



Supporting Information

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Multi-Arm PEG/Peptidomimetic Conjugate Inhibitors of DR6/APP

Interaction Block Hematogenous Tumor Cell Extravasation

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Table S1. Amino acid sequences of twenty-four peptides and the corresponding K_D values. ^a

ID	Sequence	$K_D \pm SEM^b$ (nM)
AHP-01	WTLSINS	1153.7 ± 7.94
AHP-02	FQFRLQS	NB ^c
AHP-03	YLPAPLP	688.5 ± 11.20
AHP-04	ANLLYES	NB
AHP-05	YYSGNNL	1096 ± 10.04
AHP-06	KLGPVCP	NB
AHP-07	YSLHASP	NB
AHP-08	HSTAPQL	NB
AHP-09	KLVPEPS	NB
AHP-10	NGSTHER	527.8 ± 12.09
AHP-11	GFSHYFP	NB
AHP-12	TIDATTP	321.8 ± 11.24
AHP-13	SHTDTNQ	1153.7 ± 7.94
AHP-14	AFMYSLP	457.3 ± 11.56
AHP-15	TIPKYER	NB
AHP-16	KFLQVTN	NB
AHP-17	HSYPYTQ	NB
AHP-18	TTYMPKM	NB
AHP-19	MPPFRHD	NB
AHP-20	LPRSPLA	NB
AHP-21	AGHFSLW	NB
AHP-22	LMTTKSD	NB
AHP-23	GGLSMTR	4458.8 ± 5.85
AHP-24	NHQRLLP	1093.5 ± 262

^a K_D values of peptides to DR6 were assessed as described in the Materials and Methods.

^b K_D values shown are the average of three independent experiments in triplicate.

^c NB = No binding.

Table S2. Amino acid sequences of D-peptides and DRI-peptides and the corresponding K_D values.

ID	Sequence	$K_D \pm \text{SEM}$ (nM)
AHP-D-03	D-(YLPAPLP)	4734.0 ± 8.80
AHP-D-10	D-(NGSTHER)	27110.0 ± 47383
AHP-D-12	D-(TIDATTP)	55130.0 ± 27585
AHP-D-13	D-(SHTDTNQ)	3174 ± 8.80
AHP-DRI-03	D-(PLPAPLY)	1219 ± 7.02
AHP-DRI-10	D-(REHTSGN)	17810 ± 37081
AHP-DRI-12	D-(PTTADIT)	225.19 ± 4.30
AHP-DRI-13	D-(QNTDTHS)	862.7 ± 6.59

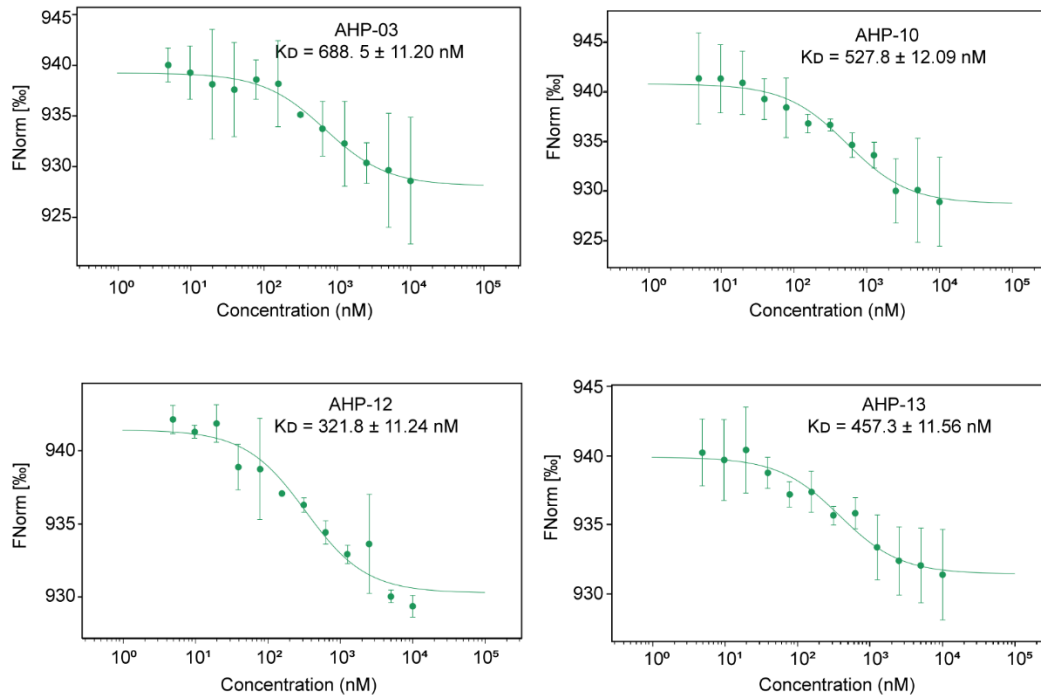
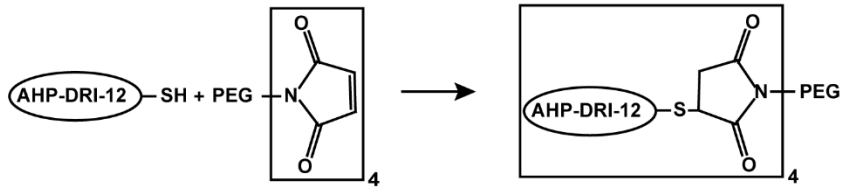
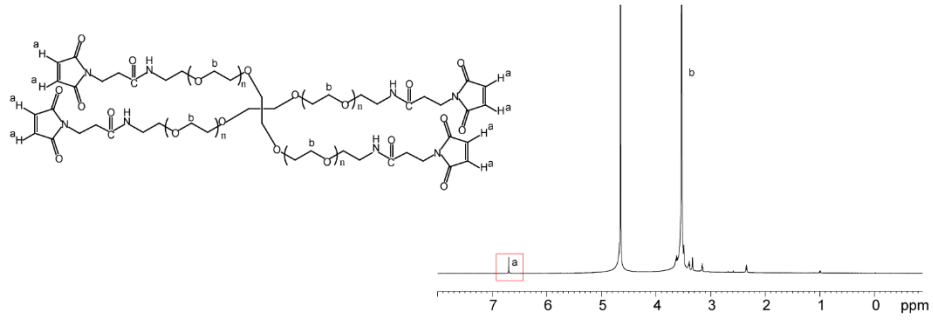


Figure S1. The binding curve of AHP-03, AHP-10, AHP-12 and AHP-13 to DR6 protein were measured by MST, respectively.

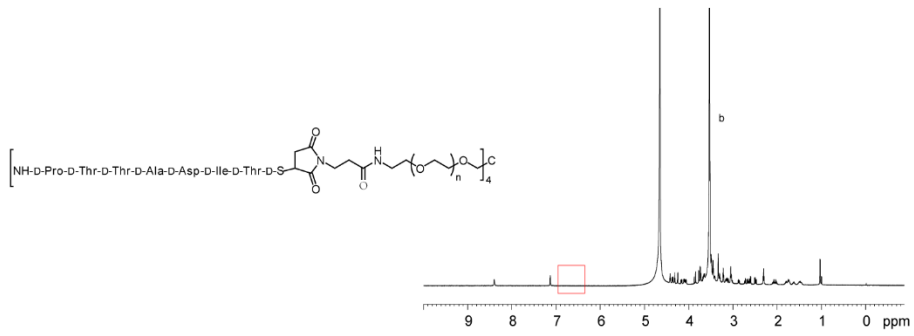
A



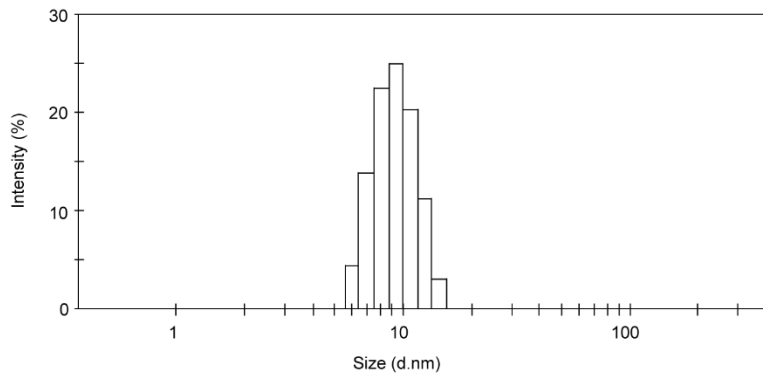
B



C



D



E

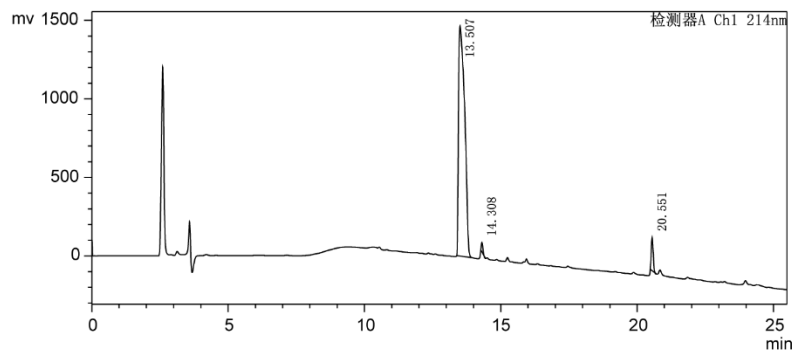
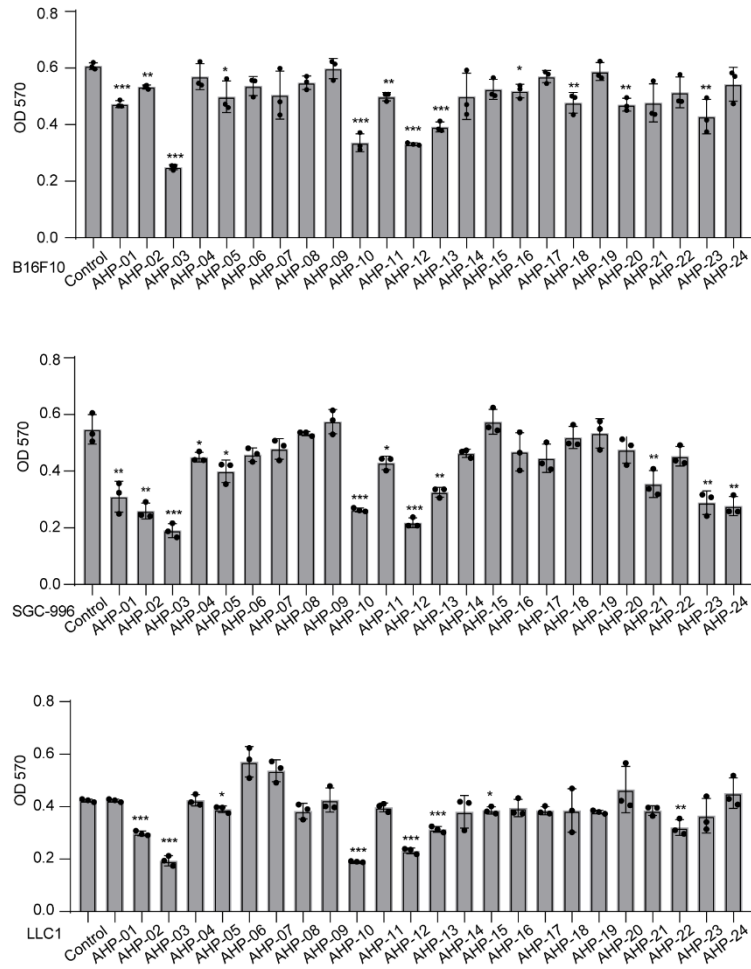
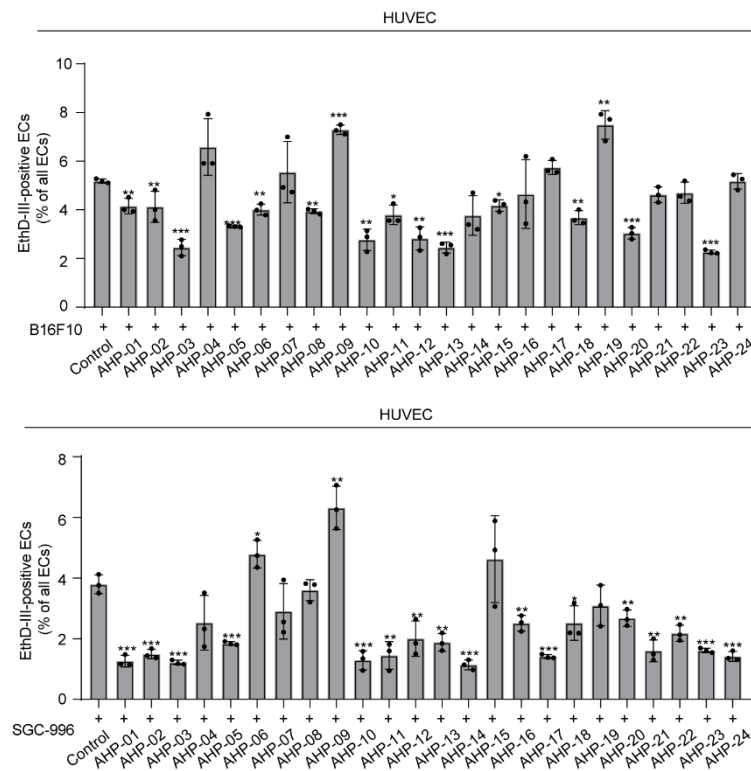


Figure S2. Synthesis of PEG-tAHP-DRI. **A)** Schematic presentation of PEG-tAHP-DRI. **B)** ^1H NMR spectra of 4-arm-PEG_{5k} in D₂O. **C)** ^1H NMR spectra of PEG-tAHP-DRI in D₂O. **D)** Size distribution of PEG-tAHP-DRI. **E)** HPLC curve of PEG-tAHP-DRI.

A



B



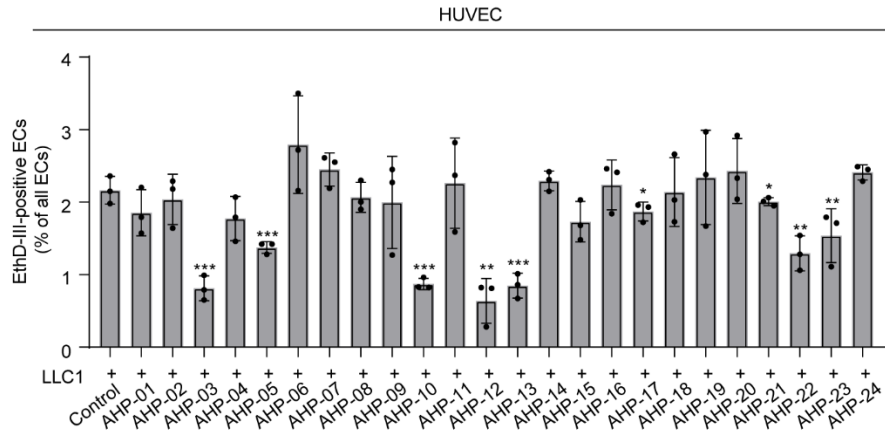
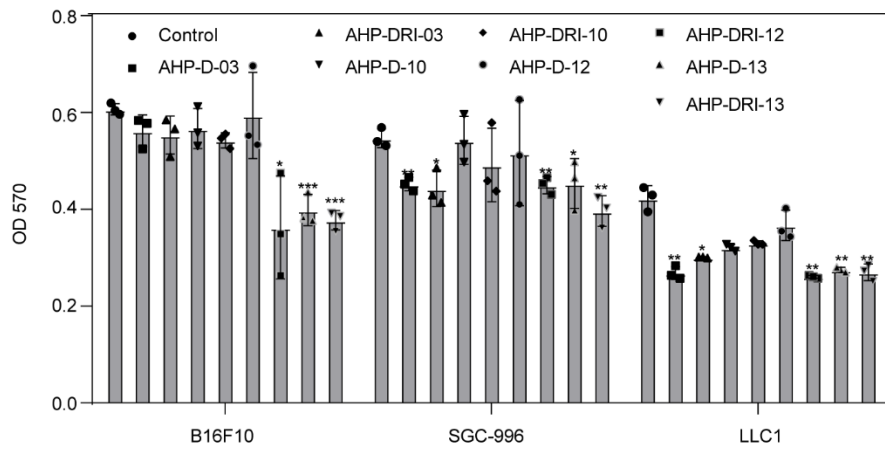
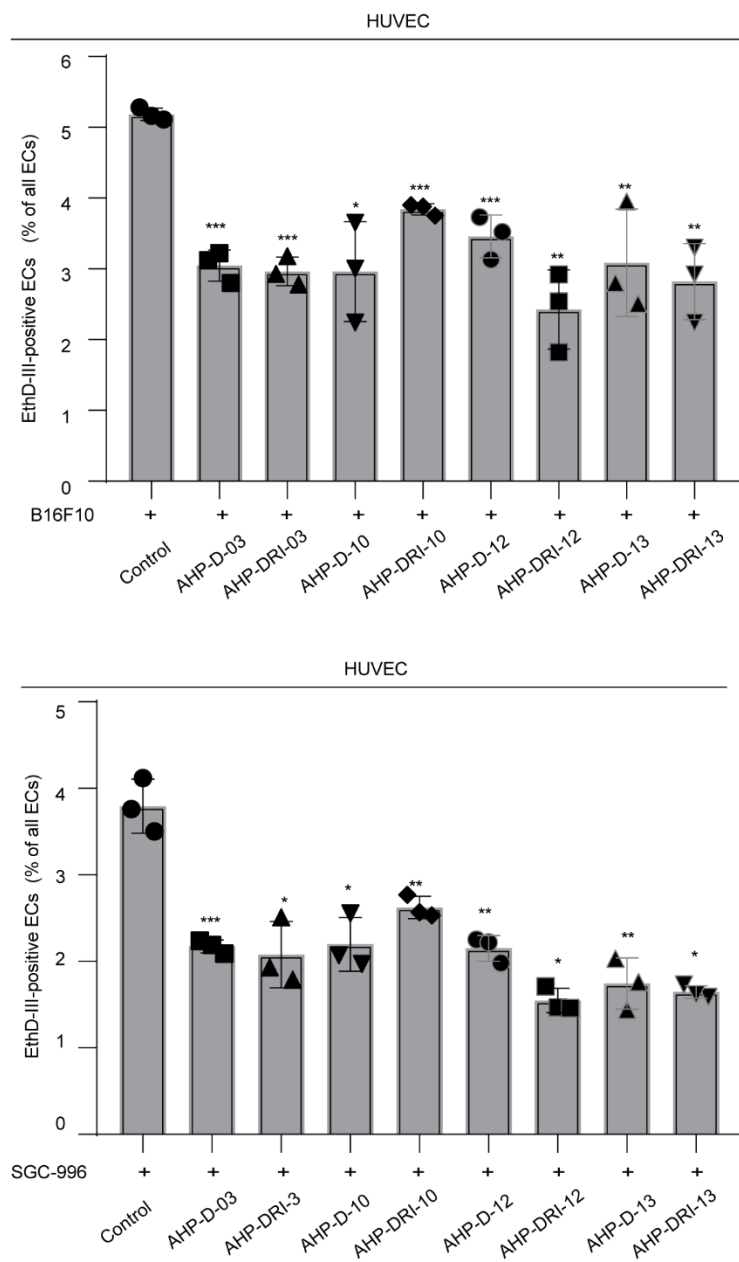


Figure S3. Screening of twenty-four DR6-targeting peptides. A) Cell attachment of different TCs to culture plate coated with DR6 protein after treated twenty-four peptides. **B)** HUVECs necroptosis when co-cultured with different TCs after treated with twenty-four peptides. Data are means \pm SD (n = 3), and analyzed with GraphPad Prism 8.0. *P < 0.05, **P < 0.01, ***P < 0.005 calculated by unpaired two-tailed Student's t-test. NS indicates P > 0.05.

A



B



C

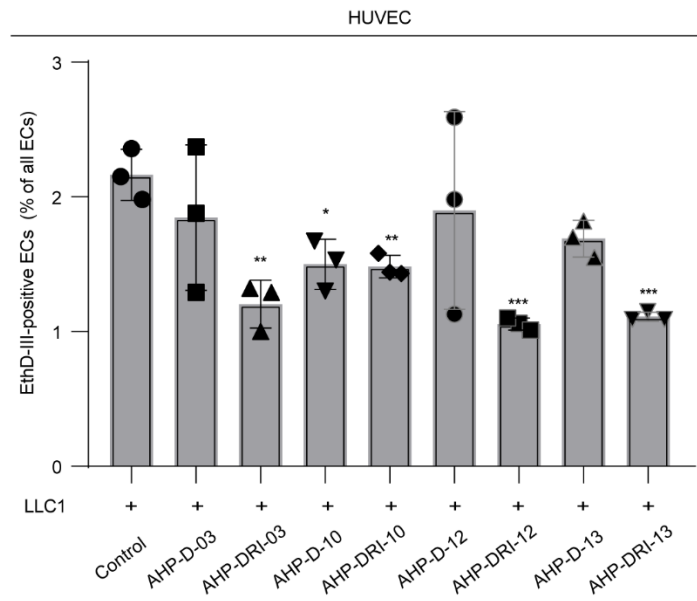


Figure S4. Screening of D-peptides and DRI-peptides. **A)** Cell attachment of different TCs to culture plate coated with DR6 protein after treated with eight peptides. **B)** HUVECs necroptosis when co-cultured with different TCs after treated with eight peptides. Data are means \pm SD ($n = 3$), and analyzed with GraphPad Prism 8.0. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.005$ calculated by unpaired two-tailed Student's t-test. NS indicates $P > 0.05$.

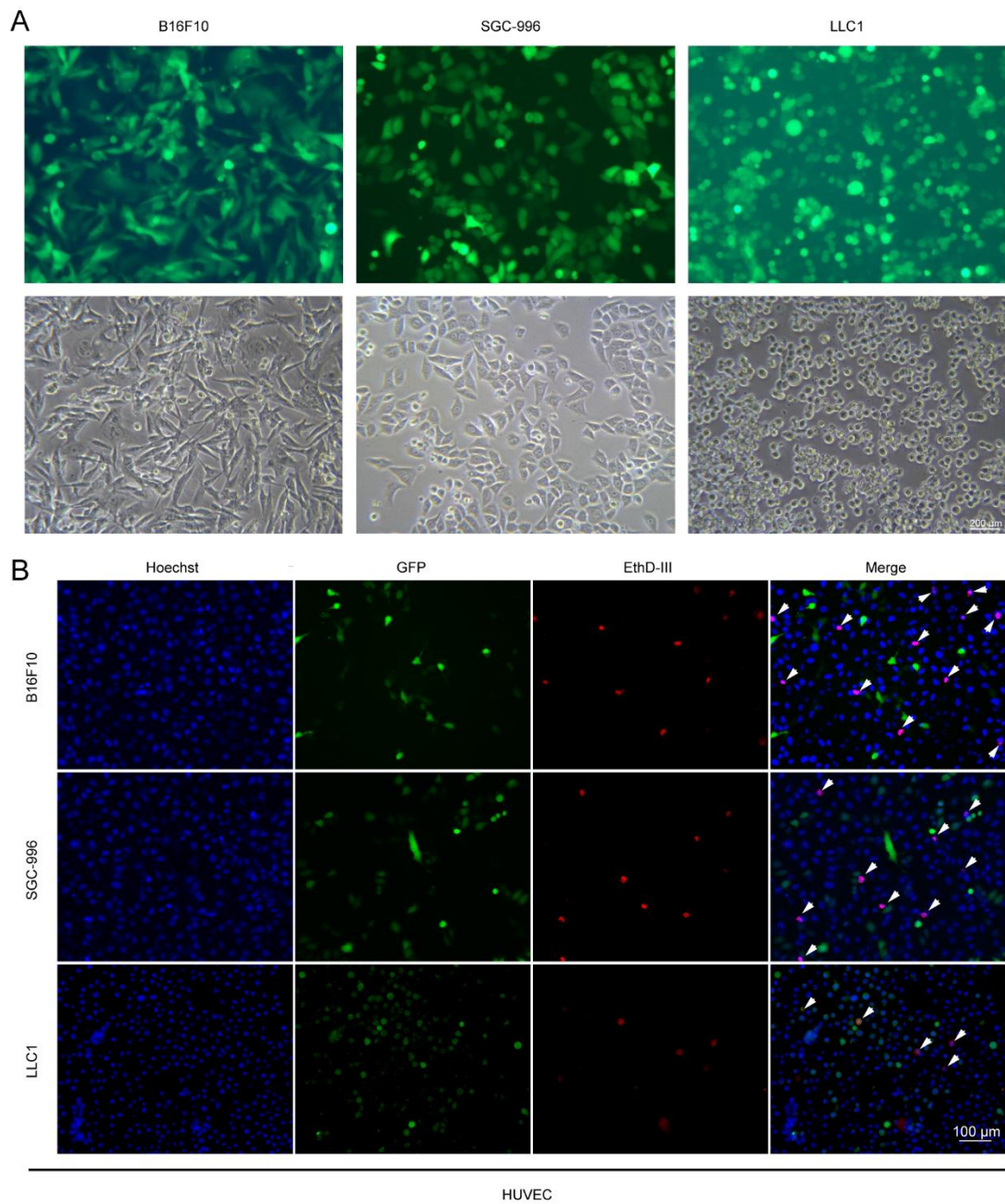


Figure S5. A) B16F10, SGC-996, LLC1 cells stably expressed GFP. **B)** Fluorescent images of HUVECs cultured in the presence of GFP-TCs (green) and stained as indicated. Arrowheads, EthD-III-positive HUVECs (red).

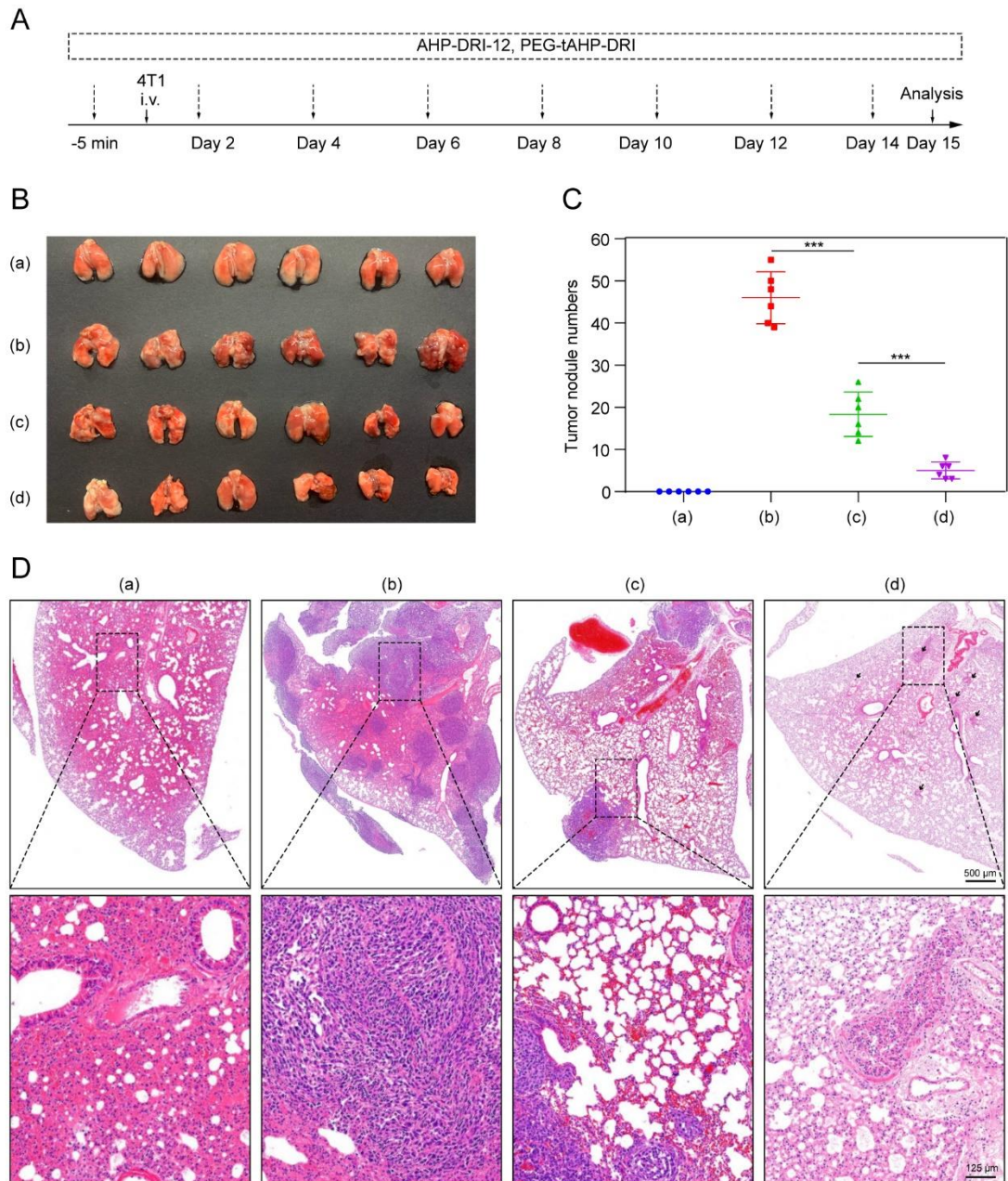


Figure S6. **A**) Experimental design. **B**) Photographs of the lungs 15 days after i.v. injection of 4T1. (a) saline, (b) 4T1 + saline (vehicle), (c) 4T1 + AHP-DRI-12, (d) 4T1 + PEG-tAHP-DRI. **C**) Metastasis nodules of lungs were quantified. **D**) H&E staining of lung sections. Data are means \pm SD ($n = 6$), and analyzed with GraphPad Prism 8.0. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.005$ calculated by unpaired two-tailed Student's t-test. NS indicates $P > 0.05$.

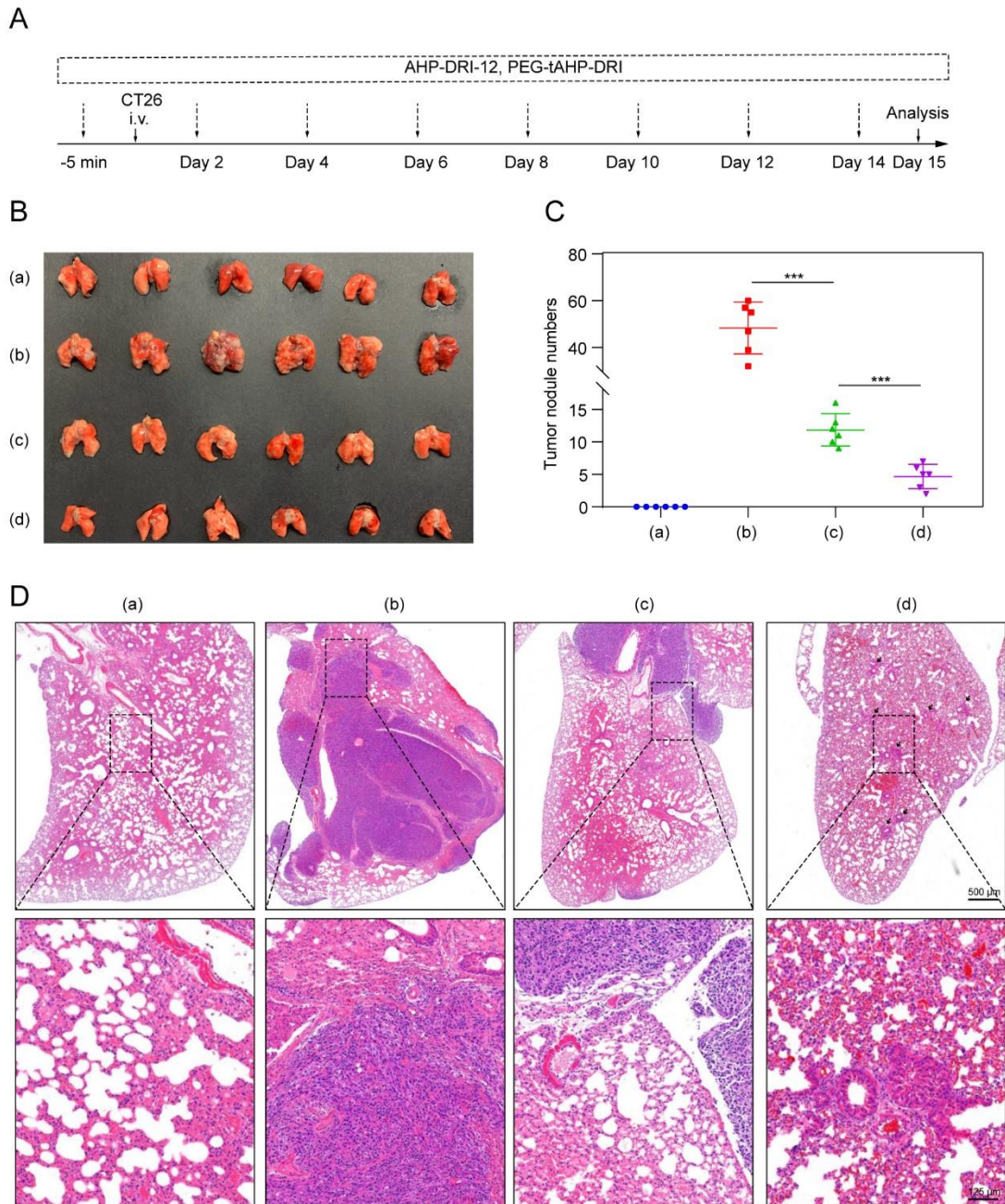


Figure S7. **A**) Experimental design. **B**) Photographs of the lungs 15 days after i.v. injection of CT26. (a) saline, (b) CT26 + saline (vehicle), (c) CT26 + AHP-DRI-12, (d) CT26 + PEG-tAHP-DRI. **C**) Metastasis nodules of lungs were quantified. **D**) H&E staining of lung sections. Data are means \pm SD ($n = 6$), and analyzed with GraphPad Prism 8.0. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.005$ calculated by unpaired two-tailed Student's t-test. NS indicates $P > 0.05$.

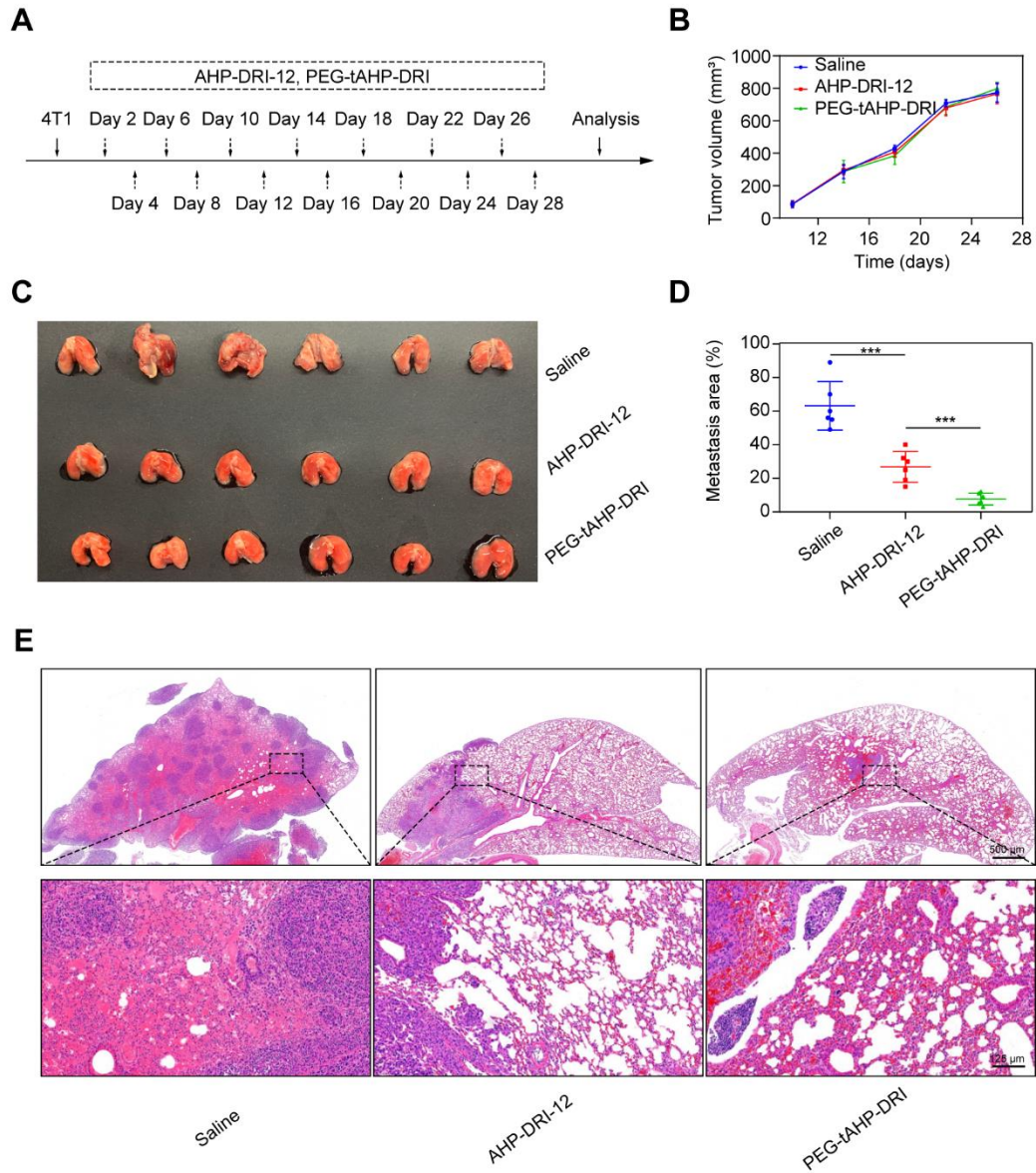
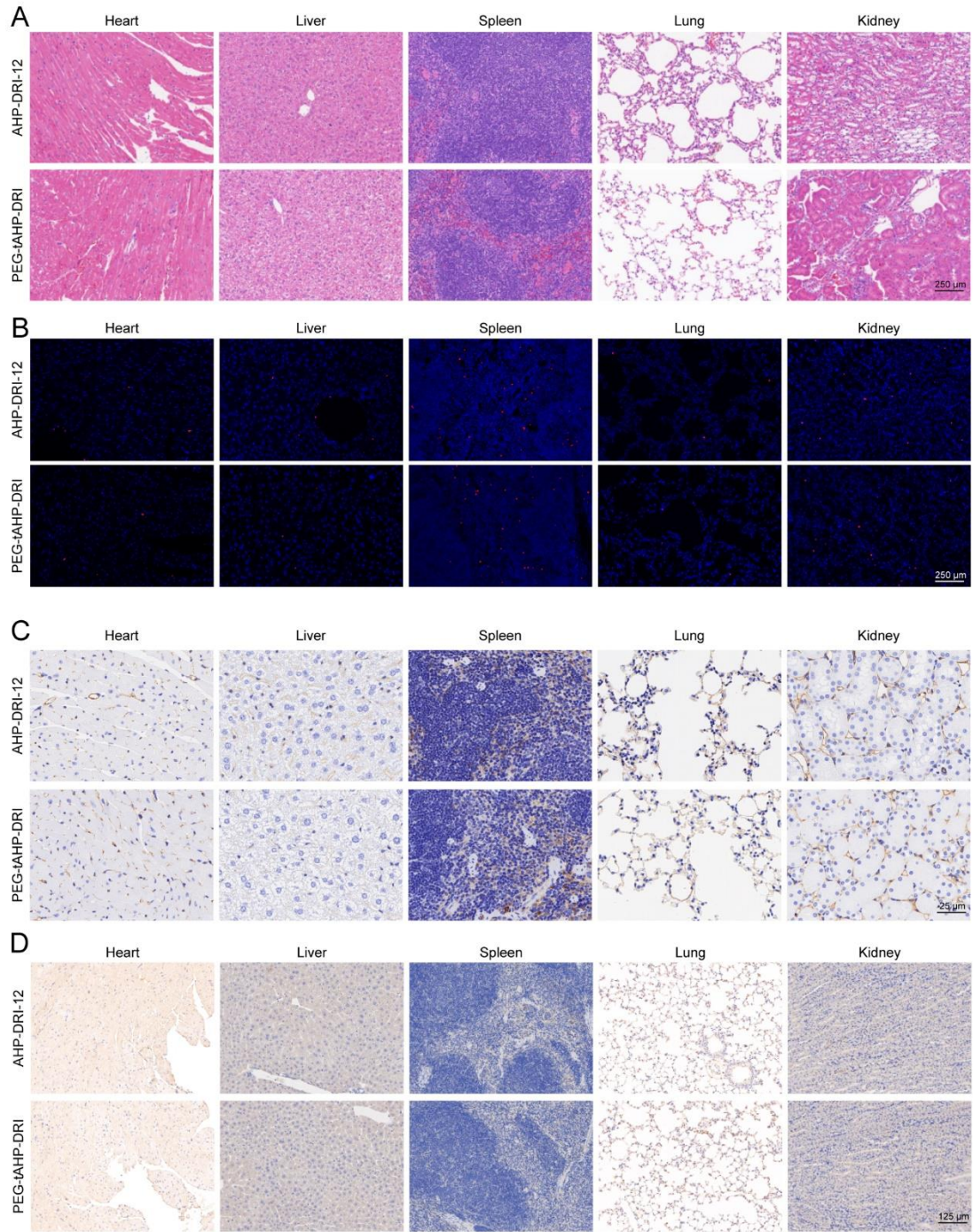


Figure S8. **A)** Experimental design. **B)** Orthotopic tumor growth of 4T1. **C)** Photographs of the lungs 30 days after 4T1 injection. **D)** Metastasis areas of lungs were quantified. **E)** H&E staining of lung sections. Data are means \pm SD ($n = 6$), and analyzed with GraphPad Prism 8.0. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.005$ calculated by unpaired two-tailed Student's t -test. NS indicates $P > 0.05$.



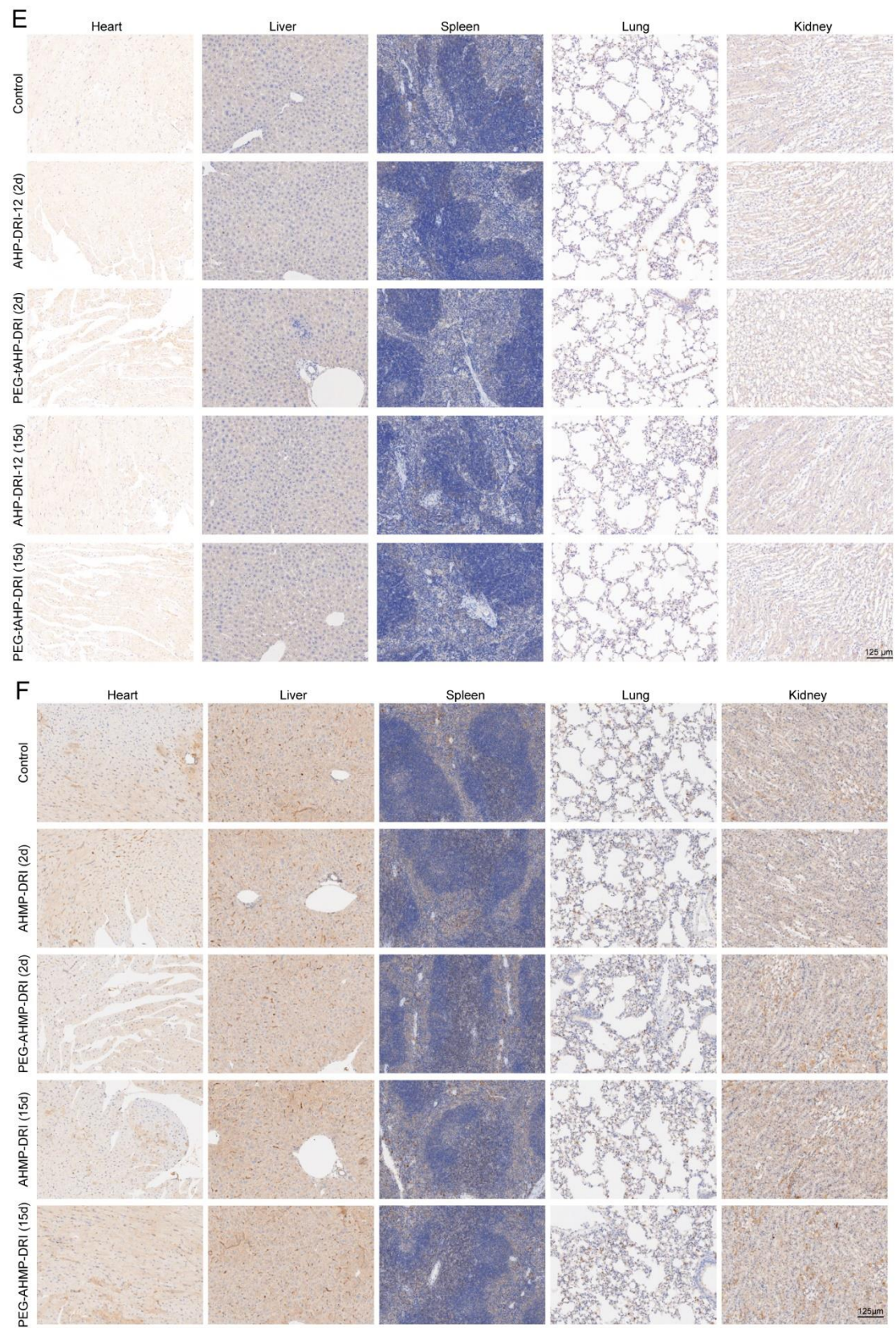


Figure S9. A, B) Tissue sections (heart, liver, spleen, lung and kidney) of healthy mice 2 days after injected with AHP-DRI-12 or PEG-tAHP-DRI were stained with H&E (**A**) and TUNEL (**B**). **C, D)** Tissue

sections (heart, liver, spleen, lung and kidney) of healthy mice 2 days after injected with AHP-DRI-12 or PEG-tAHP-DRI were stained with CD31 (C) and p-RIPK1 (D). E, F) Tissue sections (heart, liver, spleen, lung and kidney) of healthy mice 2 days and 15 days after injected with AHP-DRI-12 or PEG-tAHP-DRI were stained with p-IkB- α (E) and caspase 8 (F).

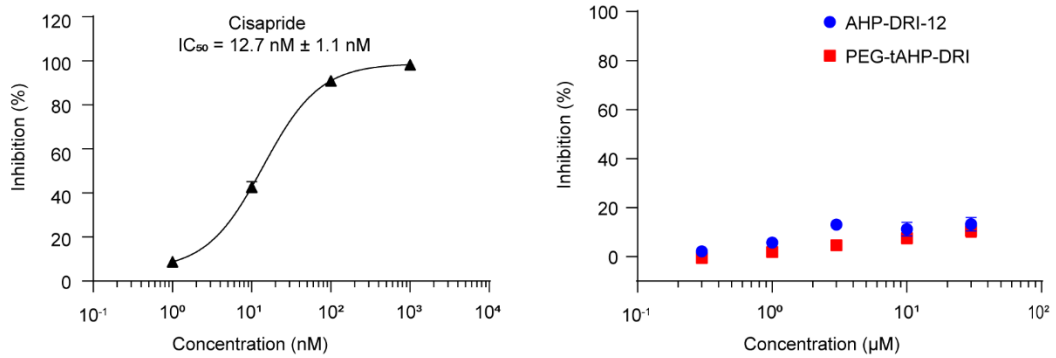


Figure S10. Current change of hERG induced by cisapride, AHP-DRI-12 or PEG-tAHP-DRI.

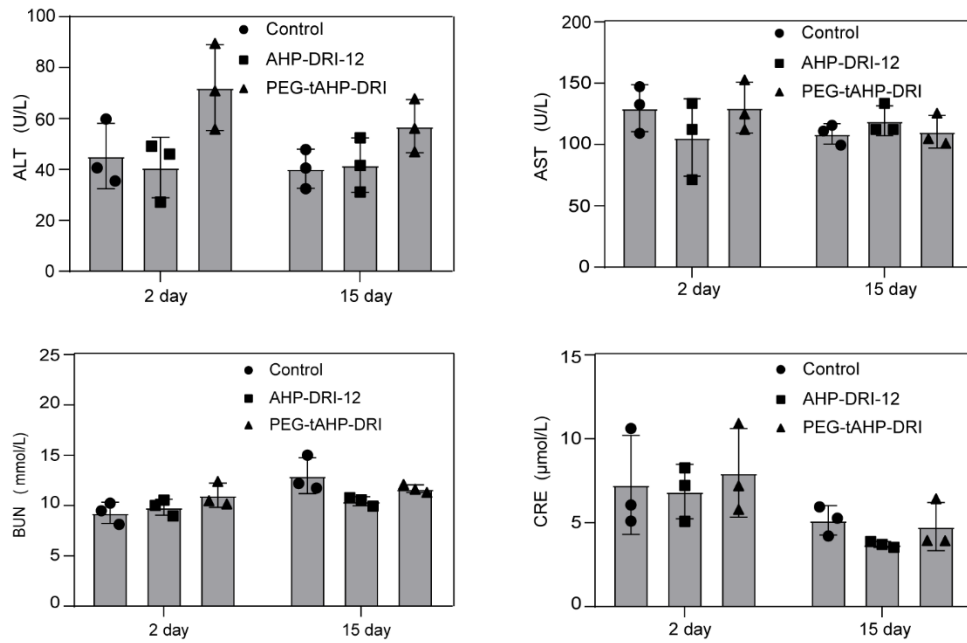
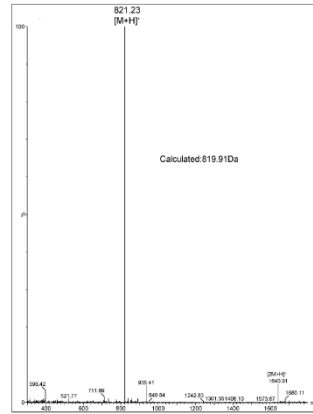
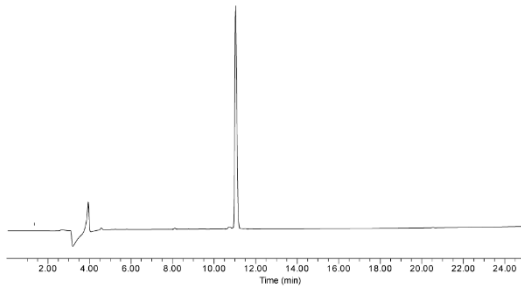
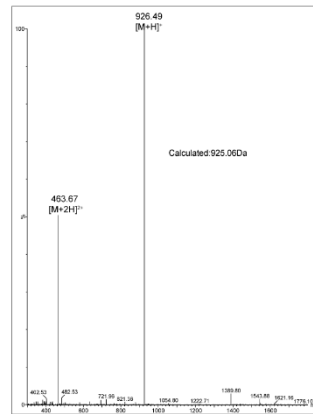
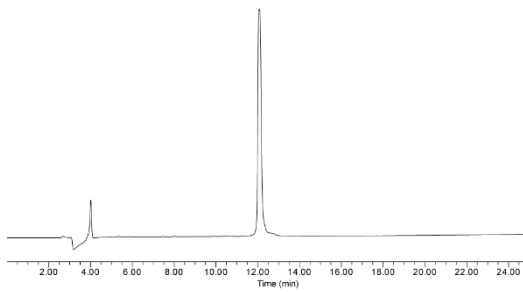


Figure S11. Serum ALT, AST, BUN and CRE levels after i.v. injection of AHP-DRI-12 and PEG-tAHP-DRI. Data are means \pm SD (n = 3), and analyzed with GraphPad Prism 8.0. *P < 0.05, **P < 0.01, ***P < 0.005 calculated by unpaired two-tailed Student's t-test. NS indicates P > 0.05.

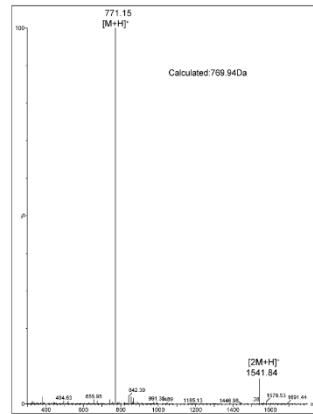
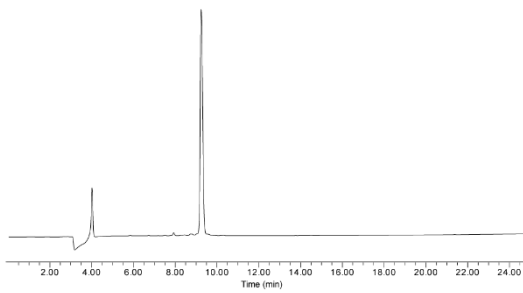
AHP-01: WTLSINS



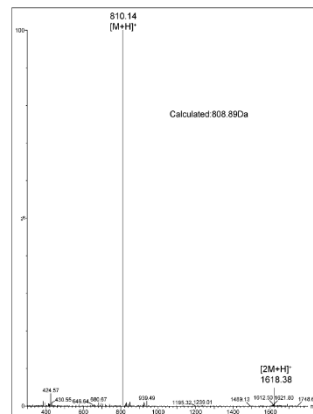
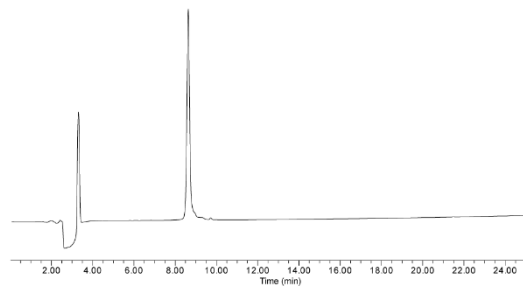
AHP-02: FQFRLQS



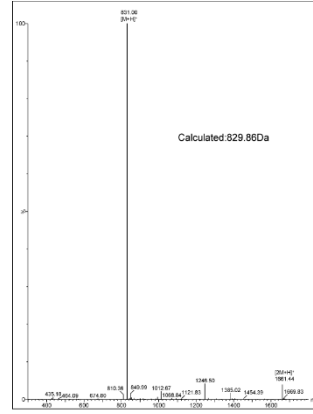
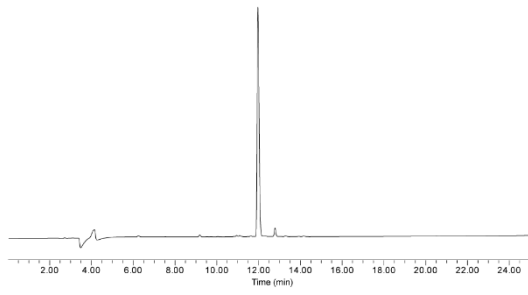
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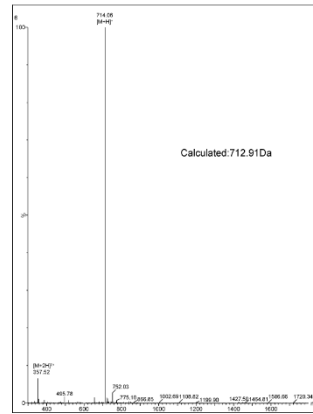
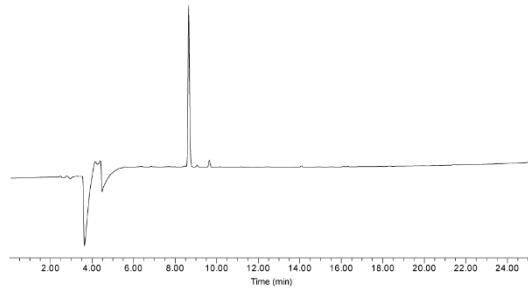
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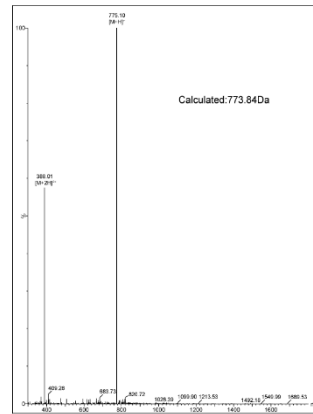
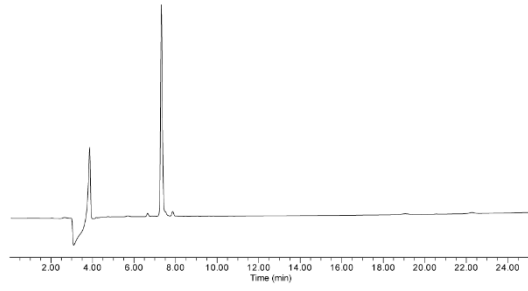
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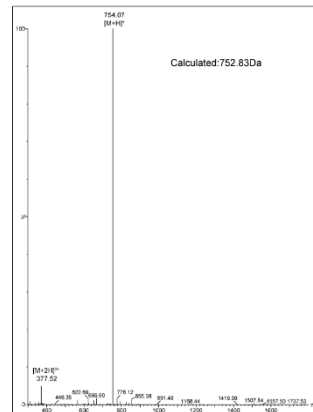
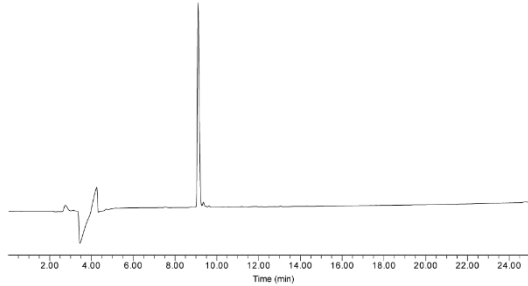
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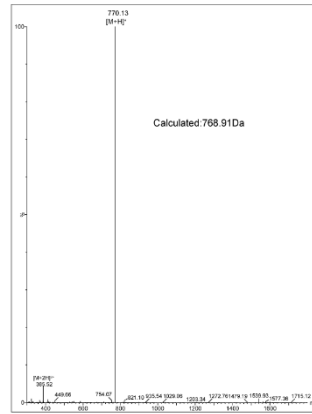
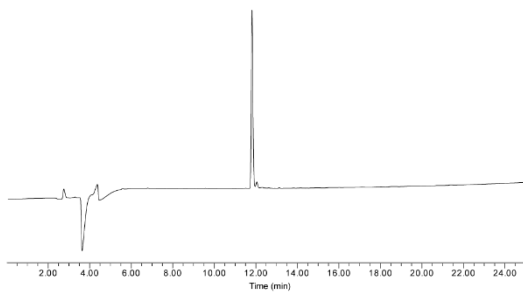
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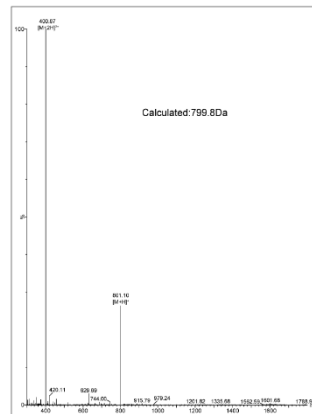
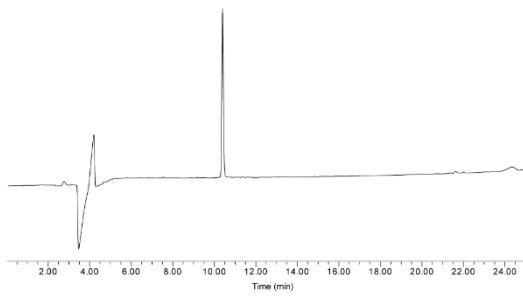
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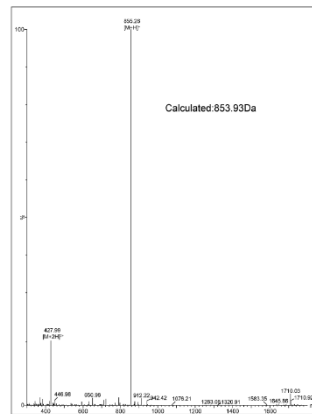
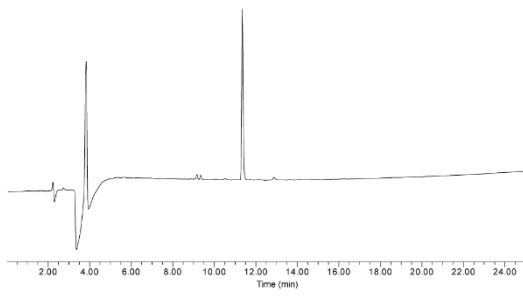
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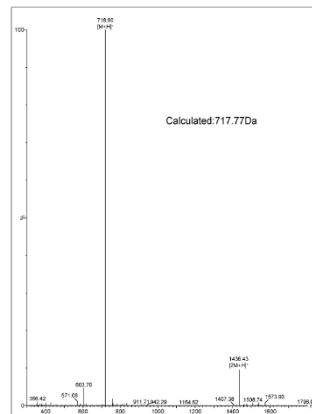
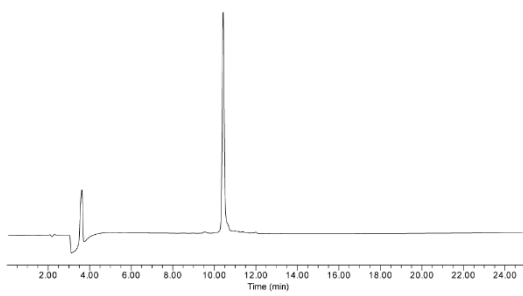
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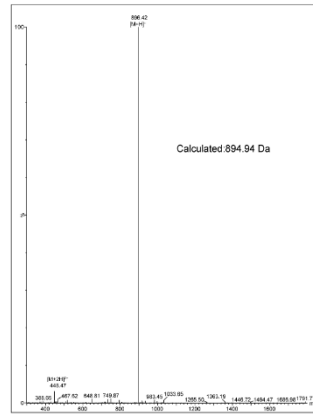
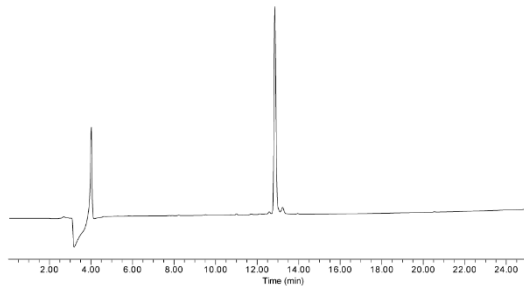
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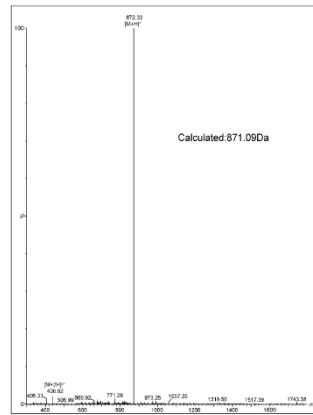
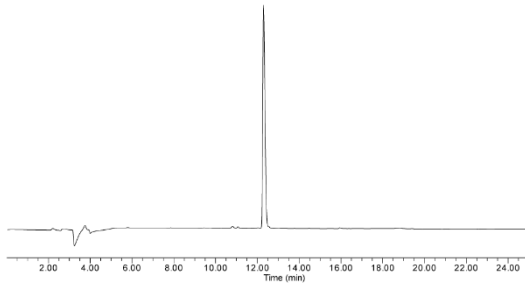
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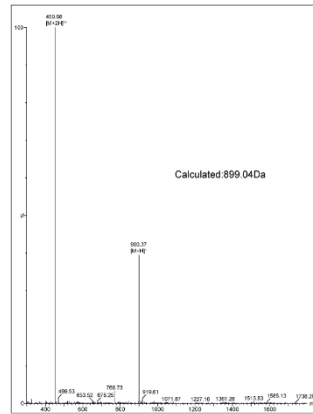
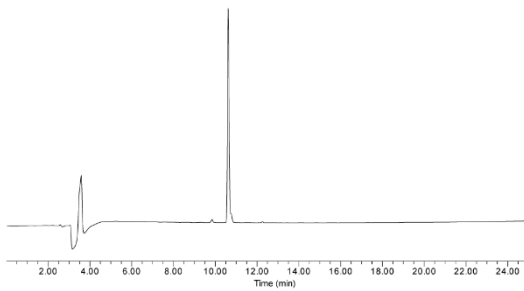
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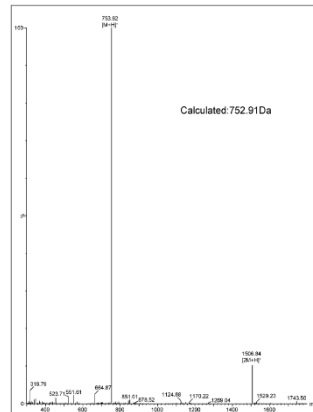
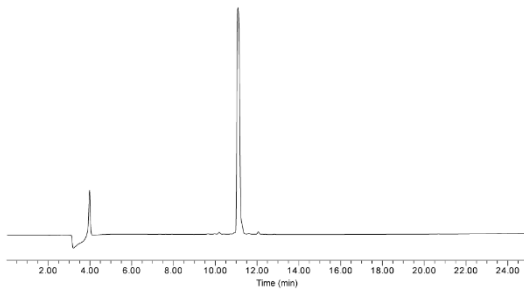
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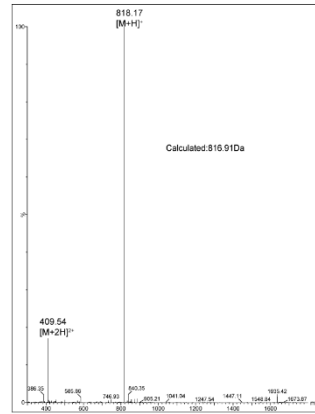
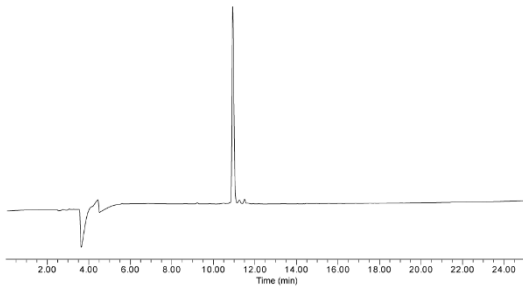
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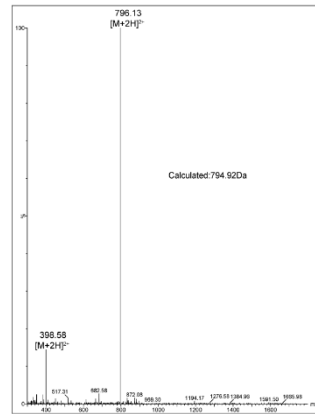
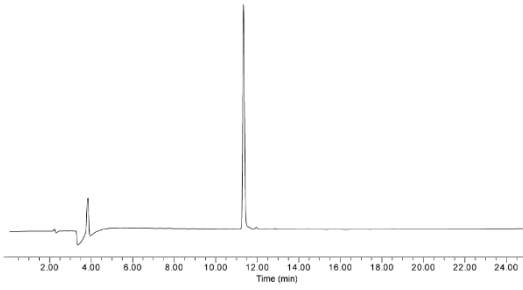
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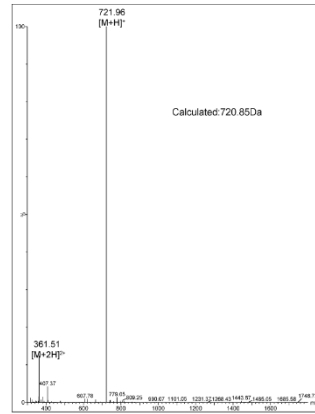
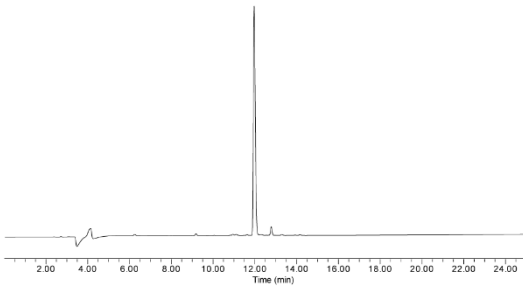
AHP-21: AGHFLW



AHP-22: LMTTKSD



AHP-23: GGLSMTR



AHP-24: NHQRLLP

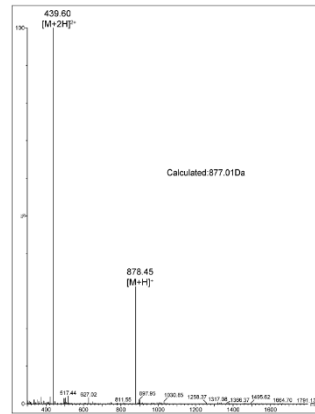
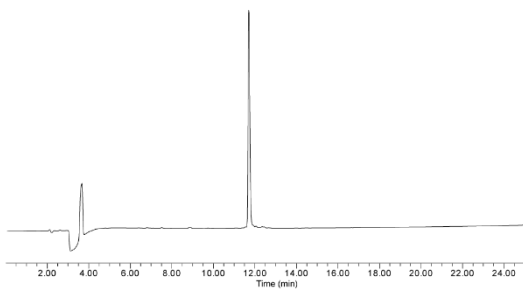
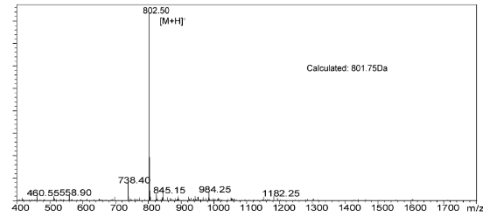
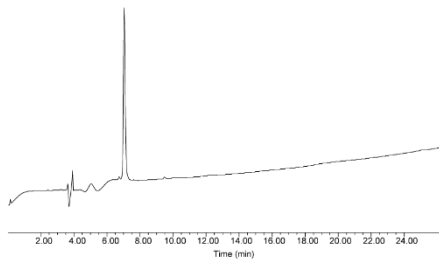
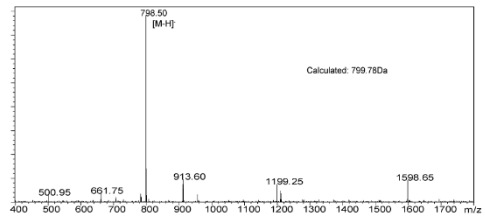
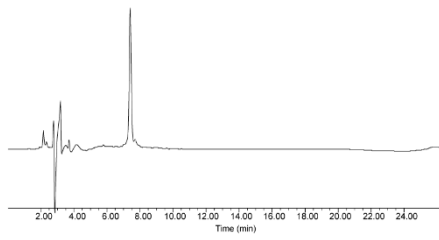


Figure S12. Characterization of twenty-four DR6-targeting peptides by HPLC ($\lambda = 220$ nm) and ESI-MS.

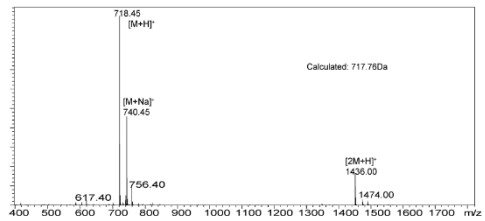
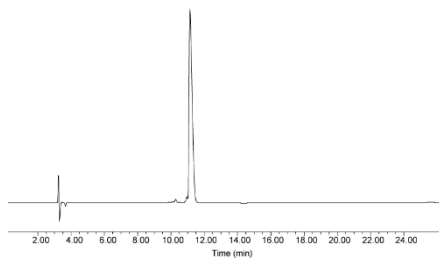
AHP-D-03: D-(YLPAPLP)



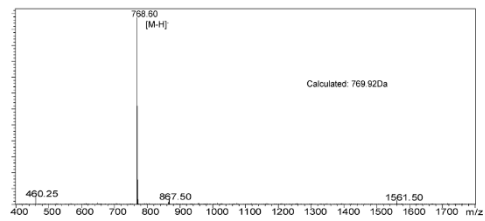
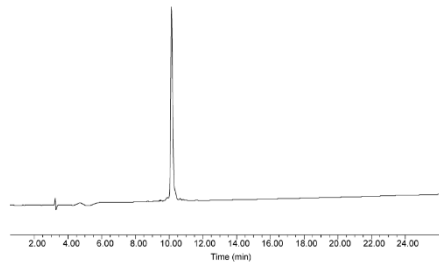
AHP-D-10: D-(NGSTHER)



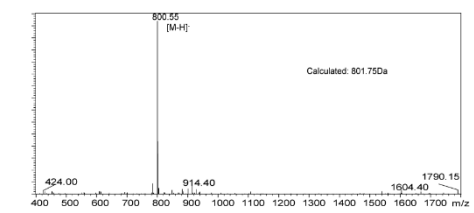
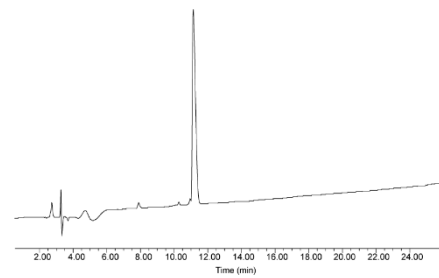
AHP-D-12: D-(TIDATTP)



AHP-D-13: D-(SHTDTNQ)



AHP-DRI-03: D-(PLPAPLY)



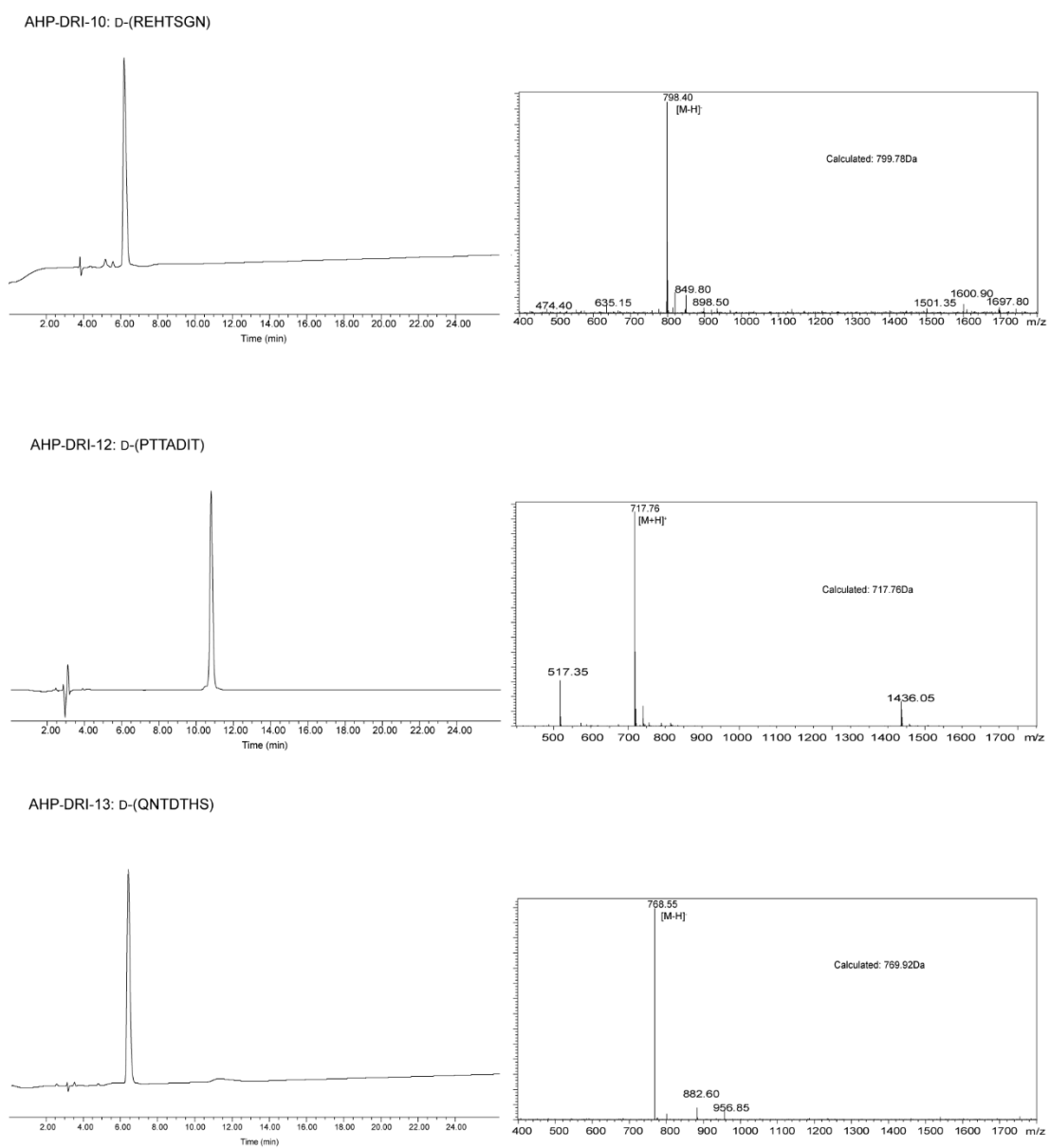


Figure S13. Characterization of D-peptides and DRI-peptides (totally composed of D amino acids) by HPLC ($\lambda = 220$ nm) and ESI-MS.