

# PROTEOMICS

## Supporting Information for Proteomics

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**A systematic analysis of the effects of increasing degrees of serum  
immunodepletion in terms of depth  
of coverage and other key aspects in top-down  
and bottom-up proteomic analyses**

**Supplementary Data 1.** Examples of proteins identified from the tryptic digest of significantly different spots in the 2D-DIGE experiment. Proteins quoted were seen in more than one preparatory gel, or no conflicting identities were apparent if seen in a single preparatory gel; multiple identities from one spot are separated by “/”; fragments of immunoglobulins observed are not listed

Increased in:	Number of spots	Examples of proteins identified
MARS14 and Prot20 compared to MARS6	19	Hemopexin, Proteoglycan 4, Inositol polyphosphate 5-phosphatase, Neutrophil defensin 1, Complement factor B (Fragment)
Prot20 compared to MARS14 and MARS6	40	Proteoglycan 4, Complement factor H, Inositol polyphosphate 5-phosphatase, Neutrophil defensin 1
Prot20 compared to MARS6	49	Hemopexin, CD5 antigen-like protein / vitamin D-binding protein precursor, Proteoglycan 4, Complement factor H, Inositol polyphosphate 5-phosphatase, Neutrophil defensin 1
Prot20 compared to MARS14	29	Hemopexin, Inositol polyphosphate 5-phosphatase
Stepwise rise (increasing progressively from MARS6 to MARS14 to Prot20)	7	Alpha-1B-glycoprotein / Inter-alpha-trypsin inhibitor heavy chain H4 / Afamin

**Supplementary Data 2: Details of immunodepleted proteins seen with LTQ-Orbitrap Velos**

**Expected depletions**

MARS 6	MARS 14	ProteoPrep20
Albumin	Albumin	Albumin
Transferrin	Transferrin	Transferrin
Fibrinogen	Fibrinogen	Fibrinogen
Alpha-1-antitrypsin	Alpha-1-antitrypsin	Alpha-1-antitrypsin
Haptoglobin	Haptoglobin	Haptoglobin
X	Alpha-2-macroglobulin	Alpha-2-macroglobulin
X	Alpha-1-acid glycoprotein	Alpha-1-acid glycoprotein
X	Apo-A-I	Apo-A-I
X	Apo-A-II	Apo-A-II
X	C3	C3
X	Transthyretin	Transthyretin
X	X	Ceruloplasmin
X	X	Apo-B
X	X	C1q
X	X	C4
X	X	Plasminogen
IgG	IgG	IgG
IgA	IgA	IgA
X	IgM	IgM
X	X	IgD

**Proteins seen in depleted sample on 1D-LC MS/MS Orbitrap Velos analysis**

(Y = evidence of the protein seen in the sample ; N= evidence of the protein not seen)

**≥2 peptide level**

MARS 6	MARS 14	ProteoPrep20
Y	Y	Y
N	Y	Y
Y	Y	Y
Y	Y	Y
Y	Y	Y
-	Y	Y
-	Y	N <sup>1</sup>
-	Y	Y
-	Y	Y
-	Y	Y
-	N	N
-	-	Y
-	-	N <sup>2</sup>
-	-	N
-	-	N
-	-	N
Y <sup>4</sup>	Y <sup>4</sup>	Y <sup>4</sup>
Y <sup>4</sup>	Y <sup>4</sup>	Y <sup>4</sup>
-	Y <sup>4</sup>	Y <sup>4</sup>
-	-	Y <sup>4</sup>

**1 peptide level**

MARS 6	MARS 14	ProteoPrep20
Y	Y	Y
N	Y	Y
Y	Y	Y
Y	Y	Y
Y	Y	Y
-	Y	Y
-	Y	Y
-	Y	Y
-	Y	Y
-	Y	N
-	-	Y
-	-	Y
-	-	Y <sup>3</sup>
-	-	N
-	-	Y
Y <sup>4</sup>	Y <sup>4</sup>	Y <sup>4</sup>
Y <sup>4</sup>	Y <sup>4</sup>	Y <sup>4</sup>
-	Y <sup>4</sup>	Y <sup>4</sup>
-	-	Y <sup>4</sup>

**Notes**

1. Alpha-1-acid glycoprotein 1 is seen in the Prot20 depleted sample at the 1+ peptide level, but not at the 2+ peptide level. Alpha-1-acid glycoprotein 2 is seen at the 2+ peptide level in all depletions
2. Refers to ApoB (IPI00894122); peptides from (either) Apo B-100 / ApoB-48 are seen at 2+ peptide level - these don't appear to be depleted
3. Complement C1q subcomponent subunit A is seen at the 1 peptide level. Subunits B and C are not seen even at that level
4. Multiple immunoglobulin fragments remain in all depletions at the 2+ peptide level. Distinguishing which are from which immunoglobulin subfamily is difficult

**Supplementary Data 3: Number of unique peptides seen in each peak 1 sample for the immunodepleted proteins.**  
 $\leq 1$  peptide detections are shown in red

Immunodepleted Proteins <sup>1</sup>	WS	MARS 6	MARS 14	Prot 20
Alpha-1-antitrypsin	31	30	31	7
Fibrinogen alpha chain	14	27	25	12
Fibrinogen beta chain	15	24	16	7
Fibrinogen gamma chain	4	13	6	4
Haptoglobin	22	17	14	12
Serotransferrin	93	0	15	8
Serum albumin	113	5	28	45
Alpha-1-acid glycoprotein 1 (Orosomucoid-1)	11	14	3	1
Alpha-1-acid glycoprotein 2 (Orosomucoid-2)	9	11	4	2
Transthyretin(Prealbumin)	5	8	1	0
Alpha-2-macroglobulin	55	73	54	56
Apolipoprotein A-I	31	58	49	41
Apolipoprotein A-II	13	15	14	10
Apolipoprotein(a)	14	33	38	13
Complement C3	22	36	2	4
Apolipoprotein B-100;Apolipoprotein B-48 <sup>2</sup>	52	154	174	157
APOB protein	2	4	4	1
Plasminogen	47	70	74	1
Ceruloplasmin	7	11	11	2
Complement C1q subcomponent subunit A	4	8	9	1
Complement C1q subcomponent subunit B	4	7	6	0
Complement C1q subcomponent subunit C	4	8	7	0
Complement component 4B	0	1	1	0

1. These are groupings of all the protein names referring to the same protein or fragments / isoforms thereof

2. The number of unique peptides seen for these suggests the immunodepletion column antibodies are not directed against these forms