REPORT Application of engine Protein Array Technology for the analysis of IgG autoantibody repertoires of five patients	•	• •	•	•				•	•	•	•	•	•	•	•			•	•	•	٠	•	•				•	•	•	•	•			•	•	•			•	•	•	•	٠	•	•		•	•	•	•	•	•		•	•	•	• •		•	•	•	•	•	•	•	• •	•	•
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Your Question. Our Service. Your Result.





BIOMARKER SCREENING with 15,000 antigens in one shot with serum samples. In a cooperation-project we have examined serum of 5 autoimmune patients with the engine UniPEx arrays (product No. 1008).

We identified 7 overlapping potential antigens for serum IgG in all 5 patients.

Did we discover new biomarkers?! We are now investigating the association between autoimmune diseases and antibodies against the following antigens:

- MAFK 8
- HNRNPUL2
- MYH14
- CREBBP
- AATF
- SDE1
- DDX51

Properties of our UniPEx Array

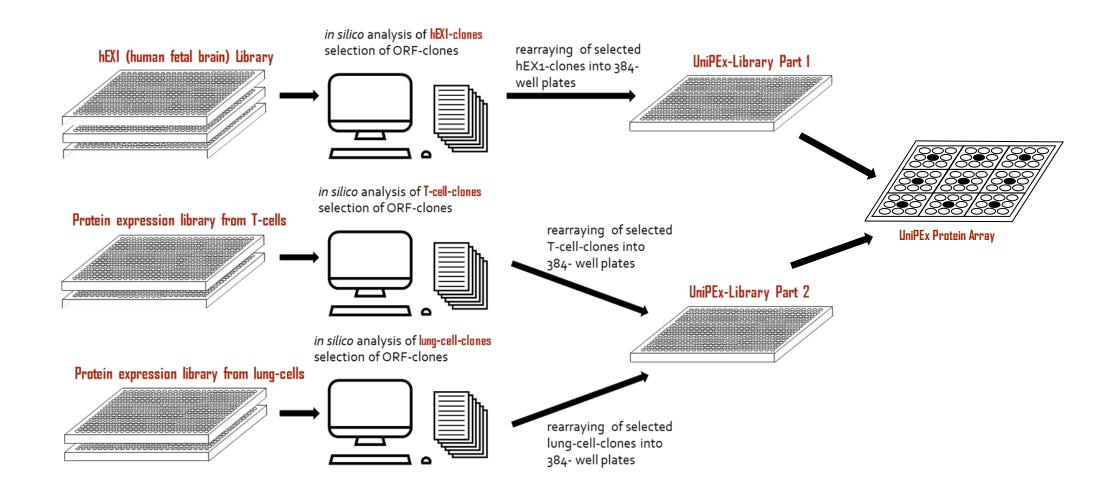


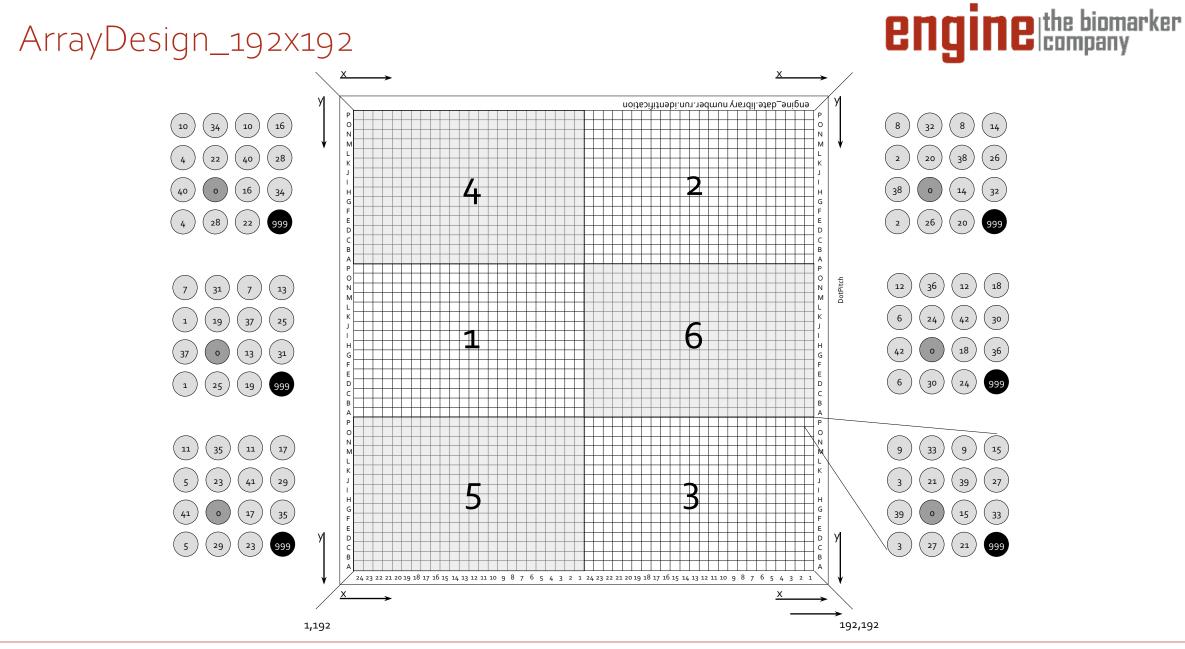


- Clones from different protein expression libraries (lung, T-cells, colon and human fetal brain)
- Mainly clones that are "in frame" are present
- Different libraries in two different vectors and expressed in different *E. coli*-strains on one array
- Spotting in 4x4 pattern
- 15,300 clones representing 7,390 different human proteins

Creation of UniPEx Array

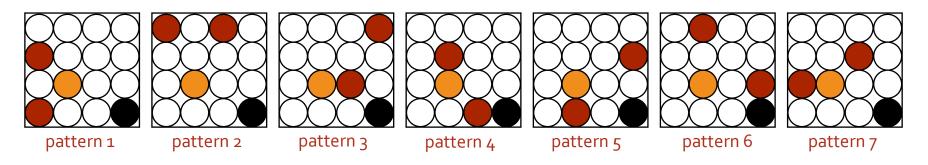




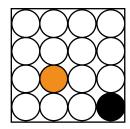


Scoring Pattern on engine Protein Arrays





- Spots higlighted in red show the corresponding pattern, in which clones were spotted
- A clone is only regarded as positive, if <u>both</u> spots can be detected



pattern 8 – guiding dot (black) and empty position (orange)

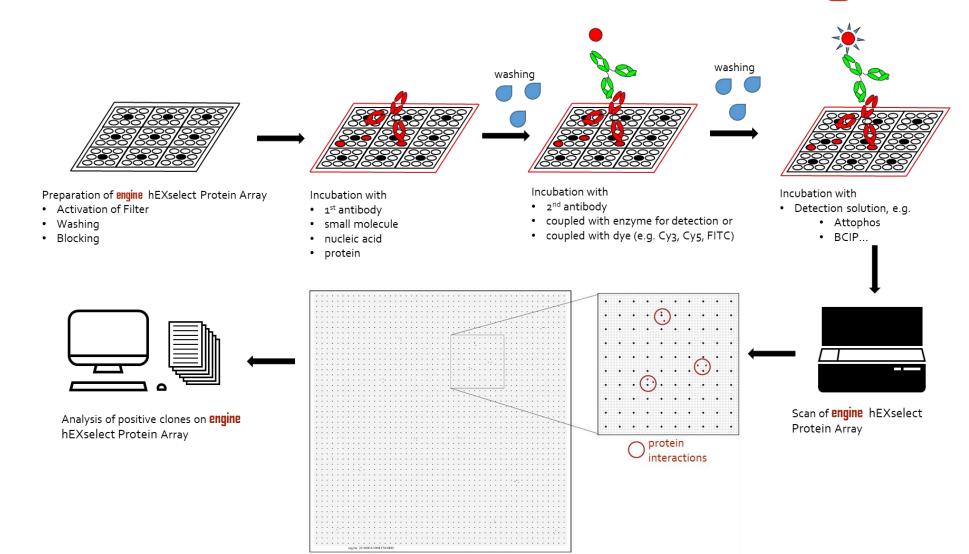
engine Protein Arrays – as simple as a Western Blot





- Removal of E. coli debris with Kimwipe tissue
- Washing
- Blocking
- Incubation with sample (serum, antibody etc.)
- Washing
- Incubation with secondary antibody (AP-labeled)
- Detection with AttoPhos® substrate
- Software-based evaluation

Detection of positives hits at a glance



Detection of positives hits at a glance



Example for Scoring:

Scoring (circle colour):

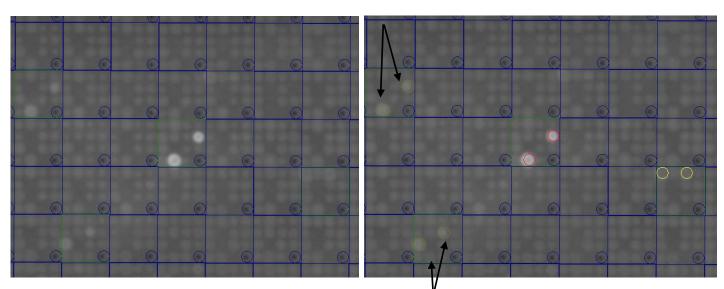
Red – 3 = strong

Brown – 2 = moderate

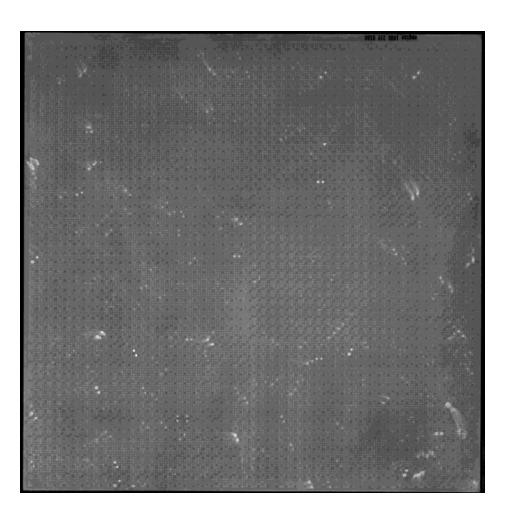
Yellow-1=low

As brown circles of score 2 are poorly visible, these are marked by arrows in addition

The same section of an Array was cropped and is shown without (left) and with (right) scoring of positive hits (bright spots), as displayed by the software.



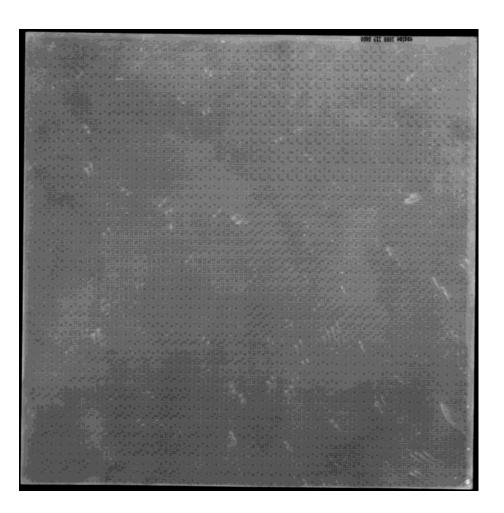
- Hits clearly detectable at a zoom of 50% in the software, used for analysis were marked with a score of "3"
- Hits clearly detectable at a zoom of 200% in the software, used for analysis were marked with a score of "2"
- Hits visible, but not standing out prominently at zoom of 200 % in the software, used for analysis were marked with a score of "1"



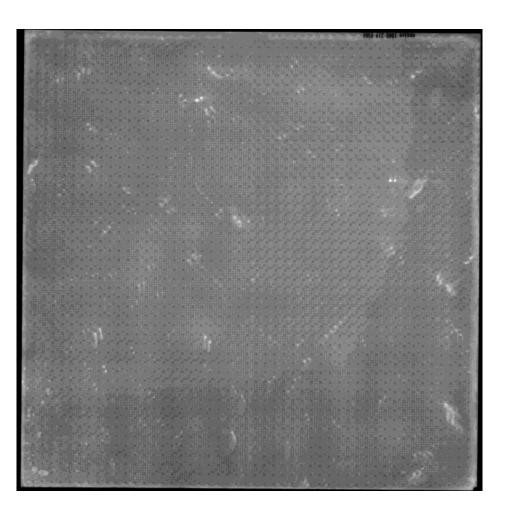
- Sample: Serum Pat1 1:1000
- Sec antibody: anti human IgG (Fc spec)-AP 1:5000
- Array: engine_1008_219_0100
- Positives total: 265
 - Intensity 1:86
 - Intensity 2: 140
 - Intensity 3: 39



- Sample: Serum Pat2 1:1000
- Sec antibody: anti human IgG (Fc spec)-AP 1:5000
- Array: engine_1008_219_0300
- Positives total: 239
 - Intensity 1: 62
 - Intensity 2: 130
 - Intensity 3: 47

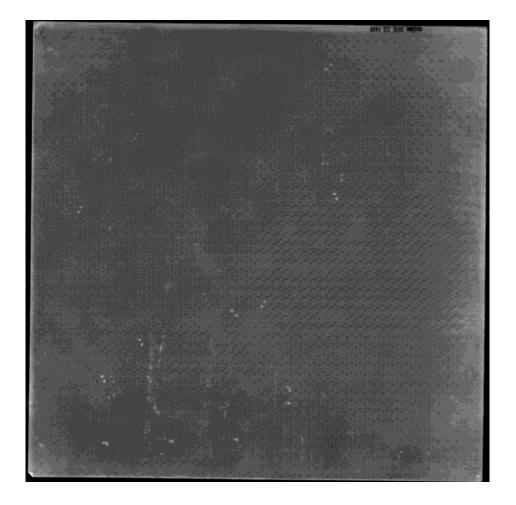


- Sample: Serum Pat3 1:1000
- Sec antibody: anti human IgG (Fc spec)-AP 1:5000
- Array: engine_1008_219_0400
- Positives total: 114
 - Intensity 1:38
 - Intensity 2: 54
 - Intensity 3: 22



- Sample: Serum Pat₄ 1:1000
- Sec antibody: anti human IgG (Fc spec)-AP 1:5000
- Array: engine_1008_219_0500
- Positives total: 242
 - Intensity 1: 75
 - Intensity 2: 115
 - Intensity 3: 52





- Sample: Serum Pat5 1:1000
- Sec antibody: anti human IgG (Fc spec)-AP 1:5000
- Array: engine_1008_221_0400
- Positives total: 61
- Intensity 1: 41
- Intensity 2: 11
- Intensity 3: 9



Summary



For all samples, antibody binding to proteins on the membrane could be detected.

Distribution of hits and intensity of hits:

			visual score	
	total	1	2	3
Patı	265	86	140	39
Pat2	239	62	130	47
Pat ₃	114	38	54	22
Pat ₄	242	75	115	52
Pat ₅	61	41	11	9

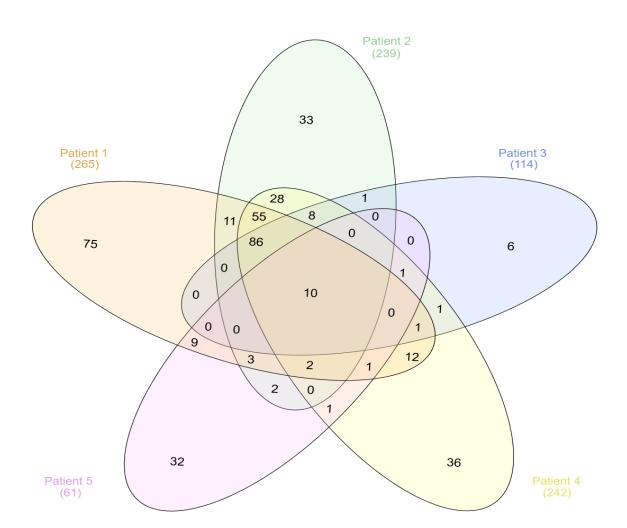
Overlap in positivity for all samples IgG

Total hits IgG

single	181
double	65
triple	69
quadruple	88
quintuple	10

Genes identified by all 5 patients excluding secondary antibody binding

Gene	Frame			Patie	Anti	# positive			
Gene	Traine	1	2	3	4	5	lgG		
MAFK	inframe	3	3	1	3	1	0	5	
HNRNPUL ₂	inframe	3	3	3	3	1	0	5	
MYH14	inframe	3	2	2	3	1	0	5	
CREBBP	inframe	2	3	2	2	1	0	5	
AATF	inframe	3	3	3	3	1	0	5	
CSDE1	inframe	3	2	3	3	1	0	5	
DDX51	inframe	3	3	3	3	1	0	5	



engine the biomarker company

ENGINE REPORT || screening of 5 patients







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