



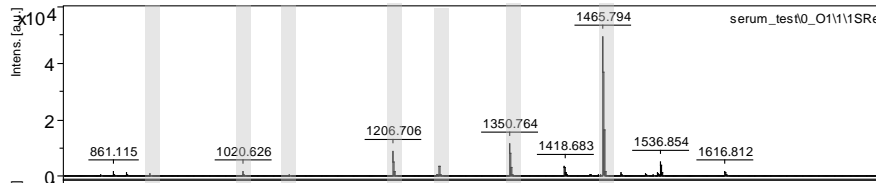
Decay-markers for quality assesment of plasma- und serum specimens

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Medical faculty Mannheim of the
University of Heidelberg

Limitations of MALDI-TOF MS Profiling

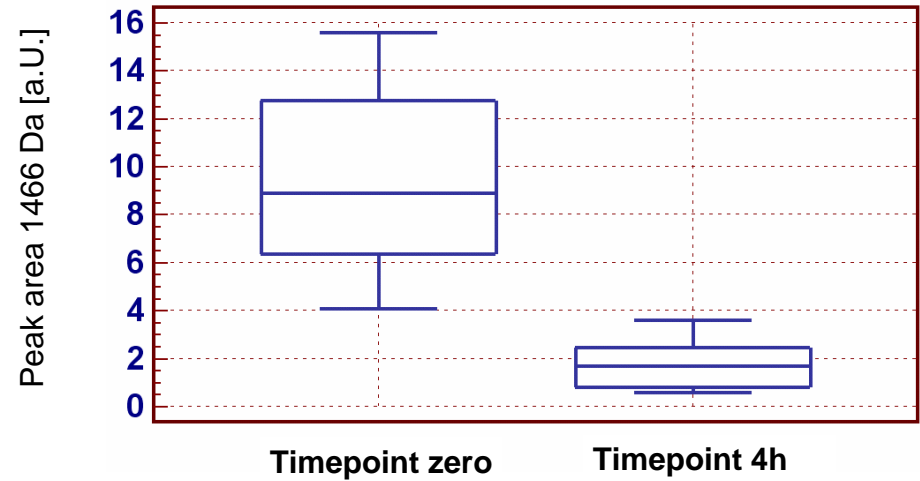
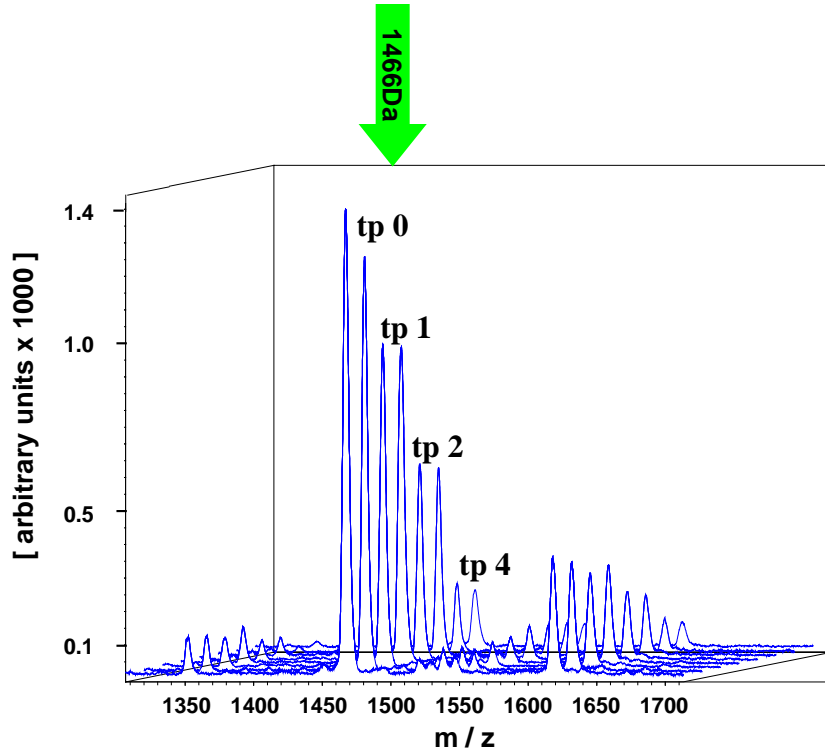
1h



1466:	DSGEGDFLAEGGGVR
1351:	SGEGDFLAEGGGVR
1263:	GEGDFLAEGGGVR
1207:	EGDFLAEGGGVR
1077:	GDFLAEGGGVR
1021:	DFLAEGGGVR
906:	FLAEGGGVR

Findeisen et al. Clin Chem 2005.51(12):2409-11

Preanalytical classification



Findeisen et al, Clin.Chem. 2005

11 Plasmaproteins

Selection of 62 endogenous decay markers

$\mu\text{mol/L}$		n	n
500-800 ^a	Alb	009	001
2-5 ^f	C4A	005	000
5-10 ^a	C3	005	002
0.2-0.4 ^e	F13	004	002
3 – 5 ^b	F2	006	004
6-12 ^a	FGA	100	031
- // -	FGB	019	013
2 - 4 ^a	ITIH4	015	003
1 - 3 ^d	KNG1	008	003
0.3-0.6 ^c	TMSB4	012	002
	FUK	003	001
	Σ	186	62

→
selection*

*highS/N ratio
highly reproducible ionisation status

^aHortin et al. *Clin Chem.* 2006 Jul;52(7):1223-37

^bKoomen et al. *J Proteome Res.* 2005 May-Jun;4(3):972-81.

^cHannappel *J Chromatogr.* 1987 Jun 26;397:279-85.

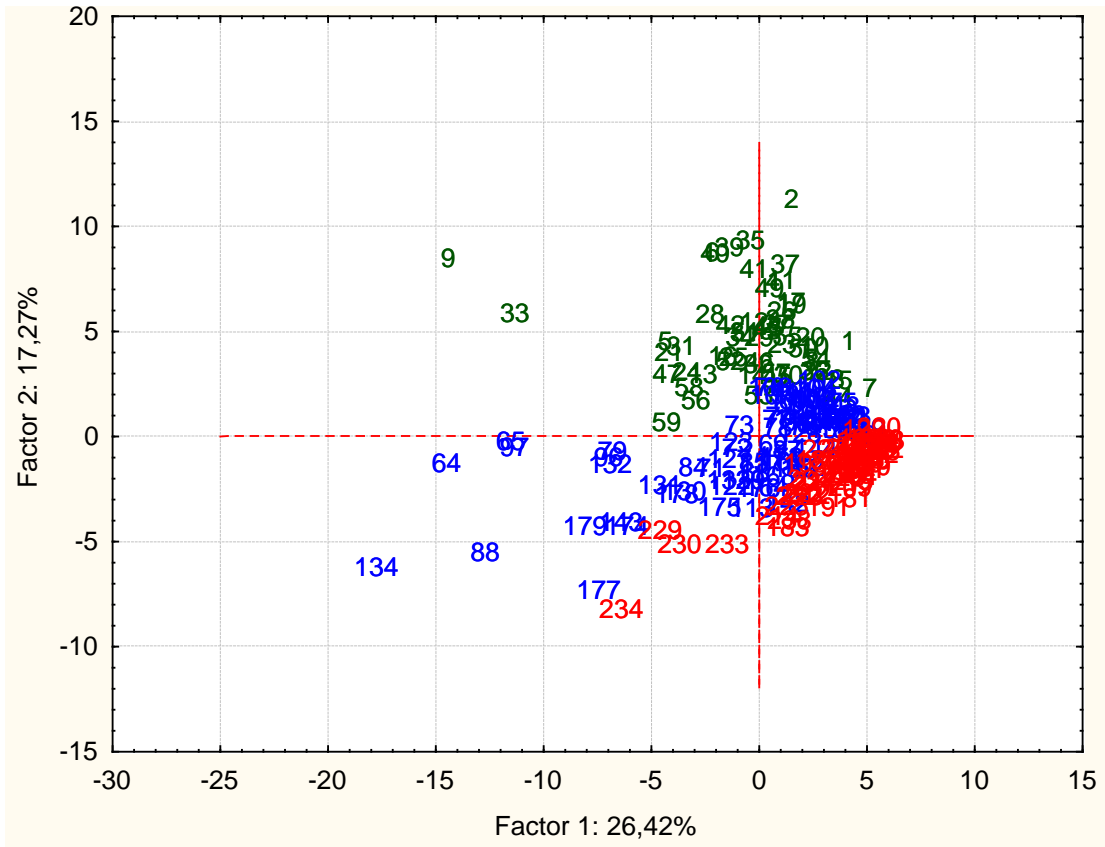
^dKleniewski et al. *Thromb Haemost.* 1979 Oct 31;42(3):1046-55.

^eKatona et al. *Thromb Haemost.* 2000 Feb;83(2):268-73.

^fGorski et al. *J Immunol Methods.* 1981;47(1):61-73.

Principle Component Analysis

Hauptkomponentenanalyse
62 Variablen (Marker); 244 Proben, 3 Qualitätsklassen



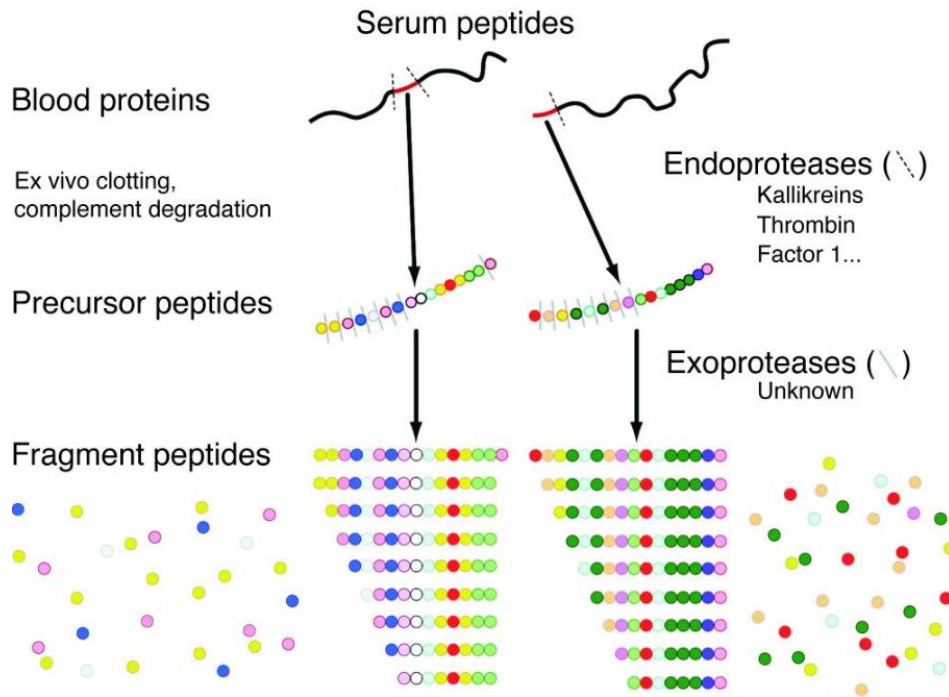
High: 0-2h
Medium: 4-9h
Low: 22-30h



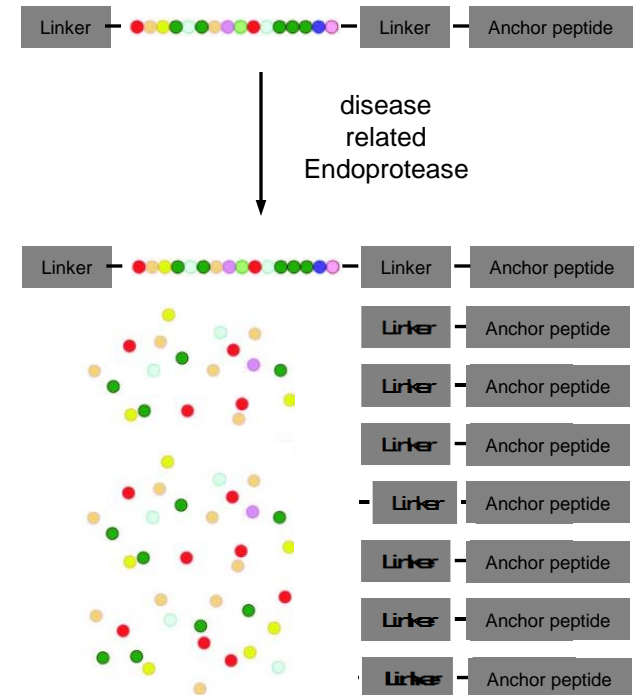
Monitoring proteolytic activity

endogenous versus exogenous decay markers

Endogenous reporter peptides

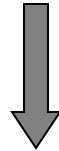


Exogenous reporter peptides



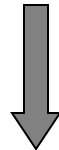
Study Design

15 Healthy Controls
15 CRC Tumor Patients



Serum and EDTA Plasma

Reporter Peptide was added prior to blood withdrawal



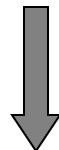
Storage at room temperature

2h

5h

8h

24h



Liquid chromatography / mass spectrometry

Quantification of RP-fragments

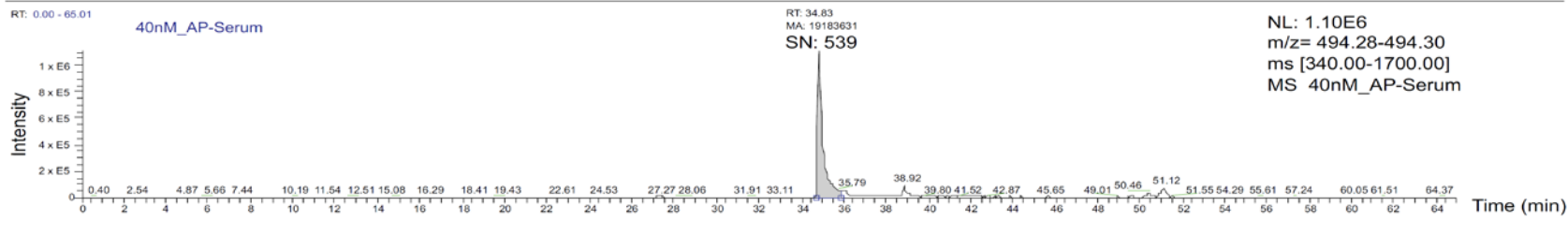
Exogenous Decay Marker (Reporter Peptide)

The `proteomic degradation clock`

Name	Sequence	m/z [M+2H] ²⁺
Reporter Peptide (RP)	LSLQKPRL-Ahx-ateelkal	
	SLQKPRL-Ahx-ateelkal	
	LQKPRL-Ahx-ateelkal	
	QKPRL-Ahx-ateelkal	
Intermediate Fragment (RP-4)	KPRL-Ahx-ateelkal	741,46
	PRL-Ahx-ateelkal	
	RL-Ahx-ateelkal	
	L-Ahx-ateelkal	
Anchor Peptide (AP)	Ahx-ateelkal	494,29

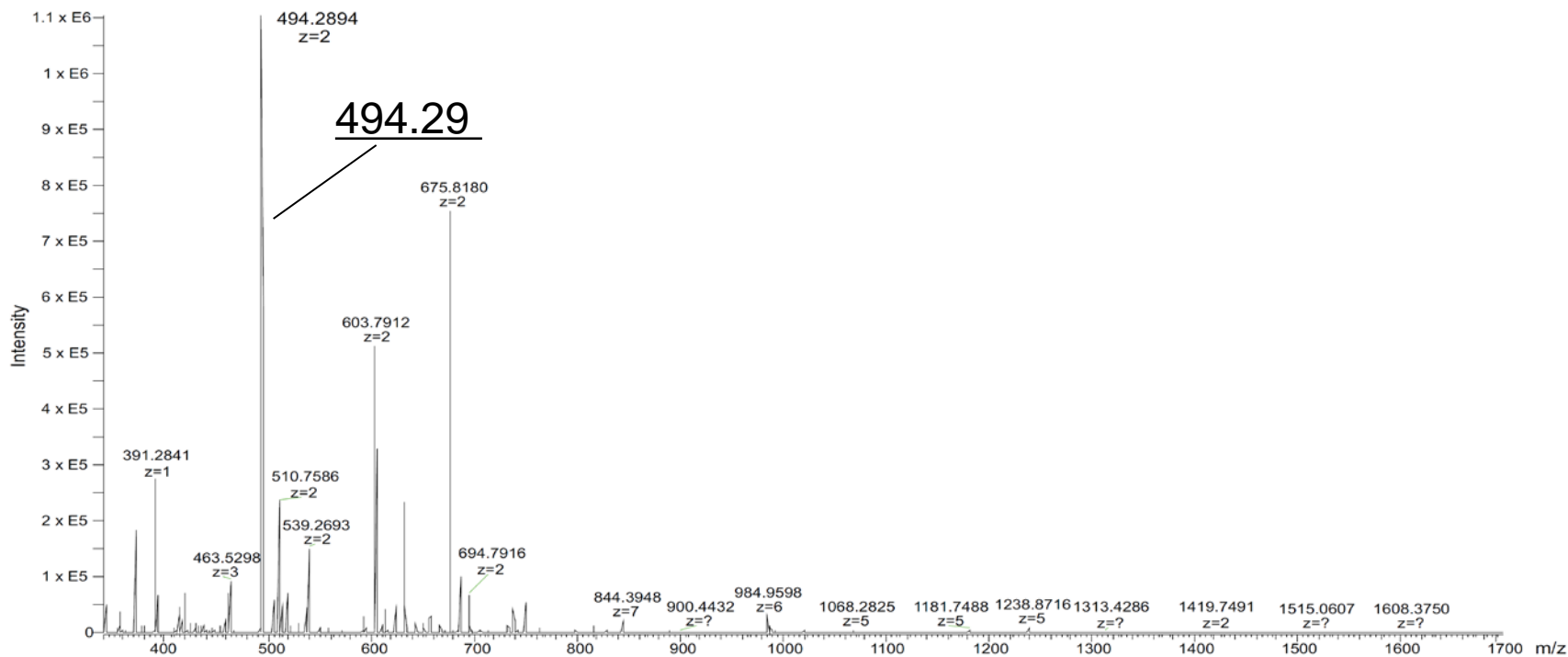
LC-MS of anchor peptide

A



B

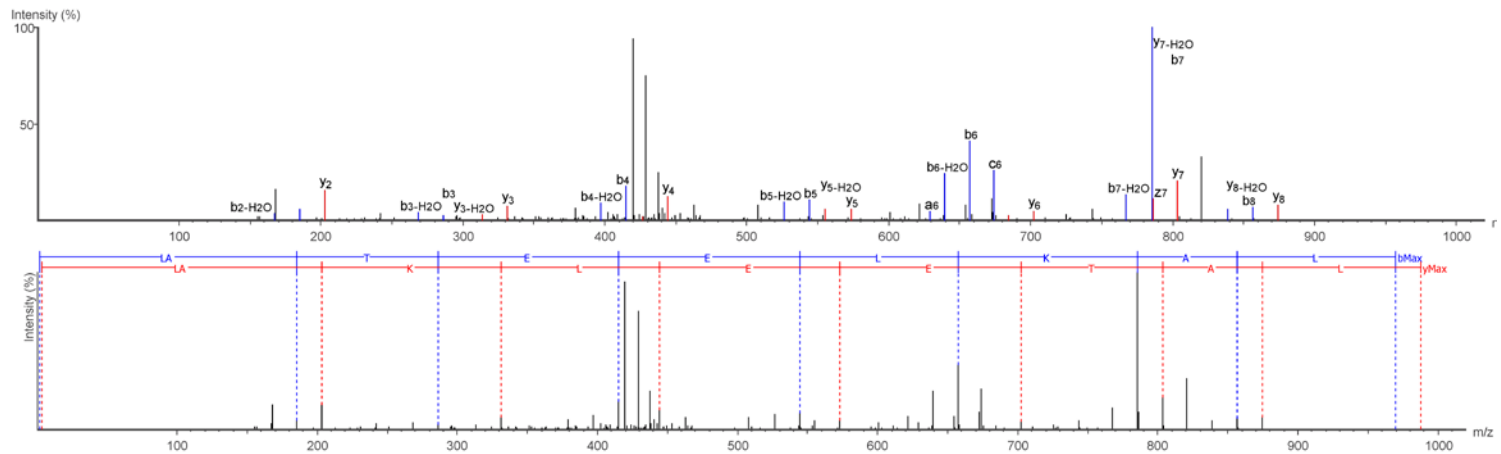
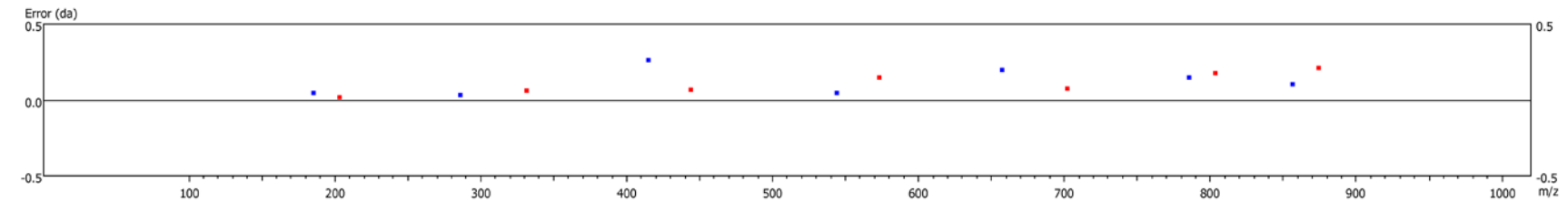
40nM_AP-Serum #2030 RT: 34.83 AV: 1 NL: 1.10E6
F: FTMS + p NSI Full ms [340.00-1700.00]



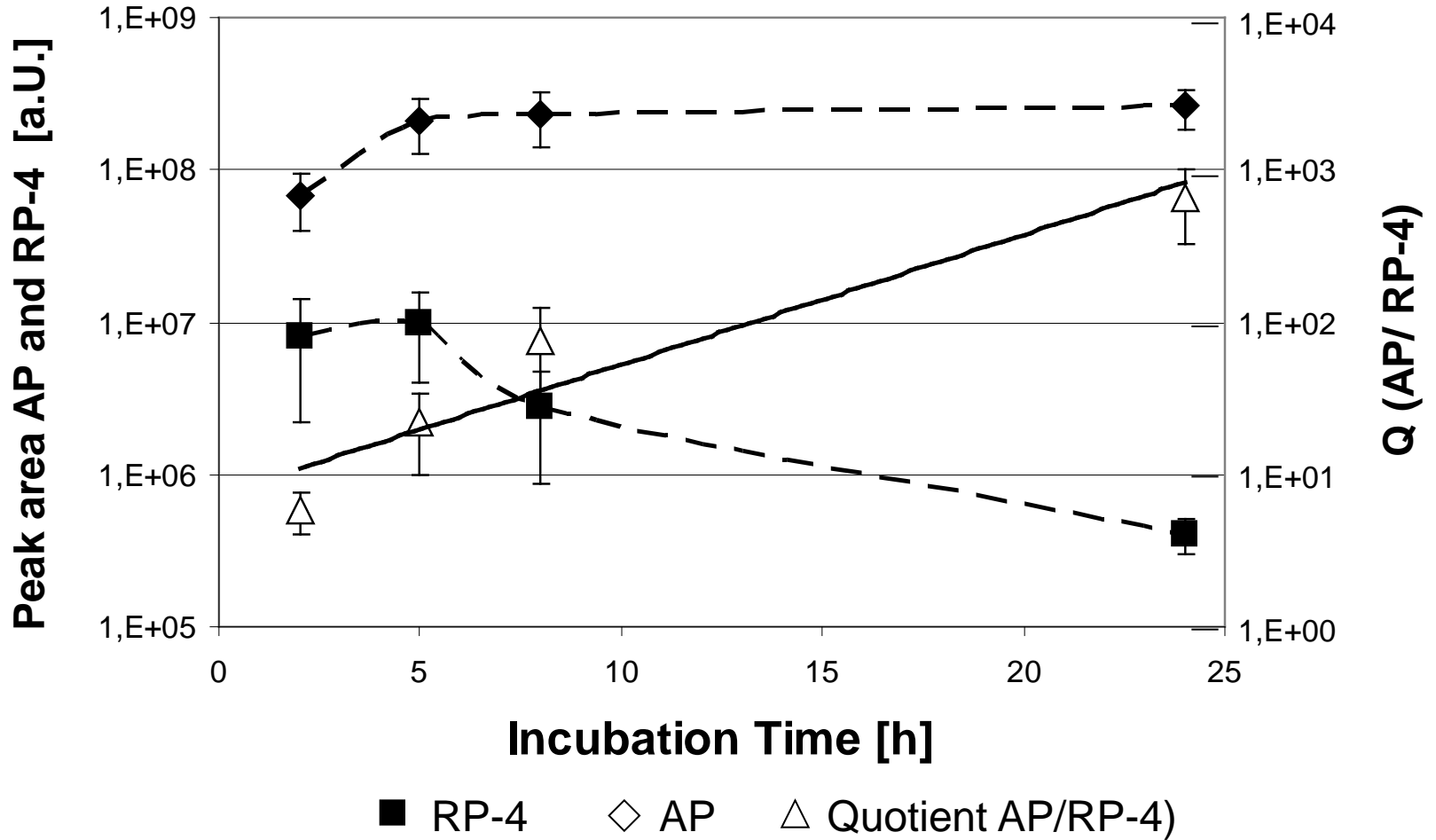
Sequence confirmation

Anchor peptide: AHX-ateelkal

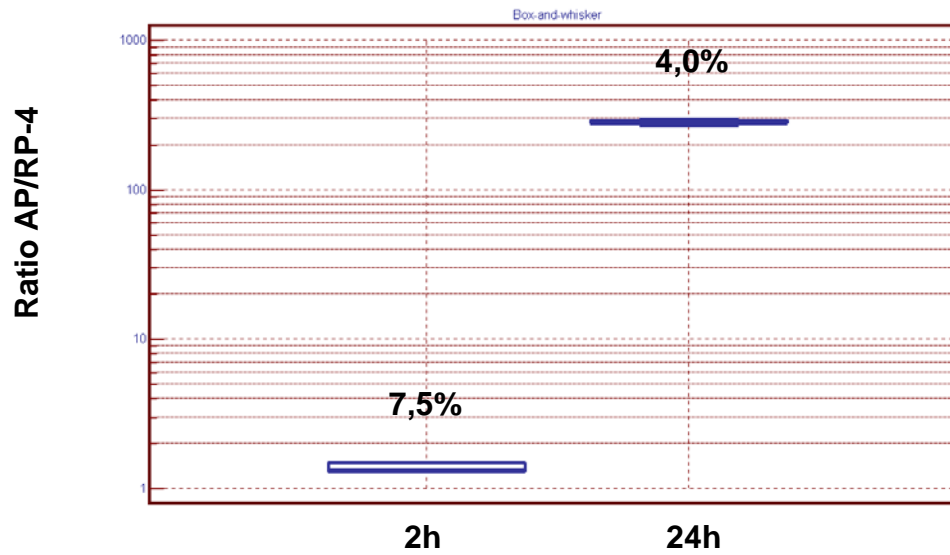
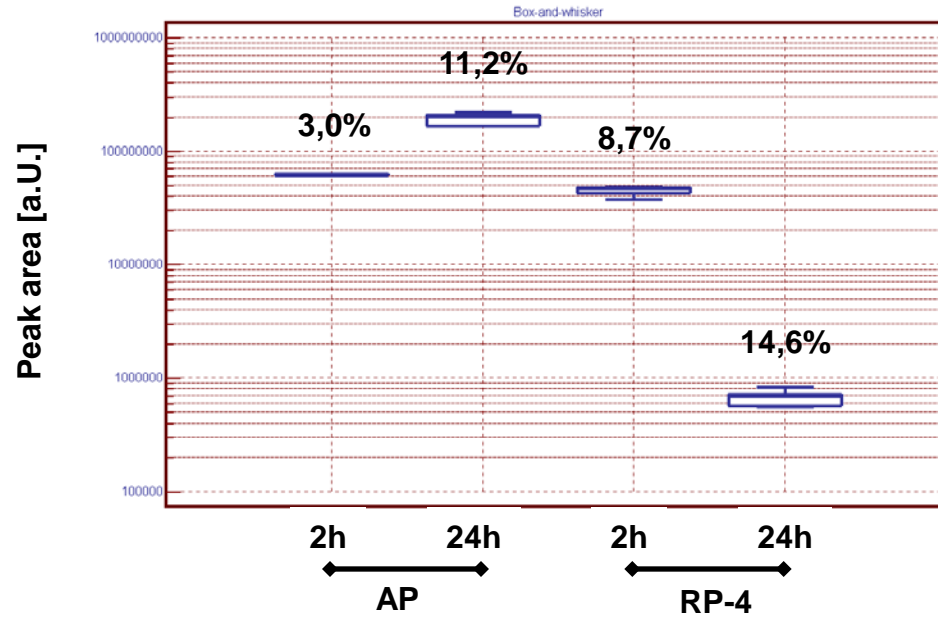
#	Immonium	b	b-H2O	a	c	Seq	y	y-H2O	z	z'	y (2+)	#
1	86.10	114.09	96.08	86.10	131.12	L						9
2	44.05	185.08	167.14	157.13	202.16	A	874.27	856.37	857.46	858.46	437.74	8
3	74.06	286.14	268.22	258.18	303.20	T	803.27	785.29	786.46	787.42	402.23	7
4	102.06	414.95	397.14	387.22	432.25	E	702.32	684.39	685.38	686.38	351.70	6
5	102.06	544.21	526.14	516.27	561.29	E	573.21	555.27	556.33	557.33	287.18	5
6	86.10	657.14	639.32	629.33	674.30	L	444.25	426.60	427.29	428.29	222.66	4
7	101.11	785.29	767.32	757.45	802.47	K	331.17	313.29	314.21	315.21	166.12	3
8	44.05	856.37	838.40	828.48	873.51	A	203.12	185.08	186.11	187.11	102.07	2
9	86.10					L	132.10	114.09	115.07	116.07	66.55	1



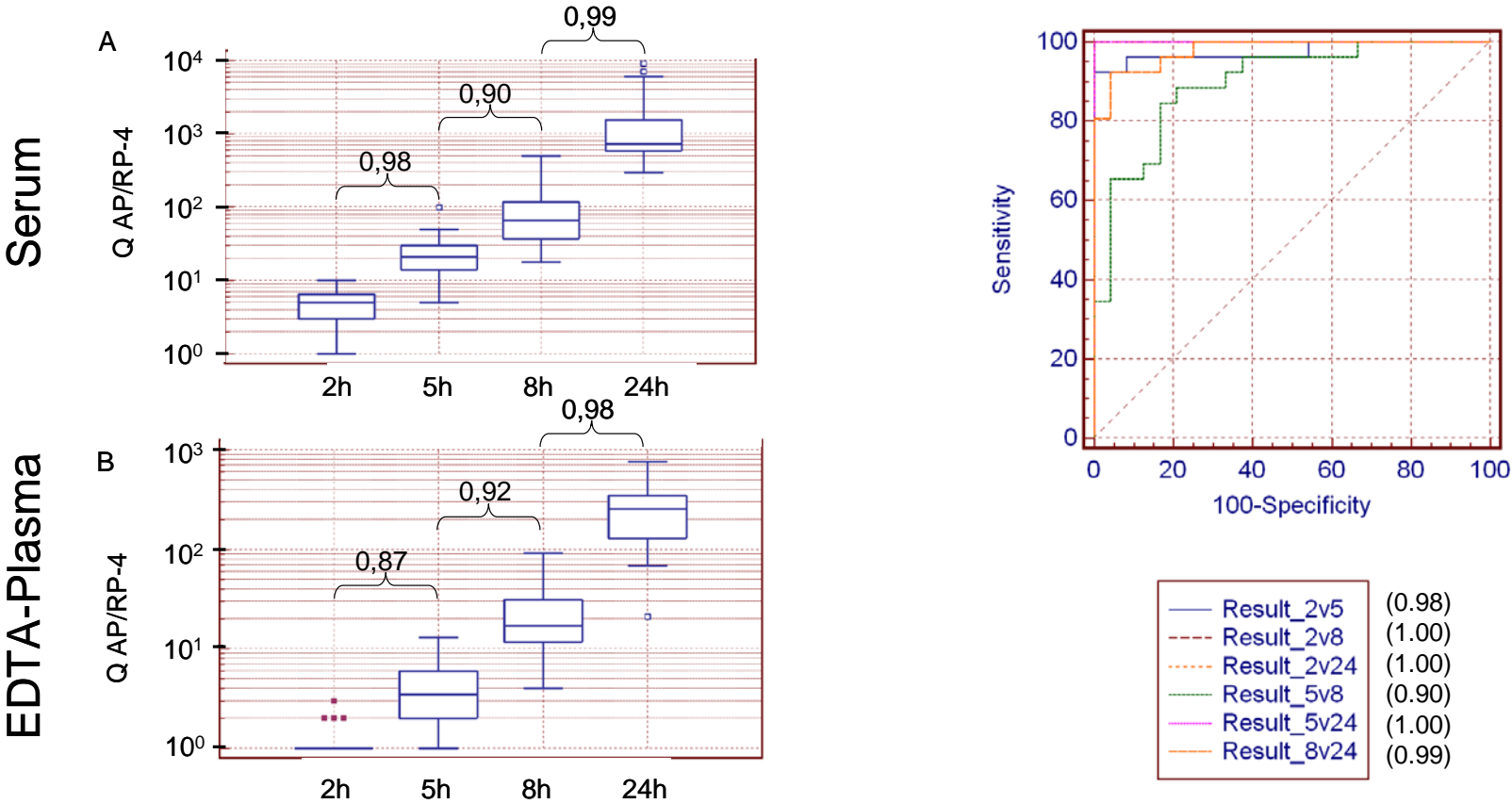
Decay Kinetics



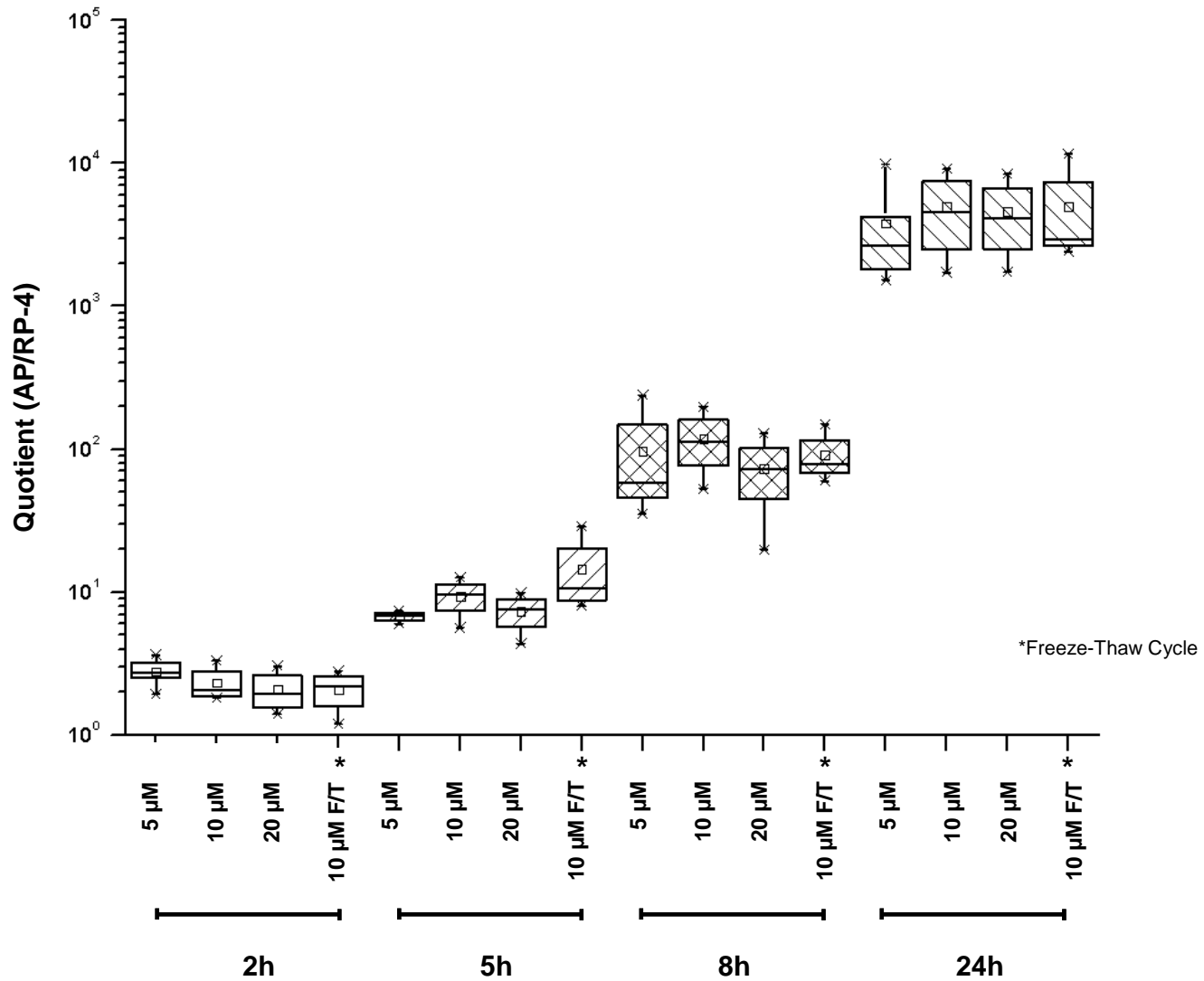
Reproducibility



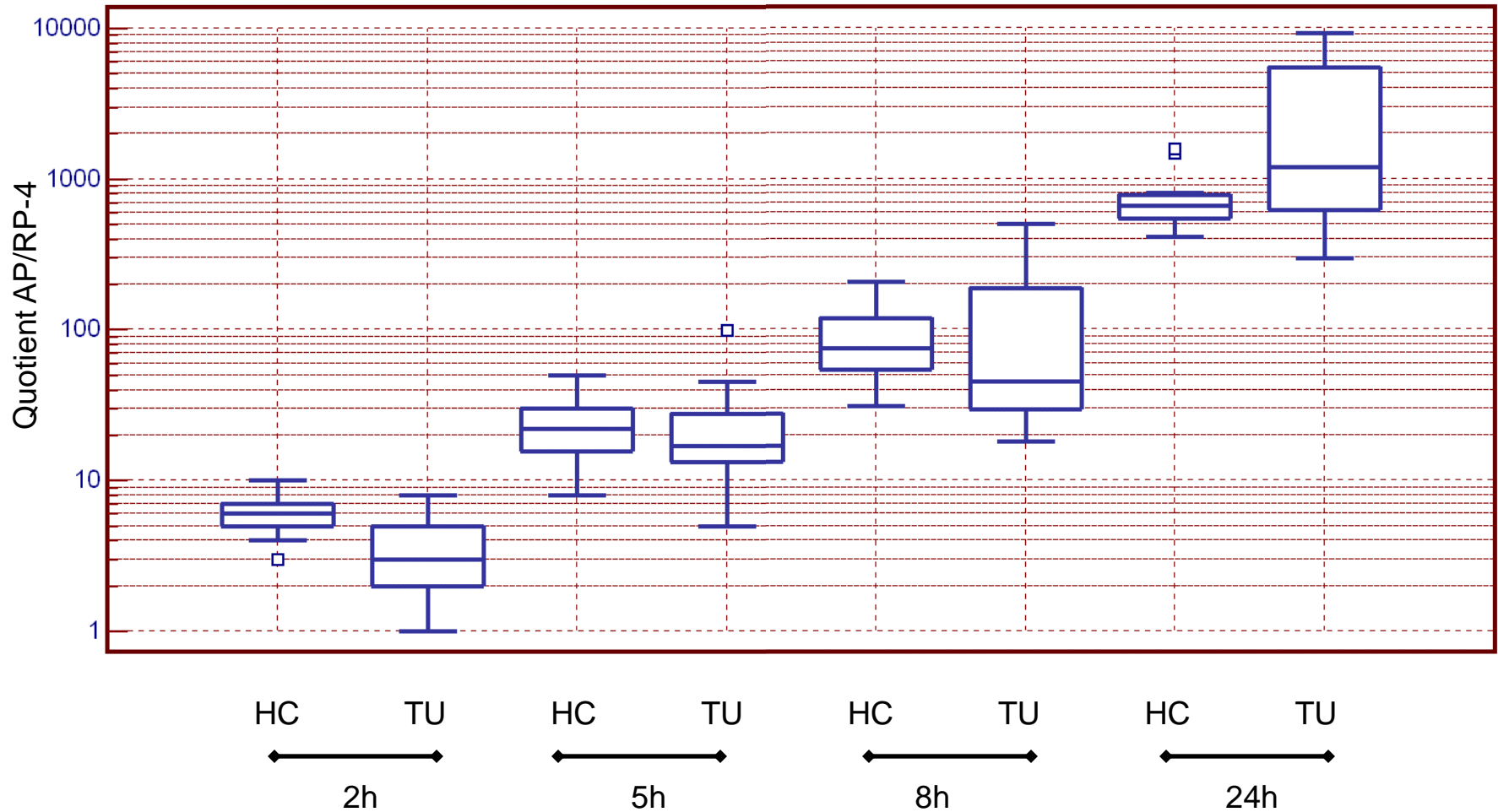
Classification



RP concentration and freeze-thaw Cycle

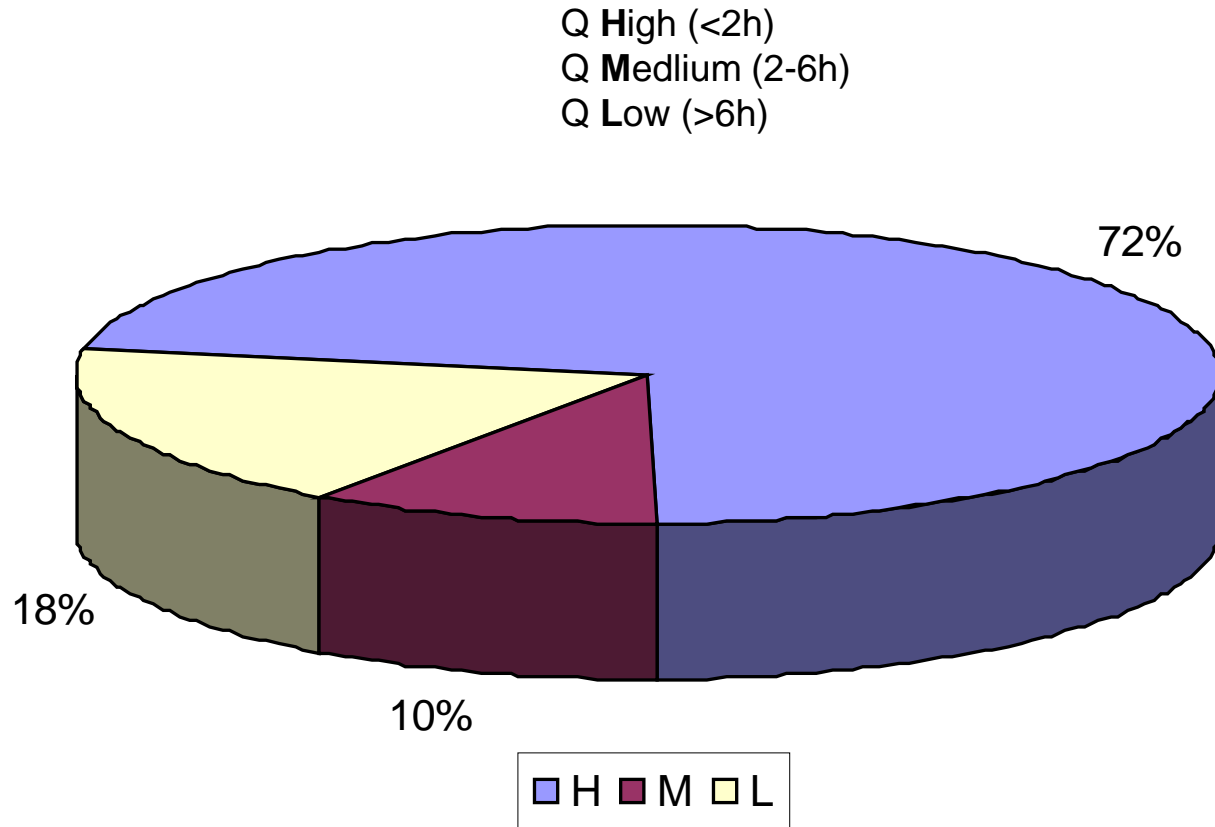


Healthy controls versus tumor patients



Exemplary application

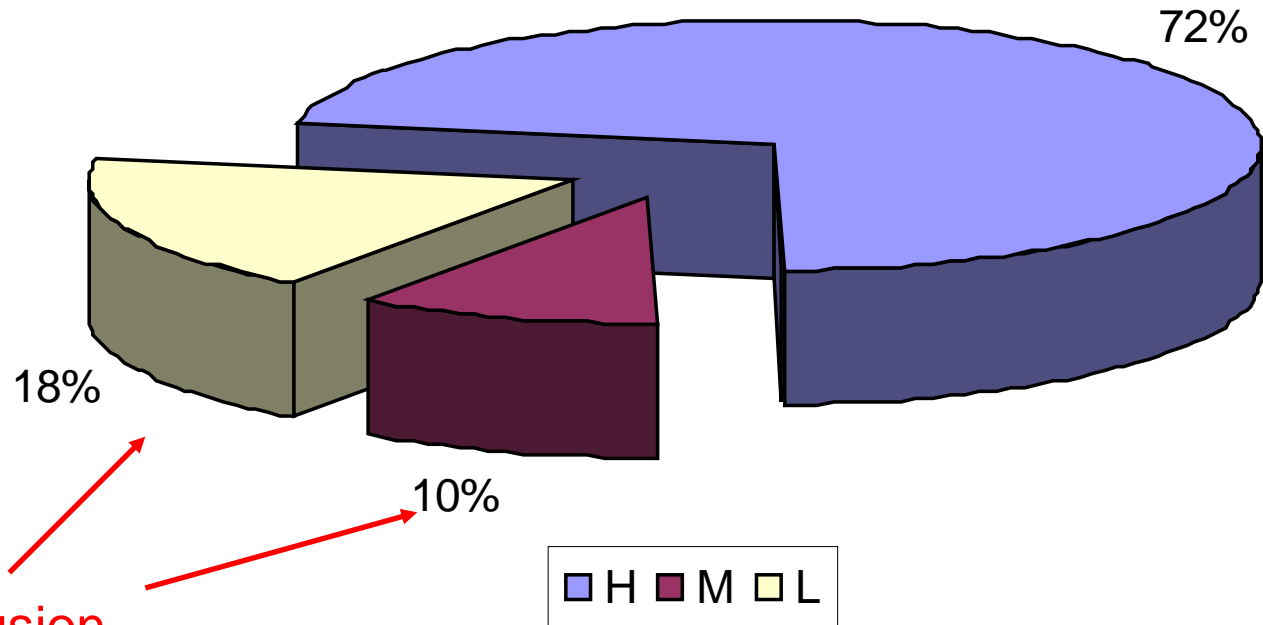
Exclusion of samples with low quality prior to IL6 measurements



Exemplary application

Exclusion of samples with low quality prior to IL6 measurements

--- Q High (<2h)
--- Q Medium (2-6h)
--- Q Low (>6h)



Exclusion

Conclusion

PROS

- Endo– and exogenous reporter peptides can be used as decay markers
- Targeted peptidomics of proteolytic fragments via LC-MS
- Preanalytical monitoring of serum and plasma specimens is feasible
- RP decay is the effect of housekeeping aminopeptidases (no relation to disease status)
- Aminopeptidases are not sensitive to freeze-thaw cycles
- External quality assessment for Biobanks

CONS

- High interindividual variability of endogenous decay markers
- Only prospective study design (RP spiking of tubes before blood withdrawal)
- Time resolution might not be sufficient
- Possible interference of exogenous RPs with downstream assays
- Long term storage?

Acknowledgement



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Dr. Kostrzewa
Dr. M. Meyer
Dr. M. Peer



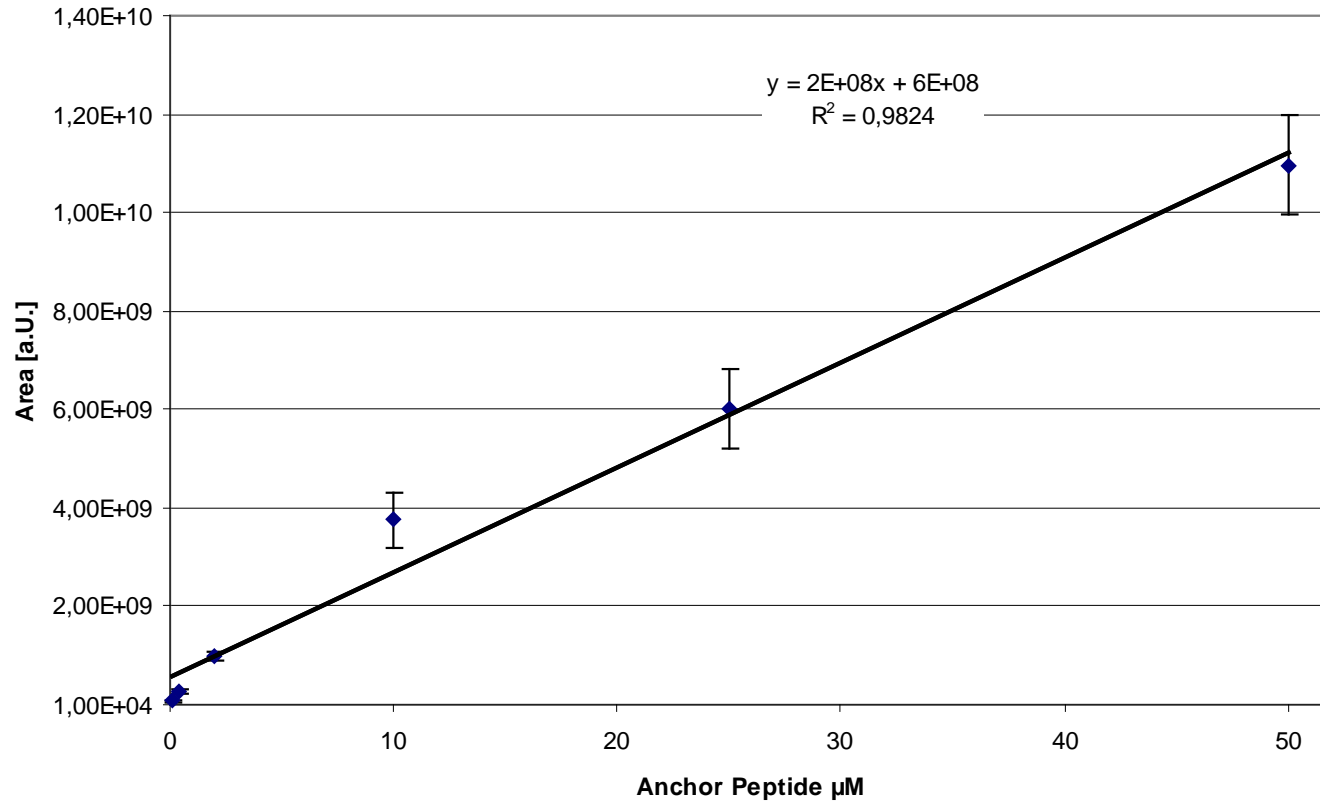
Prof. J. Rühle
Dr. T. Brandstetter
Dr. M. Rendl



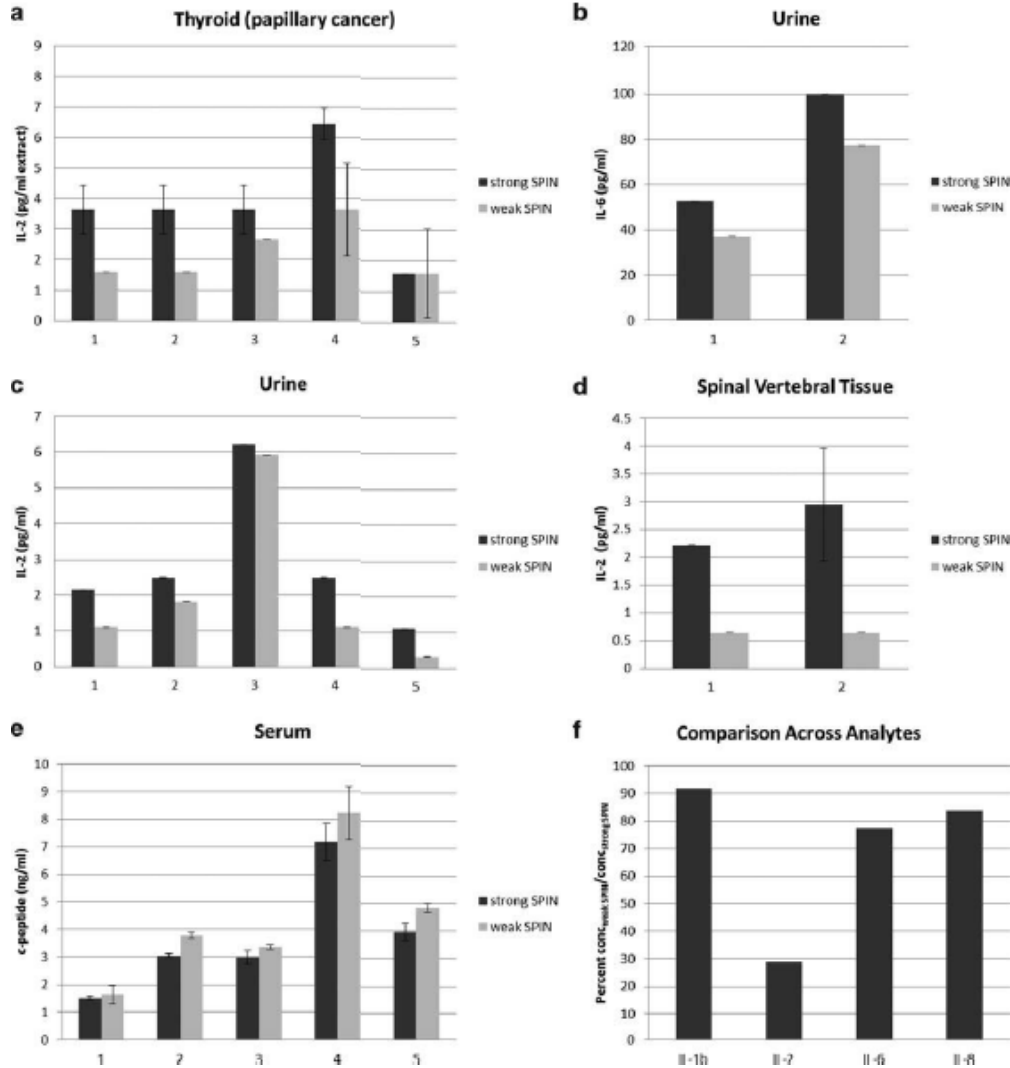
PD Dr. R. Bischoff
E. Dyrzcz



Linearity



Sample Quality ~ Analyte Concentration

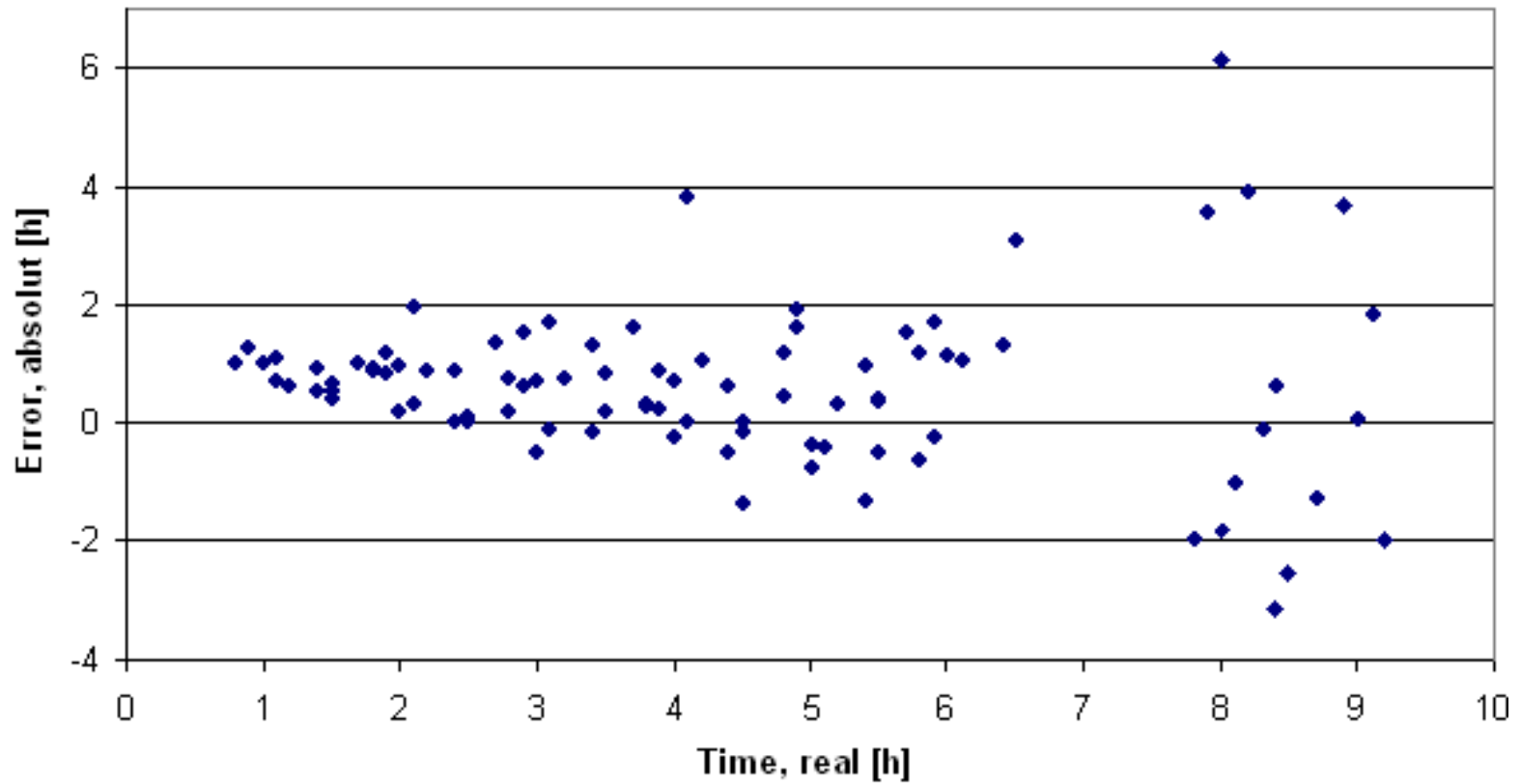


Beispiel Peptid-Degradation FGA

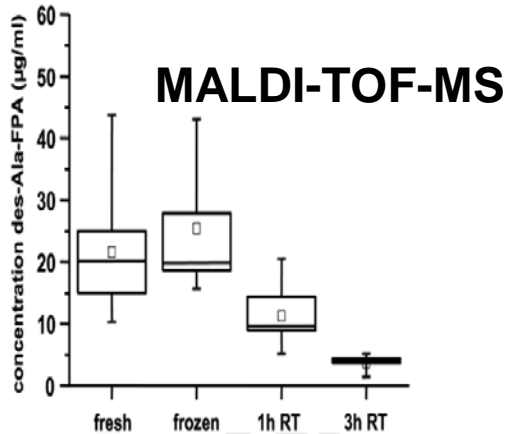
aa pos Anfang	aa pos ende	Protein	AA-Sequenz	M [Da]
21	35	FGA	A.DS G EGDFLAEGGGVR.G	1466,7
22	35	FGA	D.S G EGDFLAEGGGVR.G	1351,6
23	35	FGA	S.G E GDFLAEGGGVR.G	1264,6
24	35	FGA	G.E G DFLAEGGGVR.G	1207,6
26	35	FGA	T.G D FLAEGGGVR.G	1078,5
27	35	FGA	N-Terminus G.D F LAEGGGVR.G	1021,5
28	35	FGA	D.F L AEGGGVR.G	906,5
29	35	FGA	F.LA E GGGVR.G	759,4

576	598	FGA	K.SSSYSKQFTSSTSYNRGDSTFES.K	2555,1
576	595	FGA	K.SSSYSKQFTSSTSYNRGDST.F	2192,0
576	594	FGA	K.SSSYSKQFTSSTSYNRGDS.T	2090,9
576	593	FGA	K.SSSYSKQFTSSTSYNRGD.S	2002,9
576	592	FGA	K.SSSYSKQFTSSTSYNRG.D	1887,9
576	590	FGA	K.SSSYSKQFTSSTSYN.R	1674,7
576	589	FGA	K.SSSYSKQFTSSTSY.N	1560,7
576	588	FGA	K.SSSYSKQFTSSTS.Y	1397,6
576	587	FGA	K.SSSYSKQFTSST.S	1310,6
576	586	FGA	K.SSSYSKQFTSS.T	1209,5
576	585	FGA	K.SSSYSKQFTS.S	1122,5
576	583	FGA	K.SSSYSKQF.T	934,4

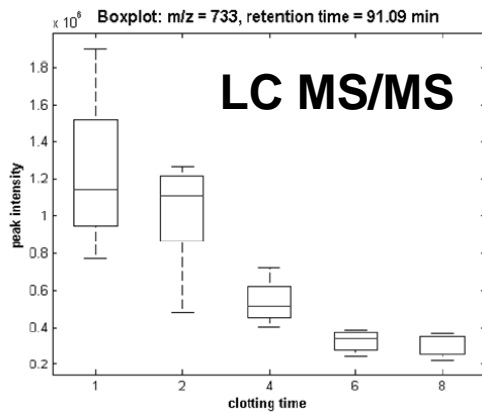
Cross Validation



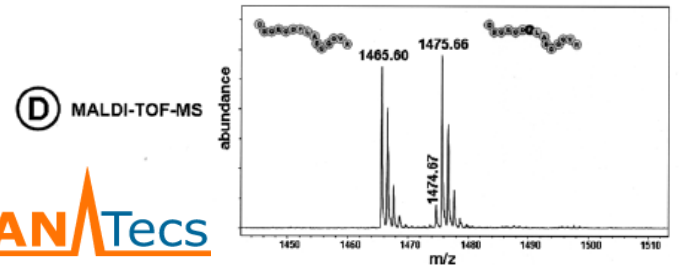
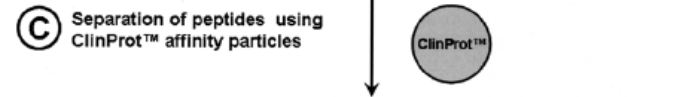
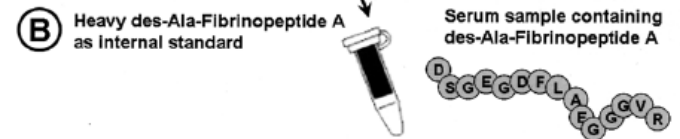
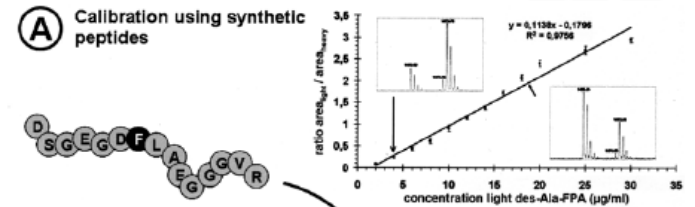
Validierung des endogenen Decay Markers



Baechle et al. Clin Proteomics 2007



Govorukhina et al. RCMS 2009



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 bioanalytical solutions

