## Supplementary information

# Interactions between $\boldsymbol{\beta}$-cyclodextrin as a carrier for anti-cancer drug delivery: A molecular dynamics simulation study 

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Table S1: The components of each simulated system in this study

|  | Components <br> Systems | Number of Drug molecule | Number of $\beta C D$ molecule | Number of Water molecule |
| :---: | :---: | :---: | :---: | :---: |
| 1 | $\beta$ CD + 5-Fu + Water | 1 | 1 | 4000 |
| 2 | 5-Fu + Water | 1 | 0 | 4000 |
| 3 | $\beta C D+$ Ald + Water | 1 | 1 | 4000 |
| 4 | Ald + Water | 1 | 0 | 4000 |
| 5 | $\beta$ CD + TMZ + Water | 1 | 1 | 4000 |
| 6 | TMZ + Water | 1 | 0 | 4000 |
| 7 | $\beta C D+$ Water | 0 | 1 | 4000 |

Table S2: Number of water molecules in different spheres inside $\beta$-CD

| System | $\mathbf{0 - 0 . 5}$ | $\mathbf{0 . 5 - 0 . 8}$ | $\mathbf{0 . 8 - 0 . 9}$ | $\mathbf{0 . 9 - 1 . 0}$ | $\mathbf{0 - 1 . 0}$ |
| :---: | :---: | :---: | :---: | :---: | :---: |
| $\beta-\mathrm{CD}+5-\mathrm{FU}+$ water | $3.620( \pm 0.247)$ | $20.269( \pm 0.001)$ | $23.175( \pm 0.021)$ | $47.115( \pm 0.000)$ | $94.179( \pm 0.225)$ |
| $\beta-\mathrm{CD}+\mathrm{Ald}+$ water | $2.476( \pm 0.089)$ | $18.461( \pm 0.005)$ | $23.262( \pm 0.038)$ | $46.870( \pm 0.009)$ | $91.069( \pm 0.047)$ |
| $\beta-\mathrm{CD}+\mathrm{TMZ}+$ water | $1.404( \pm 0.025)$ | $19.131( \pm 0.006)$ | $23.509( \pm 0.002)$ | $46.416( \pm 0.005)$ | $90.460( \pm 0.026)$ |
| $\beta-\mathrm{CD}+$ water (reference $)$ | $9.728( \pm 0.012)$ | $20.481( \pm 0.004)$ | $22.659( \pm 0.004)$ | $47.385( \pm 0.006)$ | $100.253( \pm 0.026)$ |

## Figure S1a

(5-Fu in $\beta$ CD at 20 ns )



## Figure S1a

(5-Fu in $\beta$ CD at 60 ns )



## Figure S1a

(5-Fu in $\beta$ CD at 100 ns )


Figure S1b
(Ald in $\beta$ CD at 20 ns )


Figure S1b
(Ald in $\beta$ CD at 60 ns )


Figure S1b
(Ald in $\beta$ CD at 100 ns )



Figure S1c
(TMZ in $\beta C D$ at 60 ns )


## Figure S1c

(TMZ in $\beta \mathrm{CD}$ at 100 ns )



Figure S2: The Root mean square deviation of $\beta$-CD in different simulated and reference


Figure S3: The Root mean square deviation of drugs in different simulated and reference systems

