

# Streamlined Bead-Based Assays for the High-Throughput Analysis of Antibodies and Proteins



5th Annual SLAS Conference 2016  
San Diego

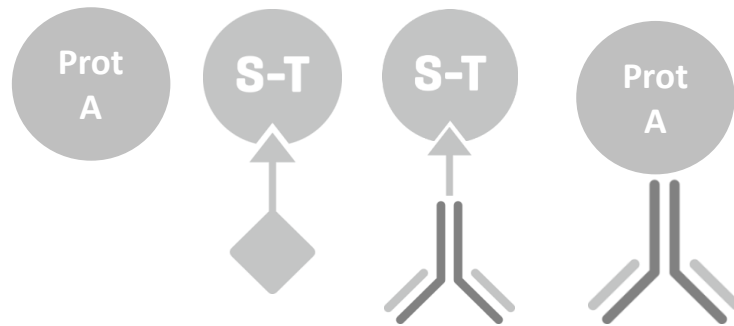
# PAIA assays in a nutshell

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- ❑ No-wash bead-based assays with fluorescence detection
- ❑ Sample volume < 10  $\mu$ L per well
- ❑ High throughput in 384-well plates
- ❑ < 60 min/plate, can be enhanced by incubation in parallel
- ❑ Incubation, bead separation and detection in the same plate
- ❑ Runs on standard fluorescence readers
- ❑ Applications: Quantification of proteins and antibodies and binding screens
- ❑ Assays kits contain *PAIAplates* with dried capture beads and a ready-to-use buffer with fluorescence marker for a specific analyte

# Assay principle

We use functionalized capture beads



to bind analytes from the sample (e.g. antibodies, proteins)



and detect the binding using fluorescence markers

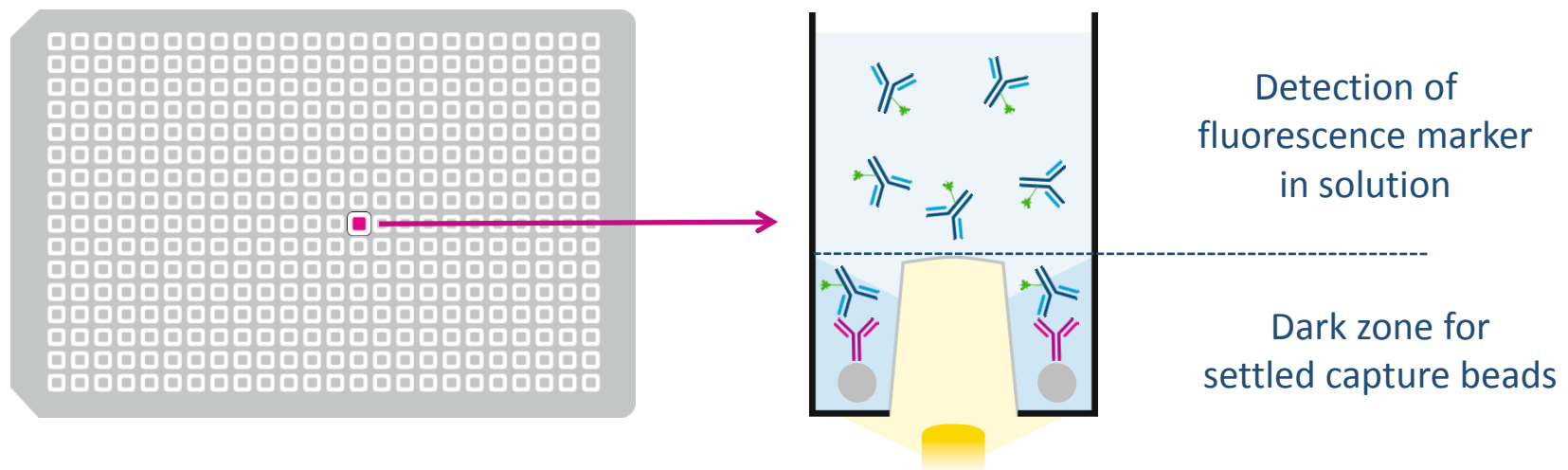


# Our innovation

- We exploit the advantages of bead based assays
- We measure the fluorescence intensity in solution (not on the beads)
- We use one microplate for incubation, bead separation & detection

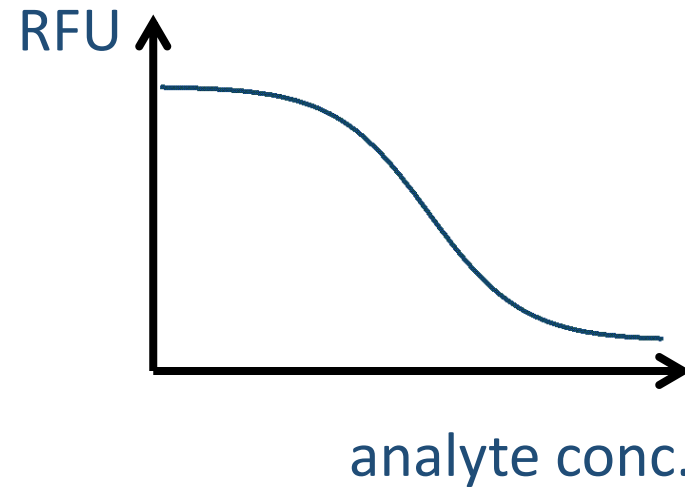
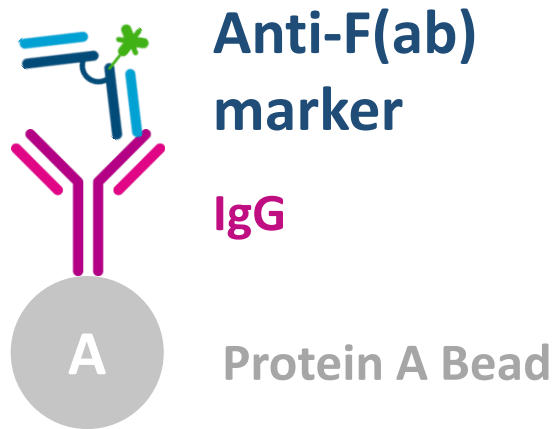
## Features of the PAIAplate

- 384-well plate with standard dimensions
- Black bottom for blocking excitation and emission of settled beads
- Transparent protrusion to direct bead settling to the well bottom and enabling fluorescence detection of unbound markers in solution

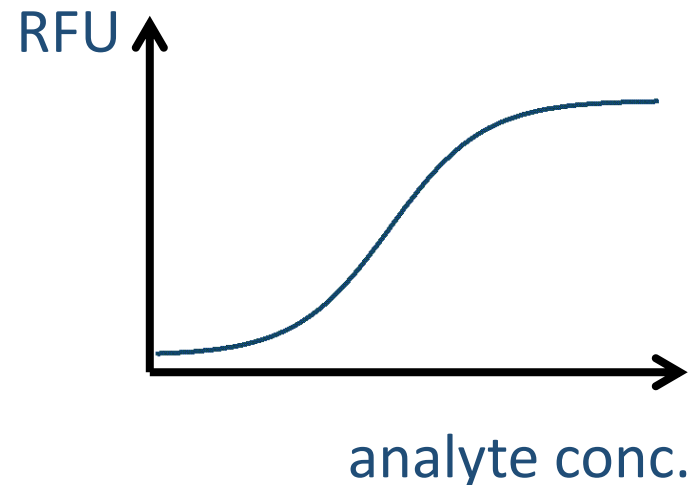
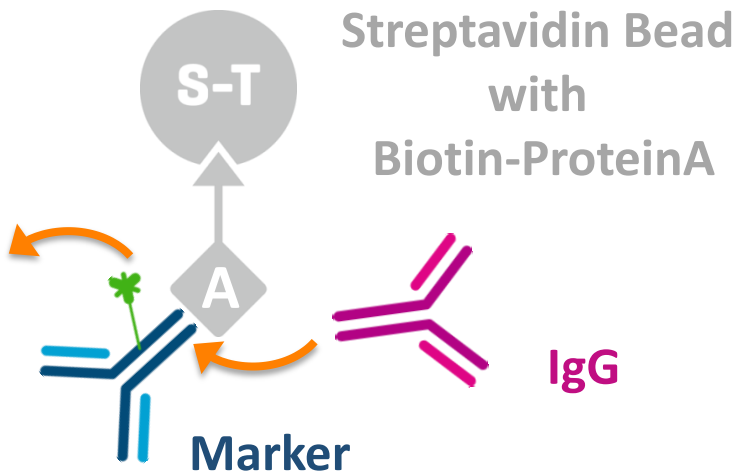


# Generic assay formats

## Direct assays (analyte binds marker to the bead)



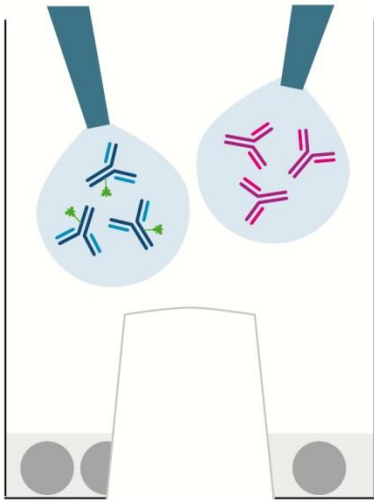
## Competitive assays (analyte replaces marker from the bead)



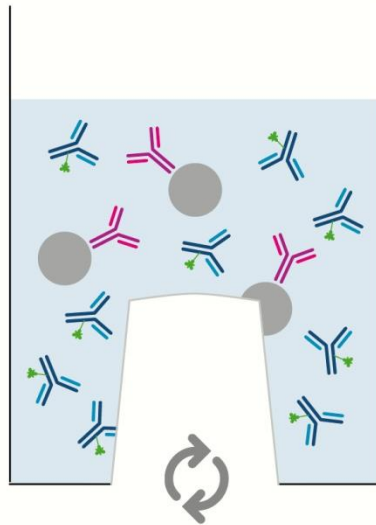
# Assay workflow

- Starts with PAIAplate containing pre-dispensed dried capture beads
- Add 50 $\mu$ L ready-to-use buffer with marker into the wells you want to use
- Add 2-10  $\mu$ L of sample containing the analyte
- Incubate the plate on orbital shaker for 30 mins, then let beads settle 15 mins.
- Measure on a fluorescence plate reader or microscope (bottom read).

1) Buffer and analyte addition

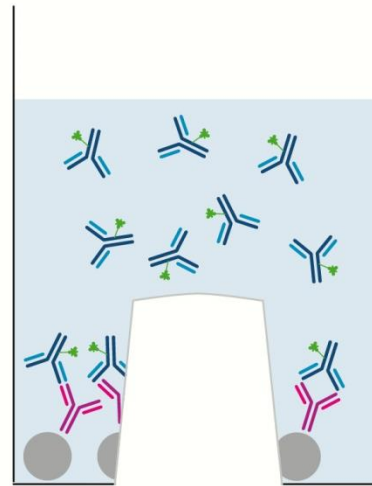


2) Analyte capture  
30 mins

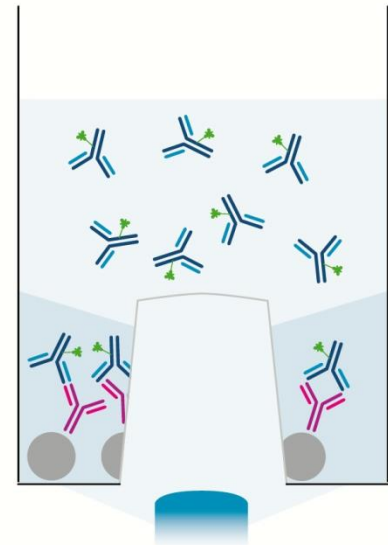


Shaking at RT

3) Separation  
15 mins



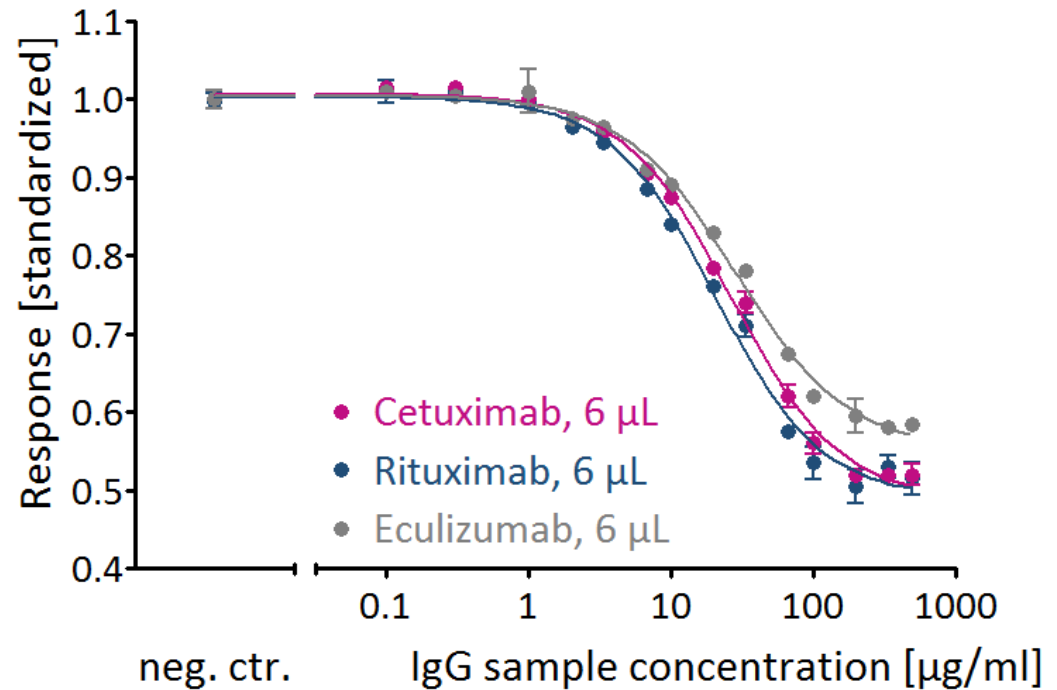
4) Measurement  
5 mins



Fluorescent read-out

# Human IgG quantification: early clone screening

- Direct assay format
- Capture via ProteinA beads, detection with human IgG Fab-specific marker



# Pilot study early clone selection (IgG1)

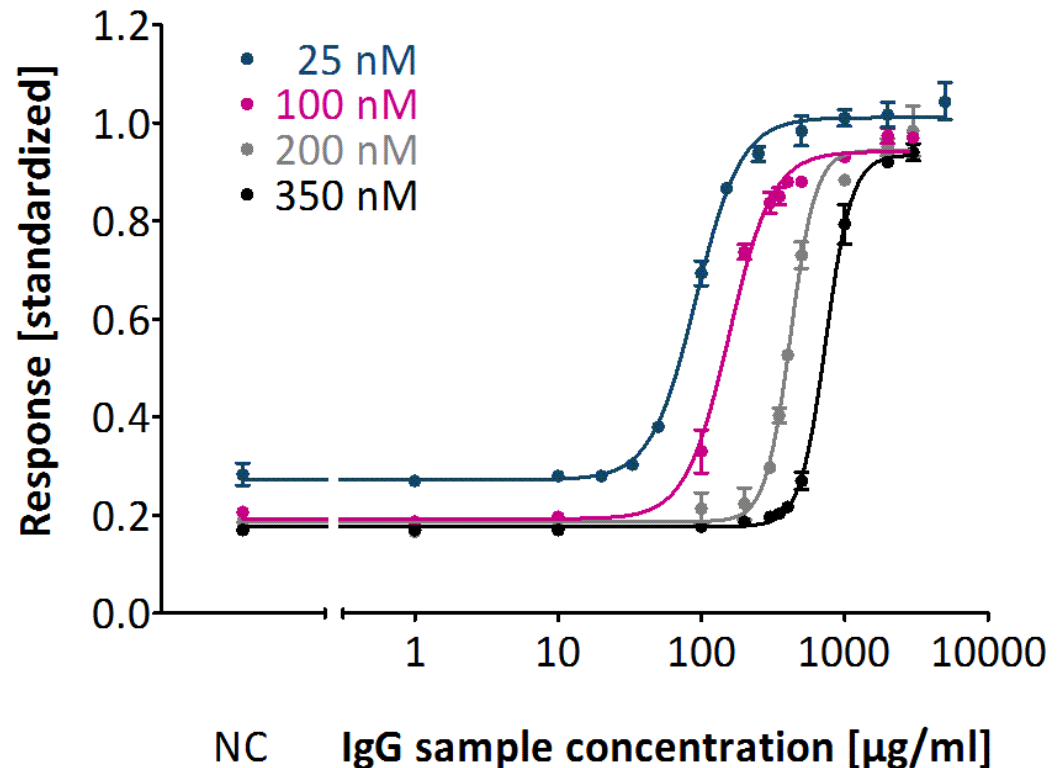
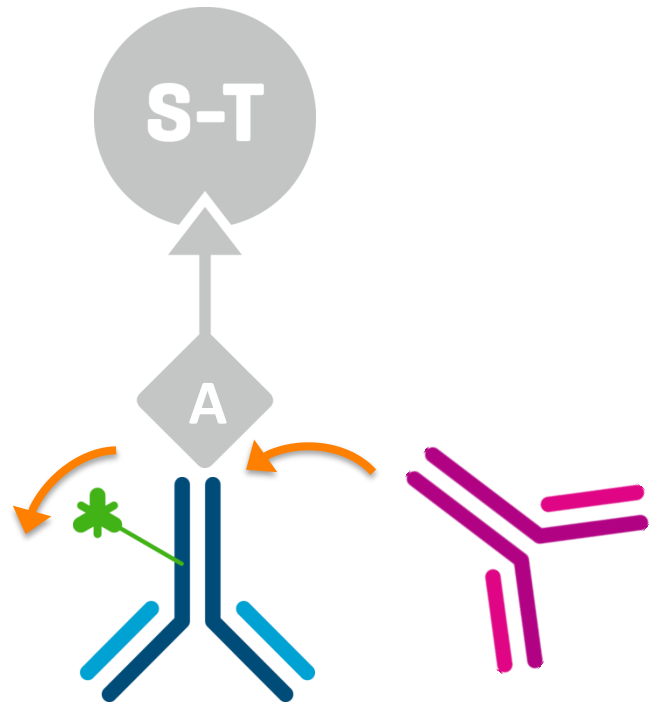
- Cell culture supernatants from CHO cells 10 days after seeding
- Four 96-well plates combined onto one PAIAplate
- Four replicate PAIAplates tested in parallel on four shakers
- All plates (1536 samples and standards) run and analyzed in 2,5 hours
- 169 of 192 top clones (88%) identical with Octet384 ranking
- Identical results for each PAIAplate
- Calibration can be done on a separate plate

	plate 1				plate 2				plate 3				plate 4			
Calibration	1	2	3	4	1	2	3	4	1	2	3	4	1	2	3	4
	9	8	6	1	9	8	6	1	3	3	2	0	4	4	2	0
	25	26	22	18	29	30	26	23	23	24	20	16	29	31	26	24
	8	7	5	1	11	10	8	3	6	5	4	0	8	7	6	1
	66	76	69	77	77	89	83	94	69	79	72	81	83	96	91	104
	52	59	52	56	63	72	65	73	58	66	59	65	58	65	58	64
	37	40	34	34	40	44	38	39	34	37	31	30	39	43	37	38
	12	11	9	4	16	15	12	8	13	12	10	5	13	13	10	5
	113	129	132	149	102	118	117	133	100	115	114	130	96	111	109	124
	105	121	121	138	101	117	116	132	99	114	112	128	97	112	109	125
	10	10	8	3	10	9	7	2	9	8	7	2	13	12	10	5
	102	117	116	132	118	135	139	157	103	119	118	134	122	139	145	163
	59	67	60	66	59	67	60	66	64	73	66	74	66	75	69	77
	117	134	138	156	115	131	135	152	106	122	122	139	99	114	113	129
	140	158	172	191	167	183	215	230	145	163	180	198	146	163	181	198
	64	73	66	74	65	75	68	76	57	64	57	63	50	57	50	53
	117	134	138	156	96	111	108	124	85	98	93	106	95	110	108	123
	10	10	8	3	12	11	9	4	8	8	6	1	13	13	10	5
	12	12	10	4	16	15	13	8	12	11	9	4	17	17	14	9
	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
	15	15	12	7	17	17	14	9	14	14	11	6	18	18	15	11
	118	135	139	157	160	176	203	219	124	141	148	167	130	147	157	175
	64	74	67	74	57	65	58	64	48	53	47	50	65	75	68	76
	84	97	92	105	94	108	105	120	89	103	99	113	87	100	96	109
	8	8	6	1	12	11	9	4	8	7	6	1	11	11	8	3
	130	147	156	175	168	184	216	230	114	130	133	151	128	145	153	172
	77	89	83	94	88	102	97	111	72	83	76	86	87	101	96	110



# IgG quantification for bioprocess development

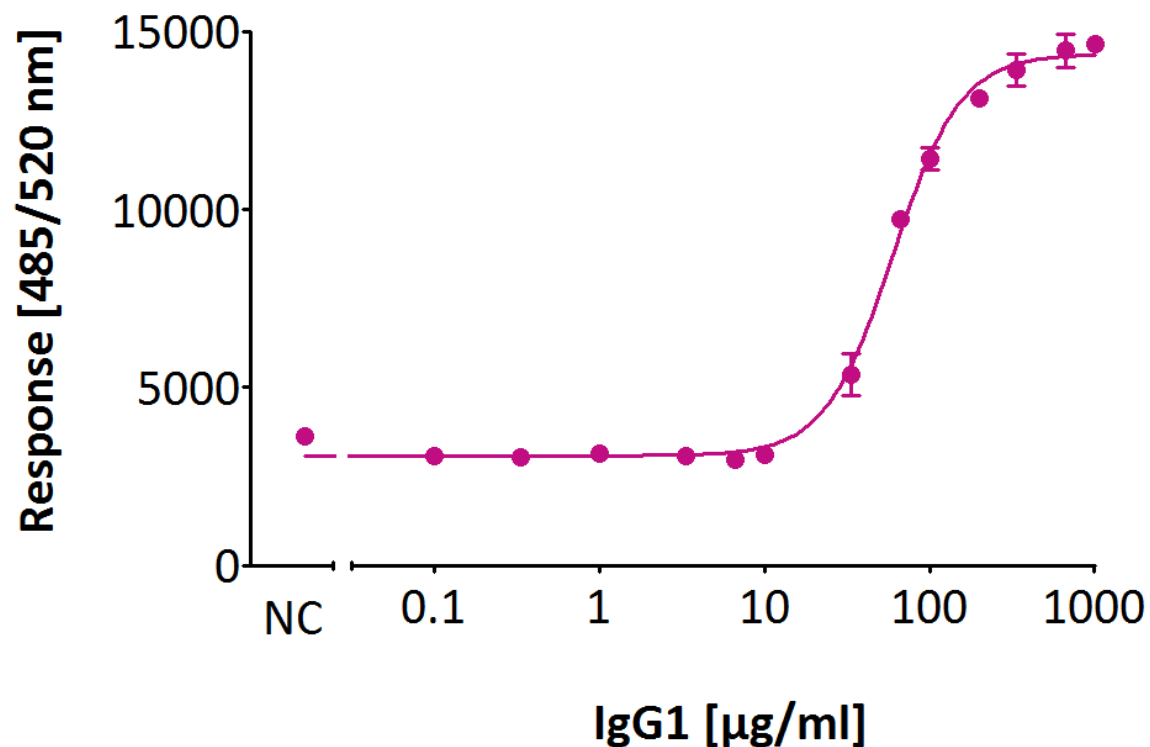
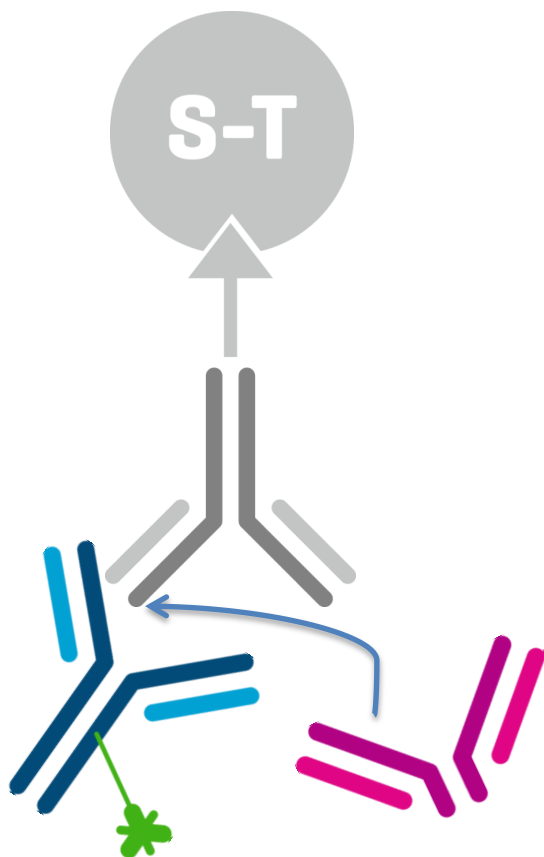
- Streptavidin capture beads with biotinylated Protein A
- Fluorescence marker: labeled human IgG
- Adjustment of dynamic range by changing the binding capacity of capture beads



# Fab Assay: CH1 Capture

- Competitive assay format
- Capture via bead-coupled anti human CH1 antibody

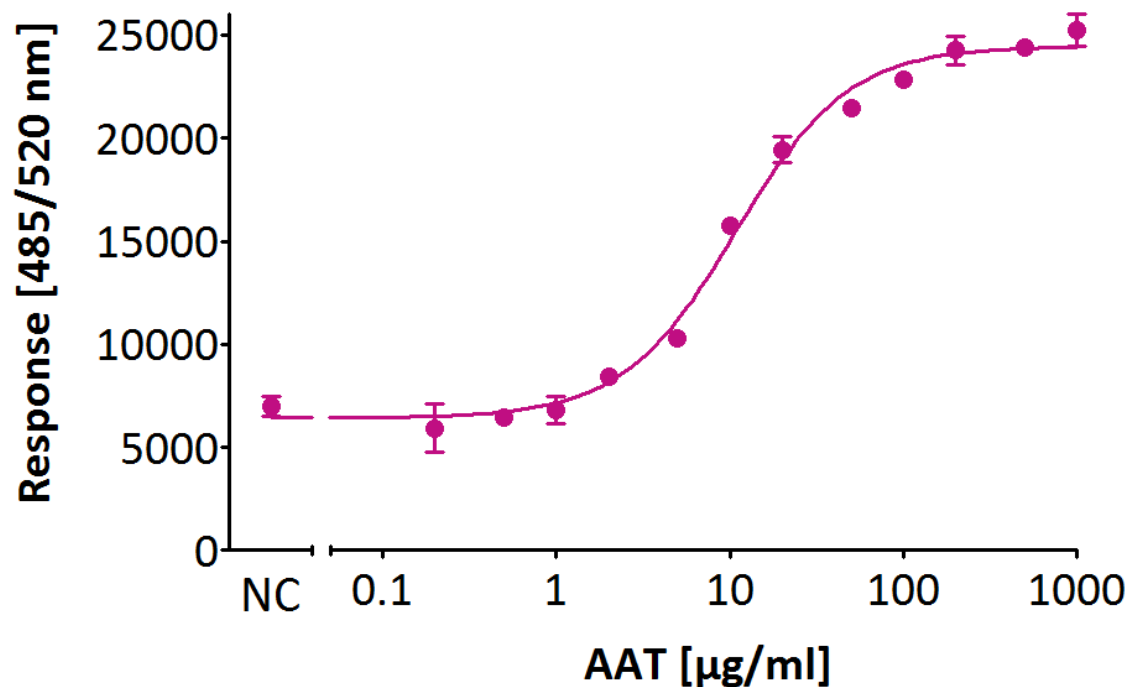
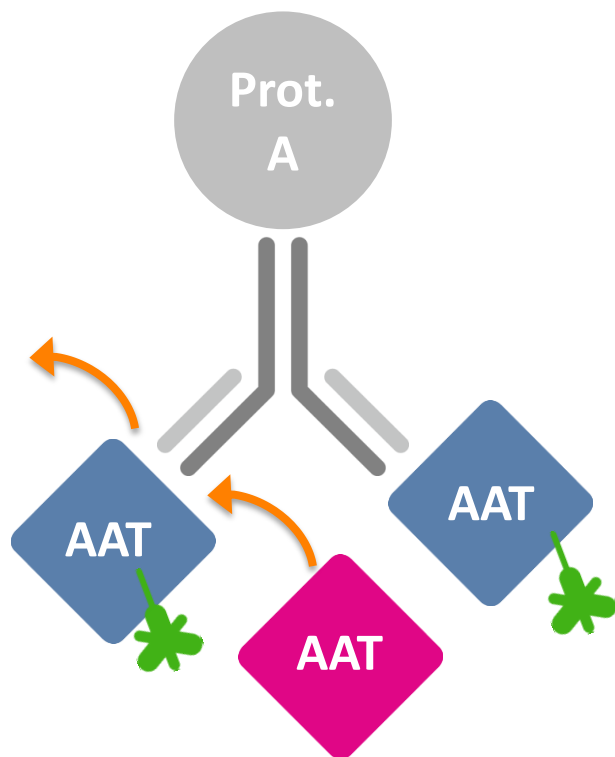
beta



# Custom Assay: human $\alpha$ -Antitrypsin (AAT)

- Competitive assay format
- Capture via bead-coupled rabbit anti-AAT antibody

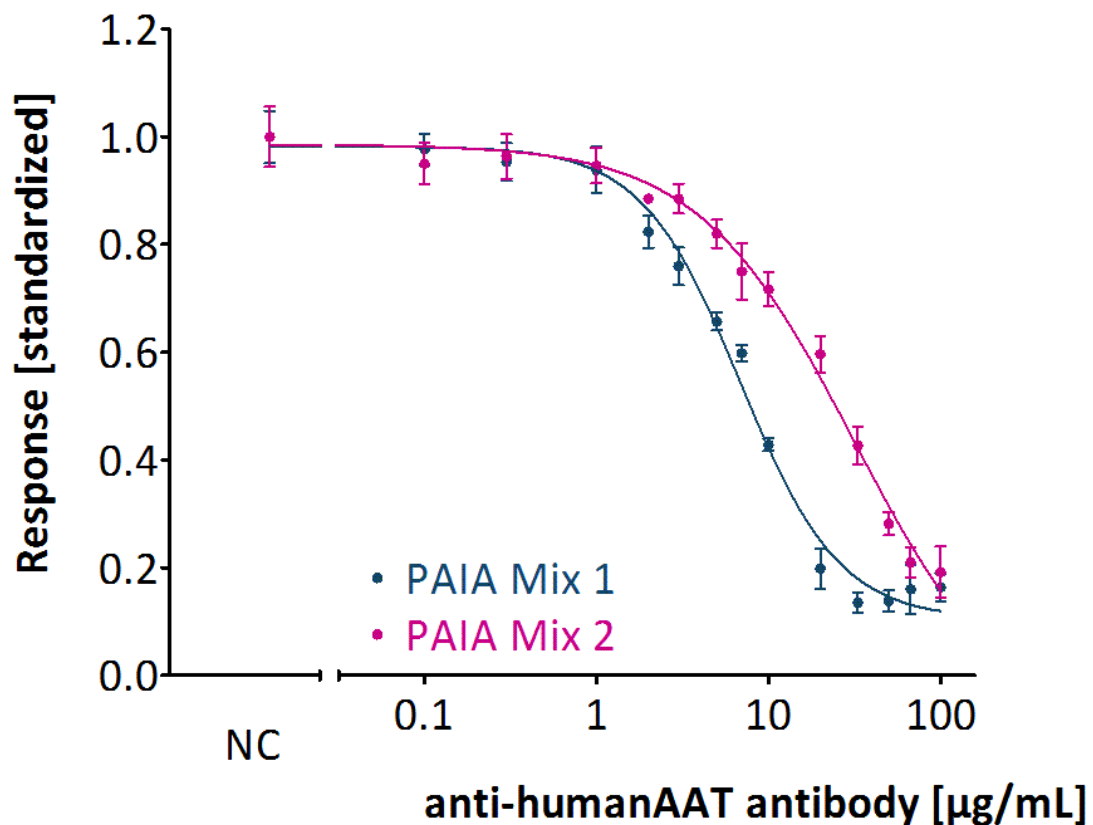
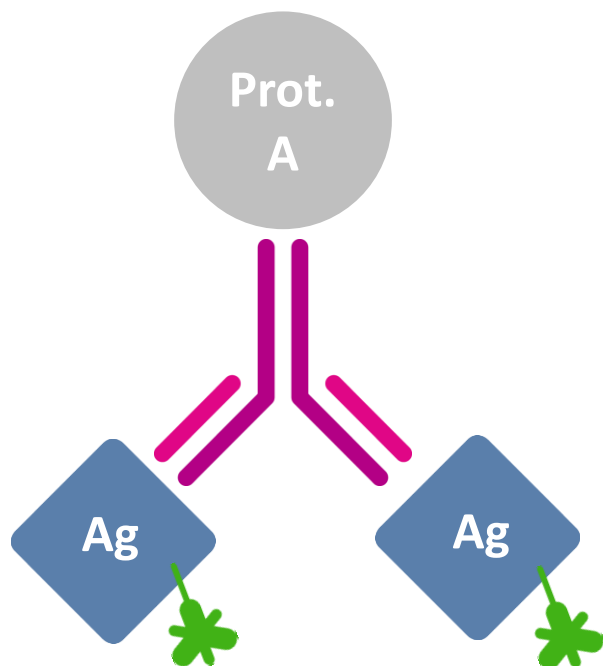
custom



# Antibody screening assay

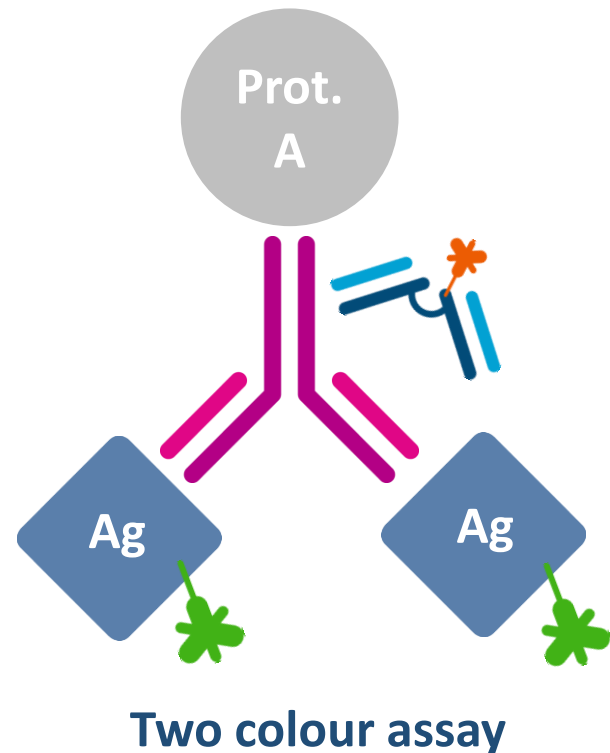
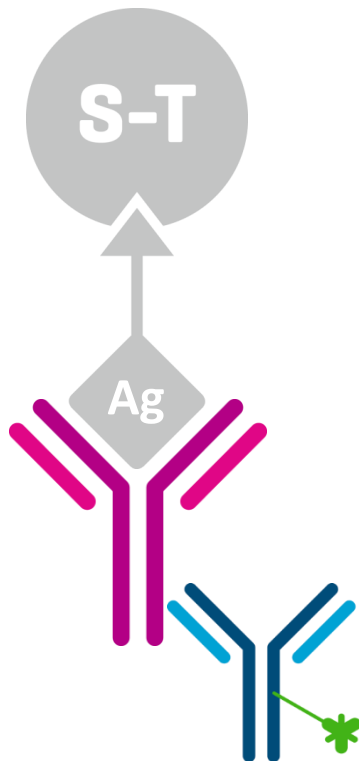
- Capture Beads: Protein A
- Fluorescence marker: labeled antigen (Ag) here: AAT

custom



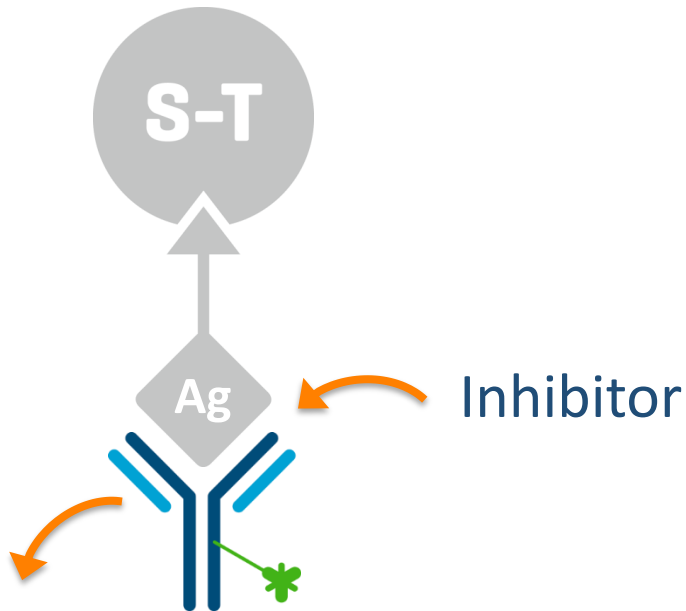
# Antibody screening assay formats

- Capture Beads: Streptavidin with biotinylated antigen
- Fluorescence marker: labeled IgG anti-Fc specific marker
- Capture Beads: ProteinA
- Fluorescence marker: labeled antigen, optional second IgG anti-Fc specific marker



# Inhibition screening assays

- Capture Beads: Streptavidin with biotinylated antigen/target
- Fluorescence marker: labeled antibody or target binding protein



Inhibition generates high fluorescence signals

# Take Home Message

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- Streamlined bead-based immunoassay platform with
- High throughput and simple, automation-friendly workflow
- No washing, no blocking, no sensor regeneration
- Sample volumes: < 10  $\mu$ L
- Fluorescence reader or fluorescence microscope (e.g. Cellavista)
- Straightforward data analysis (PAIA standalone or reader software modules)



**Time saving, usually capex-free and robust tool  
for screening applications**

# The Company

- Founded in 2014, privately owned and funded
- Research & technology development since early 2013
- Patent pending platform technology
- Based in Cologne, Germany
- Products: assay kits and custom assay development





# Thank you!

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