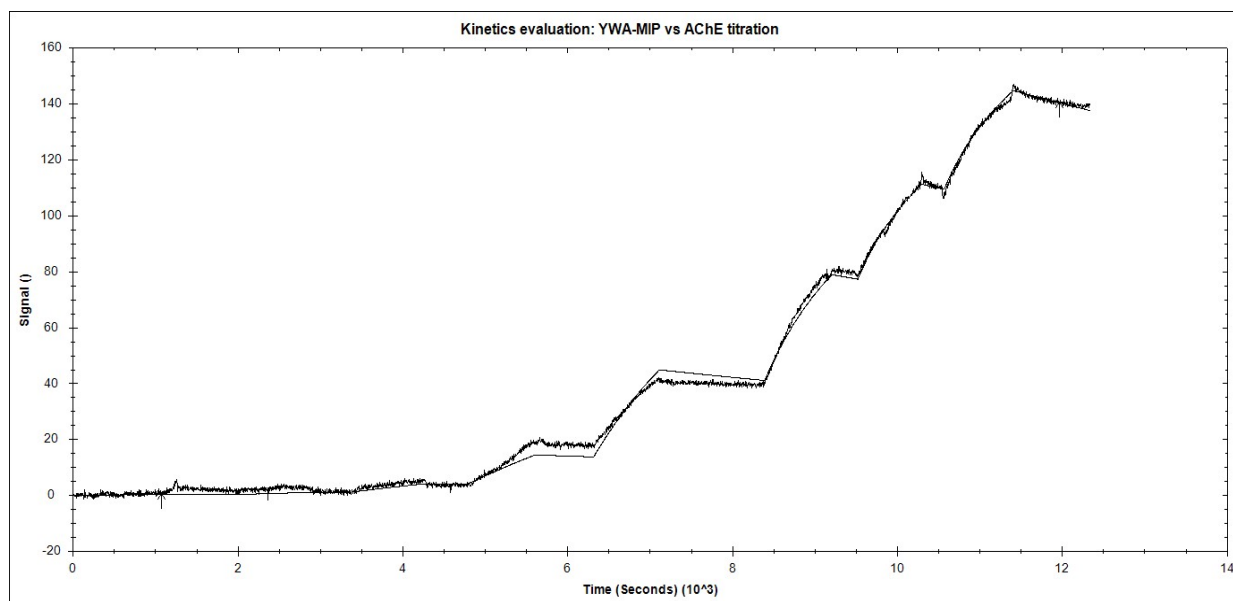
**Fig. S5.** Size distribution by intensity (top) and correlograms (bottom) for FGE-MIPs synthesized in this work (for the size and distribution values, see Table S1).



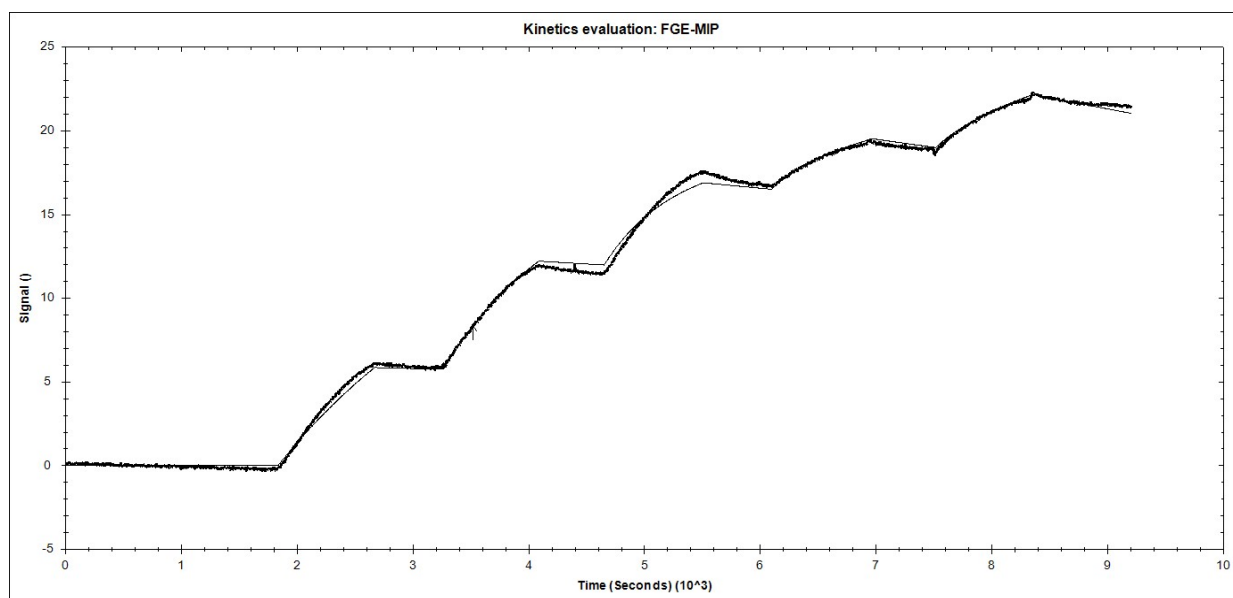
**Fig. S6.** Typical TEM image of the YWANFAR-specific MIP NPs specific for AChE.

**Table S1:** Size values and distribution for the nanoMIPs measured using DLS.

<b>MIP</b>	<b>Size by intensity (d. nm)</b>	<b>Pdi</b>	<b>Number mean (d. nm)</b>
LAL	408.4 ± 20.52	0.439 ± 0.056	393.6 ± 18.87
YWA	189.5 ± 10.52	0.304 ± 0.032	44.81 ± 2.86
FRF	484.3 ± 1.818	0.102 ± 0.047	431.1 ± 7.04
FGE	163.5 ± 9.381	0.140 ± 0.016	101.5 ± 6.76

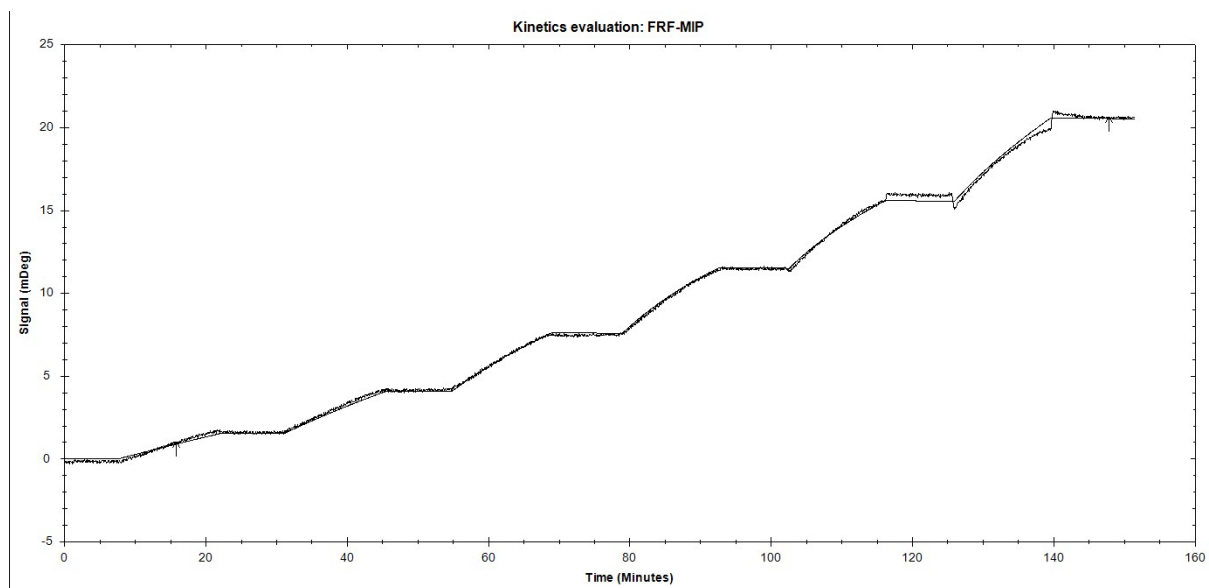


**Fig. S7.** SPR sensorgrams showing response of YWA-MIPs to injections of different concentrations of AChE. A kinetic titration injection strategy was employed for these experiments due to the difficulty of surface regeneration. AChE was injected at 5 different concentrations from 0.1 nM to 1  $\mu$ M and allowed to associate and dissociate for 14 min and 5 min respectively, before a final dissociation of 120 min (not shown). All data were reference subtracted against a control polymer of the same monomer composition, and fit to a 1:2 interaction model using Tracedrawer 1.8 software.

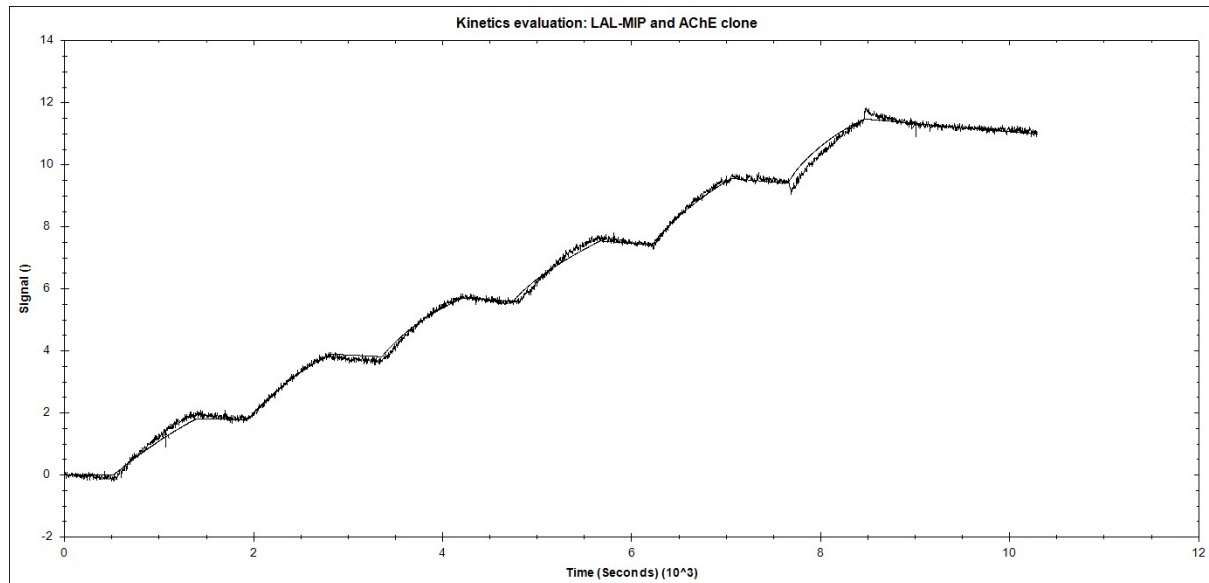


**Fig. S8.** SPR sensorgrams showing response of FGE-MIPs to injections of different concentrations of AChE. A kinetic titration injection strategy was employed for these experiments due to the difficulty of surface regeneration. AChE was injected at 5 different concentrations from 0.1 nM to 1  $\mu$ M and allowed to associate and dissociate for 14 min and 5 min respectively, before a final dissociation of 120 min (not shown). All data were reference subtracted against a control polymer of the same monomer composition, and fit to a 1:2 interaction model using Tracedrawer 1.8 software.





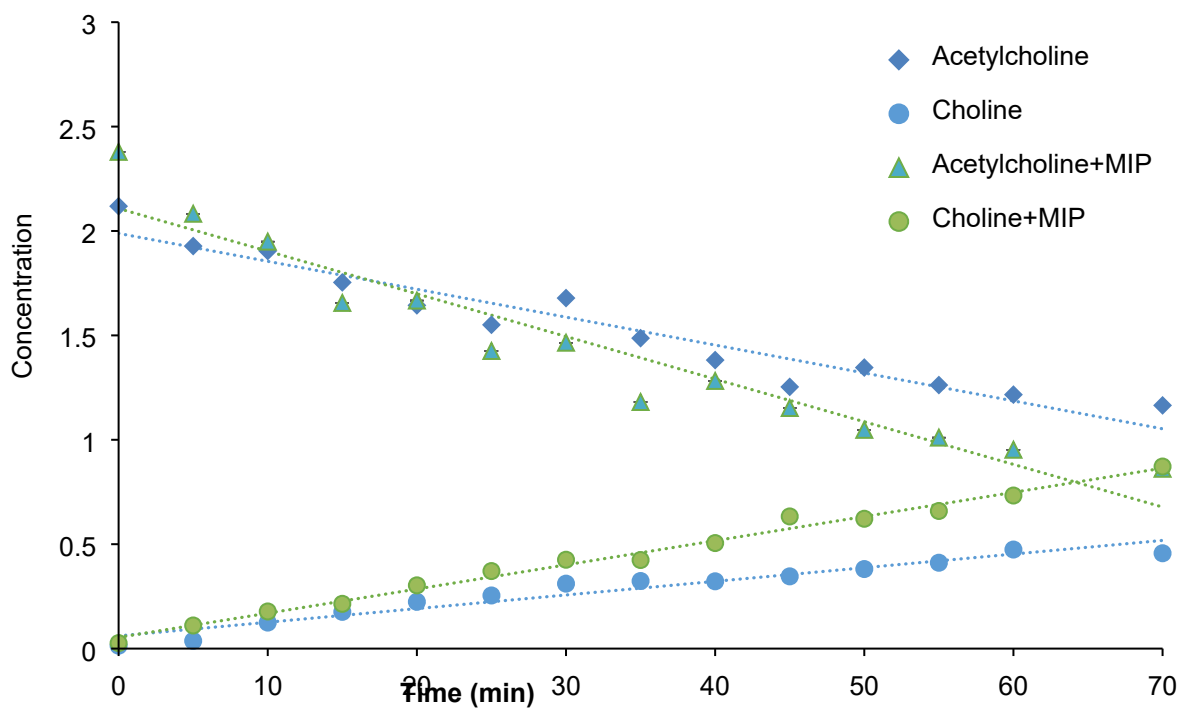
**Fig. S9.** SPR sensorgrams showing response of FRF-MIPs to injections of different concentrations of AChE. A kinetic titration injection strategy was employed for these experiments due to the difficulty of surface regeneration. AChE was injected at 5 different concentrations from 0.1 nM to 1  $\mu$ M and allowed to associate and dissociate for 14 min and 5 min respectively, before a final dissociation of 120 min (not shown). All data were reference subtracted against a control polymer of the same monomer composition, and fit to a 1:2 interaction model using Tracedrawer 1.8 software.



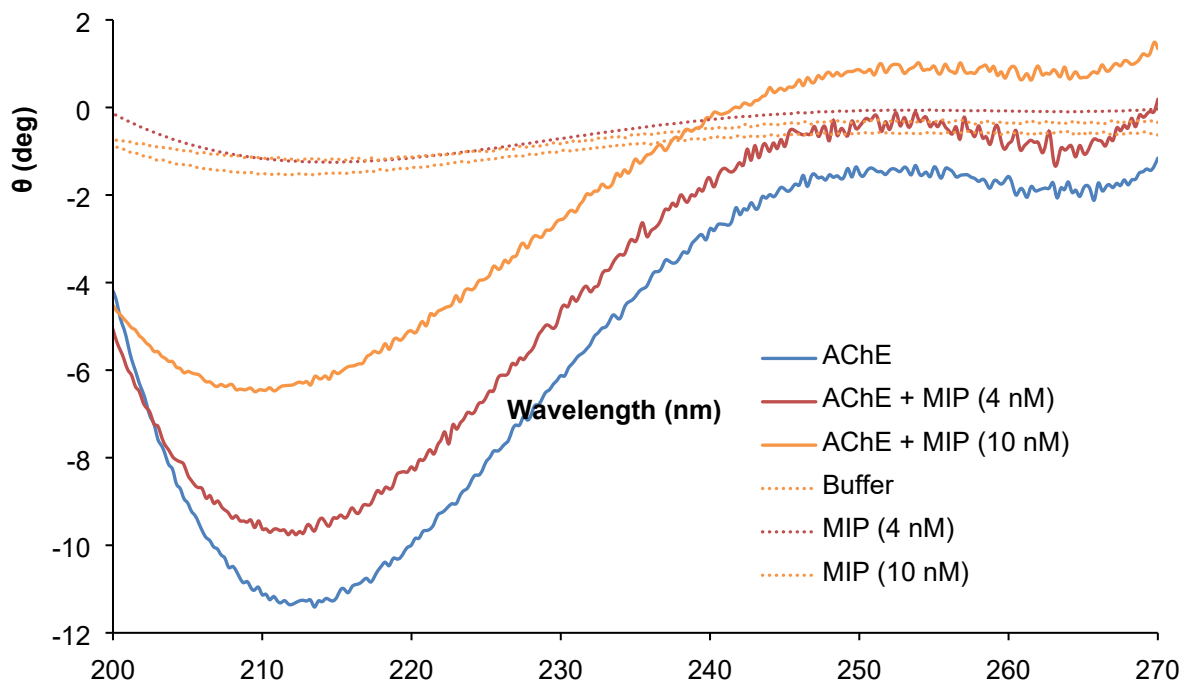
**Fig. S10.** SPR sensorgrams showing response of LAL-MIPs to injections of different concentrations of AChE. A kinetic titration injection strategy was employed for these experiments due to the difficulty of surface regeneration. AChE was injected at 5 different concentrations from 0.1 nM to 1  $\mu$ M and allowed to associate and dissociate for 14 min and 5 min respectively, before a final dissociation of 120 min (not shown). All data were reference subtracted against a control polymer of the same monomer composition, and fit to a 1:2 interaction model using Tracedrawer 1.8 software.

**Table S2.** Affinity coefficient ( $K_D$ ) of the synthesized MIP nanoparticles upon interaction with EeAChE enzyme measured using SPR.

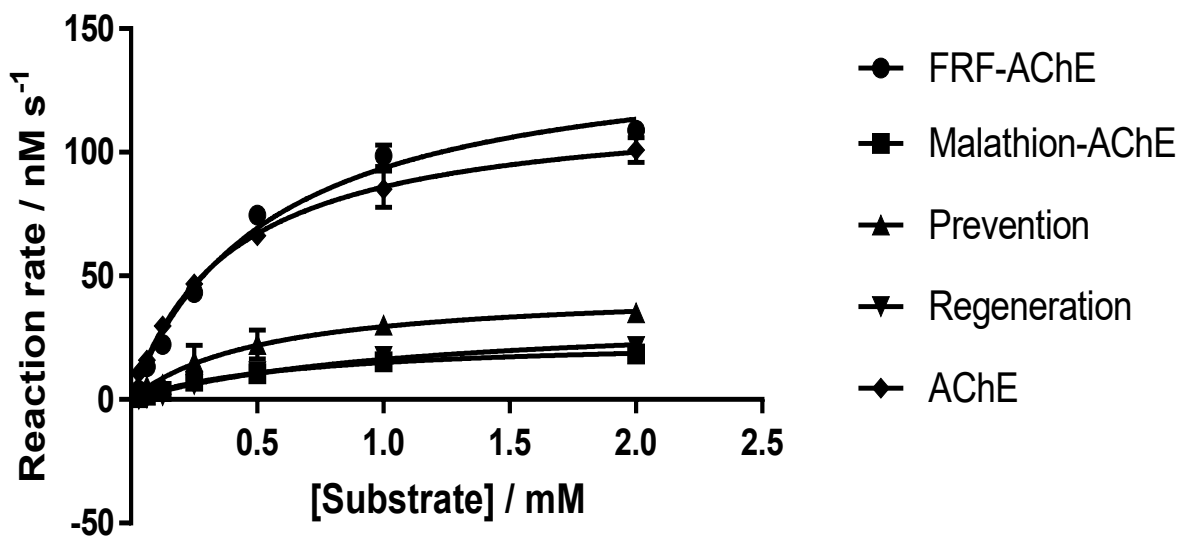
Epitope used as template	$K_D$ , nM	$\text{Chi}^2$
YWANFAR (YWA-MIP)	12.0	4.03
QVTIFGESAGAASVGM-HLLSPDSRPK (FGE-MIP)	78.6	0.06
FRFSFVPV (FRF-MIP)	0.40	0.03
LALQWVQDNIHFFGGNPK (LAL-MIP)	2.20	0.02



**Fig. S11.** Direct measure of substrate conversion by enzyme alone and in the presence of FRF-MIP.



**Fig. S12.** Circular dichroism spectra of AChE with increasing concentration of YWA-MIP. Dashed lines correspond to control measurements of MIP without protein. Note that the nanoMIP materials appear to be inert spectroscopically.



**Fig. S13.** Michaelis-Menten plot obtained for FRF-MIP. Error bars indicate SD. N=3

**Table S3.** Values obtained from Michaelis-Menten plot for FRF-MIP.

	<b>FRF-MIP- AChE</b>	<b>Malathion- AChE</b>	<b>Prevention</b>	<b>Regeneration</b>	<b>AChE</b>
<b>Michaelis-Menten</b>					
<b>Best-fit values</b>					
<b>Vmax</b>	143.9	25.01	45.12	34.92	119.7
<b>Km</b>	0.5378	0.674	0.5301	1.147	0.3928
<b>Std. Error</b>					
<b>Vmax</b>	6.829	1.315	4.506	4.047	3.342
<b>Km</b>	0.06444	0.08344	0.1343	0.2637	0.03039
<b>95% CI (profile likelihood)</b>					
<b>Vmax</b>	130.8 to 159.4	22.45 to 28.15	36.93 to 57.09	28.23 to 45.9	112.7 to 127.5
<b>Km</b>	0.4222 to 0.6895	0.5203 to 0.8828	0.3127 to 0.9282	0.7399 to 1.902	0.3313 to 0.4664
<b>Goodness of Fit</b>					
<b>Degrees of Freedom</b>	12	12	12	12	12
<b>R square</b>	0.9863	0.9867	0.9372	0.9714	0.9924
<b>Absolute Sum of Squares</b>	295.1	7.575	131.4	25.55	110.8
<b>Sy. x</b>	4.959	0.7945	3.309	1.459	3.038
<b>Constraints</b>					
<b>Km</b>	Km > 0	Km > 0	Km > 0	Km > 0	Km > 0
<b>Number of points</b>					
<b># of X values</b>	14	14	14	14	14
<b># Y values analyzed</b>	14	14	14	14	14