

Influence of the Drug Position on Bioactivity in Angiopep-2 – Daunomycin Conjugates

Lilla Pethő¹, Rita Oláh-Szabó^{1,†} and Gábor Mező^{1,2,*}

¹ ELKH-ELTE Research Group of Peptide Chemistry, 1117 Budapest, Hungary

² Institute of Chemistry, Faculty of Science, Eötvös Loránd University, 1117 Budapest, Hungary

* Correspondence: gabor.mezo@ttk.elte.hu

† Present address: Department of Genetics, Cell- and Immunobiology, Faculty of Medicine, Semmelweis University, 1089 Budapest, Hungary.

Table of contents

Analytical data of compounds 1-8	2
Analytical chromatograms and MS spectra of compounds 1-8	3
Total ion chromatograms of the lysosomal degradation of compounds 1-8	7
Detected fragments of the lysosomal degradation of compounds 1-8	11
Analytical data of compounds 9-11	17
Analytical chromatograms and MS spectra of compounds 9-11	17
Total ion chromatograms of the lysosomal degradation of compounds 9-11	19
Detected fragments of the lysosomal degradation of compounds 9-11	21

Table S1. Analytical data of the synthesized Angiopep-2 peptide and the daunomycin-Angiopep-2 conjugates

Code	Compound	R_t / min ^a	MW_{calc} / MW_{meas} ^b
1	H-TFFYGGSRGKRNNFKTEEY-OH	18.6	2300.1 / 2300.2
2	H-TFFYGGSRGKRNNFK(Dau=Aoa)TEEY-OH	20.9	2882.3 / 2882.3
3	Dau=Aoa-TFFYGGSRGKRNNFKTEEY-OH	21.9	2882.3 / 2882.3
4	H-TFFYGGSRGK(Dau=Aoa)RNNFKTEEY-OH	21.2	2882.3 / 2882.3
5	Dau=Aoa-TFFYGGSRGKRNNFK(Dau=Aoa)TEEY-OH	21.8	3464.5 / 3464.5
6	H-TFFYGGSRGK(Dau=Aoa)RNNFK(Dau=Aoa)TEEY-OH	20.6	3464.5 / 3464.5
7	Dau=Aoa-TFFYGGSRGK(Dau=Aoa)RNNFKTEEY-OH	21.8	3464.5 / 3464.5
8	Dau=Aoa-TFFYGGSRGK(Dau=Aoa)RNNFK(Dau=Aoa)TEEY-OH	21.6	4046.7 / 4046.6

^a Macherey-Nagel Nucleosil C18 column (5 μ m, 100 Å, 250 \times 4,6 mm); gradient: 0 min 2% B, 5 min 2% B, 30 min 90% B; eluent A: 0.1% TFA / distilled water, eluent B: 0.1% TFA / acetonitrile:distilled water (80:20, v/v); flow rate: 1 mL/min; detection: 214 nm.

^b ESI-MS Bruker Daltonics Esquire 3000 Plus

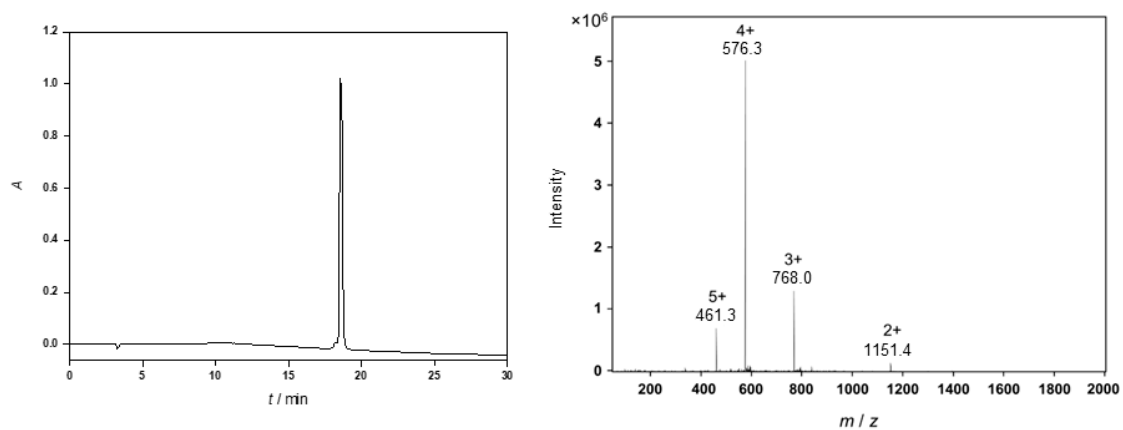


Figure S1. Analytical chromatogram and MS spectrum of Angiopep-2 (1)

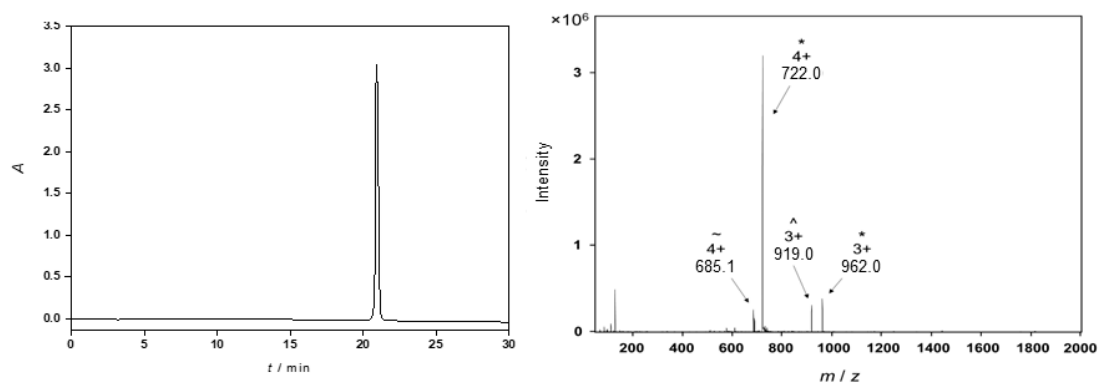


Figure S2. Analytical chromatogram and MS spectrum of H-TFFYGGSRGKRNNFK(Dau=Aoa)TEEY-OH (2). Peaks marked with - and ^ belong to the fragments resulting from the sugar loss of daunomycin occurring under general conditions of MS.

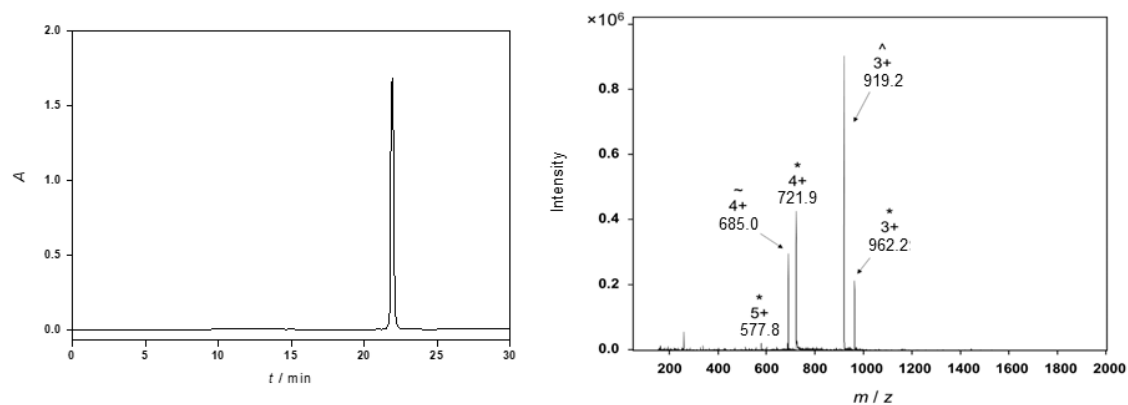


Figure S3. Analytical chromatogram and MS spectrum of Dau=Aoa-TFFYGGSRGKRNNFKTEEY-OH (3). Peaks marked with - and ^ belong to the fragments resulting from the sugar loss of daunomycin occurring under general conditions of MS.

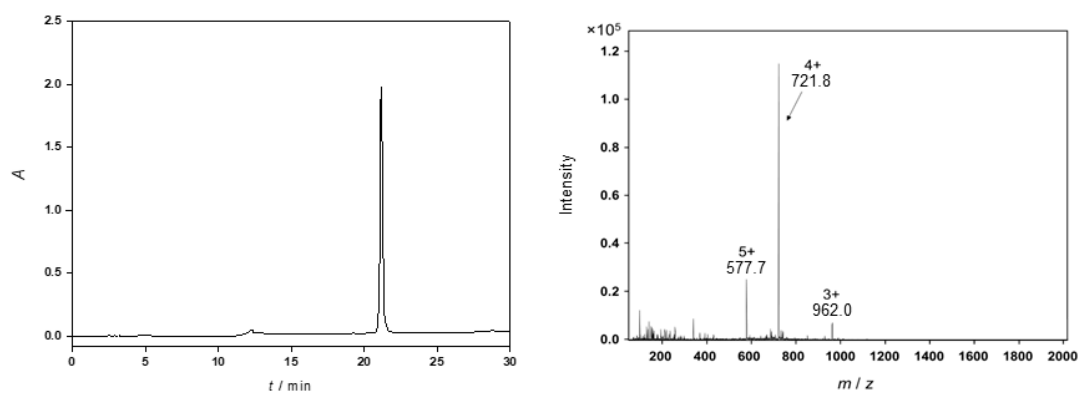


Figure S4. Analytical chromatogram and MS spectrum of H-TFFYGGSRGK(Dau=Aoa)RNNFKTEEY-OH (4)

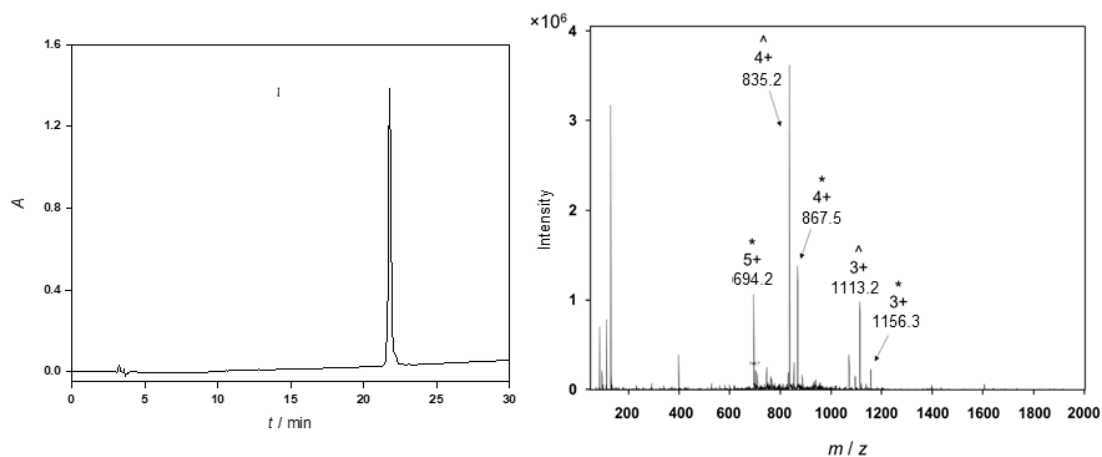


Figure S5. Analytical chromatogram and MS spectrum of Dau=Aoa-TFFYGGSRGKRNNFK(Dau=Aoa)TEEY-OH (**5**). Peaks marked with ^ belong to the fragments resulting from the sugar loss of daunomycin occurring under general conditions of MS.

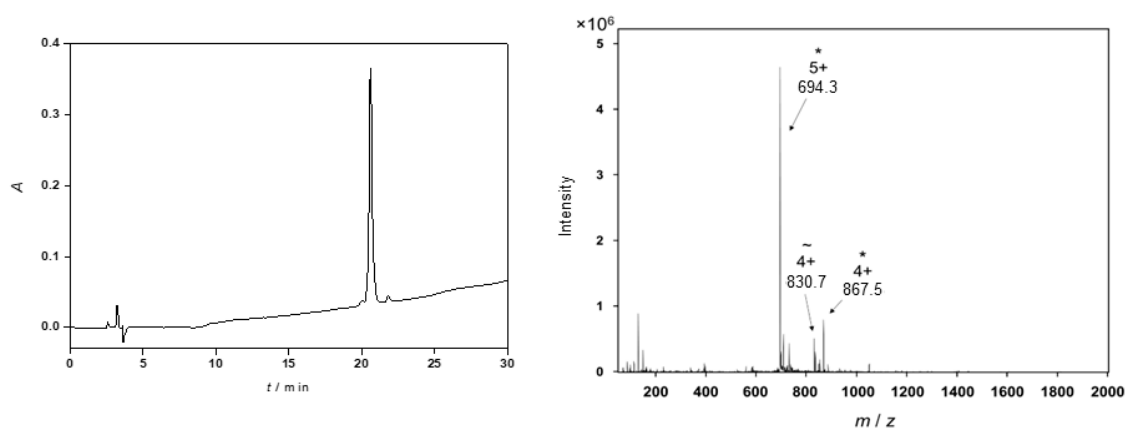


Figure S6. Analytical chromatogram and MS spectrum of H-TFFYGGSRG(Dau=Aoa)KRNNFK(Dau=Aoa)TEEY-OH (**6**). Peaks marked with ~ belong to the fragments resulting from the sugar loss of daunomycin occurring under general conditions of MS.

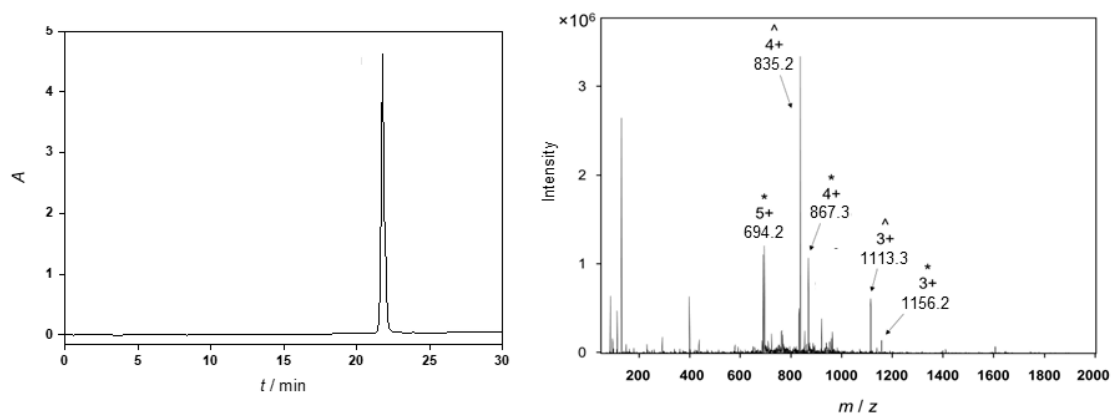


Figure S7. Analytical chromatogram and MS spectrum of Dau=Aoa-TFFYGGSRGK(Dau=Aoa)RNNFKTEEY-OH (7). Peaks marked with ^ belong to the fragments resulting from the sugar loss of daunomycin occurring under general conditions of MS.

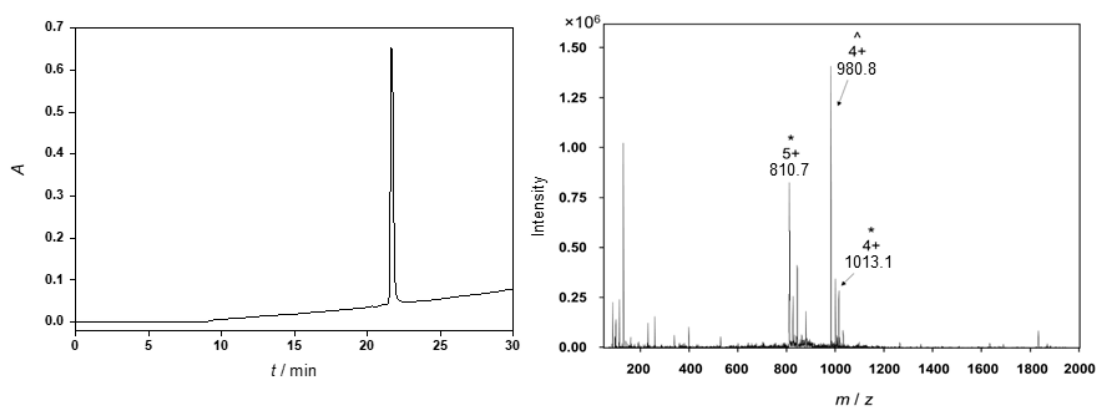


Figure S8. Analytical chromatogram and MS spectrum of Dau=Aoa-TFFYGGSRGK(Dau=Aoa)RNNFK(Dau=Aoa)TEEY-OH (8). Peaks marked with ^ belong to the fragments resulting from the sugar loss of daunomycin occurring under general conditions of MS.

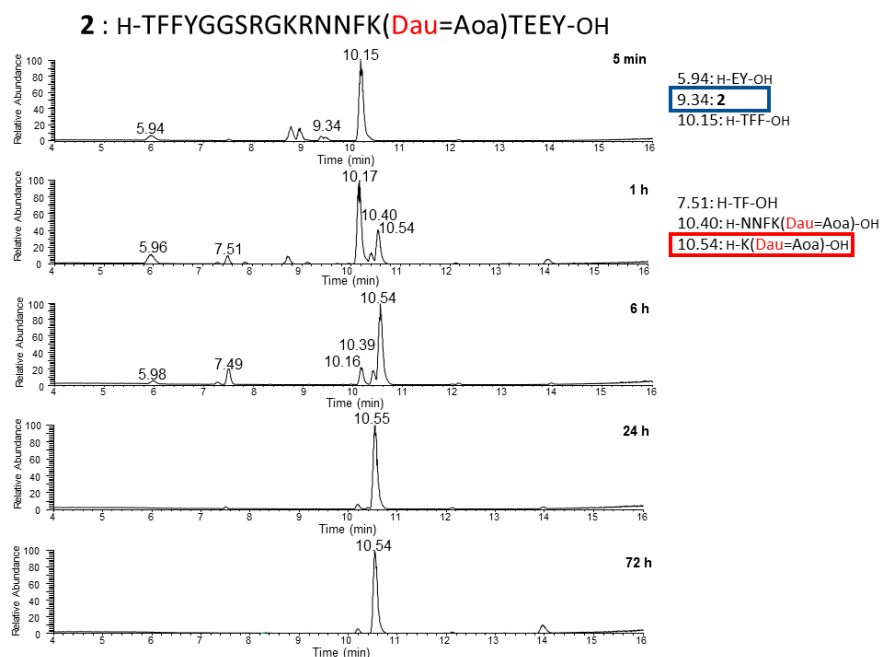


Figure S9. Detected total ion chromatograms showing the lysosomal degradation of conjugate **2**. (The blue box indicates the parent conjugate, while the red box marks the smallest drug containing metabolite.)

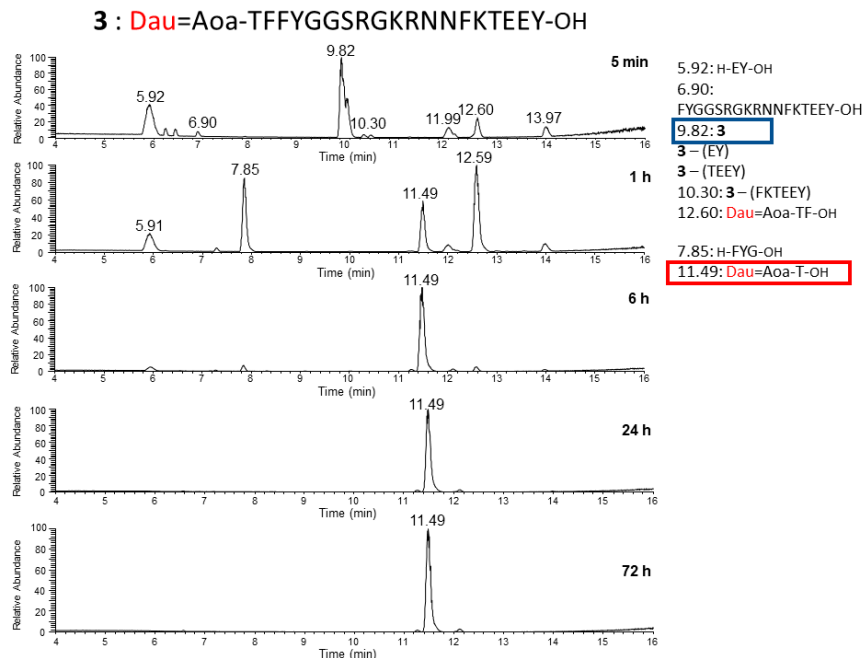


Figure S10. Detected total ion chromatograms showing the lysosomal degradation of conjugate **3**. (The blue box indicates the parent conjugate, while the red box marks the smallest drug containing metabolite.)

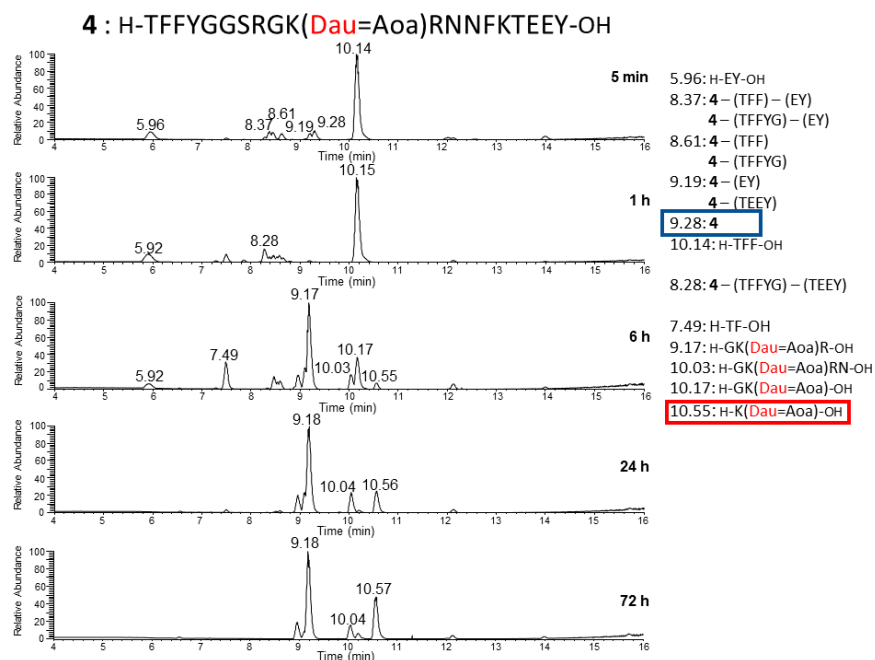


Figure S11. Detected total ion chromatograms showing the lysosomal degradation of conjugate 4. (The blue box indicates the parent conjugate, while the red box marks the smallest drug containing metabolite.)

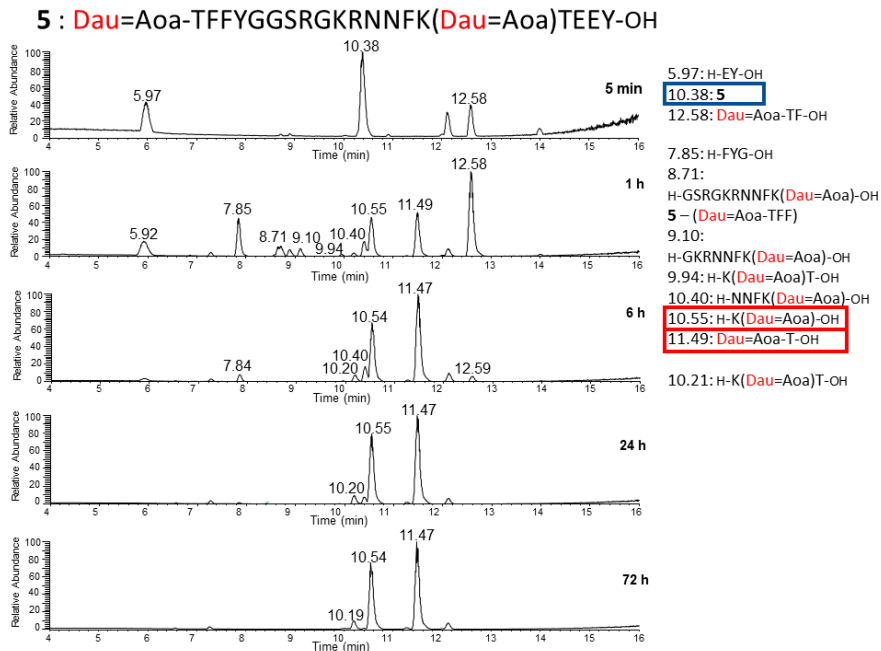


Figure S12. Detected total ion chromatograms showing the lysosomal degradation of conjugate 5. (The blue box indicates the parent conjugate, while the red boxes mark the smallest drug containing metabolites.)

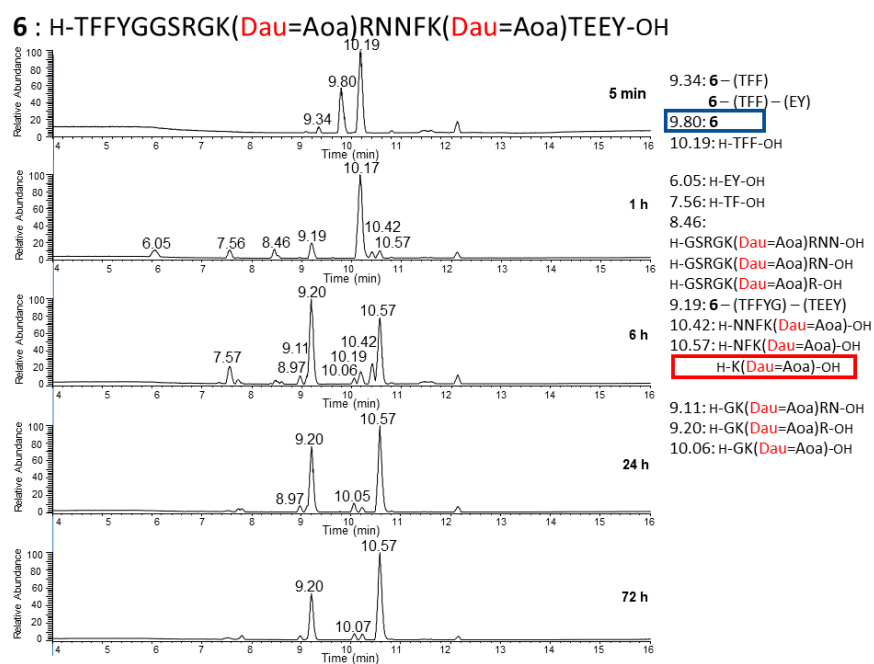


Figure S13. Detected total ion chromatograms showing the lysosomal degradation of conjugate **6**. (The blue box indicates the parent conjugate, while the red box marks the smallest drug containing metabolite.)

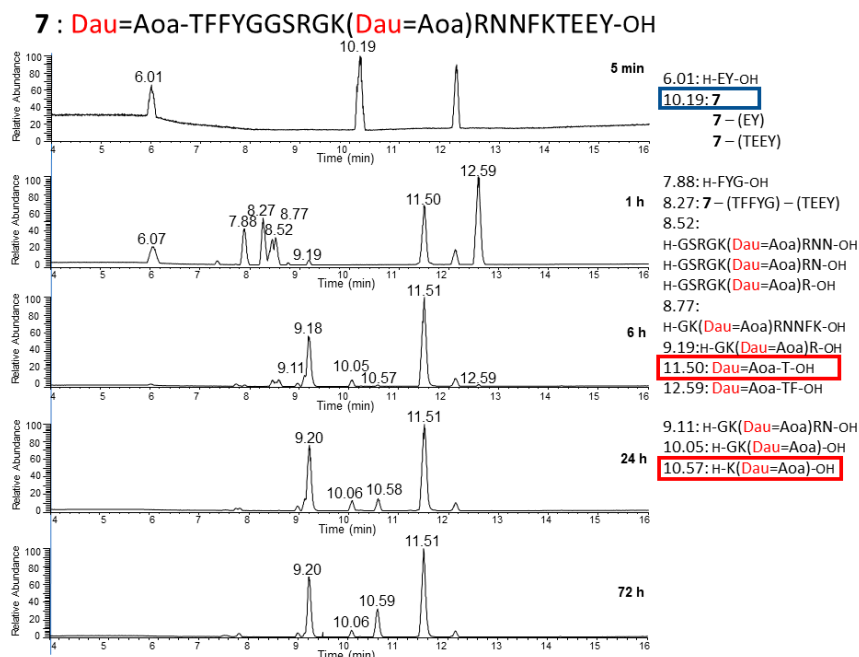


Figure S14. Detected total ion chromatograms showing the lysosomal degradation of conjugate 7. (The blue box indicates the parent conjugate, while the red boxes mark the smallest drug containing metabolites.)

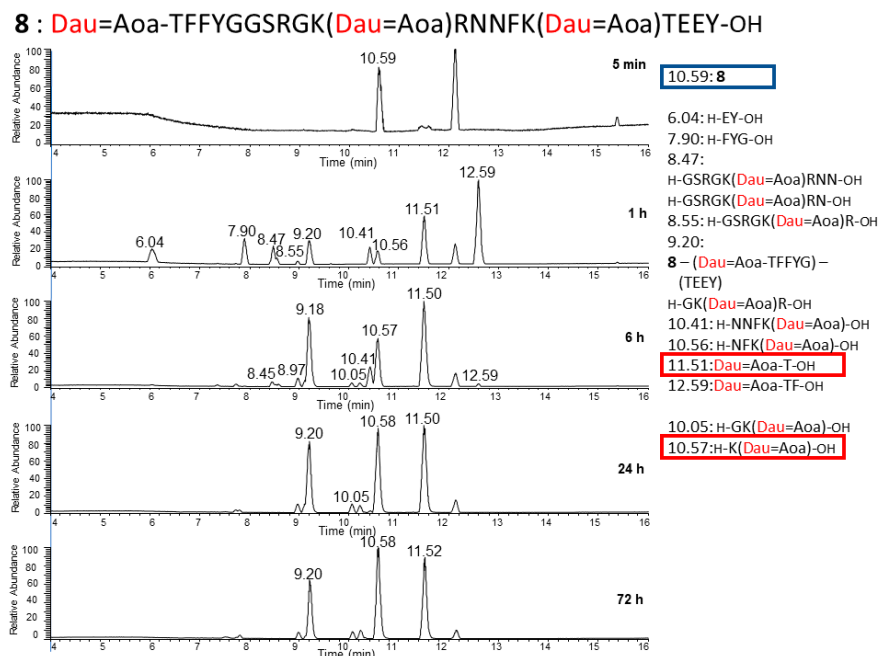


Figure S15. Detected total ion chromatograms showing the lysosomal degradation of conjugate 8. (The blue box indicates the parent conjugate, while the red boxes mark the smallest drug containing metabolites.)

Table S2. Detected fragments from the lysosomal degradation of conjugate 2.^a

	Fragment	Chemical formula	MW _{calc}	MW _{meas}	Δ ppm
1	H-TFFYGGSRGKRNNFK(Dau=Aoa)TEEY-OH	C ₁₃₃ H ₁₇₉ N ₃₁ O ₄₂	2882.2824	2882.2761	-2.19
2	H-TFFYGGSRGKRNNFK(Dau=Aoa)TE-OH	C ₁₁₉ H ₁₆₃ N ₂₉ O ₃₇	2590.1765	2590.1682	-3.20
3	H-YGGSRGKRNNFK(Dau=Aoa)TEEY-OH	C ₁₁₁ H ₁₅₄ N ₂₈ O ₃₈	2487.0979	2487.0928	-2.05
4	H-YGGSRGKRNNFK(Dau=Aoa)TE-OH	C ₉₇ H ₁₃₈ N ₂₆ O ₃₃	2194.9920	2194.9866	-2.46
5	H-YGGSRGKRNNFK(Dau=Aoa)-OH	C ₈₈ H ₁₂₄ N ₂₄ O ₂₈	1964.9017	1964.8953	-3.26
6	H-GKRNNFK(Dau=Aoa)-OH	C ₆₆ H ₉₂ N ₁₆ O ₂₁	1444.6623	1444.6582	-2.84
7	H-NNFK(Dau=Aoa)-OH	C ₅₂ H ₆₅ N ₉ O ₁₈	1103.4448	1103.4420	-2.54
8	H-K(Dau=Aoa)T-OH	C ₃₉ H ₅₁ N ₅ O ₁₅	829.3382	829.3358	-2.89
9	H-K(Dau=Aoa)-OH	C ₃₅ H ₄₄ N ₄ O ₁₃	728.2905	728.2878	-3.71
10	H-TFF-OH	C ₂₂ H ₂₇ N ₃ O ₅	413.1951	413.1937	-3.39
11	H-EY-OH	C ₁₄ H ₁₈ N ₂ O ₆	310.1165	310.1151	-4.51
12	H-TF-OH	C ₁₃ H ₁₈ N ₂ O ₄	266.1267	266.1254	-4.88

^a The grey row indicates the smallest drug containing metabolite.

Table S3. Detected fragments from the lysosomal degradation of conjugate 3.^a

	Fragment	Chemical formula	MW _{calc}	MW _{meas}	Δ ppm
1	Dau=Aoa-TFFYGGSRGKRNNFKTEEY-OH	C ₁₃₃ H ₁₇₉ N ₃₁ O ₄₂	2882.2824	2882.2767	-1.98
2	Dau=Aoa-TFFYGGSRGKRNNFKTE-OH	C ₁₁₉ H ₁₆₃ N ₂₉ O ₃₇	2590.1765	2590.1716	-1.89
3	Dau=Aoa-TFFYGGSRGKRNNFK-OH	C ₁₁₀ H ₁₄₉ N ₂₇ O ₃₂	2360.0862	2360.0796	-2.80
4	Dau=Aoa-TFFYGGSRGKRNN-OH	C ₉₅ H ₁₂₈ N ₂₄ O ₃₀	2084.9228	2084.9196	-1.53
5	H-FYGGSRGKRNNFKTEEY-OH	C ₉₁ H ₁₃₃ N ₂₇ O ₂₈	2051.9813	2051.9760	-2.58
6	Dau=Aoa-TF-OH	C ₄₂ H ₄₈ N ₄ O ₁₅	848.3116	848.3095	-2.48
7	Dau=Aoa-T-OH	C ₃₃ H ₃₉ N ₃ O ₁₄	701.2432	701.2408	-3.42
8	H-FYG-OH	C ₂₀ H ₂₃ N ₃ O ₅	385.1638	385.1624	-3.63
9	H-EY-OH	C ₁₄ H ₁₈ N ₂ O ₆	310.1165	310.1152	-4.19

^a The grey row indicates the smallest drug containing metabolite.

Table S4. Detected fragments from the lysosomal degradation of conjugate 4.^a

	Fragment	Chemical formula	MW _{calc}	MW _{meas}	Δ ppm
1	H-TFFYGGSRGK(Dau=Aoa)RNNFKTEEY-OH	C ₁₃₃ H ₁₇₉ N ₃₁ O ₄₂	2882.2824	2882.2764	-2.08
2	H-TFFYGGSRGK(Dau=Aoa)RNNFKTE-OH	C ₁₁₉ H ₁₆₃ N ₂₉ O ₃₇	2590.1765	2590.1694	-2.74
3	H-YGGSRGK(Dau=Aoa)RNNFKTEEY-OH	C ₁₁₁ H ₁₅₄ N ₂₈ O ₃₈	2487.0979	2487.0903	-3.06
4	H-TFFYGGSRGK(Dau=Aoa)RNNFK-OH	C ₁₁₀ H ₁₄₉ N ₂₇ O ₃₂	2360.0862	2360.0793	-2.92
5	H-GSRGK(Dau=Aoa)RNNFKTEEY-OH	C ₁₀₀ H ₁₄₂ N ₂₆ O ₃₅	2267.0131	2267.0070	-2.69
6	H-YGGSRGK(Dau=Aoa)RNNFKTE-OH	C ₉₇ H ₁₃₈ N ₂₆ O ₃₃	2194.9920	2194.9876	-2.00
7	H-YGGSRGK(Dau=Aoa)RNNFK-OH	C ₈₈ H ₁₂₄ N ₂₄ O ₂₈	1964.9017	1964.8970	-2.39
8	H-GSRGK(Dau=Aoa)RNNFK-OH	C ₇₇ H ₁₁₂ N ₂₂ O ₂₅	1744.8169	1744.8122	-2.69
9	H-GK(Dau=Aoa)RN-OH	C ₄₇ H ₆₅ N ₁₁ O ₁₇	1055.4560	1055.4530	-2.84
10	H-GK(Dau=Aoa)R-OH	C ₄₃ H ₅₉ N ₉ O ₁₅	941.4131	941.4102	-3.08
11	H-GK(Dau=Aoa)-OH	C ₃₇ H ₄₇ N ₅ O ₁₄	785.3120	785.3096	-3.06
12	H-K(Dau=Aoa)-OH	C ₃₅ H ₄₄ N ₄ O ₁₃	728.2905	728.2887	-2.47
13	H-TFF-OH	C ₂₂ H ₂₇ N ₃ O ₅	413.1951	413.1937	-3.39
14	H-EY-OH	C ₁₄ H ₁₈ N ₂ O ₆	310.1165	310.1152	-4.19
15	H-TF-OH	C ₁₃ H ₁₈ N ₂ O ₄	266.1267	266.1255	-4.51

^a The grey row indicates the smallest drug containing metabolite.

Table S5. Detected fragments from the lysosomal degradation of conjugate **5**.^a

	Fragment	Chemical formula	MW _{calc}	MW _{meas}	Δ ppm
1	Dau=Aoa-TFFYGGSRGKRNNFK(Dau=Aoa)TEEY-OH	C ₁₆₂ H ₂₀₉ N ₃₃ O ₅₃	3464.4673	3464.4582	-2.63
2	H-YGGSRGKRNNFK(Dau=Aoa)-OH	C ₈₈ H ₁₂₄ N ₂₄ O ₂₈	1964.9017	1964.8962	-2.80
3	H-GSRGKRNNFK(Dau=Aoa)-OH	C ₇₇ H ₁₁₂ N ₂₂ O ₂₅	1744.8169	1744.8106	-3.61
4	H-GKRNNFK(Dau=Aoa)-OH	C ₆₆ H ₉₂ N ₁₆ O ₂₁	1444.6623	1444.6576	-3.25
5	H-NNFK(Dau=Aoa)-OH	C ₅₂ H ₆₅ N ₉ O ₁₈	1103.4448	1103.4419	-2.63
6	Dau=Aoa-TF-OH	C ₄₂ H ₄₈ N ₄ O ₁₅	848.3116	848.3092	-2.83
7	H-K(Dau=Aoa)T-OH	C ₃₉ H ₅₁ N ₅ O ₁₅	829.3382	829.3361	-2.53
8	H-K(Dau=Aoa)-OH	C ₃₅ H ₄₄ N ₄ O ₁₃	728.2905	728.2876	-3.98
9	Dau=Aoa-T-OH	C ₃₃ H ₃₉ N ₃ O ₁₄	701.2432	701.2411	-2.99
10	H-FYG-OH	C ₂₀ H ₂₃ N ₃ O ₅	385.1638	385.1623	-3.89
11	H-EY-OH	C ₁₄ H ₁₈ N ₂ O ₆	310.1165	310.1151	-4.51

^a The grey rows indicate the smallest drug containing metabolites.

Table S6. Detected fragments from the lysosomal degradation of conjugate **6**.^a

	Fragment	Chemical formula	MW _{calc}	MW _{meas}	Δ ppm
1	H-TFFYGGSRGK(Dau=Aoa)RNNFK(Dau=Aoa)TEEY-OH	C ₁₆₂ H ₂₀₉ N ₃₃ O ₅₃	3464.4673	3464.4564	-3.15
2	H-YGGSRGK(Dau=Aoa)RNNFK(Dau=Aoa)TEEY-OH	C ₁₄₀ H ₁₈₄ N ₃₀ O ₄₉	3069.2828	3069.2739	-2.90
3	H-YGGSRGK(Dau=Aoa)RNNFK(Dau=Aoa)TE-OH	C ₁₂₆ H ₁₆₈ N ₂₈ O ₄₄	2777.1769	2777.1666	-3.71
4	H-GSRGK(Dau=Aoa)RNNFK(Dau=Aoa)-OH	C ₁₀₆ H ₁₄₂ N ₂₄ O ₃₆	2327.0019	2326.9952	-2.88
5	H-GSRGK(Dau=Aoa)RNN-OH	C ₆₂ H ₉₁ N ₁₉ O ₂₃	1469.6535	1469.6480	-3.74
6	H-GSRGK(Dau=Aoa)RN-OH	C ₅₈ H ₈₅ N ₁₇ O ₂₁	1355.6106	1355.6060	-3.39
7	H-GSRGK(Dau=Aoa)R-OH	C ₅₄ H ₇₉ N ₁₅ O ₁₉	1241.5677	1241.5640	-2.98
8	H-NNFK(Dau=Aoa)-OH	C ₅₂ H ₆₅ N ₉ O ₁₈	1103.4447	1103.4414	-2.99
9	H-GK(Dau=Aoa)RN-OH	C ₄₇ H ₆₅ N ₁₁ O ₁₇	1055.4560	1055.4522	-3.60
10	H-NFK(Dau=Aoa)-OH	C ₄₈ H ₅₉ N ₇ O ₁₆	989.4018	989.3988	-3.03
11	H-GK(Dau=Aoa)R-OH	C ₄₃ H ₅₉ N ₉ O ₁₅	941.4131	941.4094	-3.93
12	H-GK(Dau=Aoa)-OH	C ₃₇ H ₄₇ N ₅ O ₁₄	785.3120	785.3088	-4.07
13	H-K(Dau=Aoa)-OH	C ₃₅ H ₄₄ N ₄ O ₁₃	728.2905	728.2880	-3.43
14	H-TFF-OH	C ₂₂ H ₂₇ N ₃ O ₅	413.1951	413.1930	-5.08
15	H-EY-OH	C ₁₄ H ₁₈ N ₂ O ₆	310.1165	310.1149	-5.16
16	H-TF-OH	C ₁₃ H ₁₈ N ₂ O ₄	266.1267	266.1253	-5.26

^a The grey row indicates the smallest drug containing metabolite.

Table S7. Detected fragments from the lysosomal degradation of conjugate 7.^a

	Fragment	Chemical formula	MW _{calc}	MW _{meas}	Δ ppm
1	Dau=Aoa-TFFYGGSRGK(Dau=Aoa)RNNFKTEEY-OH	C ₁₆₂ H ₂₀₉ N ₃₃ O ₅₃	3464.4673	3464.4546	-3.67
2	Dau=Aoa-TFFYGGSRGK(Dau=Aoa)RNNFKTE-OH	C ₁₄₈ H ₁₉₃ N ₃₁ O ₄₈	3172.3614	3172.3488	-3.97
3	Dau=Aoa-TFFYGGSRGK(Dau=Aoa)RNNFK-OH	C ₁₃₉ H ₁₇₉ N ₂₉ O ₄₃	2942.2712	2942.2614	-3.33
4	H-GSRGK(Dau=Aoa)RNNFK-OH	C ₇₇ H ₁₁₂ N ₂₂ O ₂₅	1744.8169	1744.8093	-4.36
5	H-GSRGK(Dau=Aoa)RNN-OH	C ₆₂ H ₉₁ N ₁₉ O ₂₃	1469.6535	1469.6476	-4.01
6	H-GK(Dau=Aoa)RNNFK-OH	C ₆₆ H ₉₂ N ₁₆ O ₂₁	1444.6623	1444.6576	-3.25
7	H-GSRGK(Dau=Aoa)RN-OH	C ₅₈ H ₈₅ N ₁₇ O ₂₁	1355.6106	1355.6052	-3.98
8	H-GSRGK(Dau=Aoa)R-OH	C ₅₄ H ₇₉ N ₁₅ O ₁₉	1241.5677	1241.5628	-3.95
9	H-GK(Dau=Aoa)RN-OH	C ₄₇ H ₆₅ N ₁₁ O ₁₇	1055.4560	1055.4522	-3.60
10	H-GK(Dau=Aoa)R-OH	C ₄₃ H ₅₉ N ₉ O ₁₅	941.4131	941.4101	-3.19
11	H-GK(Dau=Aoa)-OH	C ₃₇ H ₄₇ N ₅ O ₁₄	785.3120	785.3090	-3.82
12	Dau=Aoa-TF-OH	C ₄₂ H ₄₈ N ₄ O ₁₅	848.3116	848.3084	-3.77
13	H-K(Dau=Aoa)-OH	C ₃₅ H ₄₄ N ₄ O ₁₃	728.2905	728.2880	-3.43
14	Dau=Aoa-T-OH	C ₃₃ H ₃₉ N ₃ O ₁₄	701.2432	701.2402	-4.28
15	H-FYG-OH	C ₂₀ H ₂₃ N ₃ O ₅	385.1638	385.1619	-4.93
16	H-EY-OH	C ₁₄ H ₁₈ N ₂ O ₆	310.1165	310.1148	-5.48

^a The grey rows indicate the smallest drug containing metabolites.

Table S8. Detected fragments from the lysosomal degradation of conjugate 8.^a

	Fragment	Chemical formula	MW _{calc}	MW _{meas}	Δ ppm
1	Dau=Aoa-TFFYGGSRGK(Dau=Aoa)RNNFK(Dau=Aoa)TEEY-OH	C ₁₉₁ H ₂₃₉ N ₃₅ O ₆₄	4046.6523	4046.6460	-1.56
2	H-GSRGK(Dau=Aoa)RNNFK(Dau=Aoa)-OH	C ₁₀₆ H ₁₄₂ N ₂₄ O ₃₆	2327.0019	2326.9950	-2.97
3	H-GSRGK(Dau=Aoa)RNN-OH	C ₆₂ H ₉₁ N ₁₉ O ₂₃	1469.6535	1469.6478	-3.88
4	H-GSRGK(Dau=Aoa)RN-OH	C ₅₈ H ₈₅ N ₁₇ O ₂₁	1355.6106	1355.6058	-3.54
5	H-GSRGK(Dau=Aoa)R-OH	C ₅₄ H ₇₉ N ₁₅ O ₁₉	1241.5677	1241.5634	-3.46
6	H-NNFK(Dau=Aoa)-OH	C ₅₂ H ₆₅ N ₉ O ₁₈	1103.4447	1103.4409	-3.44
7	H-NFK(Dau=Aoa)-OH	C ₄₈ H ₅₉ N ₇ O ₁₆	989.4018	989.3984	-3.44
8	H-GK(Dau=Aoa)R-OH	C ₄₃ H ₅₉ N ₉ O ₁₅	941.4131	941.4094	-3.93
9	Dau=Aoa-TF-OH	C ₄₂ H ₄₈ N ₄ O ₁₅	848.3116	848.3082	-4.01
10	H-GK(Dau=Aoa)-OH	C ₃₇ H ₄₇ N ₅ O ₁₄	785.3120	785.3090	-3.82
11	H-K(Dau=Aoa)-OH	C ₃₅ H ₄₄ N ₄ O ₁₃	728.2905	728.2877	-3.84
12	Dau=Aoa-T-OH	C ₃₃ H ₃₉ N ₃ O ₁₄	701.2432	701.2401	-4.42
13	H-FYG-OH	C ₂₀ H ₂₃ N ₃ O ₅	385.1638	385.1619	-4.93
14	H-EY-OH	C ₁₄ H ₁₈ N ₂ O ₆	310.1165	310.1149	-5.16

^a The grey rows indicate the smallest drug containing metabolites.

Table S9. Analytical data of the enzyme labile spacer containing Angiopep-2 – daunomycin conjugates

Code	Conjugate	R_t / min ^a	MW_{calc} / MW_{meas} ^b
9	H-TFFYGGSRGK(Dau=Aoa-GFLG)RNNFKTEEY-OH	22.3	3256.5 / 3256.5
10	H-TFFYGGSRGK(Dau=Aoa-VA)RNNFKTEEY-OH	21.5	3052.4 / 3052.4
11	H-TFFYGGSRGK(Dau=Aoa-VAGG)RNNFKTEEY-OH	21.5	3166.4 / 3166.4

^a Macherey-Nagel Nucleosil C18 column (5 μ m, 100 Å, 250 \times 4,6 mm); gradient: 0 min 2% B, 5 min 2% B, 30 min 90% B; eluent A: 0.1% TFA / distilled water, eluent B: 0.1% TFA / acetonitrile:distilled water (80:20, v/v); flow rate: 1 mL/min; detection: 214 nm.

^b ESI-MS Bruker Daltonics Esquire 3000 Plus

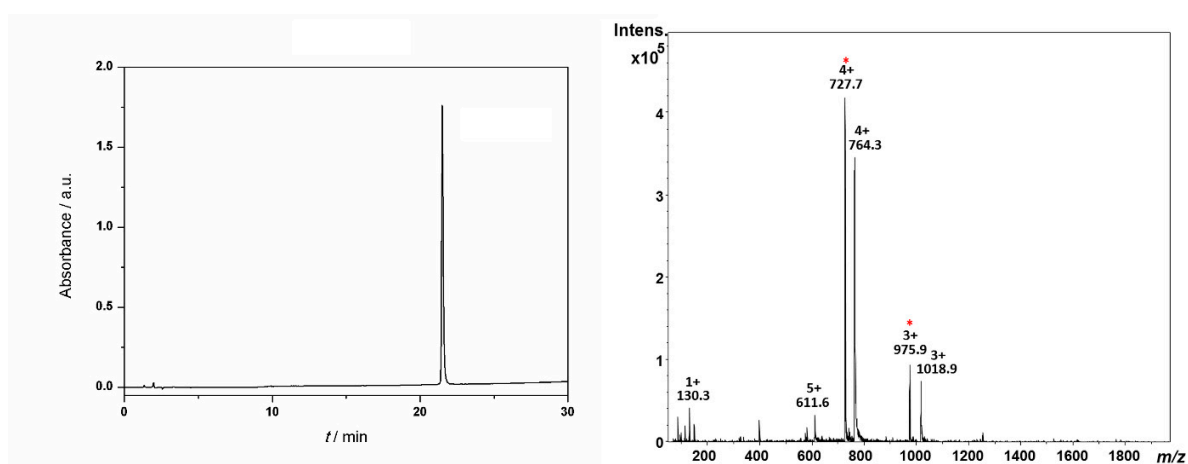


Figure S16. Analytical chromatogram and MS spectrum of H-TFFYGGSRGK(Dau=Aoa-GFLG)RNNFKTEEY-OH (9). Peaks marked with * belong to the fragments resulting from the sugar loss of daunomycin occurring under general conditions of MS.

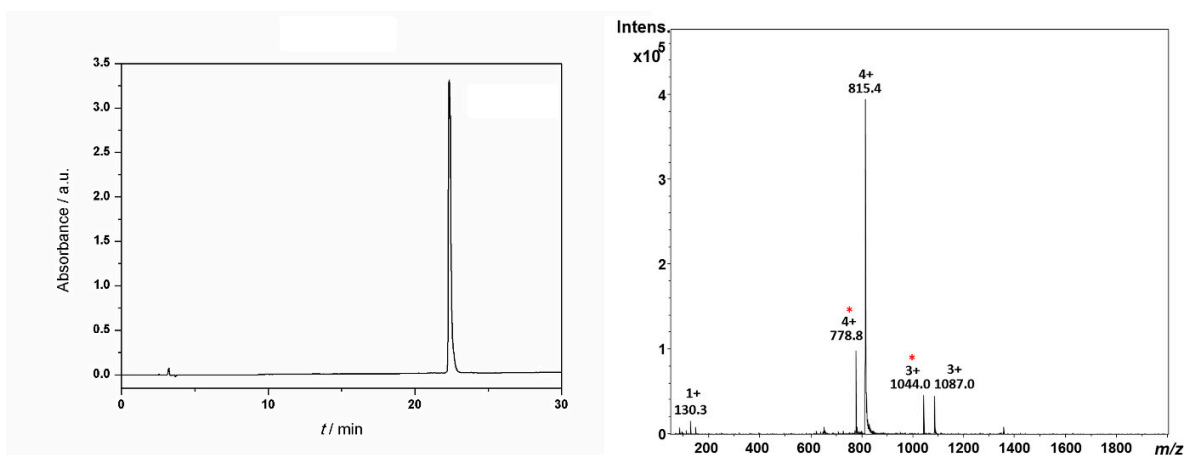


Figure S17. Analytical chromatogram and MS spectrum of H-TFFYGGSRGK(Dau=Aoa-VA)RNNFKTEEY-OH (**10**). Peaks marked with * belong to the fragments resulting from the sugar loss of daunomycin occurring under general conditions of MS.

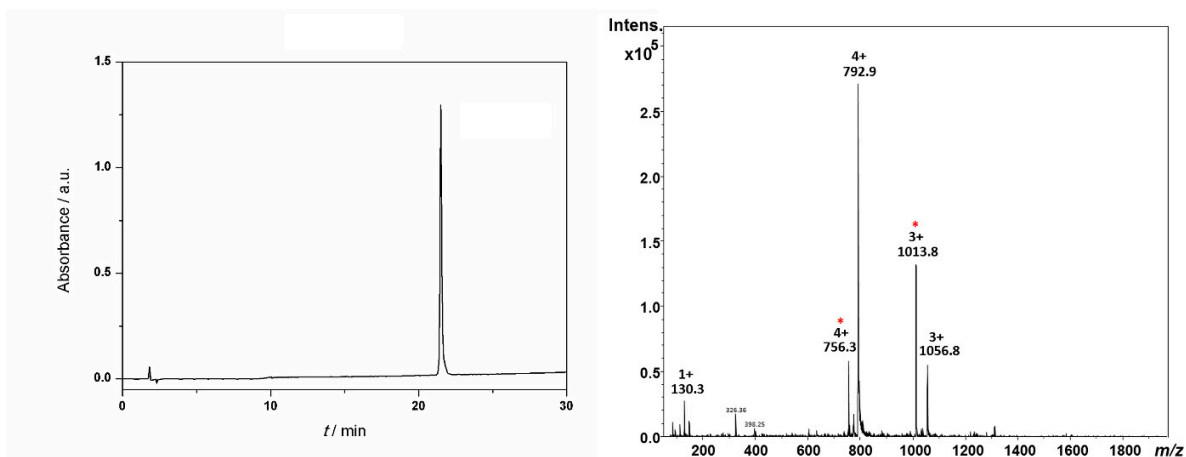


Figure S18. Analytical chromatogram and MS spectrum of H-TFFYGGSRGK(Dau=Aoa-VAGG)RNNFKTEEY-OH (**11**). Peaks marked with * belong to the fragments resulting from the sugar loss of daunomycin occurring under general conditions of MS.

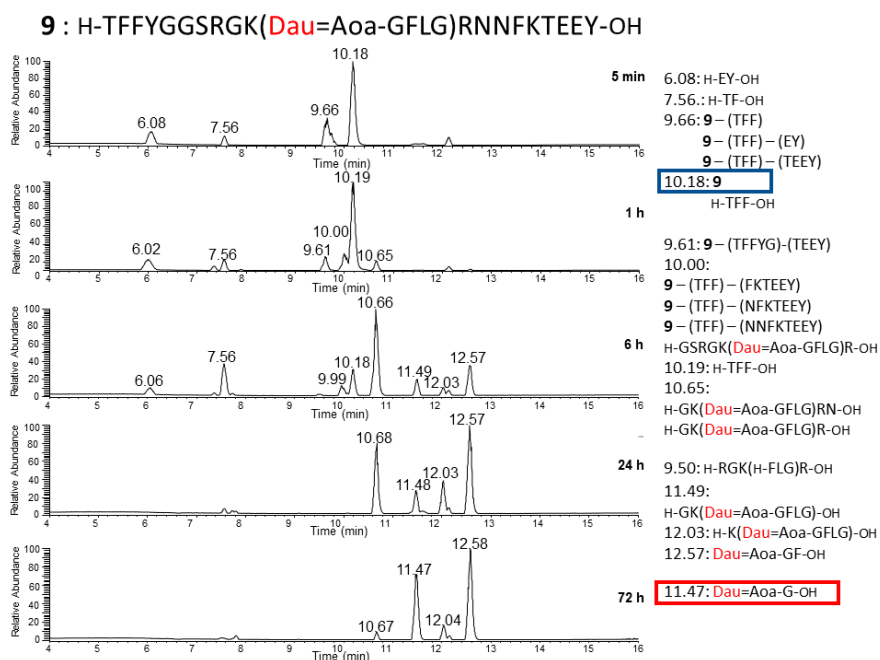


Figure S19. Detected total ion chromatograms showing the lysosomal degradation of conjugate **9**. (The blue box indicates the parent conjugate, while the red box marks the smallest drug containing metabolite.)

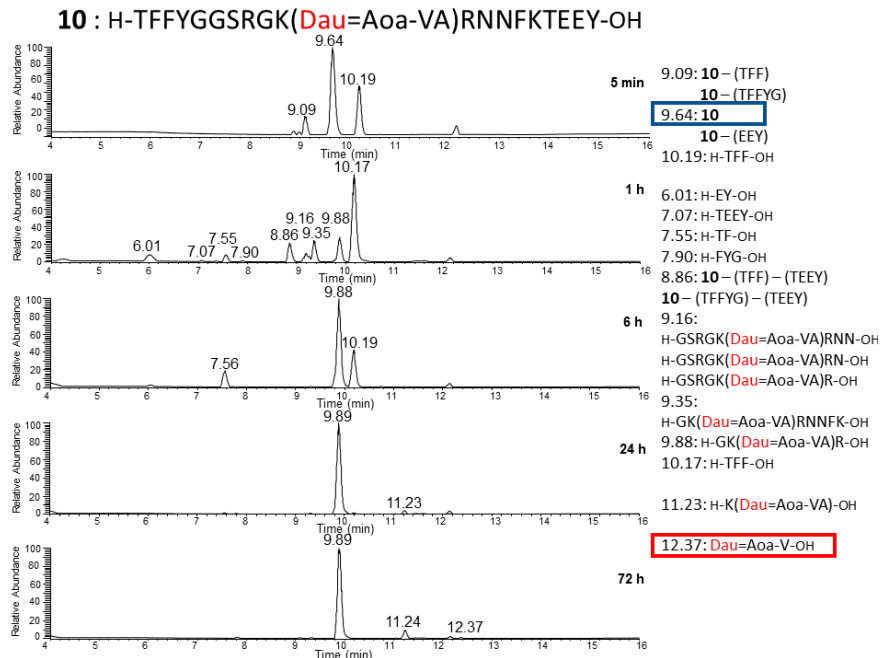


Figure S20. Detected total ion chromatograms showing the lysosomal degradation of conjugate **10**. (The blue box indicates the parent conjugate, while the red box marks the smallest drug containing metabolite.)

11 : H-TFFYGGSRGK(Dau=Aoa-VAGG)RNNFKTEEY-OH

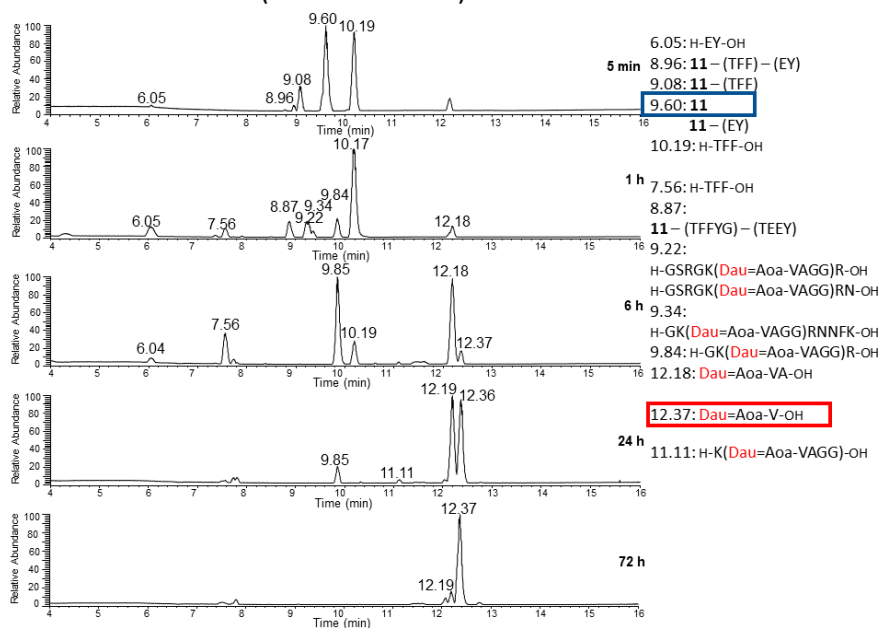


Figure S21. Detected total ion chromatograms showing the lysosomal degradation of conjugate **11**. (The blue box indicates the parent conjugate, while the red box marks the smallest drug containing metabolite.)

Table S10. Detected fragments from the lysosomal degradation of conjugate **9**.^a

	Fragment	Chemical formula	MW _{calc}	MW _{meas}	Δ ppm
1	H-TFFYGGSRGK(Dau =Aoa-GFLG)RNNFKTEEY-OH	C ₁₅₂ H ₂₀₅ N ₃₅ O ₄₆	3256.4778	3256.4668	-3.38
2	H-YGGSRGK(Dau =Aoa-GFLG)RNNFKTEEY-OH	C ₁₃₀ H ₁₈₀ N ₃₂ O ₄₂	2861.2933	2861.2815	-4.12
3	H-YGGSRGK(Dau =Aoa-GFLG)RNNFKTE-OH	C ₁₁₆ H ₁₆₄ N ₃₀ O ₃₇	2569.1874	2569.1775	-3.85
4	H-YGGSRGK(Dau =Aoa-GFLG)RNNFK-OH	C ₁₀₇ H ₁₅₀ N ₂₈ O ₃₂	2339.0971	2339.0868	-4.40
5	H-GSRGK(Dau =Aoa-GFLG)RNNFK-OH	C ₉₆ H ₁₃₈ N ₂₆ O ₂₉	2119.0123	2119.0026	-4.58
6	H-YGGSRGK(Dau =Aoa-GFLG)RNN-OH	C ₉₂ H ₁₂₉ N ₂₅ O ₃₀	2063.9337	2063.9258	-3.83
7	H-YGGSRGK(Dau =Aoa-GFLG)RN-OH	C ₈₈ H ₁₂₃ N ₂₃ O ₂₈	1949.8908	1949.8828	-4.10
8	H-YGGSRGK(Dau =Aoa-GFLG)R-OH	C ₈₄ H ₁₁₇ N ₂₁ O ₂₆	1835.8479	1835.8404	-4.09
9	H-GGSRGK(Dau =Aoa-GFLG)RN-OH	C ₇₉ H ₁₁₄ N ₂₂ O ₂₆	1786.8275	1786.8196	-4.42
10	H-GGSRGK(Dau =Aoa-GFLG)R-OH	C ₇₅ H ₁₀₈ N ₂₀ O ₂₄	1672.7845	1672.7776	-4.12
11	H-GSRGK(Dau =Aoa-GFLG)R-OH	C ₇₃ H ₁₀₅ N ₁₉ O ₂₃	1615.7631	1615.7556	-4.64
12	H-GK(Dau =Aoa-GFLG)RN-OH	C ₆₆ H ₉₁ N ₁₅ O ₂₁	1429.6514	1429.6454	-4.20
13	H-GK(Dau =Aoa-GFLG)R-OH	C ₆₂ H ₈₅ N ₁₃ O ₁₉	1315.6085	1315.6038	-3.57
14	H-GK(Dau =Aoa-GFLG)-OH	C ₅₆ H ₇₃ N ₉ O ₁₈	1159.5074	1159.5041	-2.85
15	H-K(Dau =Aoa-GFLG)-OH	C ₅₄ H ₇₀ N ₈ O ₁₇	1102.4859	1102.4827	-2.90
16	Dau =Aoa-GF-OH	C ₄₀ H ₄₄ N ₄ O ₁₄	804.2854	804.2826	-3.48
17	Dau =Aoa-G-OH	C ₃₁ H ₃₅ N ₃ O ₁₃	657.2170	657.2145	-3.80
18	H-TFF-OH	C ₂₂ H ₂₇ N ₃ O ₅	413.1951	413.1931	-4.84
19	H-EY-OH	C ₁₄ H ₁₈ N ₂ O ₆	310.1165	310.1148	-5.48
20	H-TF-OH	C ₁₃ H ₁₈ N ₂ O ₄	266.1267	266.1253	-5.26

^a The grey row indicates the smallest drug containing metabolite.

Table S11. Detected fragments from the lysosomal degradation of conjugate 10.^a

	Fragment	Chemical formula	MW _{calc}	MW _{meas}	Δ ppm
1	H-TFFYGGSRGK(Dau=Aoa-VA)RNNFKTEEY-OH	C ₁₄₁ H ₁₉₃ N ₃₃ O ₄₄	3052.3879	3052.3761	-3.87
2	H-TFFYGGSRGK(Dau=Aoa-VA)RNNFKTE-OH	C ₁₂₇ H ₁₇₇ N ₃₁ O ₃₉	2760.2820	2760.2706	-4.13
3	H-YGGSRGK(Dau=Aoa-VA)RNNFKTEEY-OH	C ₁₁₉ H ₁₆₈ N ₃₀ O ₄₀	2657.2034	2657.1939	-3.58
4	H-GSRGK(Dau=Aoa-VA)RNNFKTEEY-OH	C ₁₀₈ H ₁₅₆ N ₂₈ O ₃₇	2437.1186	2437.1088	-4.02
5	H-YGGSRGK(Dau=Aoa-VA)RNNFK-OH	C ₉₆ H ₁₃₈ N ₂₆ O ₃₀	2135.0072	2134.9980	-4.31
6	H-GSRGK(Dau=Aoa-VA)RNNFK-OH	C ₈₅ H ₁₂₆ N ₂₄ O ₂₇	1914.9224	1914.9144	-4.18
7	H-GSRGK(Dau=Aoa-VA)RNN-OH	C ₇₀ H ₁₀₅ N ₂₁ O ₂₅	1639.7590	1639.7522	-4.15
8	H-GK(Dau=Aoa-VA)RNNFK-OH	C ₇₄ H ₁₀₆ N ₁₈ O ₂₃	1614.7678	1614.7622	-3.47
9	H-GSRGK(Dau=Aoa-VA)RN-OH	C ₆₆ H ₉₉ N ₁₉ O ₂₃	1525.7161	1525.7100	-4.00
10	H-GSRGK(Dau=Aoa-VA)R-OH	C ₆₂ H ₉₃ N ₁₇ O ₂₁	1411.6732	1411.6678	-3.83
11	H-GK(Dau=Aoa-VA)R-OH	C ₅₁ H ₇₃ N ₁₁ O ₁₇	1111.5186	1111.5147	-3.51
12	H-K(Dau=Aoa-VA)-OH	C ₄₃ H ₅₈ N ₆ O ₁₅	898.3960	898.3929	-3.45
13	Dau=Aoa-V-OH	C ₃₄ H ₄₁ N ₃ O ₁₃	699.2639	699.2615	-3.43
14	H-TEEY-OH	C ₂₃ H ₃₂ N ₄ O ₁₁	540.2068	540.2051	-3.15
15	H-TFF-OH	C ₂₂ H ₂₇ N ₃ O ₅	413.1951	413.1931	-4.84
16	H-FYG-OH	C ₂₀ H ₂₃ N ₃ O ₅	385.1638	385.1624	-3.63
17	H-EY-OH	C ₁₄ H ₁₈ N ₂ O ₆	310.1165	310.1149	-5.16
18	H-TF-OH	C ₁₃ H ₁₈ N ₂ O ₄	266.1267	266.1253	-5.26

^a The grey row indicates the smallest drug containing metabolite.

Table S12. Detected fragments from the lysosomal degradation of conjugate **11**.^a

	Fragment	Chemical formula	MW _{calc}	MW _{meas}	Δ ppm
1	H-TFFYGGSRGK(Dau=Aoa-VAGG)RNNFKTEEY-OH	C ₁₄₅ H ₁₉₉ N ₃₅ O ₄₆	3166.4308	3166.4211	-3.06
2	H-TFFYGGSRGK(Dau=Aoa-VAGG)RNNFKTE-OH	C ₁₃₁ H ₁₈₃ N ₃₃ O ₄₁	2874.3249	2874.3138	-3.86
3	H-YGGSRGK(Dau=Aoa-VAGG)RNNFKTEEY-OH	C ₁₂₃ H ₁₇₄ N ₃₂ O ₄₂	2771.2463	2771.2365	-3.54
4	H-YGGSRGK(Dau=Aoa-VAGG)RNNFKTE-OH	C ₁₀₉ H ₁₅₈ N ₃₀ O ₃₇	2479.1404	2479.1313	-3.67
5	H-GSRGK(Dau=Aoa-VAGG)RNNFK-OH	C ₈₉ H ₁₃₂ N ₂₆ O ₂₉	2028.9654	2028.9580	-3.65
6	H-GK(Dau=Aoa-VAGG)RNNFK-OH	C ₇₈ H ₁₁₂ N ₂₀ O ₂₅	1728.8107	1728.8036	-4.11
7	H-GSRGK(Dau=Aoa-VAGG)RN-OH	C ₇₀ H ₁₀₅ N ₂₁ O ₂₅	1639.7590	1639.7516	-4.51
8	H-GSRGK(Dau=Aoa-VAGG)R-OH	C ₆₆ H ₉₉ N ₁₉ O ₂₃	1525.7161	1525.7096	-4.26
9	H-GK(Dau=Aoa-VAGG)R-OH	C ₅₅ H ₇₉ N ₁₃ O ₁₉	1225.5615	1225.5562	-4.32
10	H-GK(Dau=Aoa-VAGG)-OH	C ₄₉ H ₆₇ N ₉ O ₁₈	1069.4604	1069.4572	-2.99
11	H-K(Dau=Aoa-VAGG)-OH	C ₄₇ H ₆₄ N ₈ O ₁₇	1012.4389	1012.4358	-3.06
12	Dau=Aoa-VA-OH	C ₃₇ H ₄₆ N ₄ O ₁₄	770.3011	770.2986	-3.25
13	Dau=Aoa-V-OH	C ₃₄ H ₄₁ N ₃ O ₁₃	699.2639	699.2613	-3.72
14	H-TFF-OH	C ₂₂ H ₂₇ N ₃ O ₅	413.1951	413.1931	-4.84
15	H-EY-OH	C ₁₄ H ₁₈ N ₂ O ₆	310.1165	310.1150	-4.84
16	H-TF-OH	C ₁₃ H ₁₈ N ₂ O ₄	266.1267	266.1252	-5.64

^a The grey row indicates the smallest drug containing metabolite.