# Supporting Information

## Hetero-modification of TRAIL trimer for improved drug delivery and in vivo antitumor activities

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## **Preparation of TRAIL-vcMMAE and PEG-TRAIL-vcMMAE conjugates.**

Steps:

Number	Reduction	First Step	Second step
		mPEG-MAL	vcMMAE
1	TCEP (10 eq) 1 h, 37 °C	(2 eq/ SH group)	(4 eq / SH group)
		40 min	40 min
2		vcMMAE	mPEG-MAL
		(4 eq / SH group)	(2 eq/ SH group)
		40 min	40 min
3		mPEG-MAL was added 30 seconds ahead of	
		vcMMAE, and incubated together for 40 min at RT	

## Characterization of N109C-vcMMAE

1) Q-TOF MS (mass spectrometry)

TRAIL<sub>114-281</sub>, N109C and N109C-vcMMAE were diluted by dd  $H_2O$  to a final concentration of 1 mg/mL, and centrifuged (12,000 rpm, 15 min). Supernatant was analyzed by Q-TOF MS.

Instrument: Waters UPLC Acquity Bio H Class, Waters Xevo G2-S Q TOF

Column: MassPREP Desalting column

Column temp: 80°C

A: 0.1% Formic Acid in H<sub>2</sub>O

B: 0.1% Formic Acid in ACN

Gra	di	ent	:
oru	ur	one	•

Time(min)	Flow Rate(mL/min)	%A	%B
 0	0.200	90.0	10.0
2.00	0.200	90.0	10.0
13.00	0.200	20.0	80.0
13.50	0.200	10.0	90.0
15.50	0.200	10.0	90.0
16.00	0.200	90.0	10.0
18.00	0.200	90.0	10.0

MS parameters:

Capillary (kV): 2.5 Sampling Cone (V): 60 Source Temperature (°C): 100 Desolvation Temperature (°C): 500 Desolvation Gas Flow (L/Hr): 800.0 Mass range(m/z): 400 to 4,000

Results:

Subject	Expected MW (Da)	Calculated MW (Da)
TRAIL <sub>114-281</sub> monomer	19491.85	19497.97
N109C monomer	21652.23	21657.97
N109C dimer	43302.46	43316.8
N109C-vcMMAE monomer	22966.86	22969.8

Due to the high column temperature, all the trimeric TRAIL and its conjugates were depolymerized in monomers (Figure S1). And because PEG is mixture of polymers with different molecular weight (MW), so we did not determine its MW through Q-TOF MS. These results indicated that all the proteins were expressed in *E.coli* correctly, and TRAIL mutant N109C was successfully conjugated with MMAE.



Figure S1. Q-TOF MS results of TRAIL<sub>114-281</sub>, N109C and N109C-vcMMAE

#### 1) Size-exclusion chromatography (SEC)

A silica-based G3000SW<sub>XL</sub> size exclusion column (7.8 mm×30 cm dimension, 5  $\mu$ m particle size, 250 Å pore size) (Tosoh Bioscience LLC, Montgomeryville, PA, USA) was used to determine the polymerization of N109C and its conjugates. The separation was performed using size exclusion column with mobile phase (50 mM NaH<sub>2</sub>PO<sub>4</sub> (pH 6.7), 300 mM NaCl, 0.05% NaN<sub>3</sub>) at flow rate of 0.6 mL/min. The injection volume was 20  $\mu$ L.



**Figure S2**. Size-exclusion chromatography analysis of the tertiary structure of N109C and its conjugates.

### In vitro antitumor activity of PEG-TRAIL-vcMMAE conjugates

NCI-H460 is human large cell lung cancer cell line (TRAIL sensitive); K562 is human erythromyeloblastoid leukemia cell line (TRAIL resistant); MCF-7 is human breast adenocarcinoma cell line (TRAIL resistant). All tumor cells were seeded at a density of  $3 \times 10^3$  cells/100 µL/well in 96-well microtiter plates (Costar, Corning). Twenty-four hours post-seeding, the cultures were washed twice with PBS and then exposed to various concentrations (1.6~250 nM) of TRAIL<sub>114-281</sub>, N109C-vcMMAE and PEG-N109C-vcMMAE for 96 h. The PBS (pH 7.4) was used as negative control. Cell viability was determined by Cell Counting Kit-8 (Dojindo, Osaka, Japan). The absorbance of 450 nm was measured by BioRad Model 680 Microplate Reader. The results were expressed as mean ±standard deviation (S.D.) (n =3).



Figure S3. In vitro antitumor activities of TRAIL and its conjugates on differents tumor cell lines.

#### In vivo antitumor activities of TRAIL and its conjugates

Balb/c athymic nude mice (Slaccas Laboratory Animal Co., Ltd., Shanghai, China) were used for in vivo antitumor activities of TRAIL and its conjugates. Fourty-four athymic nude mice (female, 6-8 weeks) were divided into 7 groups [saline (6 mice), MMAE (390 µg/kg) (4 mice), BSAvcMMAE (10 mg/kg, 390 µg/kg MMAE) (3 mice), TRAIL<sub>114-281</sub> 10 mg/kg (4 mice), PEG-N109CvcMMAE (4mg/kg, 65 µg/kg MMAE) (5 mice), PEG-N109C-vcMMAE (24mg/kg, 390 µg/kg MMAE) (6 mice), N109C-vcMMAE (10 mg/kg, 390 µg/kg MMAE) (6 mice)]. To build the tumorbearing mice model, freshly harvested NCI-H460 cells ( $5 \times 10^6$  cells per mouse) were inoculated subcutaneously in the right flank of athymic nude mice. When tumor volume reached 50 mm<sup>3</sup>, mice were treated with above samples once every two days for four times  $(q2d\times4)$  intravenously. Tumor volumes were continuously monitored until the end of the experiment and calculated by the formula  $V = (L^*W^2)/2$ , L and W refer to longitudinal and transverse tumor diameters respectively. The body weight of mice was also monitored once every two days (Figure S4). In vivo cell apoptosis was evaluated by terminal deoxynucleotidyl transferase-mediated dUTP nick end labeling (TUNEL) assays (In Situ Cell Death Detection kit, Roche). The tumor tissues for TUNEL assay were obtained from 6 groups (except BSA-vcMMAE group) above randomly (one mouse per group), 24 h after the fourth administration of samples. After fixed with 10% neutral-buffered formalin and paraffin-embedded, sections (5 µm) were cut from tumor tissues, developed by TUNEL reagents after route dewaxing and washing steps and imaged by a light microscope(**Figure S5**). Lung, Liver and kidney tissues were also obtained from the same mice and immediately fixed by 10% neutral-buffered formalin for the following toxicity evaluation. Acute liver or kidney toxicity was studied through H&E (Hematoxylin and Eosin) staining for the morphological examination of tissue cells. In brief, the lung, liver and kidney tissues obtained were fixed, paraffin-embedded and stained by H&E according to above-mentioned steps. The sections were examined by light microscopes at the magnification  $400 \times$  (lung and kidney) or  $100 \times$  (liver).



Figure S4. Body weight monitoring of mice treated with different sampls.



Figure S5. TUNEL assays for the in situ cell death within tumor tissues. Magnification 100×.