

Innovative bioconjugation technology for antibody-drug conjugates: proof of concept in a CD30-positive lymphoma mouse model

Ludovic Juen,^{§*} Christine B. Baltus,[§] Camille Gély,[§] Ofelia Feuillâtre,^{§,‡} Audrey Desgranges,[§]
Marie-Claude Viaud-Massuard,^{‡,‡} Camille Martin[§]

[§] McSAF, 1 rue Claude Thion 37000 Tours, France

[#] University of Tours, GICC, Team IMT EA7501, 31 avenue Monge 37200 Tours, FRANCE

[‡] McSAF, cofounder, 1 rue Claude Thion 37000 Tours, France

*corresponding author: ludovic.juen@mcsaf.fr; +33247241254.

Contents:

S1. Linker 7 mass spectrometry analysis.	2
S2. MF-BTX-MMAE native mass spectrometry analysis.	3
S3. MF-BTX-MMAE HIC analysis.	4
S4. MF-BTX-MMAE denaturing mass spectrometry analysis.	5
S5. HPLC SEC-UV analysis of MF-BTX-MMAE and Adcetris®.	6
S6. Stability of MF-BTX-MMAE upon storage at 4 °C.	7
S7. Antigen-binding analysis data by FACS on Karpas-299 (CD30-positive).	8
S8. Antigen-binding analysis by high content analysis on Karpas-299 (CD30-positive).	9
S9. Median tumor volume curves of CB17-SCID mice.	10
S10. Tumor growth inhibition (T/C%) results.	11
S11. Individual tumor volumes of CB17-SCID mice.	12
S12. Mean body weight curves of CB17-SCID mice.	13
S13. Relative body weight change curves of Sprague-Dawley rats.	14-15

Supporting Information

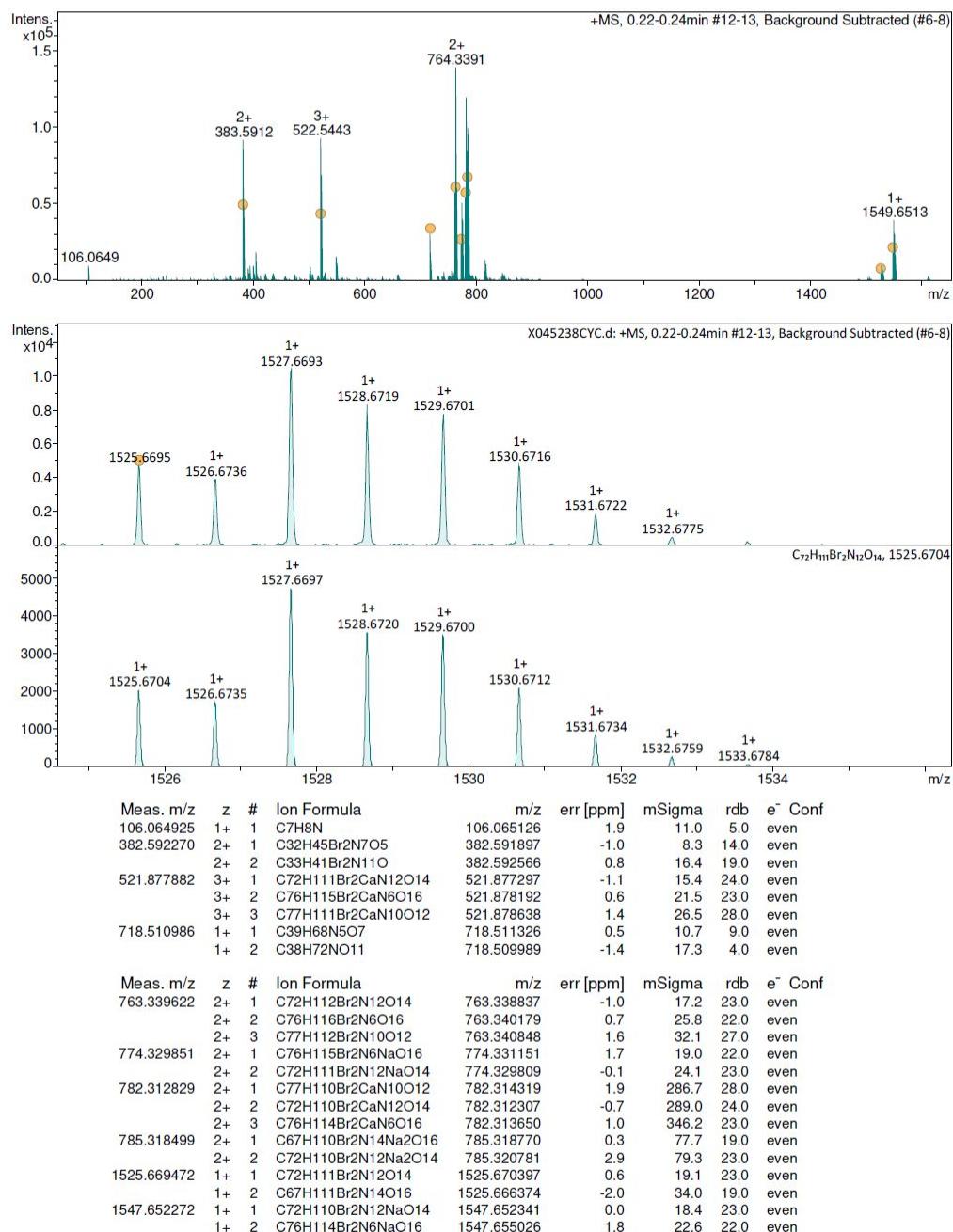
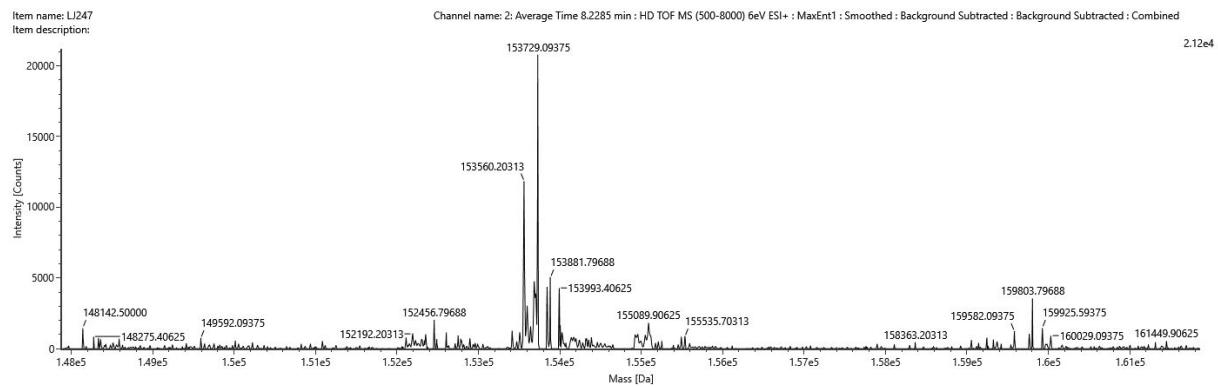


Figure S1. Linker 7 mass spectrometry analysis.

Supporting Information



MF-BTX-MMAE	DAR 0	DAR 1	DAR 2	DAR 3	DAR 4	DAR 5
Expected MW (Da)¹	148083	149451	150819	152187	153555	154923
Measured MW (Da)¹	n.o. ²	n.o. ²	n.o. ²	152194	153561	154932
Measurement error (ppm)	-	-	-	-48.17	-38.8	-58.35
Area	-	-	-	77042	560913	59337
Area (%)	-	-	-	11.1	80.4	8.5
Average DAR	4.0					

¹Only the first glycosylation peak (GOF) was considered.

²Not observed.

The mass increment of 1368 Da expected for the conjugated linker is confirmed.

Figure S2. MF-BTX-MMAE native mass spectrometry analysis.

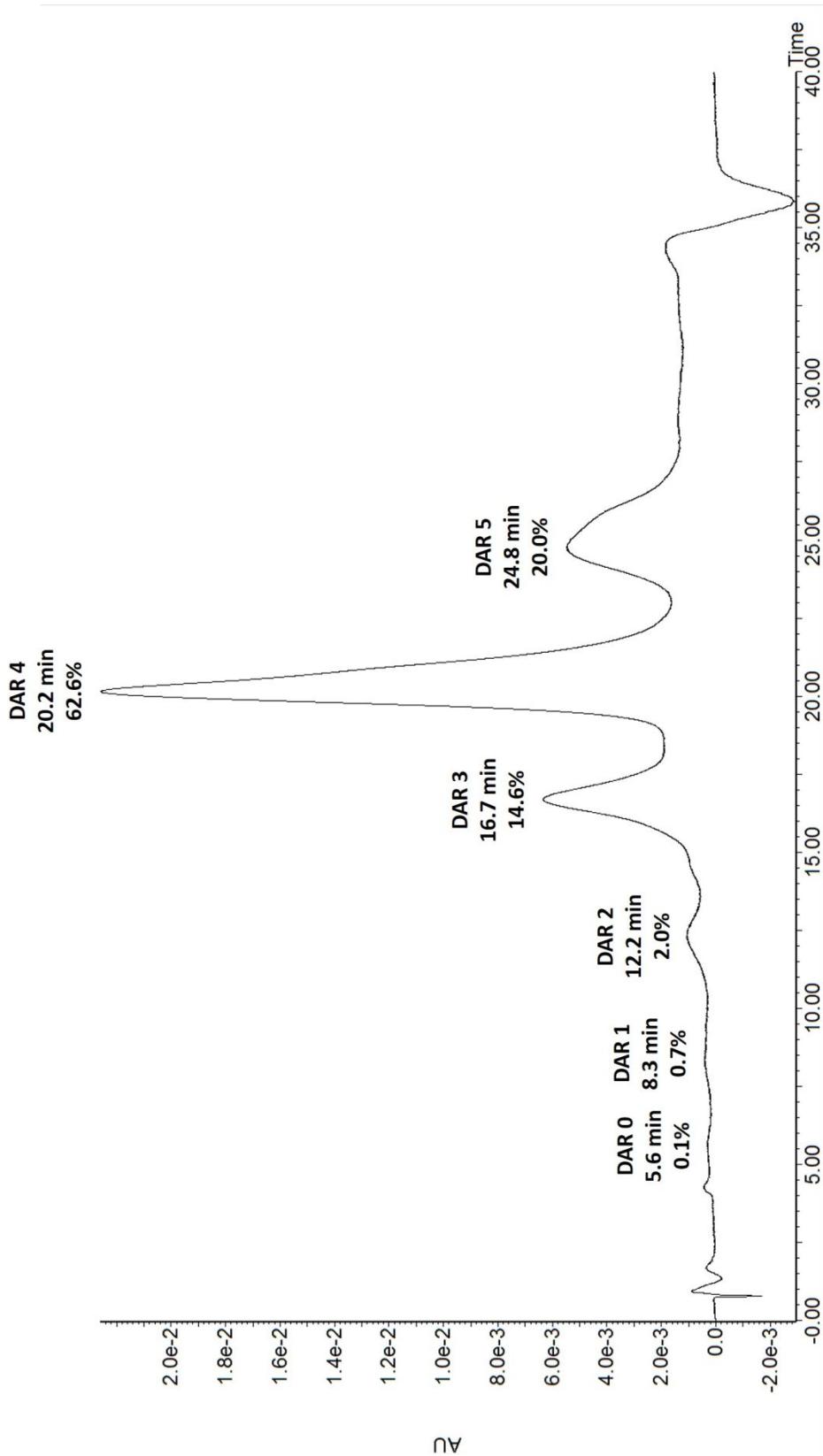
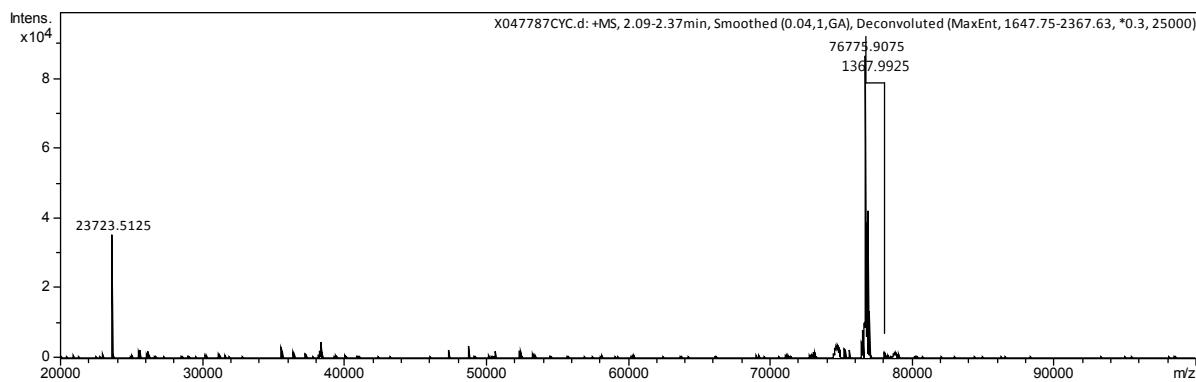
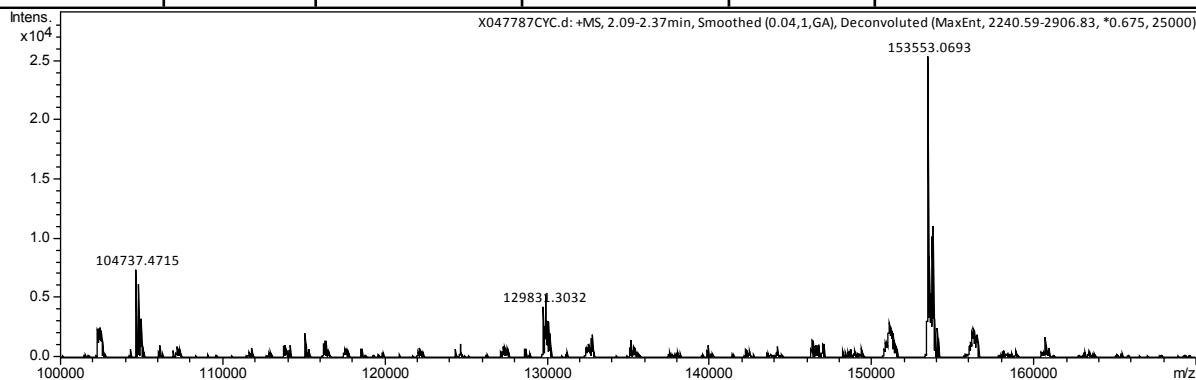


Figure S3. MF-BTX-MMAE HIC analysis.

Supporting Information



Fragments observed	DAR observed	Expected MW (Da) ¹	Measured MW (Da) ¹	Intensity	Relative intensity (%)
L	0	23723	23723	35296	41.0
LH	2	76779	76776	86140	100.0
LH	3	78147	78144	1614	1.9



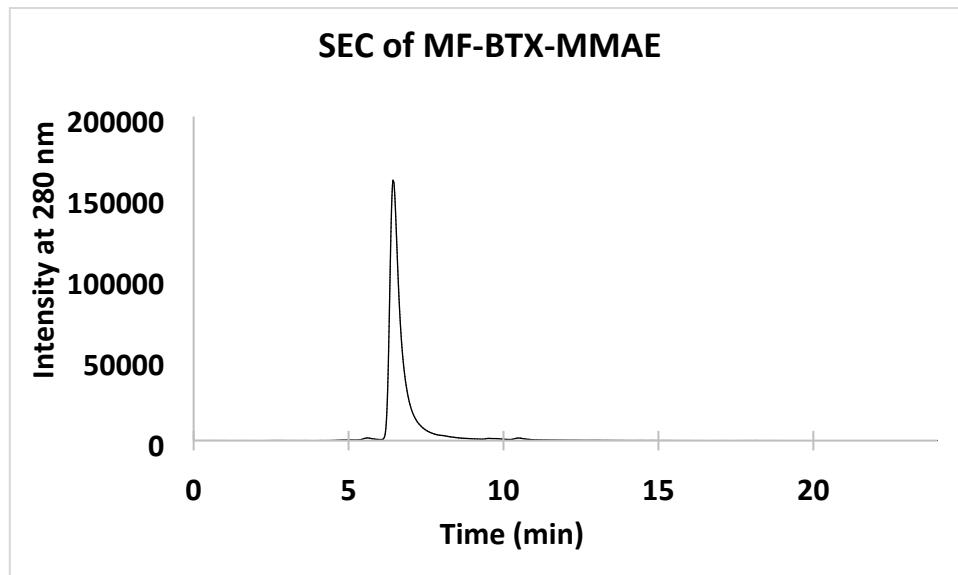
Fragments observed	DAR observed	Expected MW (Da) ¹	Measured MW (Da) ¹	Intensity	Relative intensity (%)
HH	3	104744	104737	7408	29.3
LHH	4	129832	129831	4272	16.9
LHHL	4	153555	153553	622804	100.00

Fragments	LHHL	LHH	HH	LH	H	L
Average DAR	4	4	3	2.02	-	0

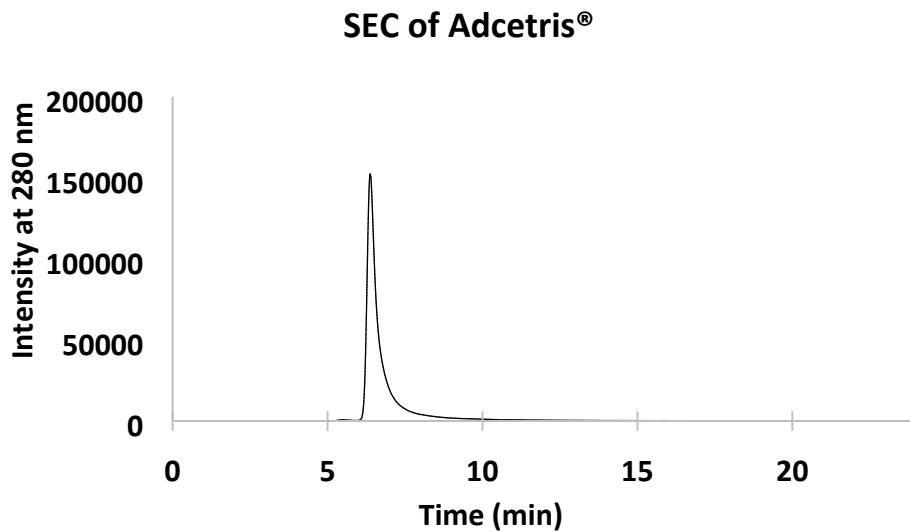
¹Only the first glycosylation peak (GOF) was considered.

The mass increment of 1368 Da expected for the conjugated linker is confirmed.

Figure S4. MF-BTX-MMAE denaturing mass spectrometry analysis.



MF-BTX-MMAE	Aggregates	Oligomers	Monomer
Retention time(min)	2.71	5.59	6.44
Area (%)	0.05	1.4	98.55



Adcetris®	Aggregates	Oligomers	Monomer
Retention time(min)	2.77	5.49	6.38
Area (%)	0.04	0.45	99.51

Figure S5. HPLC SEC-UV analysis of **MF-BTX-MMAE** and **Adcetris®**.

Supporting Information

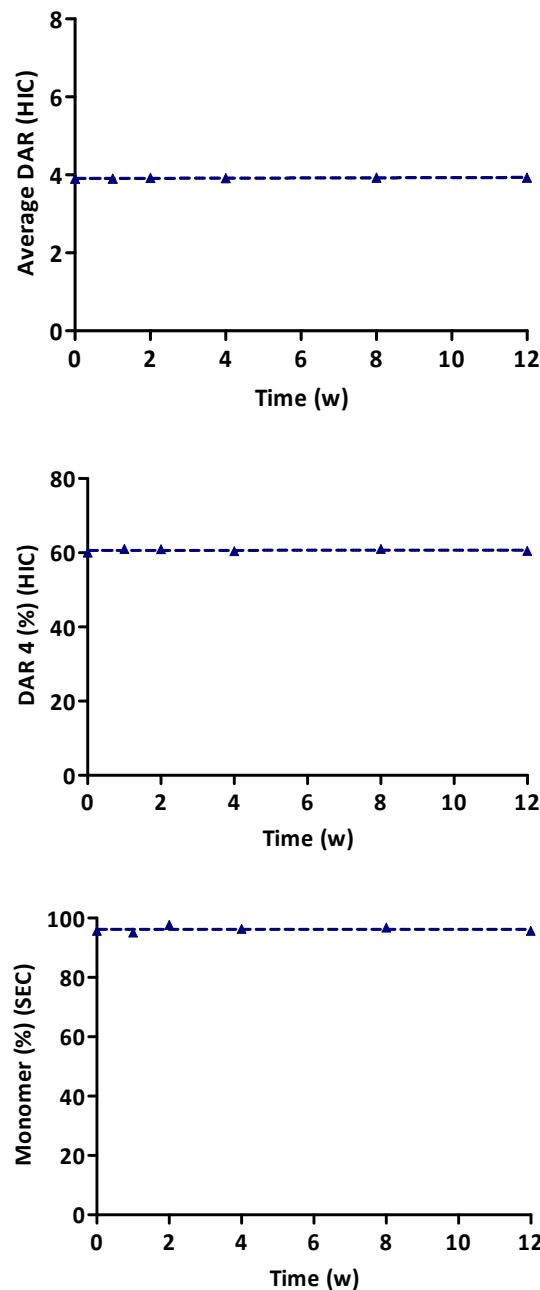


Figure S6. Stability of **MF-BTX-MMAE** upon storage at 4 °C in PBS pH 7.4 for 12 weeks ($N = 1$).

Supporting Information

Name	Events	% of Vis	Mean	GeoMean	Median	CV
Hilary Ab only PE anti-human	8650	100.00	3.96	3.36	3.19	115.50
Isotype control ADC PE anti-human	8815	100.00	5.09	4.08	3.85	284.87
Adcetris® PE anti-human	8809	100.00	2829.78	2512.67	2617.99	45.10
MF-BTX-MMAE PE anti-human	8595	100.00	2693.90	2396.49	2525.48	44.85

Figure S7. Antigen-binding analysis data by FACS on Karpas-299 (CD30-positive).

Supporting Information

High Content Analysis (Operetta Images)

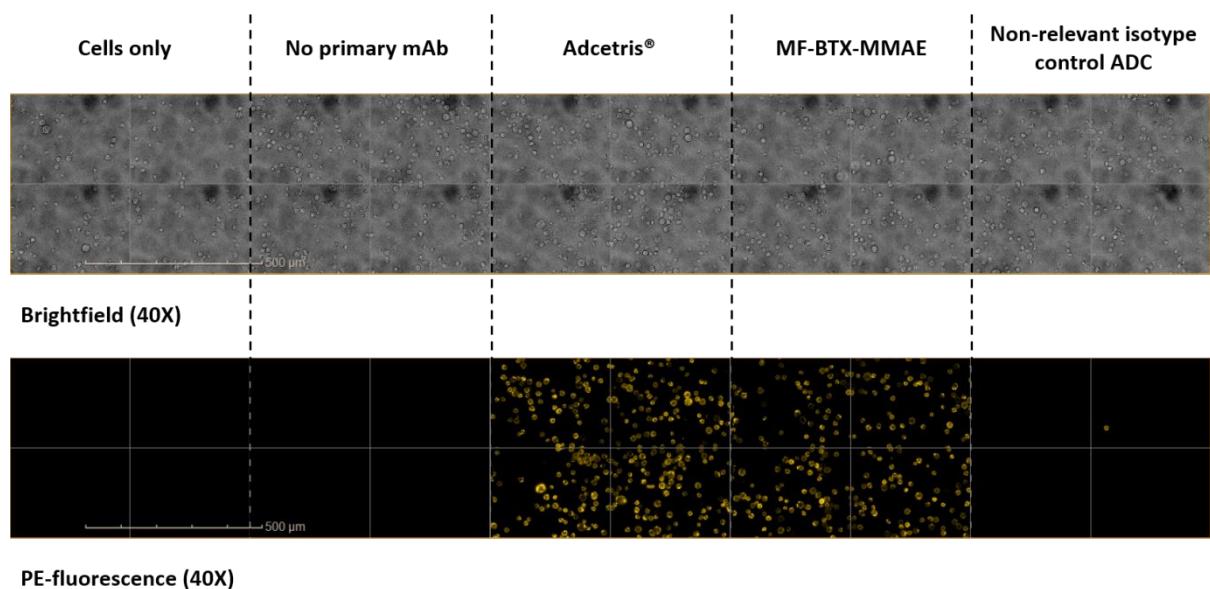


Figure S8. Antigen-binding analysis by high content analysis on Karpas-299 (CD30-positive).

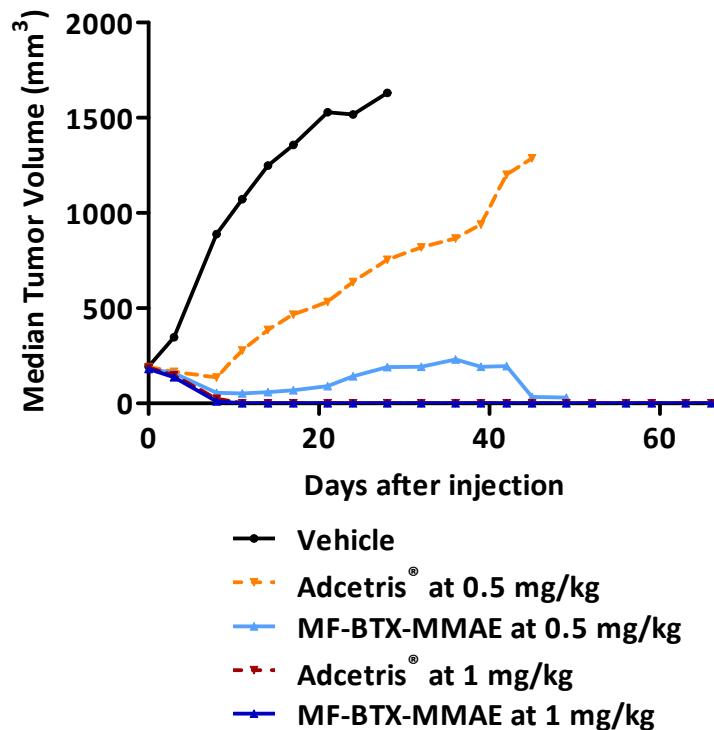


Figure S9. Median tumor volume curves of CB17-SCID mice bearing subcutaneous Karpas-299 human lymphoma tumors. Each point represents the median of the recorded tumor volume per group. Animals were randomized and treated on D0. Last mice were sacrificed on D66. Graphs are shown when at least 80% of the animals are alive.

Group	Data	D0	D3	D8	D11	D14	D17	D21	D24	D28	D32	D36
Vehicle	Median	194	347	889	1072	1250	1358	1529	1517	1631	1521	1401
	T/C%	100	100	100	100	100	100	100	100	100	100	100
Adcetris® 0.5 mg/kg	Median	191	166	135	277	385	465	532	637	755	819	865
	T/C%	98	48	15	26	31	34	35	42	46	54	62
Adcetris® 1 mg/kg	Median	187	147	24	0	0	0	0	0	0	0	0
	T/C%	96	42	3	0	0	0	0	0	0	0	0
MF-BTX-MMAE 0.5 mg/kg	Median	183	159	56	52	60	69	91	143	191	193	230
	T/C%	94	46	6	5	5	5	6	9	12	13	16
MF-BTX-MMAE 1 mg/kg	Median	181	137	10	0	0	0	0	0	0	0	0
	T/C%	93	39	1	0	0	0	0	0	0	0	0

Figure S10. Tumor growth inhibition (T/C%) results in treated CB17-SCID mice bearing subcutaneous Karpas 299 human lymphoma tumors. Animals were randomized and treated on D0. Last mice were sacrificed on D66. No further calculations were possible after D36 since there were less than four mice remaining in the vehicle group.

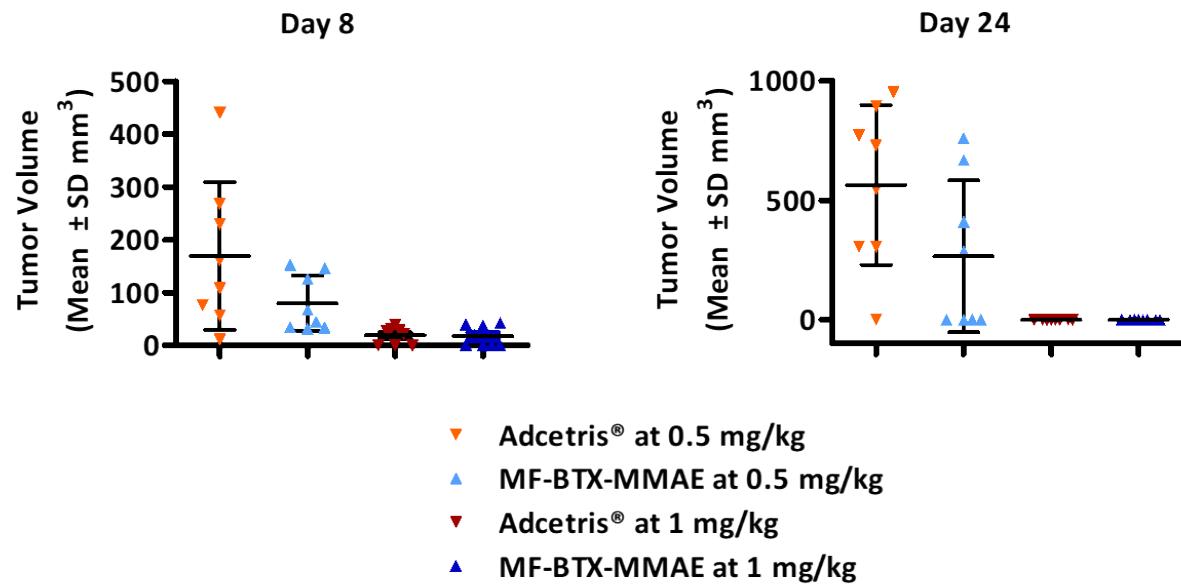


Figure S11. Individual tumor volumes of CB17-SCID mice bearing subcutaneous Karpas 299 human lymphoma tumors on D8 and D24. Animals were randomized and treated on D0. Last mice were sacrificed on D66.

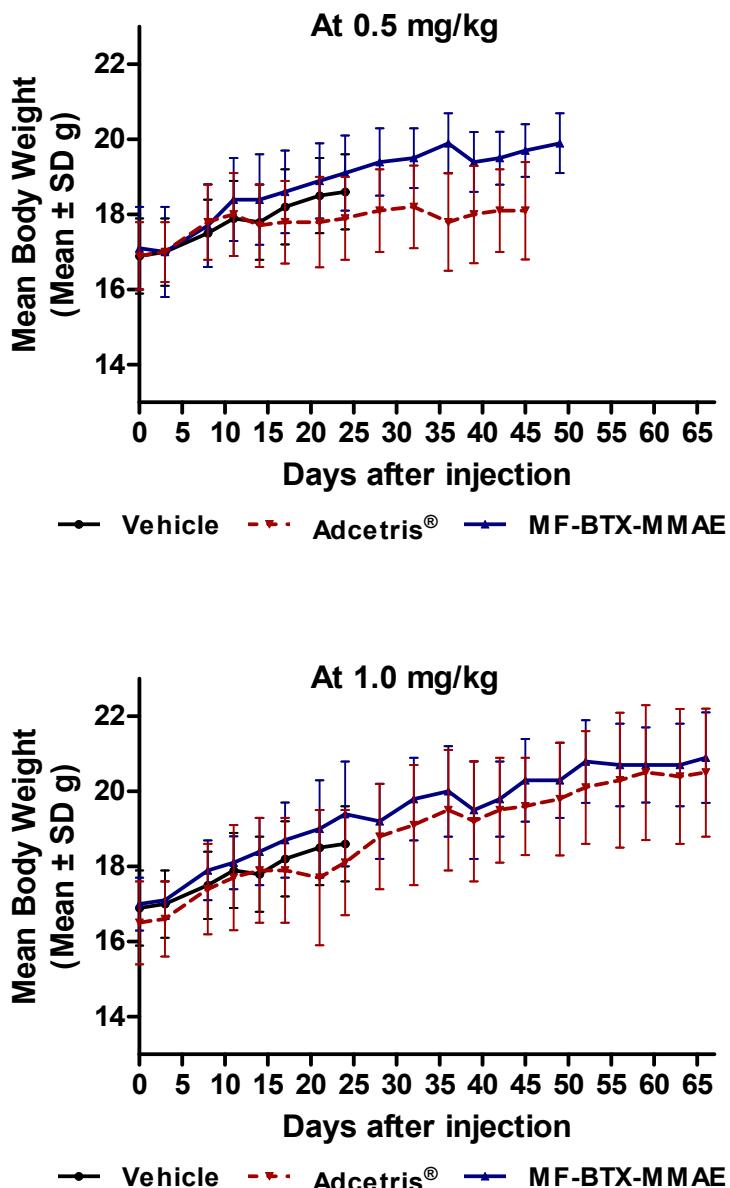
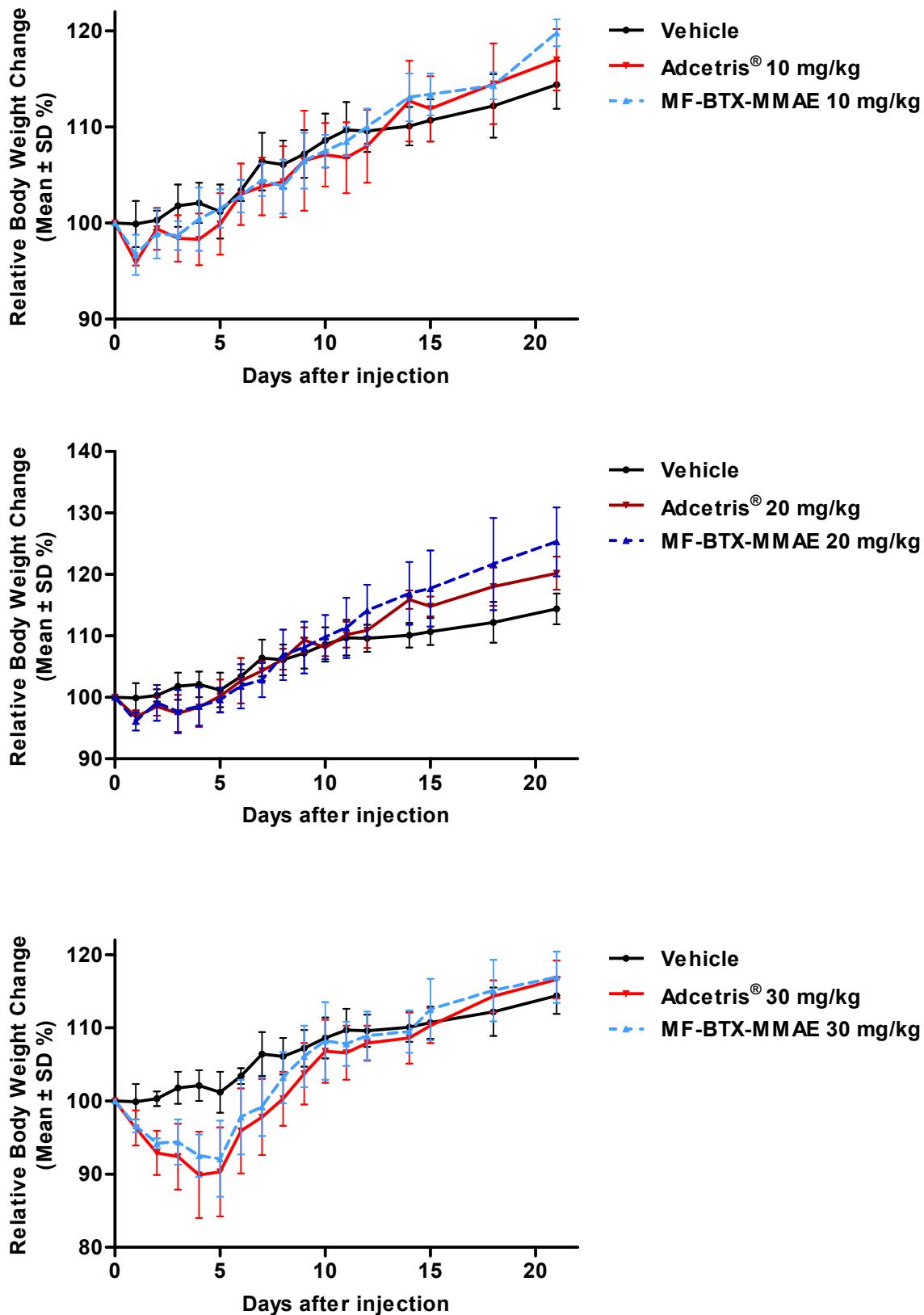


Figure S12. Mean body weight curves of CB17-SCID mice bearing subcutaneous Karpas 299 human lymphoma tumors treated with a single dose on day 0 either with vehicle, Adcetris® or MF-BTX-MMAE at 0,5 and 1 mg/kg. Each point represents the mean of the recorded body weight per group. Graphs are shown when more than 80% of the animals are alive. Data are mean ± SD, N=16 for vehicle and N=8 for other groups.

Supporting Information



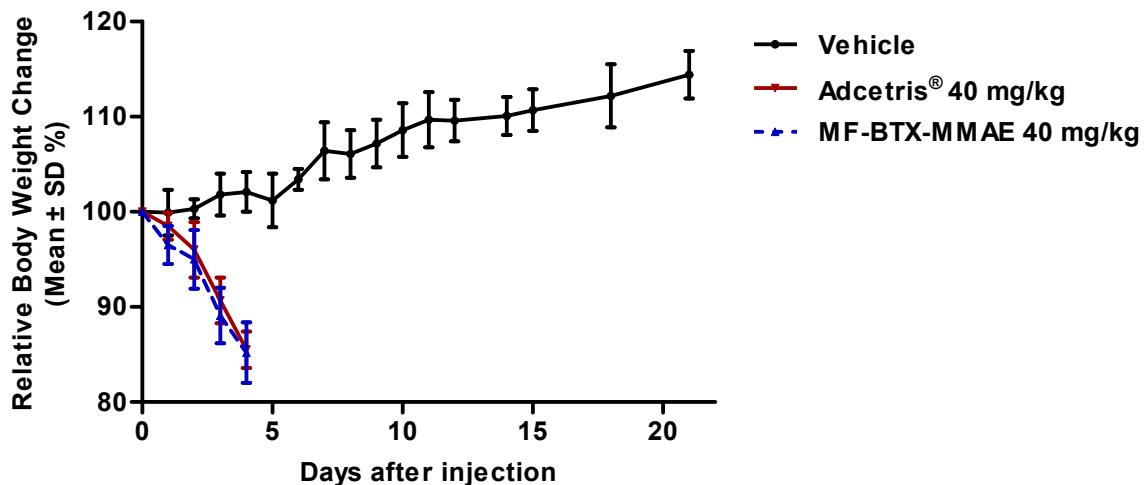


Figure S13. Relative body weight change curves of Sprague-Dawley rats treated with a single dose on day 0 either with vehicle, Adcetris® or **MF-BTX-MMAE** at 10, 20, 30 and 40 mg/kg. Each point represents the mean of the recorded relative body weight change per group. Data are mean \pm SD, N=3 for vehicle, N=9 for **MF-BTX-MMAE** at 40 mg/kg and N=5 for other groups.