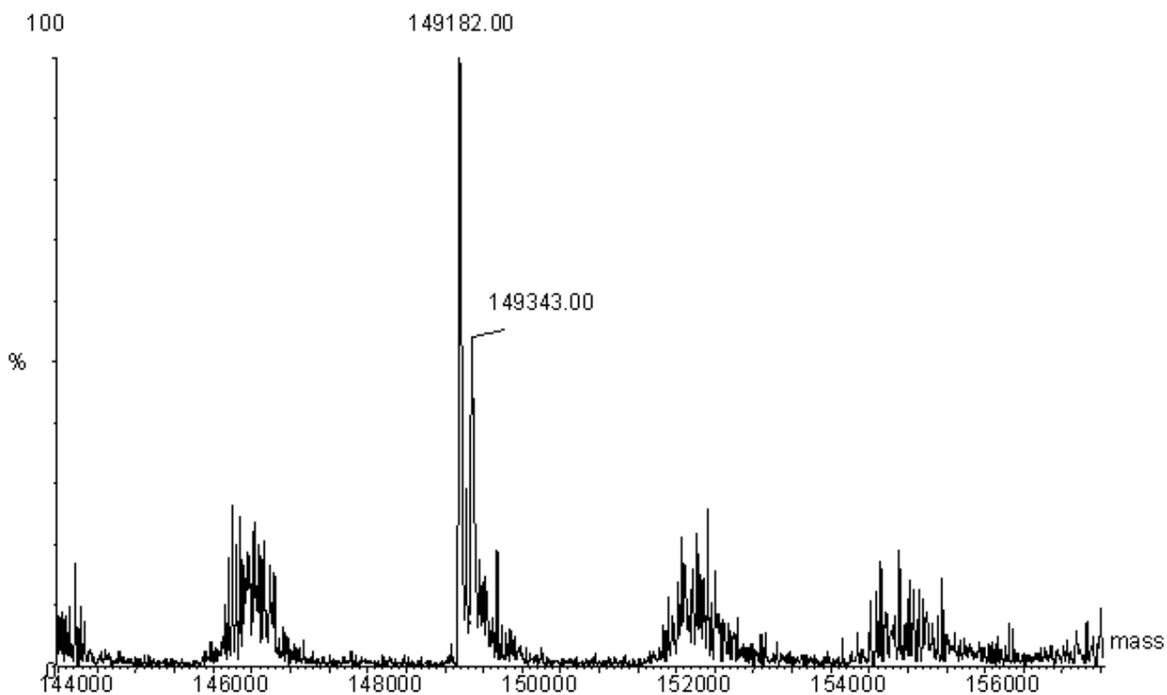


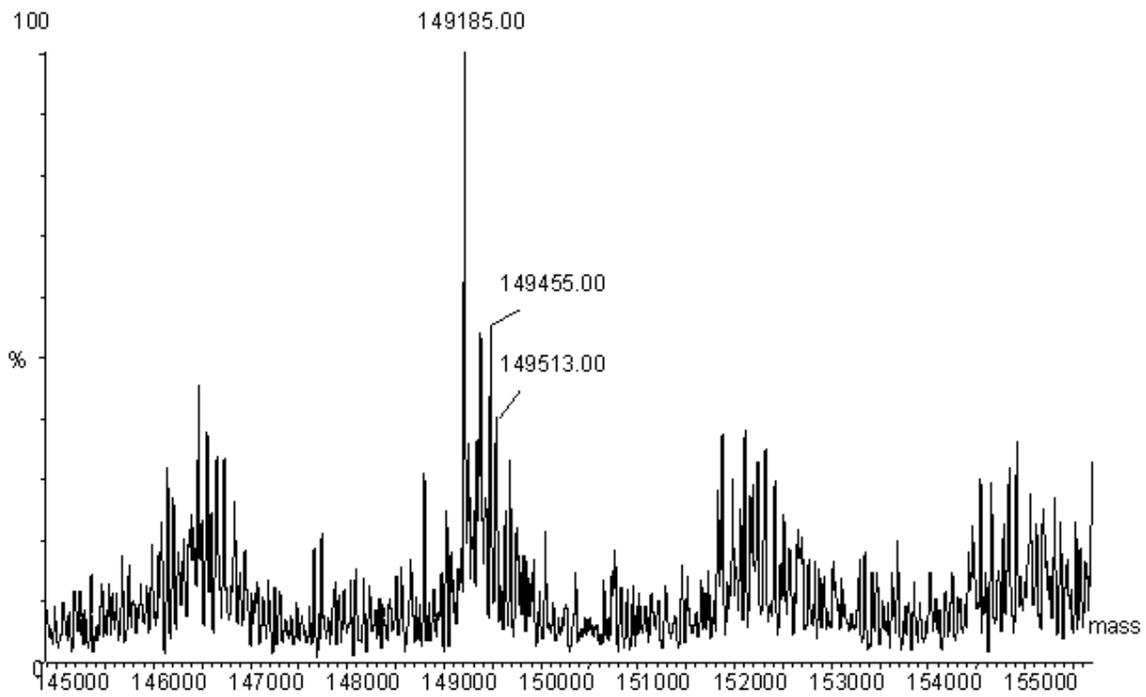
Supporting information

Figure S1. Mass spectra of *ex in vivo* Bevacizumab samples after incubation in vitreous humour from (a) day 0, (b) day 7 to (c) day 14. Structural heterogeneity peaks were found along with time.

(a)



(b)



(c)

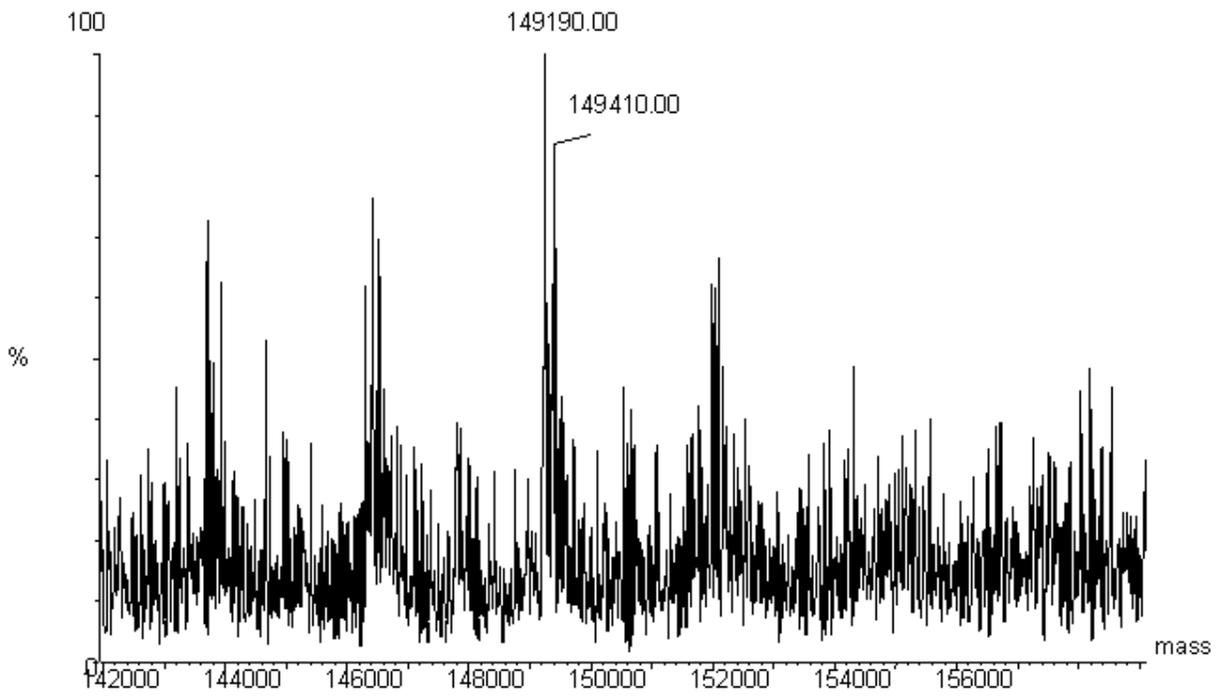
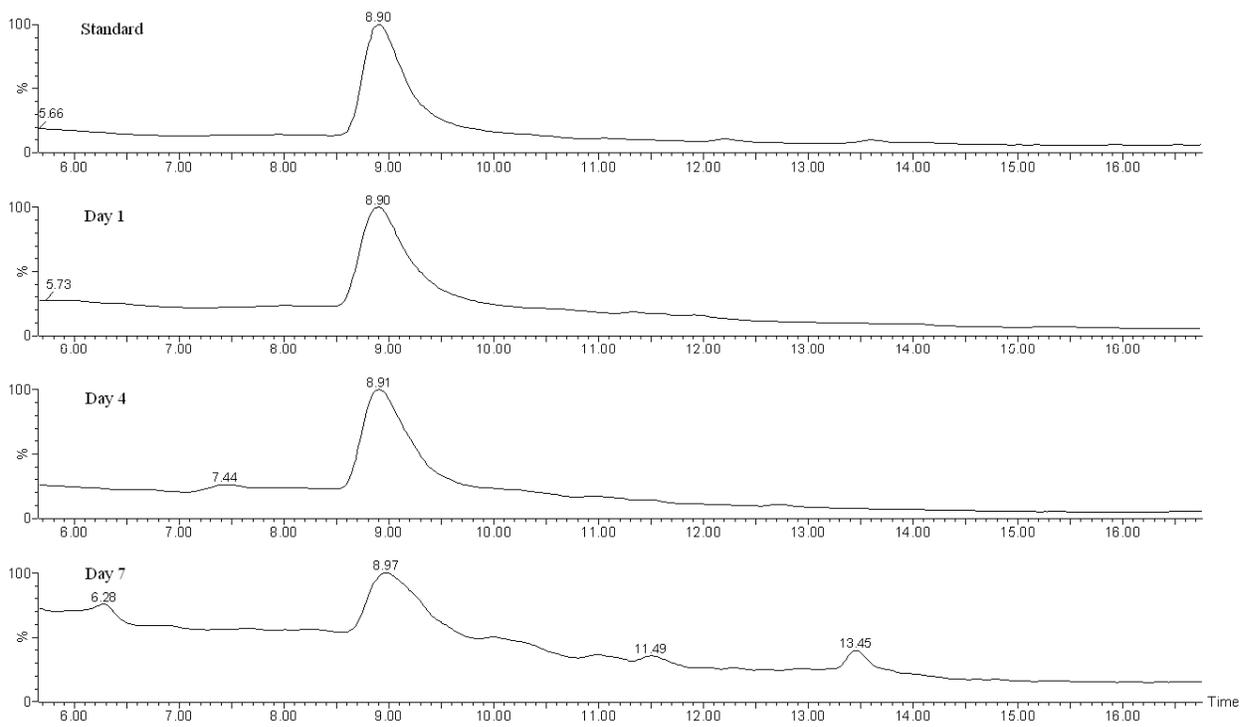
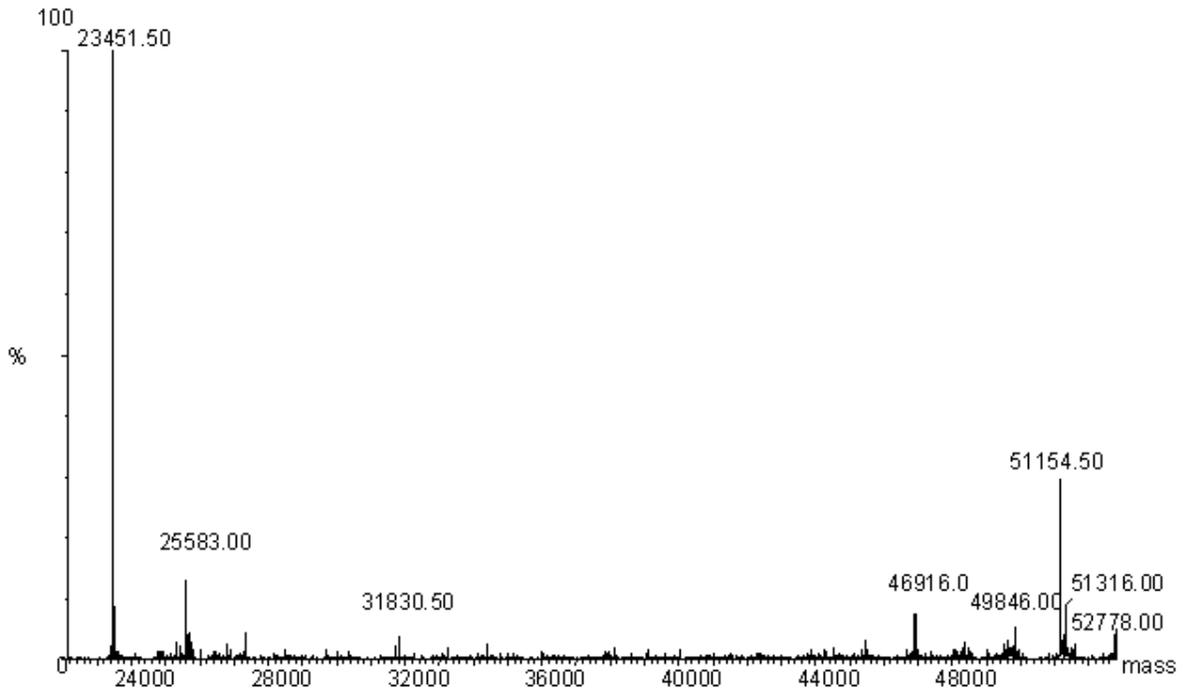


Figure S2. Chromatograms (a) and mass spectra of standard Bevacizumab spiked in vitreous humor and vitreous samples from 1, 4 and 7 days after intra-vitreous injection following isolation and DTT reduction. The heavy and light chains were co-eluted. (b-e) They showed the respective mass spectrum of the Bevacizumab from spiked standard, day 1 to day 7. (b) It showed mass 23451.5 m/z was the mass of light chain, the mass of 51154.5 m/z was the mass of heavy chain and the mass 51316 m/z was the mass of heavy chain with an additional galactose on the glycan chain. (c) It showed similar mass spectrum whereas the heavy was becoming heterogeneous from (d – e).

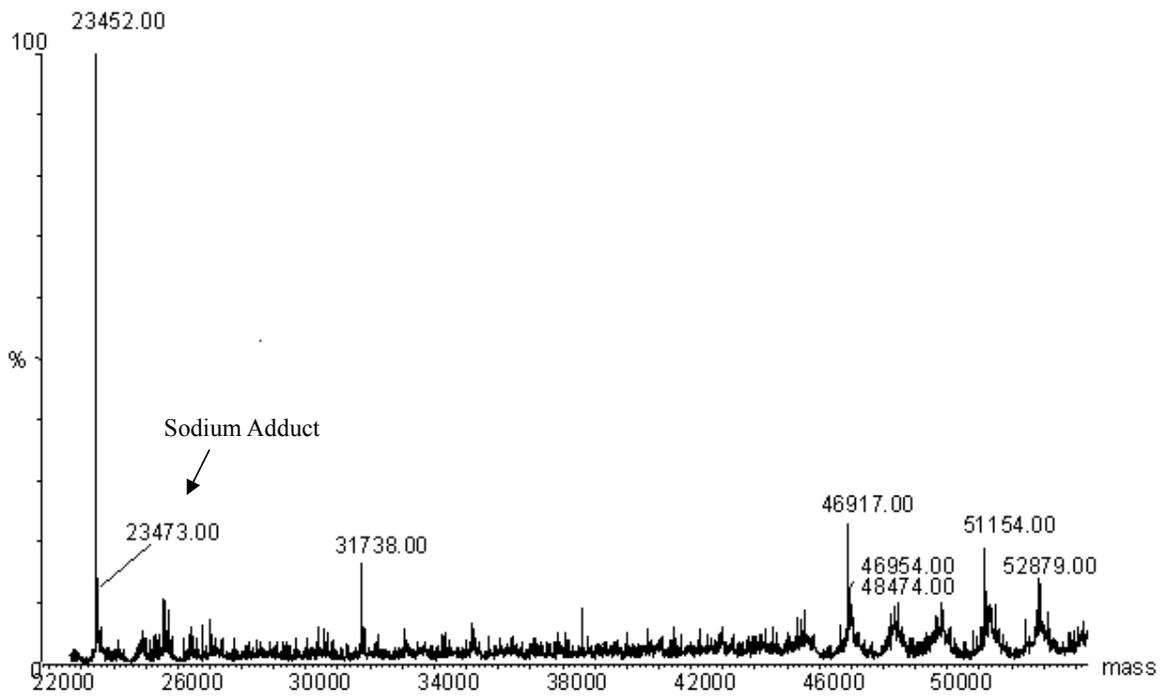
(a)



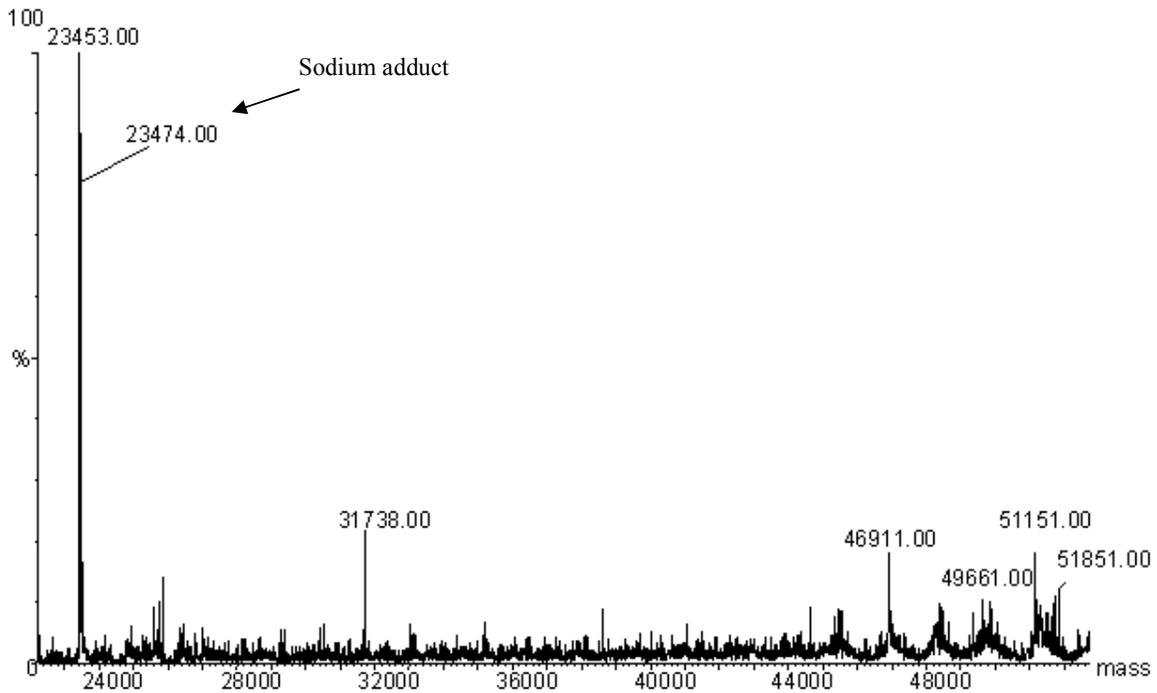
(b)



(c)



(d)



(e)

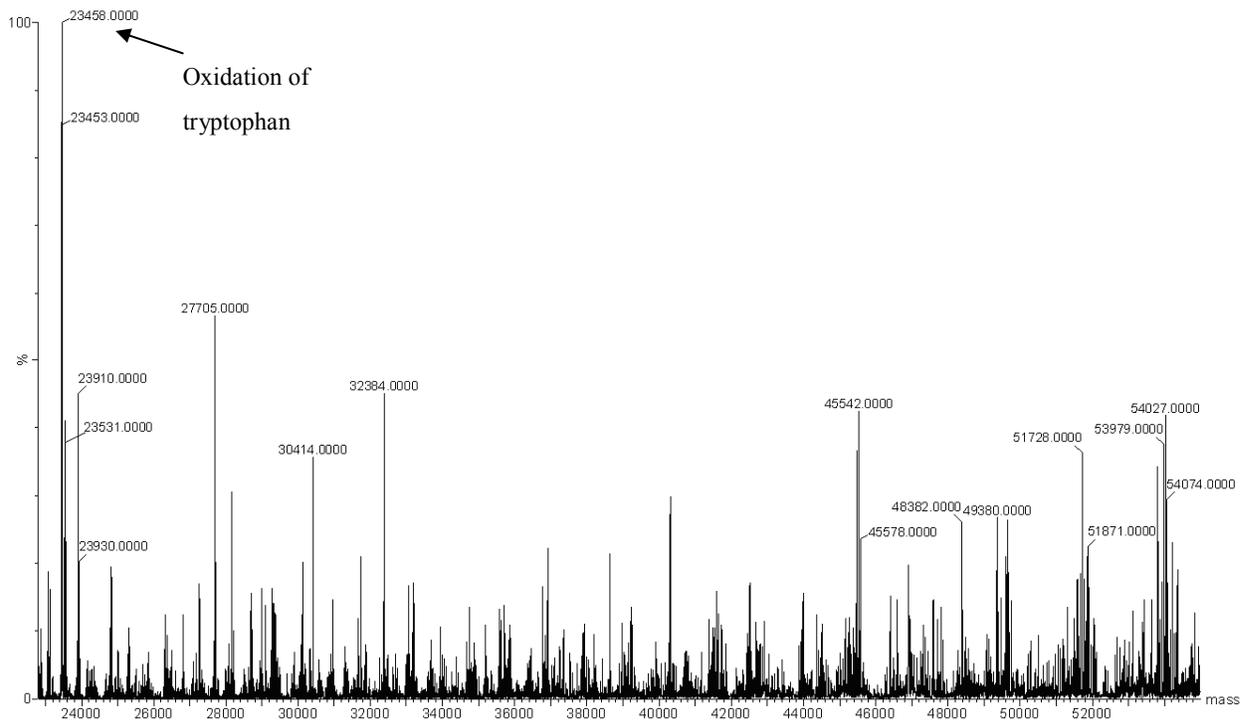


Figure S3. LCMS chromatograms of spiked Bevacizumab in vitreous humor and 1, 4 and 7 days after intravitreal injection following Poros G/20 isolation, deglycosylated and DTT reduction.

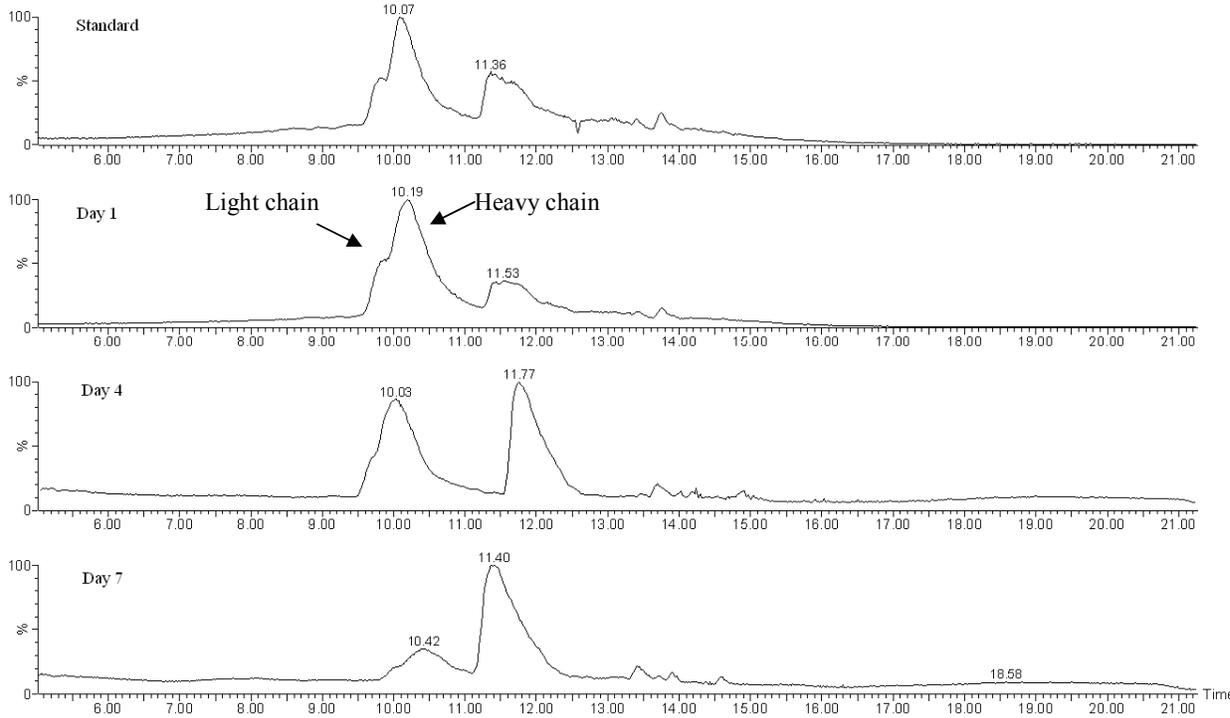


Figure S4. Mass spectrum of spiked Bevacizumab in vitreous humor after deglycosylation by PNGase F. The mass was the mass of intact Bevacizumab minus two times of Go glycan.

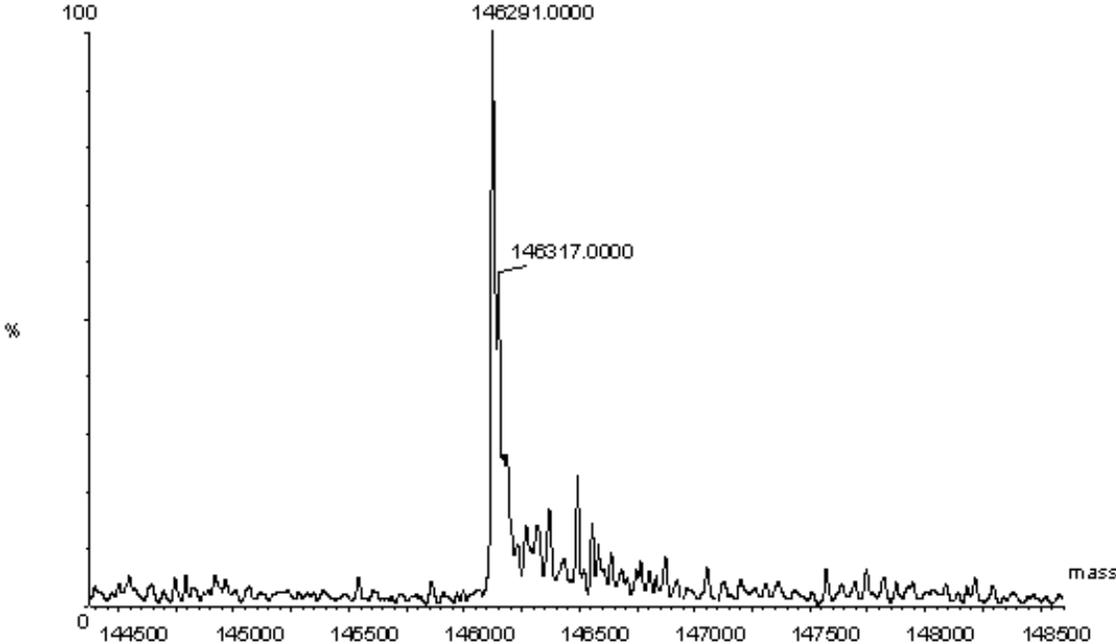


Figure S5. Difference of relative Ka at day 1, 7, and 21 respected to the relative Ka at 4 hours after intra-vitreous injection of three rabbit eyes. The symbol * shows significant difference between the results of day 21 to day 1 and day 7 calculated by ANOVA post hoc Dunnett test ($p < 0.05$).

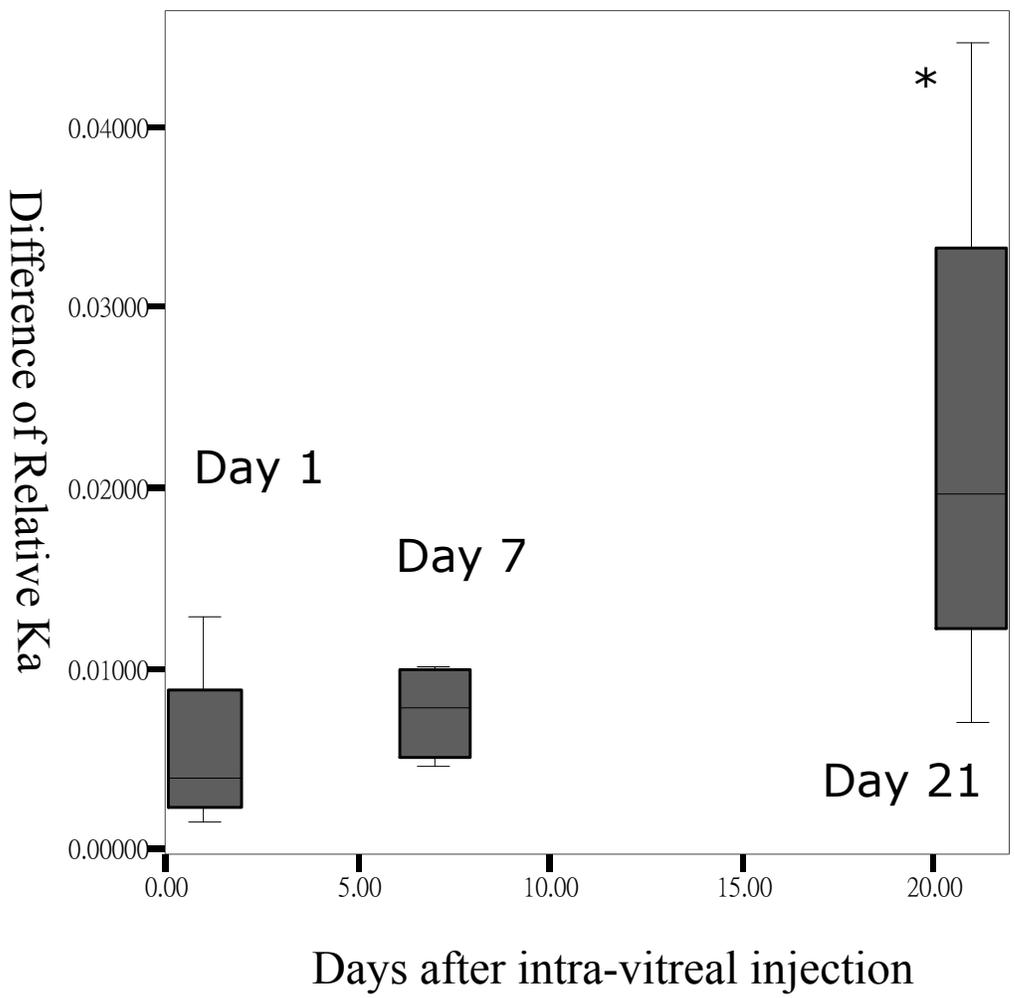


Figure S6. LCMS chromatograms of Bevacizumab after purified by Poros G/20 affinity column and trypsin digestion treatment. The samples were Bevacizumab standards and vitreous samples after different days of intra-vitreous injection.

