

## Supporting Information

### All-stage precisional glioma targeted therapy enabled by a well-designed D-peptide

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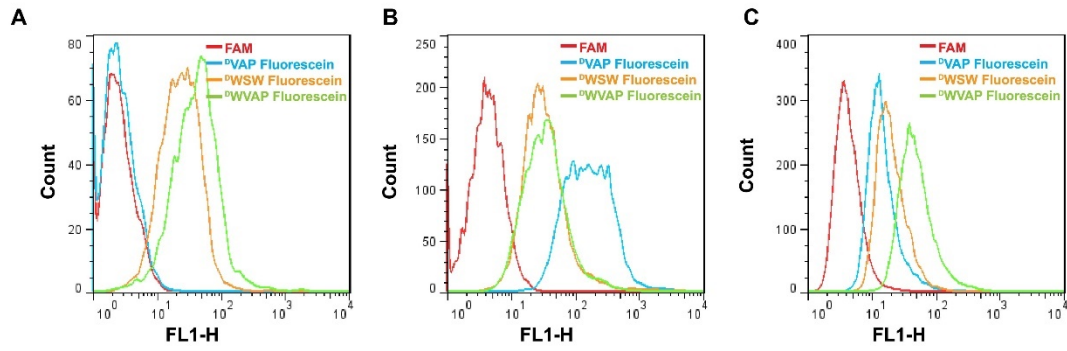
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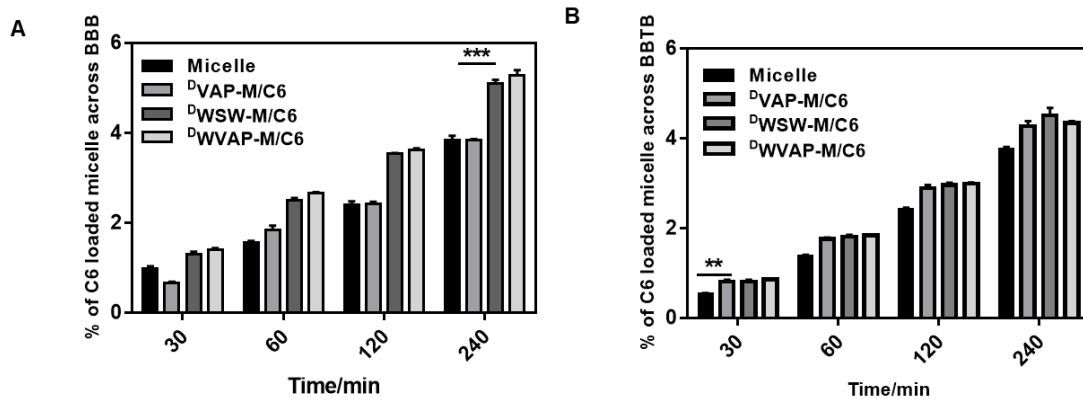
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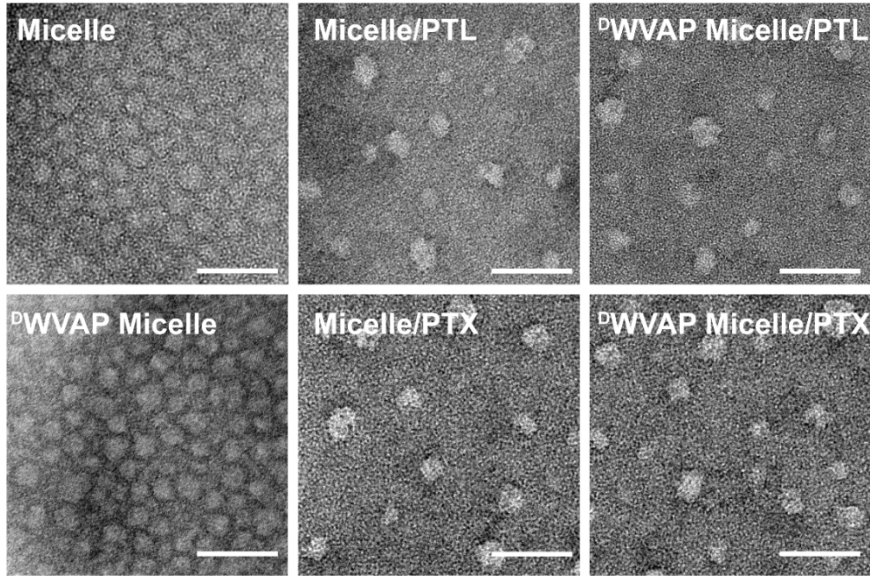
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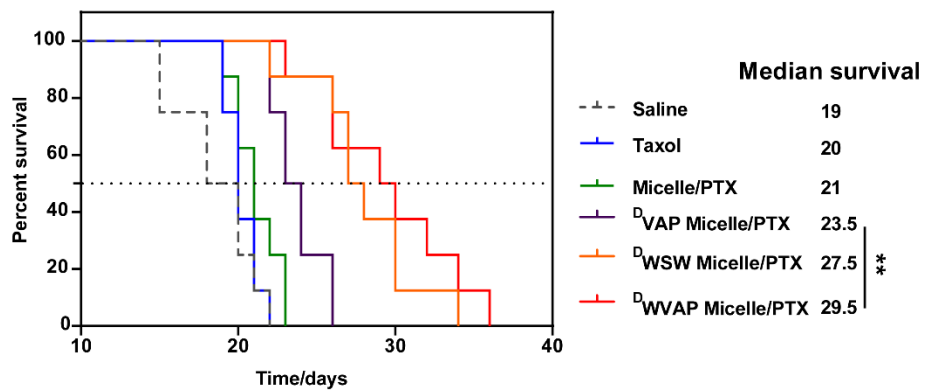
**Figure S1.** Quantitative cellular uptake was measured by flow cytometry in (A) BCEC cells, (B) U87 cells and (C) HUVEC cells.



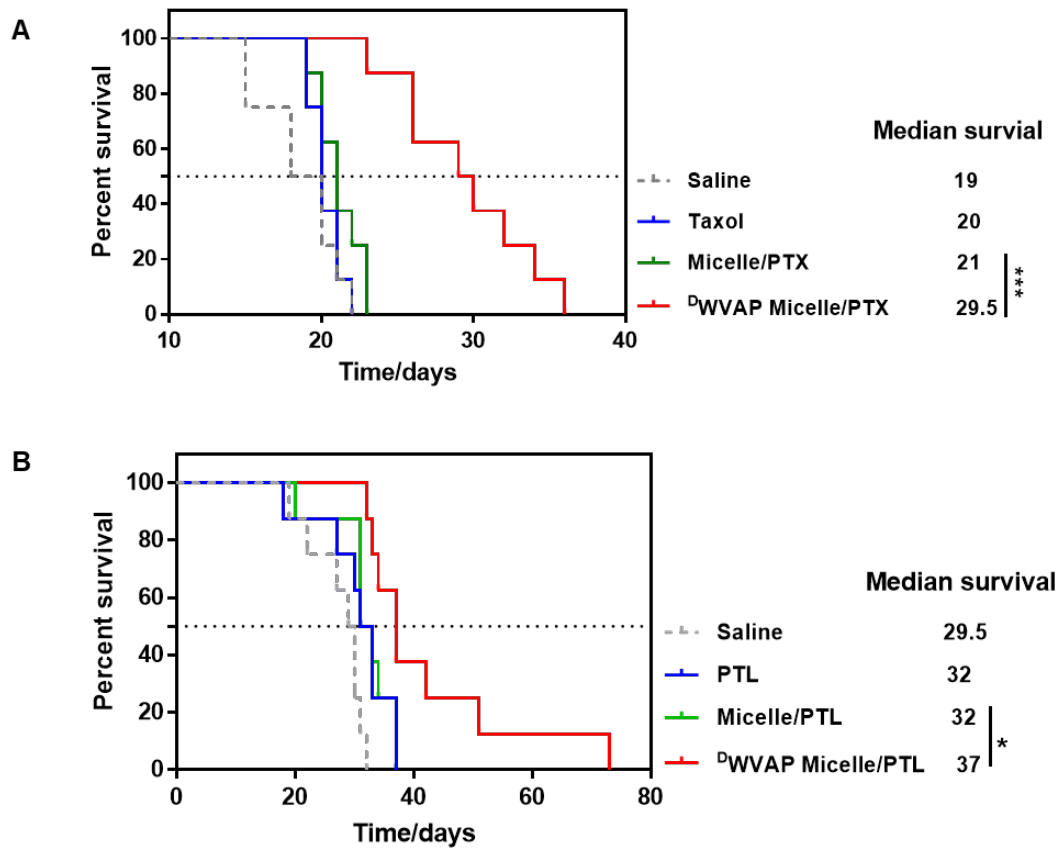
**Figure S2.** Evaluation of the BBB and BBTB penetrating capacity *in vitro* by using BCECs or HUVEC monolayers. (A) Transcytosis efficiency of Coumarin-6-loaded plain Micelles, <sup>125</sup>I-VAP Micelles, <sup>125</sup>I-WSW Micelles and <sup>125</sup>I-WVAP Micelles on *in vitro* BBB model. (B) Transcytosis efficiency of Coumarin-6-loaded plain Micelles, <sup>125</sup>I-VAP Micelles, <sup>125</sup>I-WSW Micelles and <sup>125</sup>I-WVAP Micelles on *in vitro* BBTB model. Data were presented as mean  $\pm$  SD, n = 3, \*\* p < 0.01, \*\*\* p < 0.001.



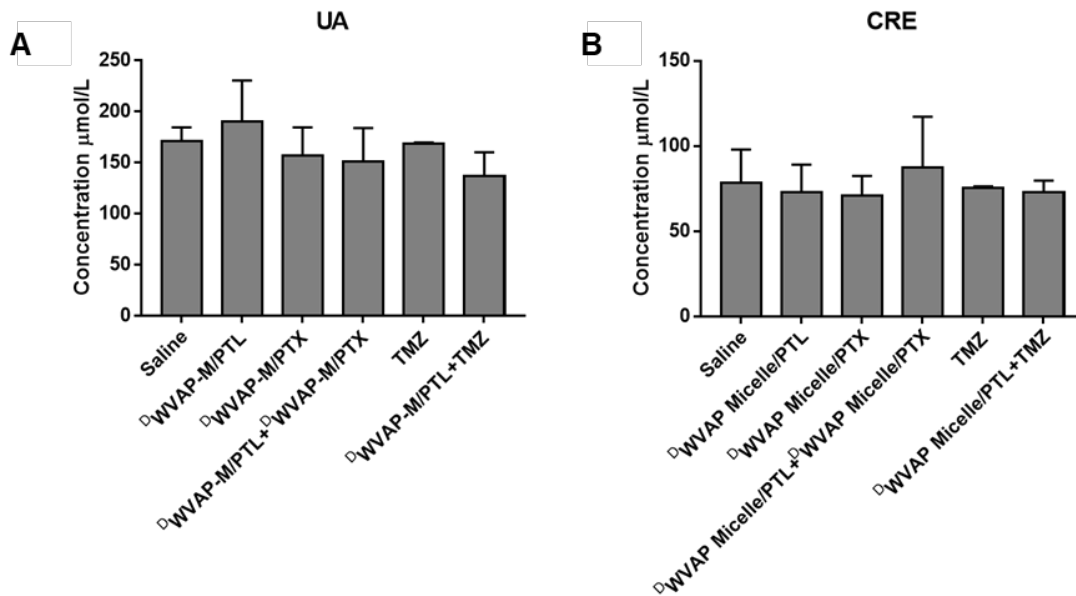
**Figure S3.** TEM images of <sup>D</sup>WVAP peptide modified micelles. (Bar = 50 nm)



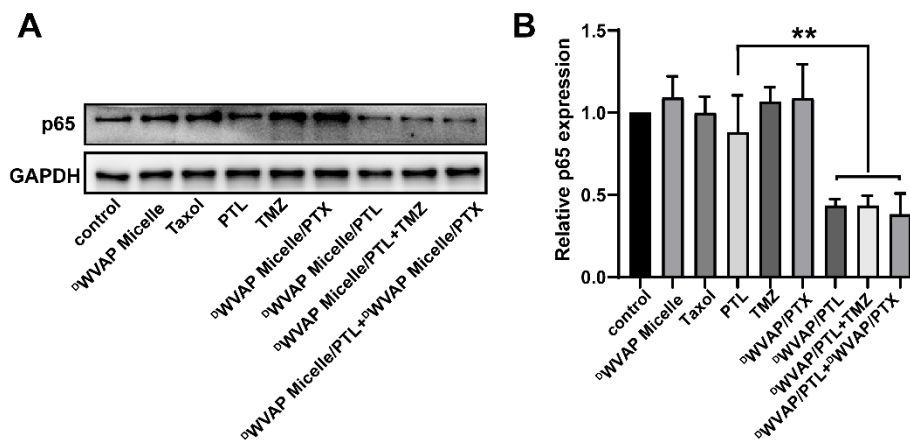
**Figure S4.** Kaplan–Meier survival curves of intracranial glioblastoma-implanted nude mice treated with different regimens. Mice (n = 8) were treated with four injections (6 mg/kg each time) of equal PTX dose (at 6, 9, 12 and 15 days after glioblastoma implantation). \*\*p < 0.01



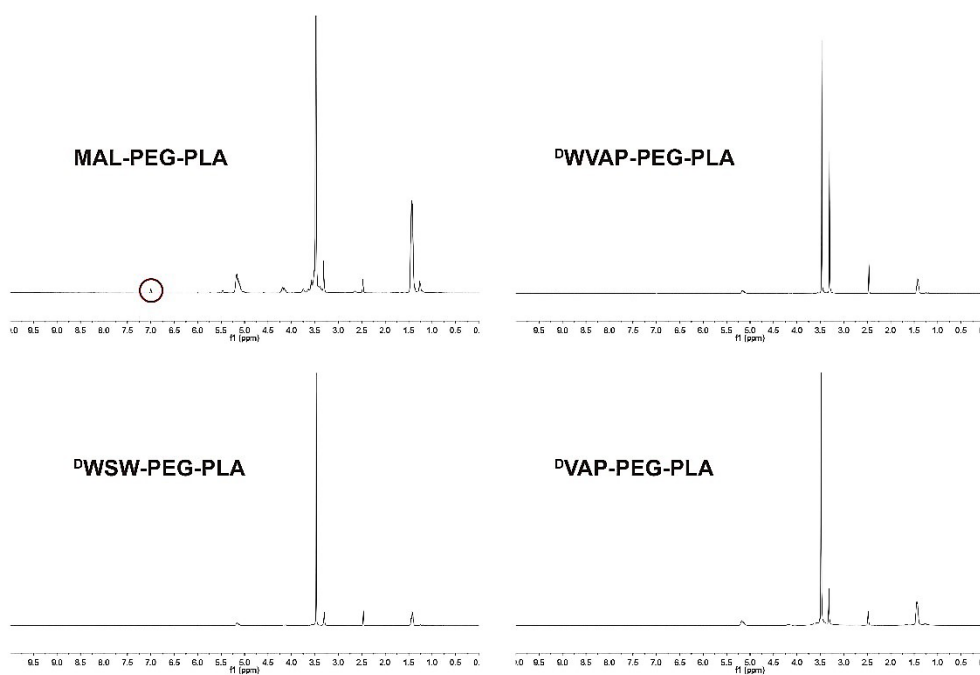
**Figure S5.** Kaplan–Meier survival curves of intracranial glioblastoma-implanted nude mice treated with different regimens. (A) Mice (n = 8) were treated with four injections (6 mg/kg each time) of equal PTX dose (at 6, 9, 12 and 15 days after glioblastoma implantation). (B) Mice (n = 8) were treated with five injections (5 mg/kg each time) of equal PTL dose (at 6, 8, 10, 12 and 14 days after glioblastoma implantation). \* $p < 0.05$ , \*\*\* $p < 0.001$



**Figure S6.** Biochemical analysis of blood samples from various formulation treated groups. UA (Uric acid) and CRE (Creatinine) were used to evaluate the kidney functions (A and B).



**Figure S7.** P65 protein expression in the nucleus of U87 cells analyzed by (A) Western blotting, (B) relative protein expression were quantified using ImageJ software. Results were presented as mean  $\pm$  SD, n= 3, \*\*p < 0.01.



**Figure S8.** The NMR spectra of Mal-PEG<sub>3000</sub>-PLA<sub>2000</sub>, <sup>D</sup>WVAP-PEG<sub>3000</sub>-PLA<sub>2000</sub>, <sup>D</sup>VAP-PEG<sub>3000</sub>-PLA<sub>2000</sub> and <sup>D</sup>WSW-PEG<sub>3000</sub>-PLA<sub>2000</sub>. The Mal group presented a sharp peak around 7 ppm in NMR spectrum of Mal-PEG<sub>3000</sub>-PLA<sub>2000</sub>, which disappeared in that of <sup>D</sup>WVAP-PEG<sub>3000</sub>-PLA<sub>2000</sub>, <sup>D</sup>VAP-PEG<sub>3000</sub>-PLA<sub>2000</sub> and <sup>D</sup>WSW-PEG<sub>3000</sub>-PLA<sub>2000</sub>.

**Table S1.** Characterization of different micelles (n = 3).

Formulation	Size (nm)	Zeta Potential (mV)	Encapsulation efficiency (%)	Drug loading capacity (%)
Micelle/PTX	29.11 ± 2.59	-2.82 ± 1.10	88.36 ± 1.84	20.75 ± 0.76
Micelle/PTL	25.69 ± 1.97	-2.86 ± 1.21	87.36 ± 1.62	8.84 ± 0.51
<sup>D</sup> WVAP Micelle/PTX	29.25 ± 1.87	-0.53 ± 0.92	84.52 ± 1.82	20.65 ± 1.29
<sup>D</sup> WVAP Micelle/PTL	27.60 ± 2.42	-0.55 ± 0.85	83.09 ± 1.80	8.86 ± 0.34