

Supporting Information

MiS-MALDI: Microgel-Selected detection of protein biomarkers by MALDI-ToF mass spectrometry

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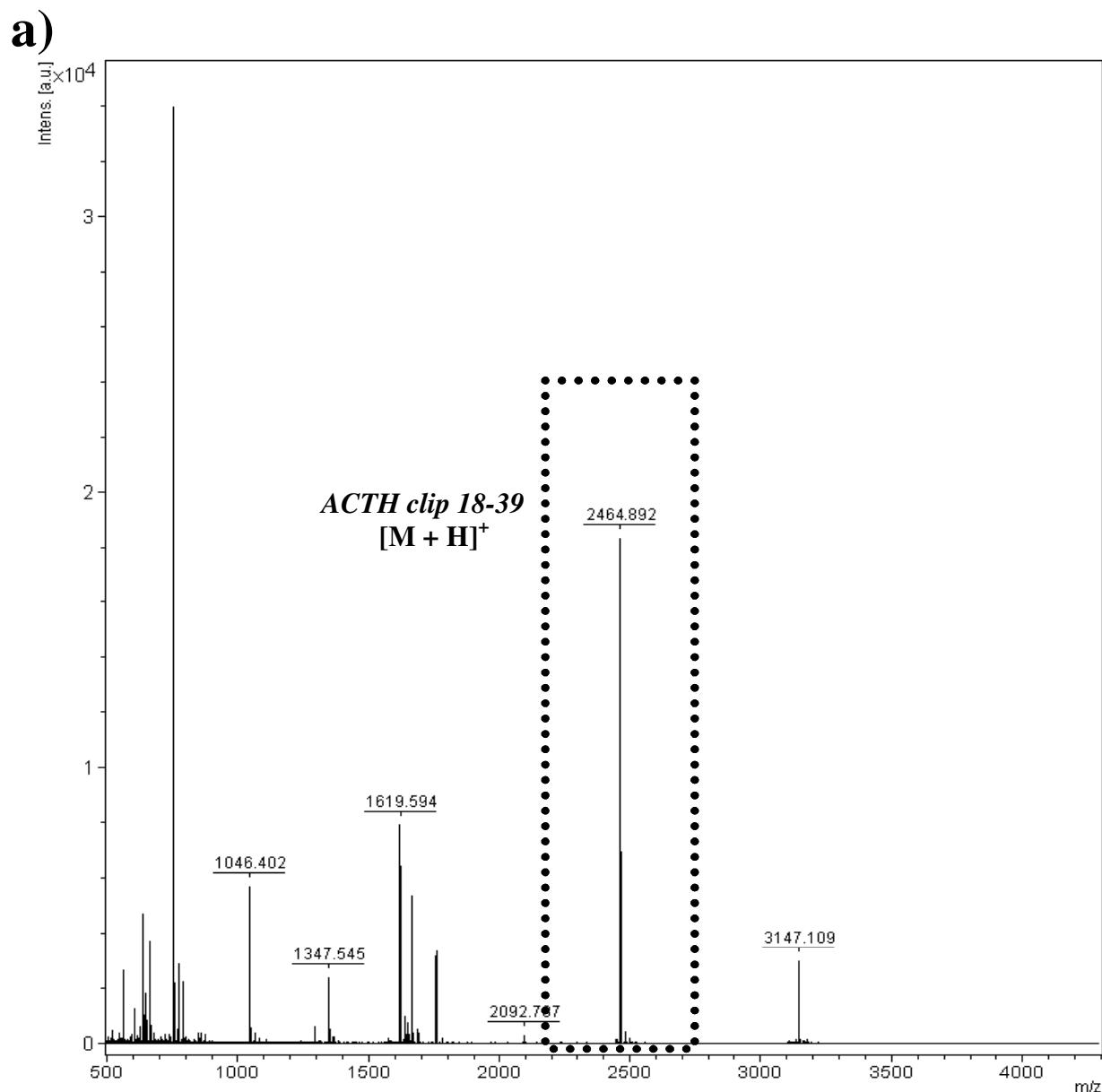
Table S1. Peptides analysed by MiS-MALDI.

Peptide calibrant	$[M+H]^+$, m/z		Sequence	Theoretical pI	Captured by microgel	Theoretical R_H (nm)
	calc.	found				
<i>Angiotensin II</i>	1046.2	1046.5	DRVYIHPF	6.74	No	0.61
<i>Angiotensin I</i>	1296.5	1296.5	DRVYIHPFHL	6.92	No	0.66
<i>Substance P</i>	1348.6	1347.7	RPKPQQFFGLM	11.00	No	0.68
<i>Bombesin</i>	1620.8	1619.8	Pyr ^a -QRLGNQWAVGHL	9.76	No	0.73
<i>ACTH clip 1-17</i>	2093.4	2093.1	SYSMEHFRWGKPVGKKR	10.45	No	0.82
<i>gp41₆₅₉₋₆₇₁</i>	1616.8	1616.8	ELLELDKWASLWN	4.14	Yes	0.73
<i>ACTH clip 18-39</i>	2465.7	2464.9	RPVKVYPNGAEDESAAEAPLEF	4.25	Yes	0.88
<i>Somatostatin 28</i>	3150.6	3149.5	SANSNPAMAPRERKAGCKNFFWKTFTSC	9.85	No	0.97
<i>Coiled coil strand A•</i>	2301.6	2302.2	ac-YGKEIAALEQEIAALEQEIAA-am	3.98	Yes	0.85
<i>Coiled coil strand B•</i>	2160.4	2161.1	ac-LEQENAALEKEIAALEQGY-am	3.98	Yes	0.83
<i>Coiled coil strand AB</i>	4361.3	4361.7	ac-YGQKIAALKVAAALKQKIAALKQKNAALKKEKA-am	10.15	No	1.12

^a pyrrolidone carboxylic acid (Pyr)

Table S2. Series of proteins analysed by MiS-MALDI used to verify the selectivity towards charge and size.

Protein	$[M+H]^+$, m/z		Theoretical pI	Captured by microgel	Theoretical R_H , (nm)
	calc.	found			
RNase B	14700	14895	9.6	No	1.88
EDQM hGH	22124	22135	4.9	Yes	2.14
Ovalbumin	42000	40081	4.5 – 5.1	No	2.97
mAb	150000	146935	5.5	No	5.07
Aldolase (subunits)	158000 (~40000)	- (~38006)	6.6 6.7 – 8.2	No	5.18



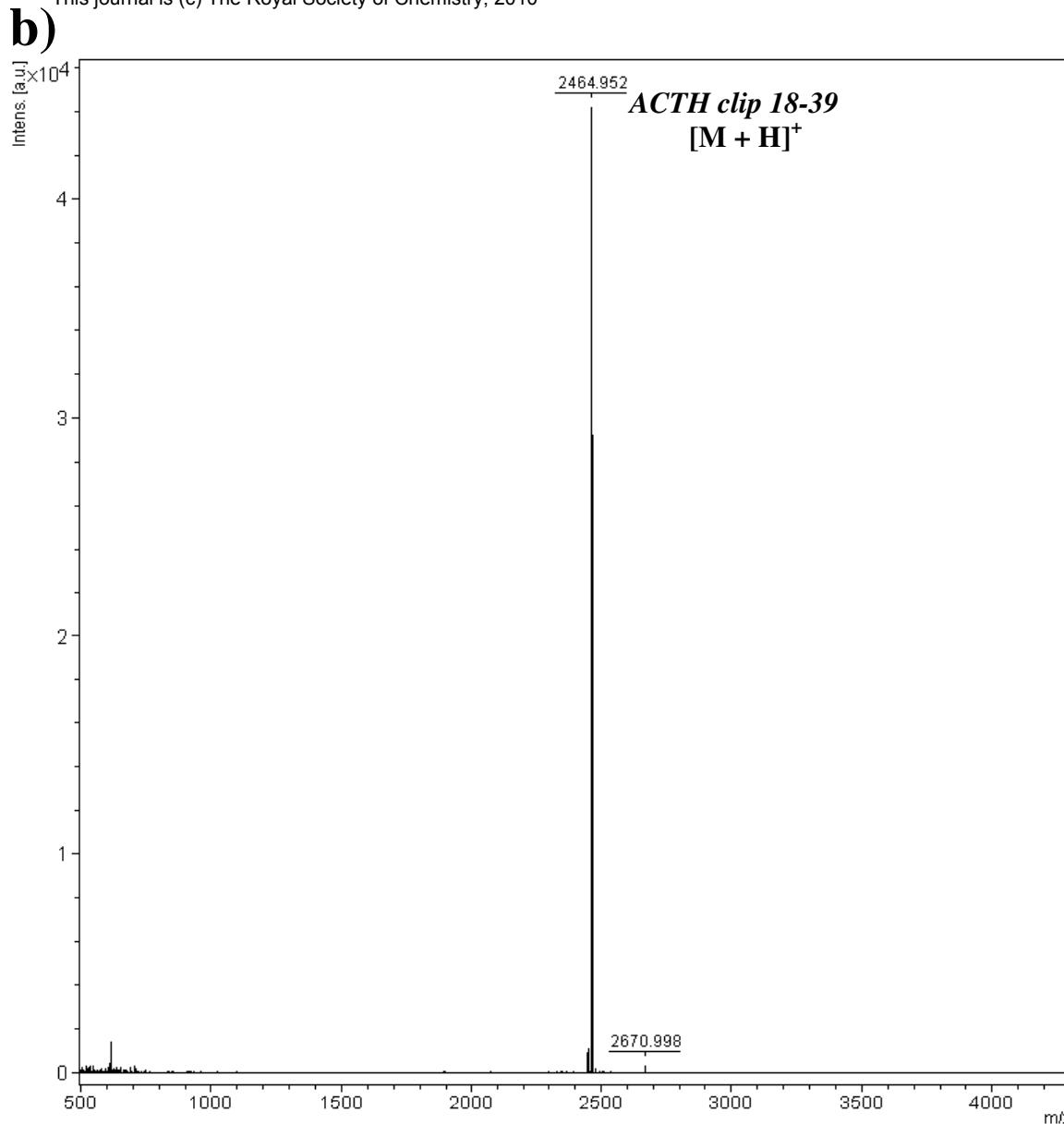
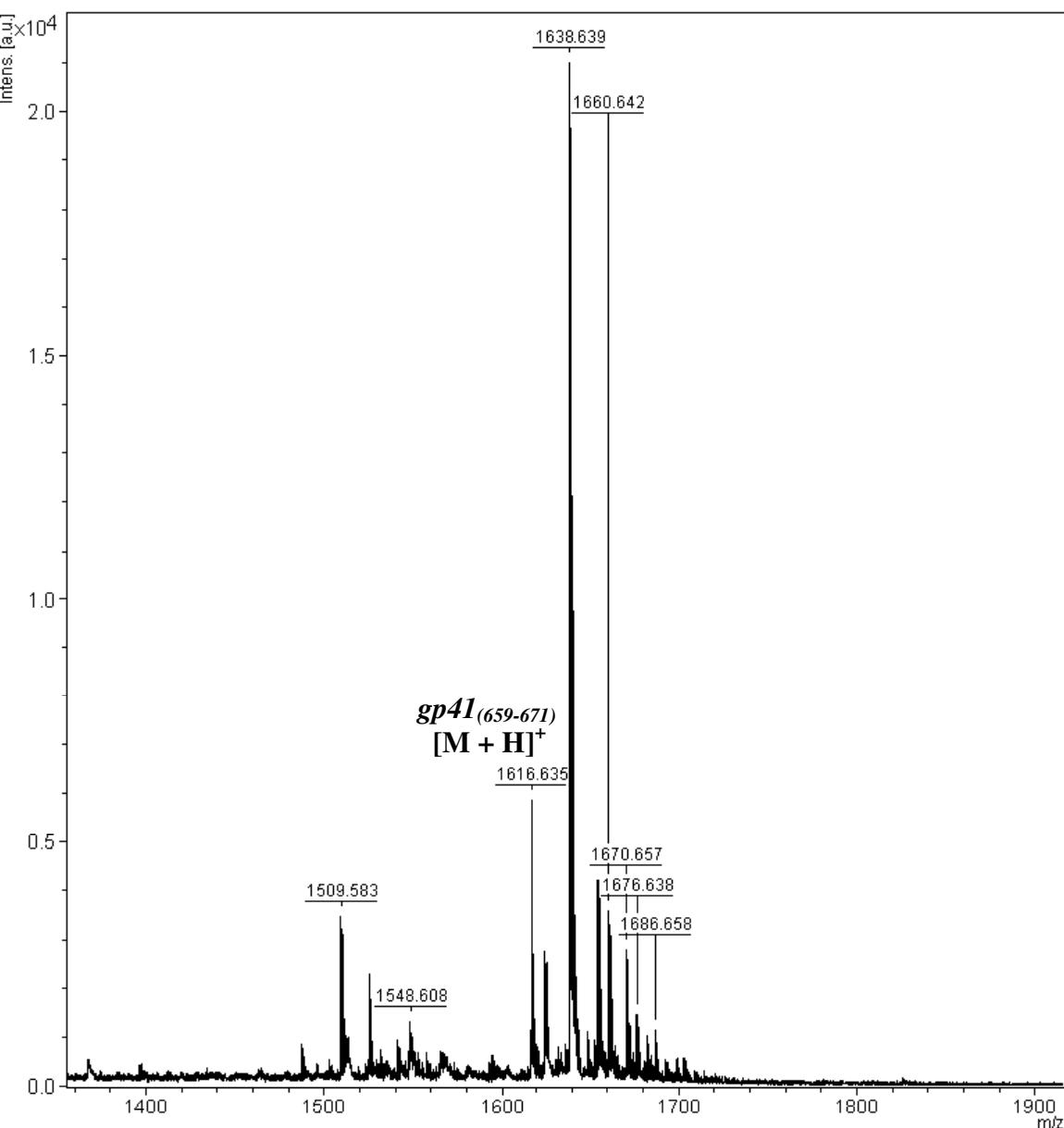


Figure S1. MALDI-ToF mass spectra of peptide calibrants obtained from solution (a) and from the microgel (b). Only ACTH clip 18-39 peptide (m/z 2465) was captured and detected in the pellet.

a)



b)

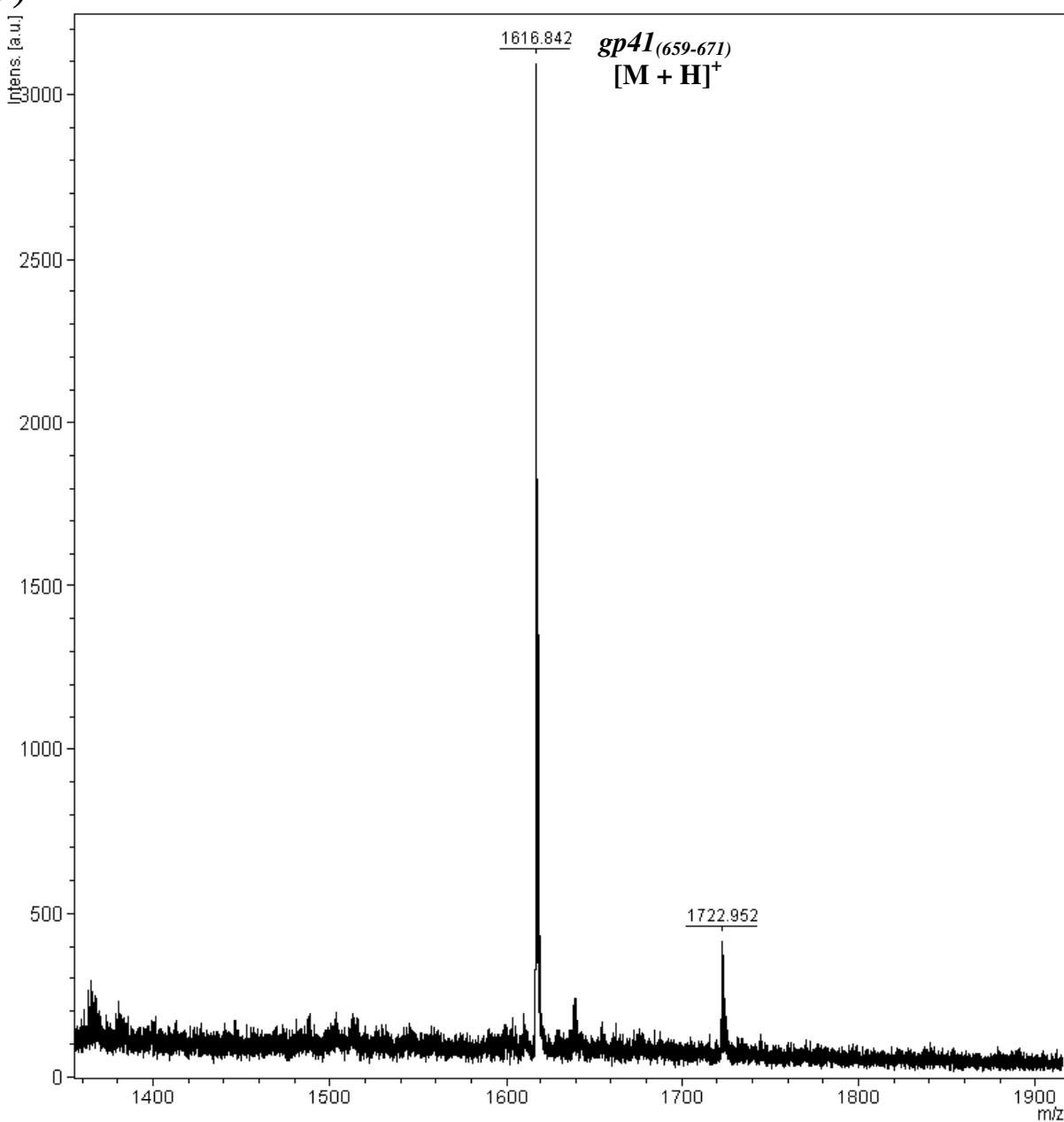


Figure S2. MALDI-ToF mass spectra for $gp41_{(659-671)}$ recorded from solution (a) and the microgel (b).

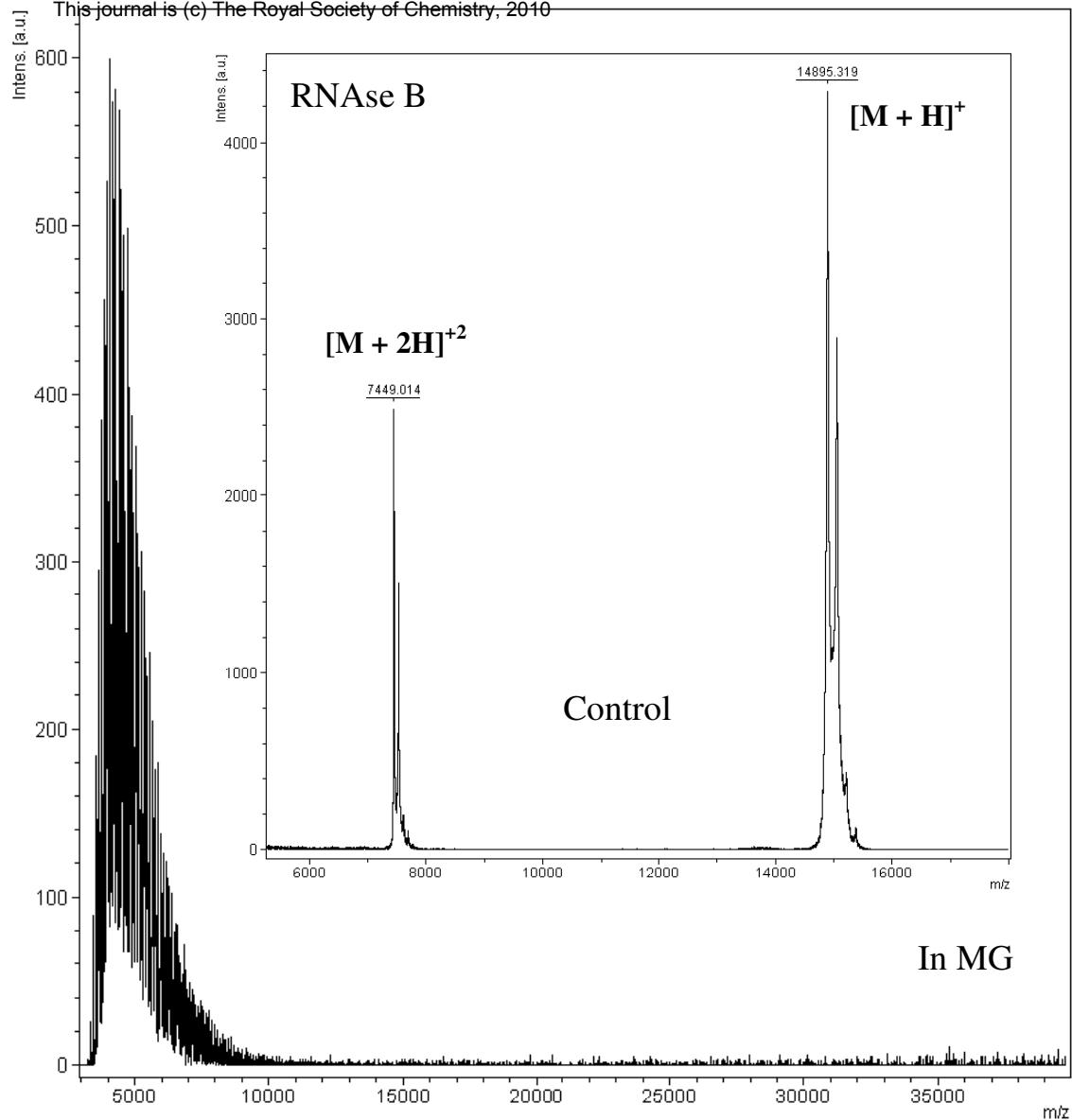


Figure S3. MALDI-ToF mass spectra of RNase B obtained from the microgel and from solution (inset) showing the charge-exclusion of the protein from the microgel.

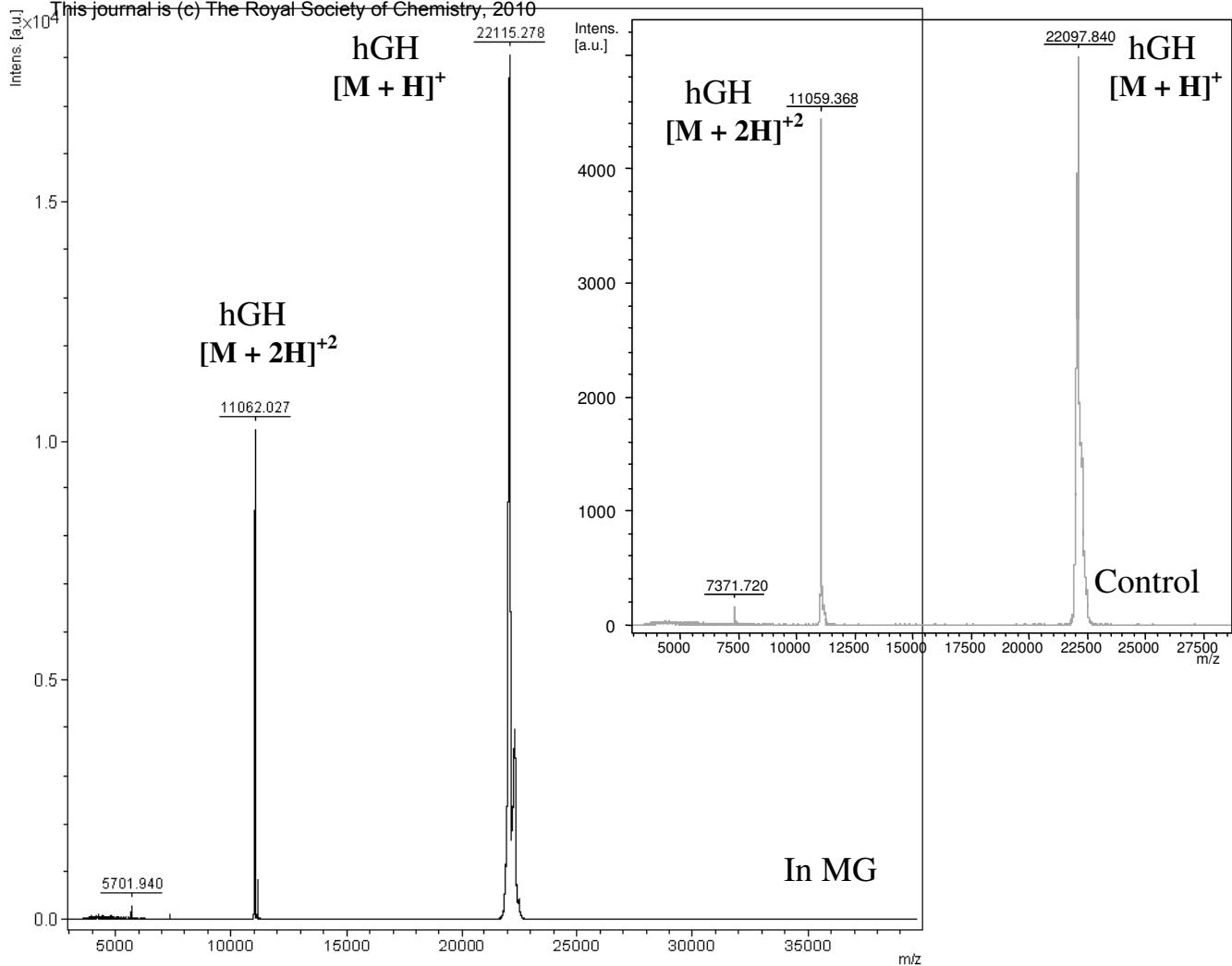


Figure S4. MALDI-ToF mass-spectra of hGH obtained from the microgel and from solution (inset) showing the inclusion of the protein in the microgel.

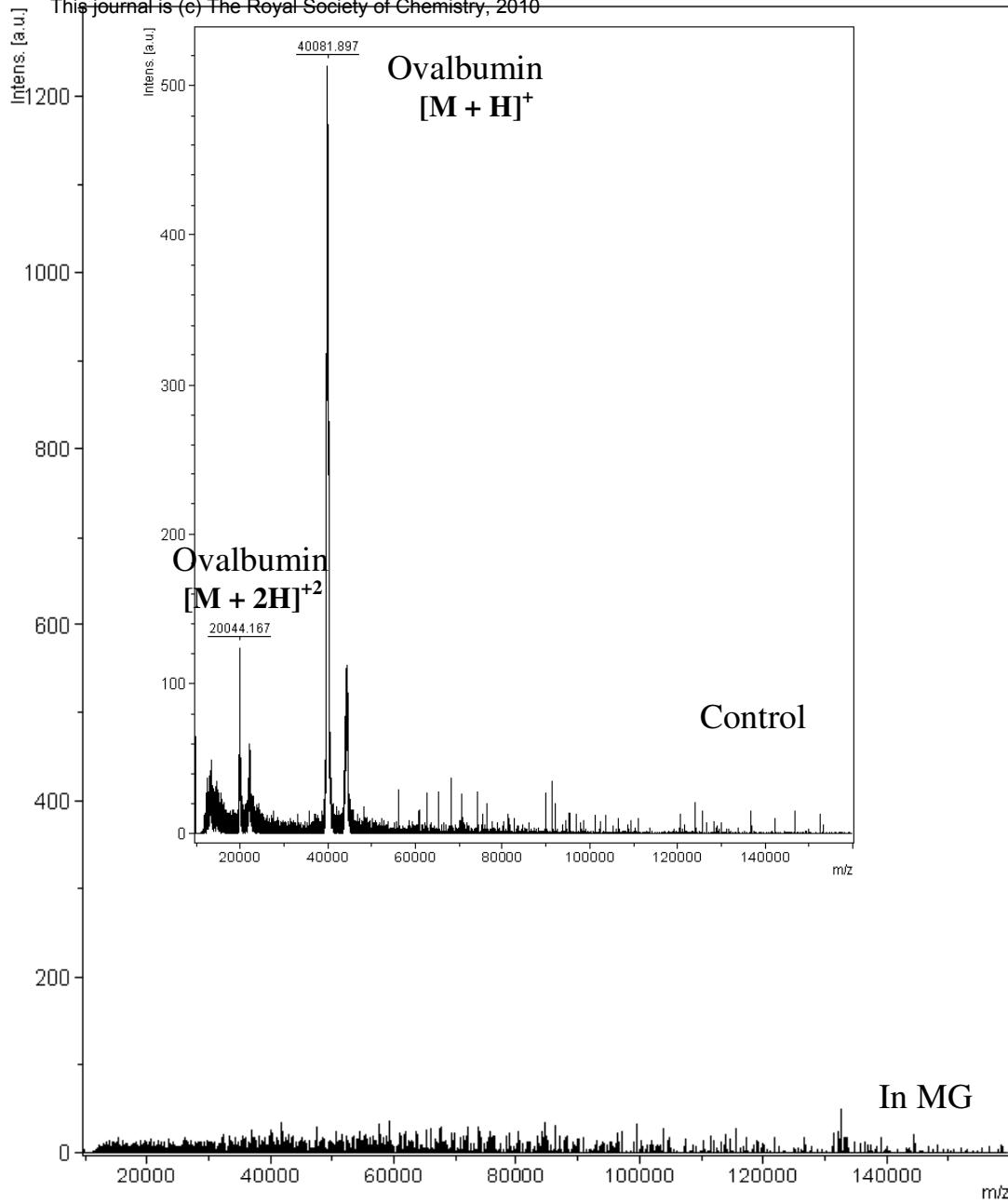


Figure S5. MALDI-ToF mass spectra of ovalbumin obtained from the microgel and from solution (inset) showing the size-exclusion of the protein from the microgel.

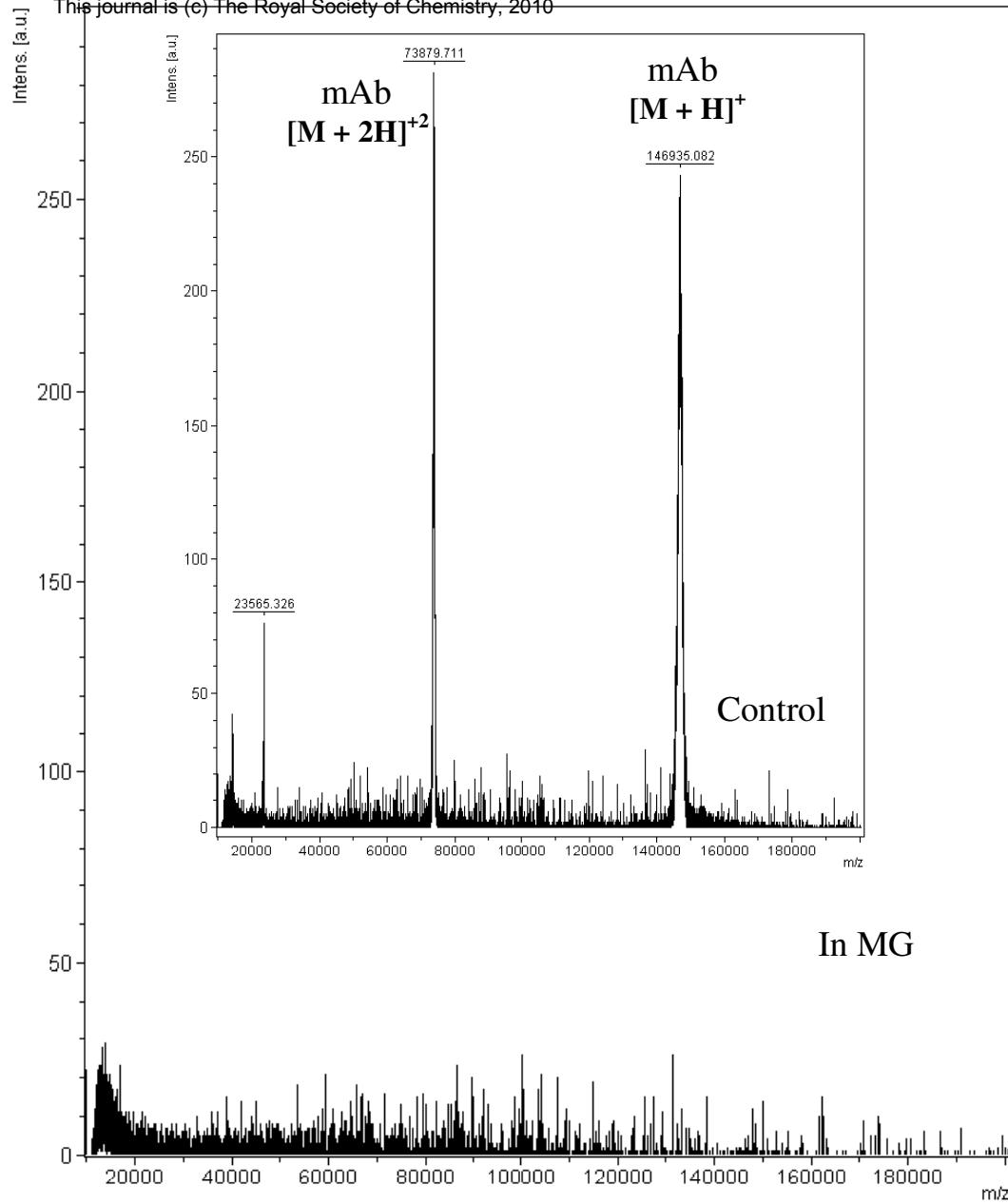


Figure S6. MALDI-ToF mass spectra of a monoclonal antibody (mAb) obtained from the microgel and from solution (inset) showing the size-exclusion of the protein from the microgel.

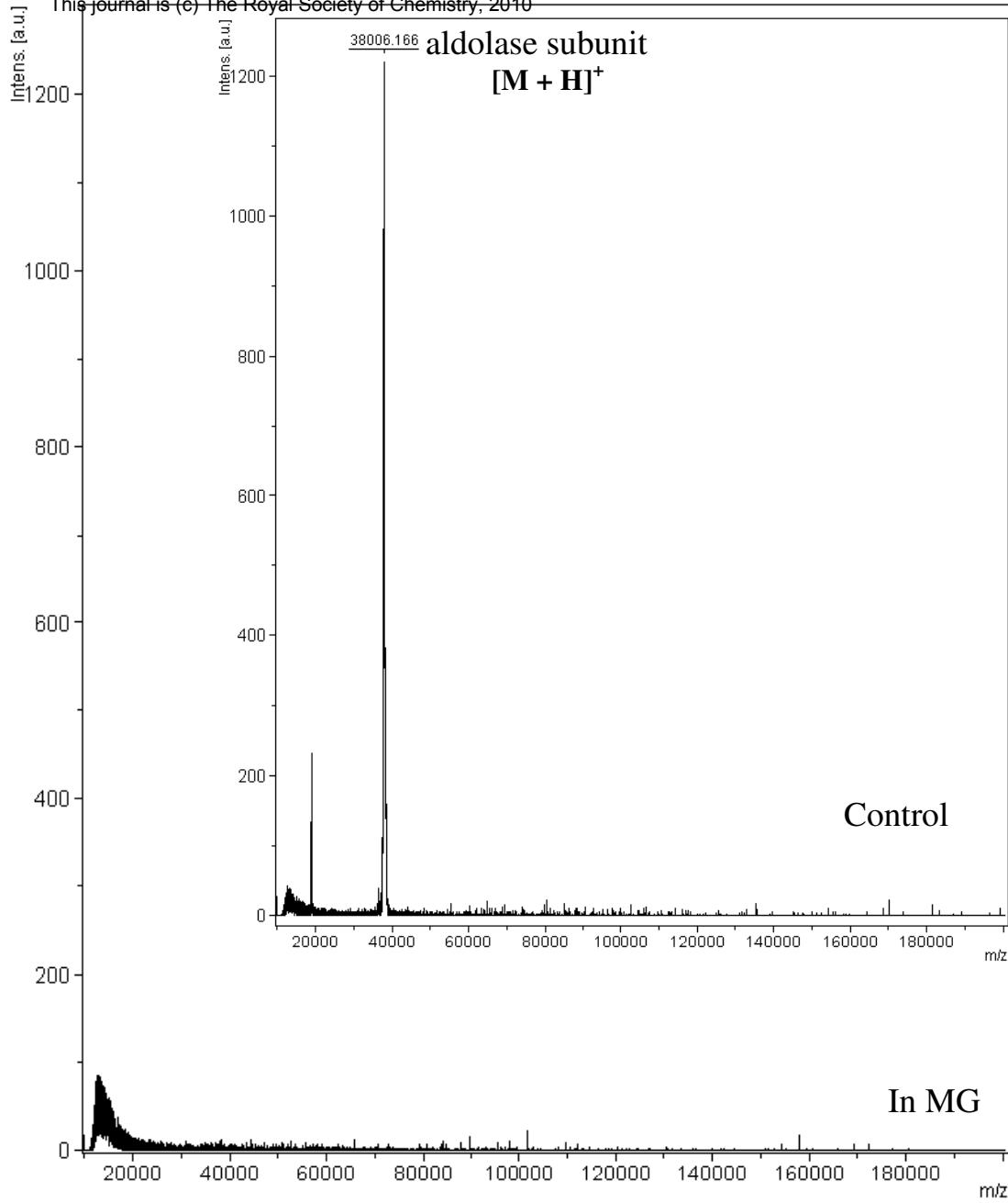
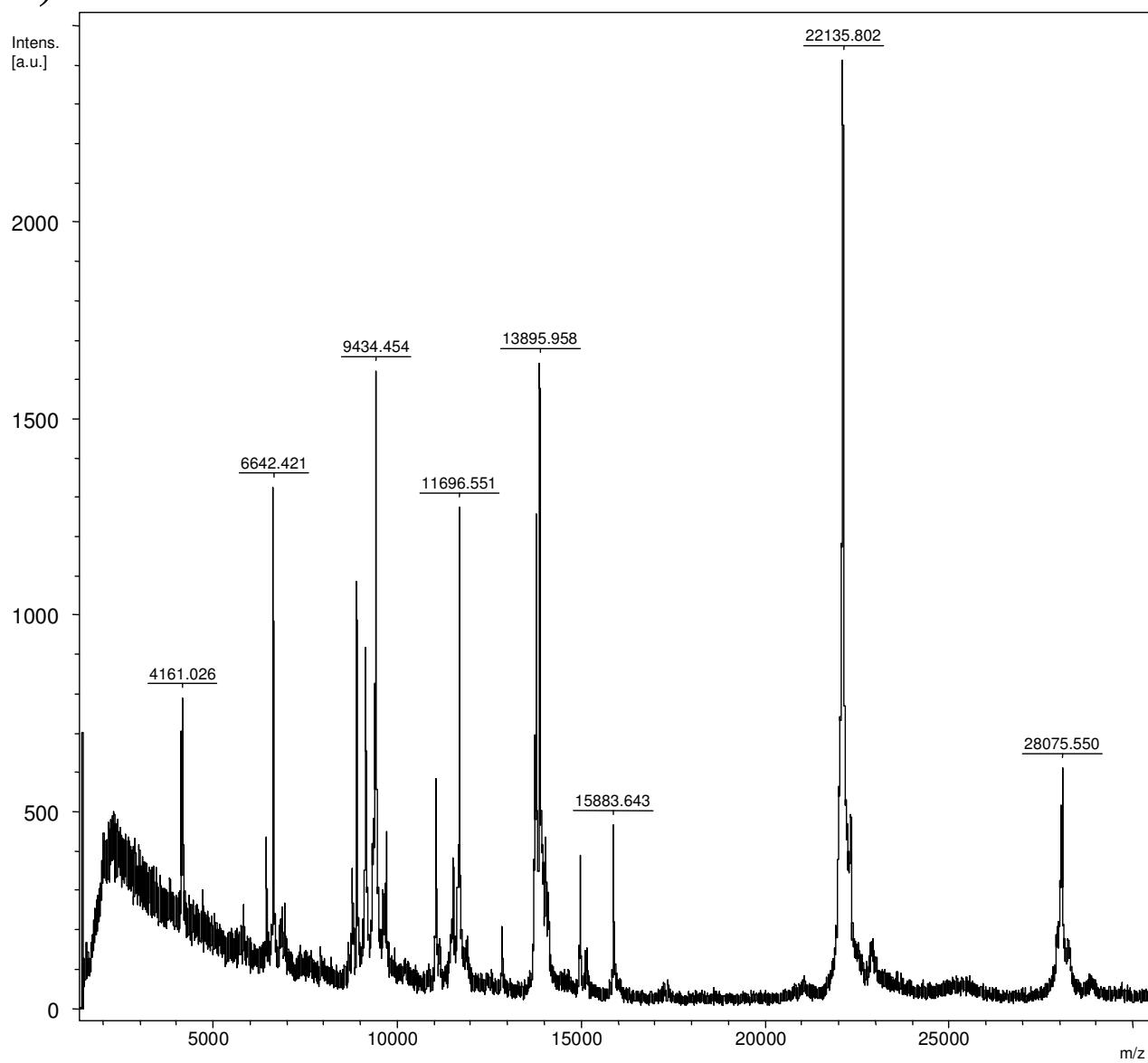
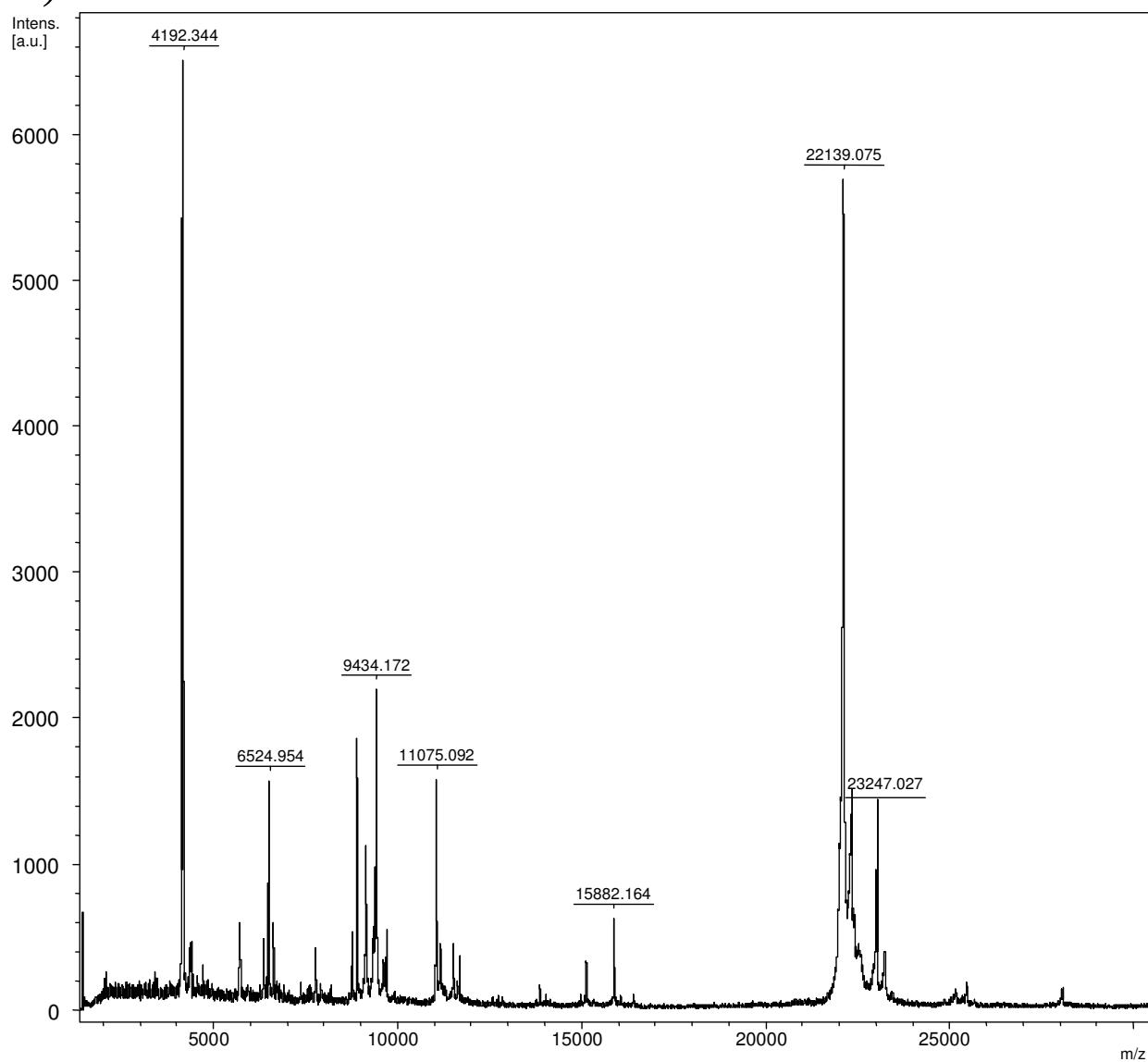


Figure S7. MALDI-ToF mass spectra of aldolase subunit obtained from the microgel and from solution (inset) showing the size-exclusion of the protein from the microgel.

a)



b)



c)

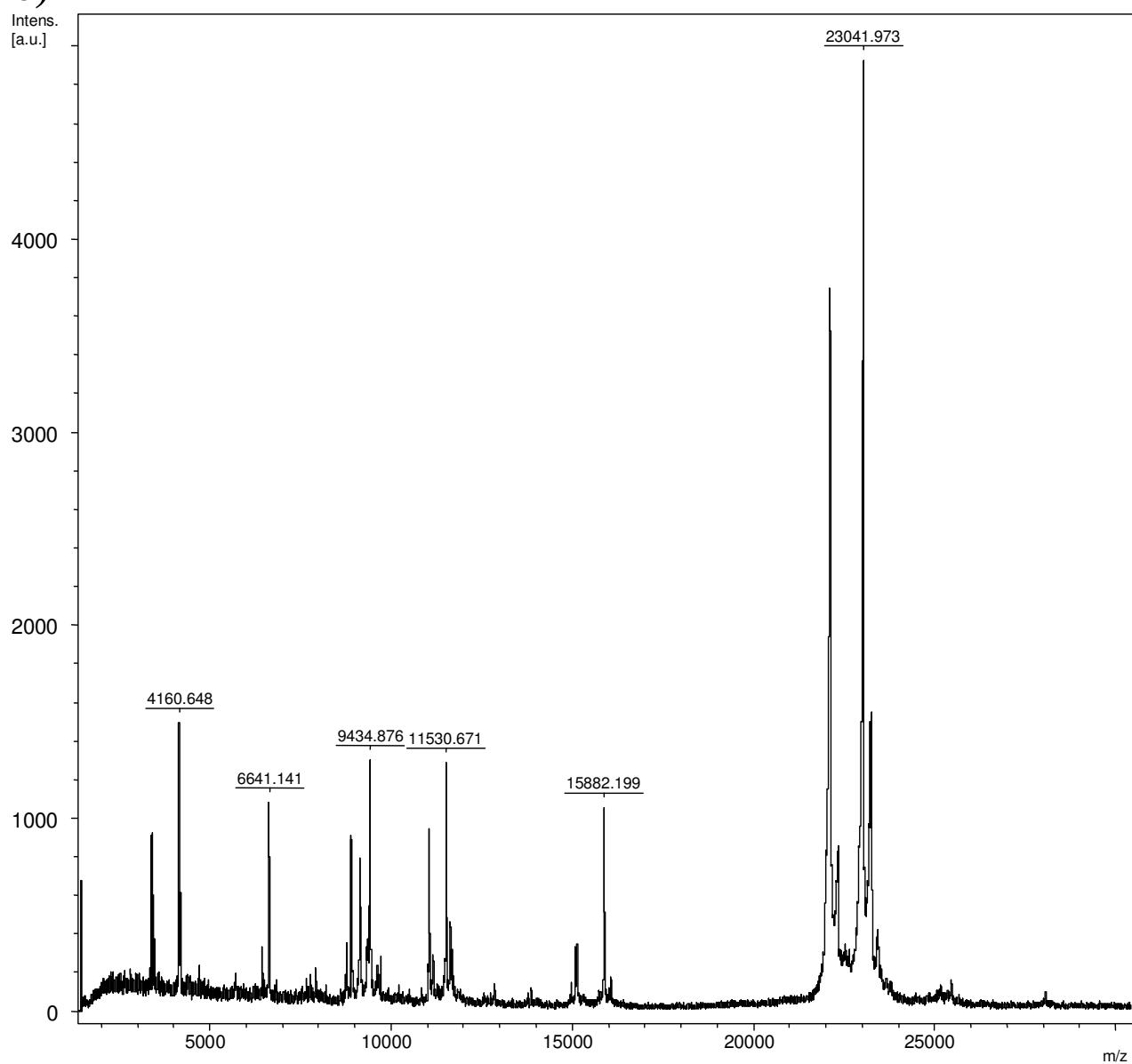
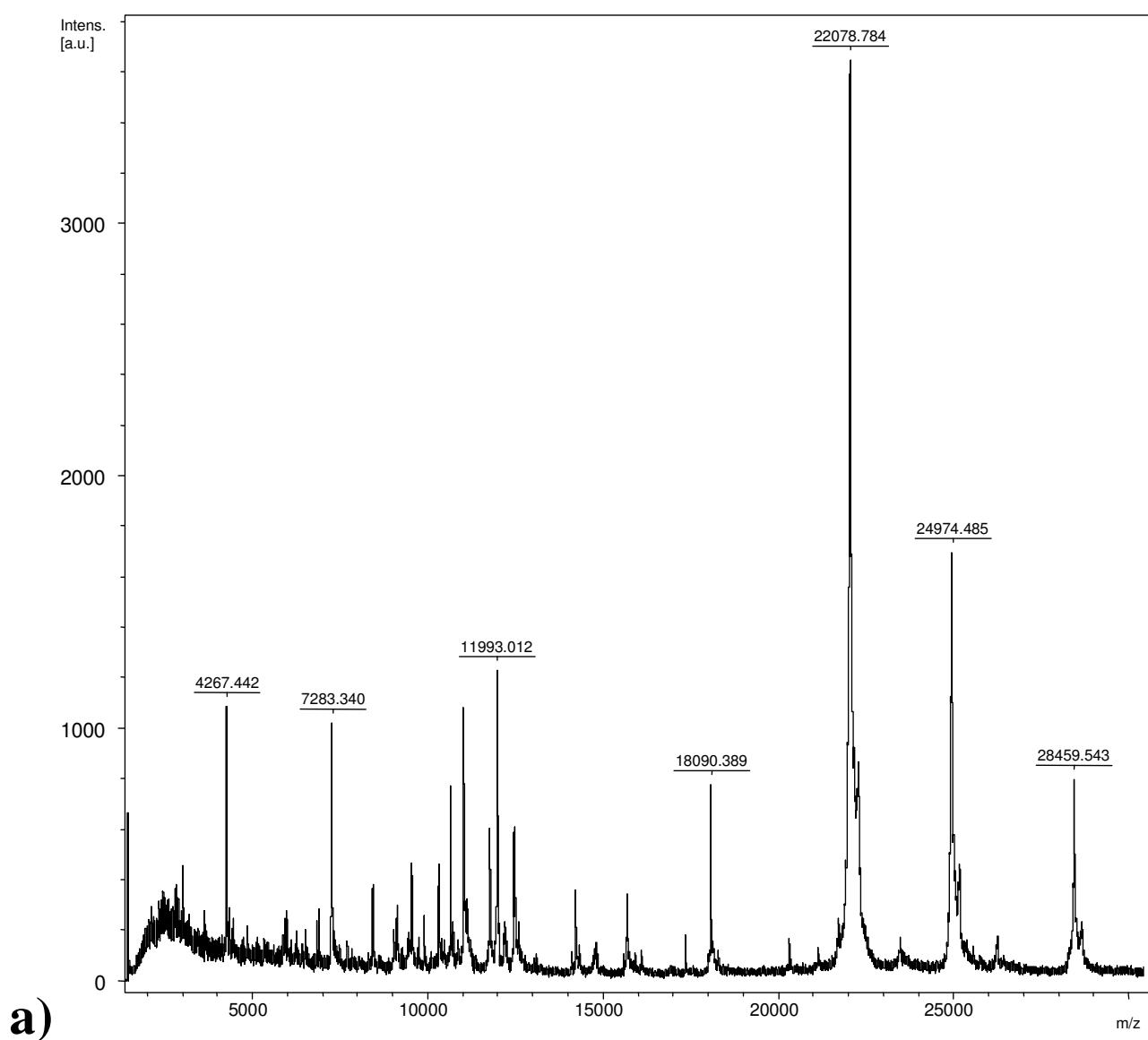


Figure S8. MALDI-ToF mass spectra of a tenfold-diluted serum with hGH spiked at 100 µg/mL (**a**), its microgel-captured fraction (**b**), and microgel-captured fraction of undiluted serum with hGH spiked at 100 µg/mL (**c**).



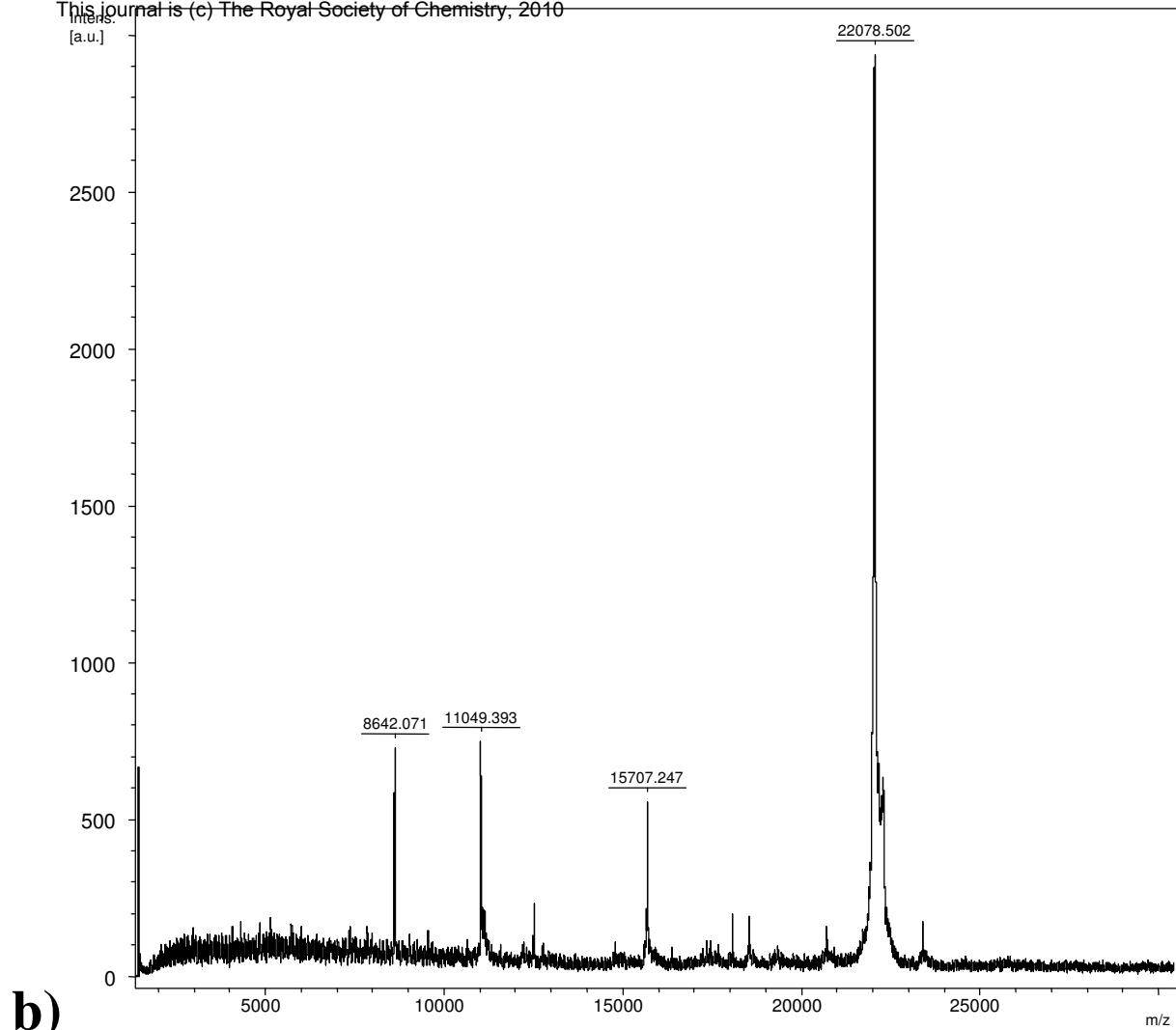


Figure S9. MALDI-ToF mass spectra of crude *E. coli* periplasm containing over-expressed hGH (a), and its microgel-captured fraction (b).

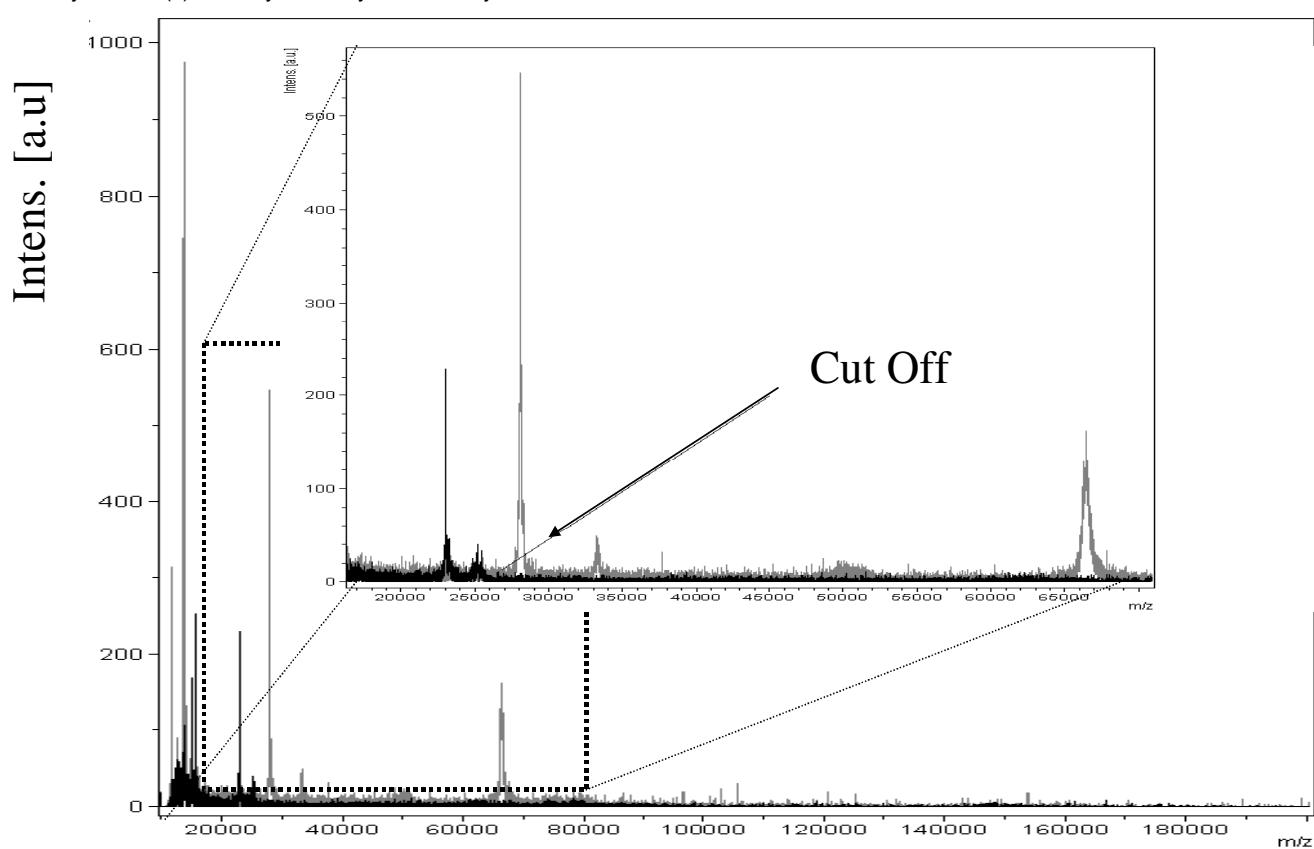


Figure S10. MALDI-ToF mass spectra of a tenfold-diluted serum obtained from the microgel (black) and from solution (grey) showing the size-selective capture of plasma proteins with MW <30 kDa (inset).

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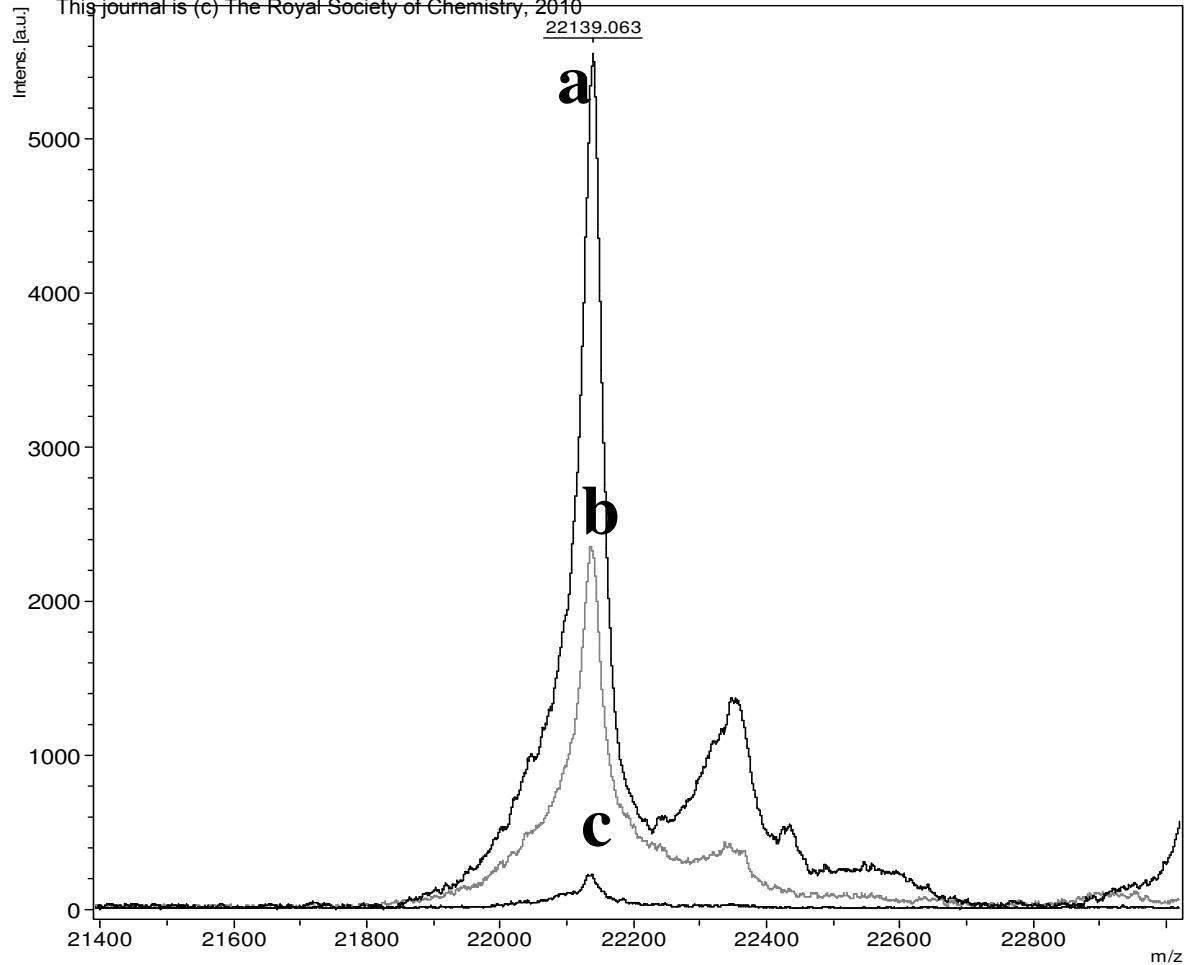


Figure S11. Concentrating effect of MiS-MALDI. Mass-spectra of hGH obtained from the microgel (**a**), from solution before incubation (**b**) and from the supernatant after incubation (**c**).

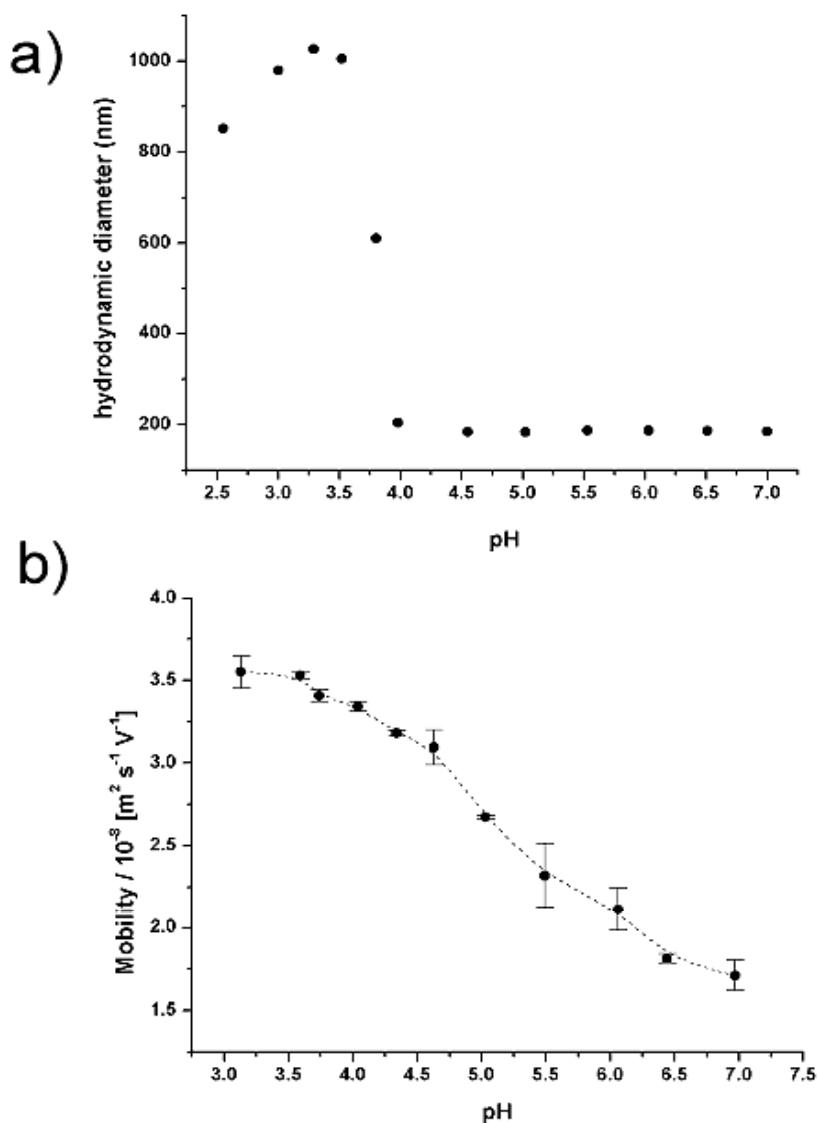


Figure S12. Hydrodynamic diameter (a) and electrophoretic mobility (b) of 1%-x-linked PVP/PEGMA MG as a function of pH.