

Multiple-Responsive Supramolecular Vesicle Based on Azobenzene-Cyclodextrin Host-Guest Interaction

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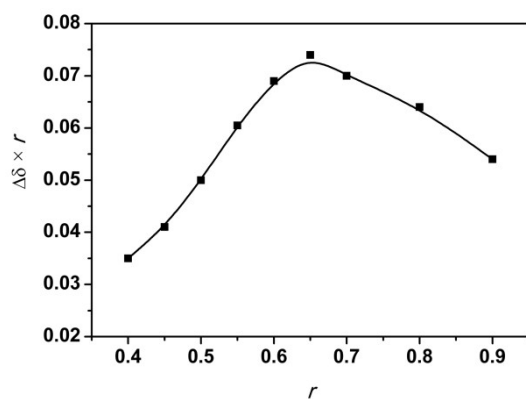


Fig. S1 ^1H NMR Job's plot corresponding to the chemical shift of H-5 of β -CD in D_2O . $[\text{HPB}] + [\beta\text{-CD}] = 1.0 \text{ mM}$.

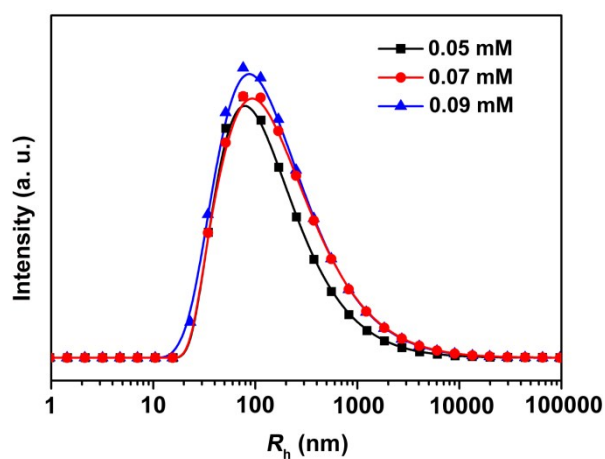


Fig. S2 Hydrodynamic radius (R_h) distribution determined by DLS for $\text{HPB}@ \beta\text{-CD}$ aqueous solution at different concentration.

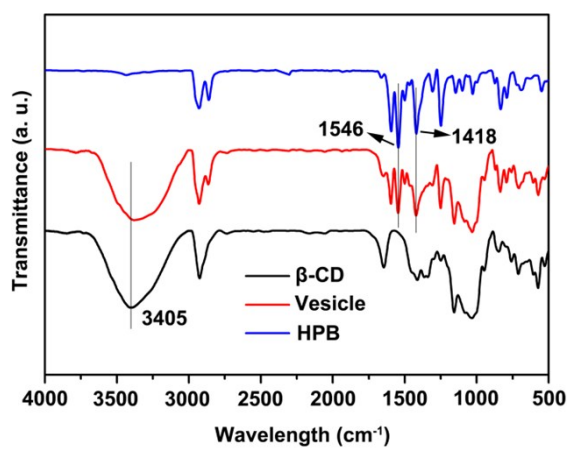


Fig. S3 FTIR spectra of pure β -CD, HPB and dehydrated vesicles from $0.1 \text{ mM HPB}@ \beta\text{-CD}$ sample.

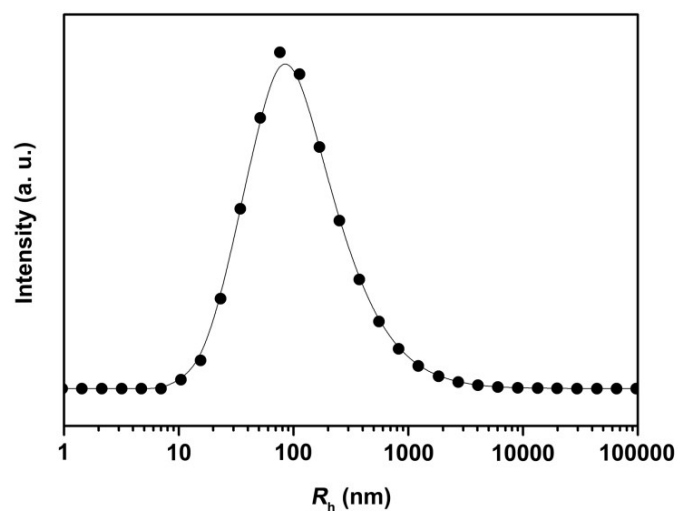


Fig. S4 Hydrodynamic radius (R_h) distribution determined by DLS for the reformed vesicles.

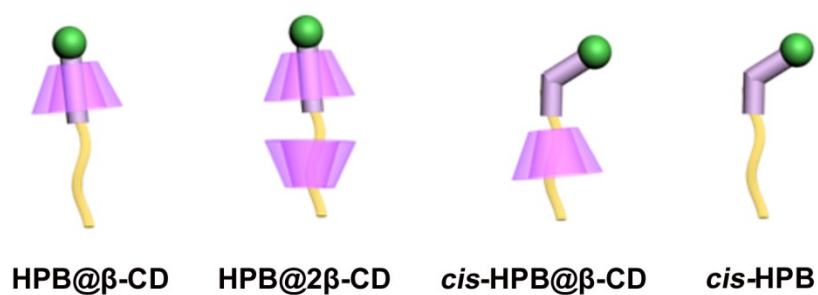


Fig. S5 Molecular structure models of HPB@β-CD, HPB@2β-CD, *cis*-HPB@β-CD and *cis*-HPB.

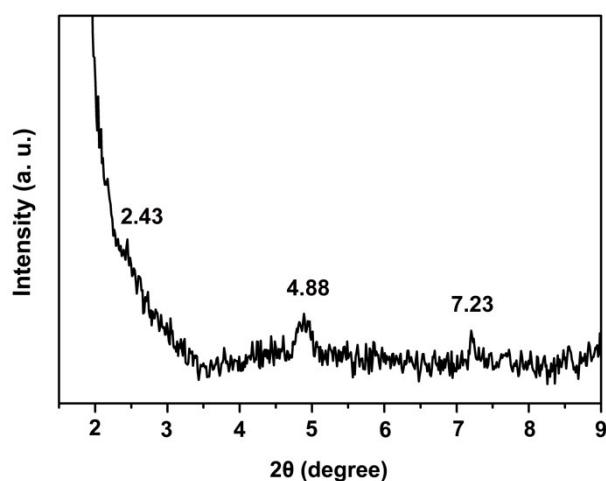


Fig. S6 XRD pattern of microbelts from 0.1 mM HPB@β-CD sample at pH = 4.5. The sample was dried in vacuum at room temperature.

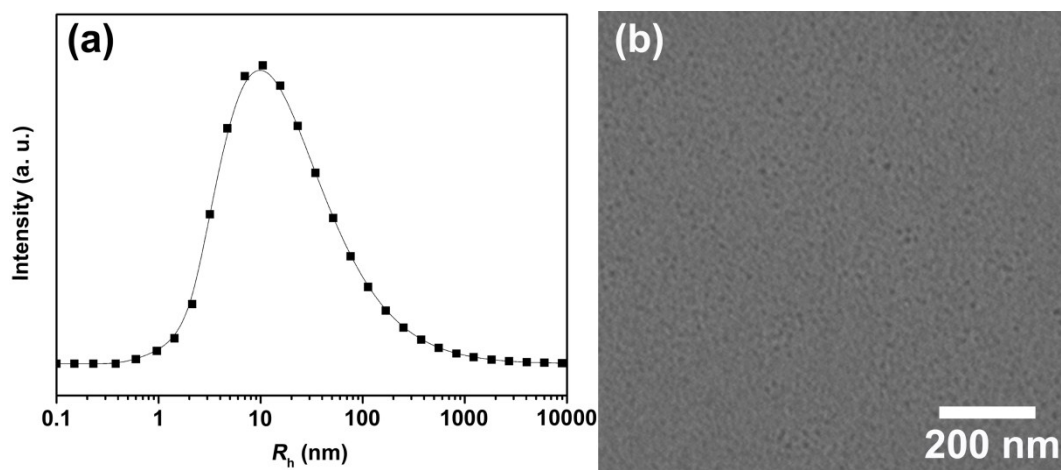


Fig. S7 Hydrodynamic radius (R_h) distribution (a) determined by DLS and TEM (b) for micelles from 0.1 mM HPB@ β -CD aqueous solution after adding ADA at 25 °C.

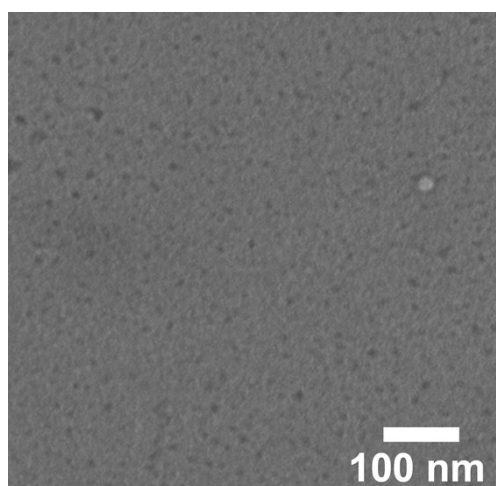


Fig.S8 TEM (b) of micelles formed in 0.1 mM HPB@ β -CD sample after adding α -amylase.