## ELECTRONIC SUPPLEMENTARY INFORMATION

## Limpid hydrogels from $\boldsymbol{\beta}$-turn motif-connected tandem repeats of $\mathbf{A} \boldsymbol{\beta}_{16-22}$

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## Transmission Electron Microscopy (TEM)

TEM samples were prepared by drop-casting $5 \mu \mathrm{l}$ of the gel samples ( 24 h old) on 200 mesh Formvar/Carbon coated copper grids. After 2 minutes, the excess amount was blotted out from the periphery of the grid by Whatman I filter paper. The grids were then stained with saturated uranyl acetate negative stain solution which was blotted out after 30 seconds. The samples were dried and images recorded on Field Emission Transmission Electron Microscopy instrument (Make: JEOL, Model: 2100F).
(A)

(D)

(B)

(E)

(C)

(F)


Figure S1. TEM images of A $\beta$ FF-NG (A), A $\beta$ FF-pG (B), A $\beta$ FF-Up (C), A $\beta F Y-N G(D), A \beta F Y-$ pG (E), and AßFY-Up (F) gel samples


Figure S2. Mechano-responsive behaviour of the $5 \mathrm{mM} A \beta F F-N G$ and $A \beta F Y-N G$ gels. The shearthinned samples form gels upon 12 h standing.


Figure S3. Hydrogelation by $A \beta F F-N G$ and $A \beta F Y-N G$ at pH 2 and 4. The pH of the peptide solutions was adjusted using HCl .


Figure S4. Fraction doxorubicin release from $5 \mathrm{mM} A \beta F F-p G$ (panel A) and A $\beta$ FF-Up (panel B) gel samples.

Table S1. The sequences of the PCR primers used for GAPDH and insulin-1 cDNA amplification

| GAPDH primers |
| :---: |
| 5'-ATGGAGAAGGCTGGGGCTCA-3' <br> 5'-GTTGTCATGGATGACCTTGGC-3' |
| Insulin-1 primers |
| 5'-GCACCTTTGTGGTCCTCACCT-3' |
| 5'-GCCTCCACCCAGCTCCAGTT-3' |

