



BioPhase 8800 system with NF

Year: 2025

TRKE **CHARGE**

Maintain all existing capabilities on the BioPhase 8800 system.

Benefit from the addition of native fluorescence detection:

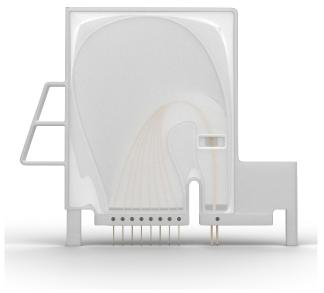
- 1. Protein detection with improved sensitivity compared to traditional UV detection.
- 2. Faster, more confident peak integration than traditional UV due to significantly reduced noise and flattened baseline.
- 3. No need for protein-dye conjugation during sample preparation, speeding up sample preparation.

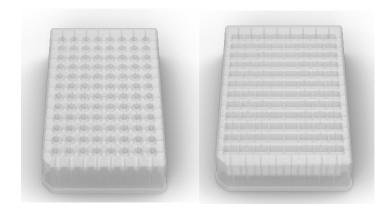




Features - hardware

- Factory-built cartridge with push-in load: Reduce user error
- a capillaries parallel processing: Speed up your sample analysis
- Built-in electrodes, interface block, and aperture: Reduce maintenance
- Smart chip with SN, usage tracking: Improve traceability
- Precise temperature capillary control: Maximize data reproducibility
- Plate-based sample/reagent: Rutomate processes
- Software-driven detector selection: Simplify detector changes and enable a single sequence with multiple detector types
- Reduced stray light and higher transmission: Gain higher signal intensity







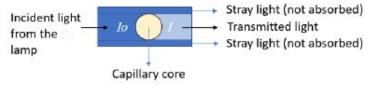


Figure 3. Schematic diagram of the stray light.



Features - software

The user can choose to set up either in a local or a network configuration



Intuitive touchscreen interface



Drag and drop method programming



BioPhase 8800 Driver for Empower software & documentation + 003





Native fluorescence of proteins: A new way of detection

The native fluorescence of proteins originates from residues of three aromatic amino acids: **tryptophan** , tyrosine, and phenylalanine

Tryptophan : $\lambda_{\rm ex}/\lambda_{\rm em}$ =280/350 nm

-- High emission intensity, the primary native fluorescence source in proteins

Tyrosine : $\lambda_{\rm ex}/\lambda_{\rm em}$ =204/304 nm & $\lambda_{\rm ex}/\lambda_{\rm em}$ =220/304 nm

-- High emission intensity, but quenched by its interaction with the peptide chain via energy transfer

Phenylalanine : $\lambda_{px}/\lambda_{pm} = 270/280 \text{ nm}$

-- Low emission intensity

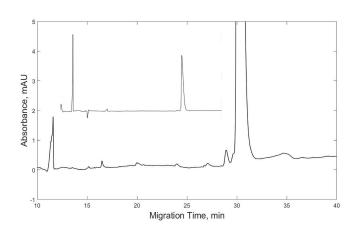
No Need to rely on low-wavelength amide bond or fluorescent tag labeling for protein detection





BioPhase 8800 system with NF: non-reduced CE-SDS

Traditional UV 220 nm detection

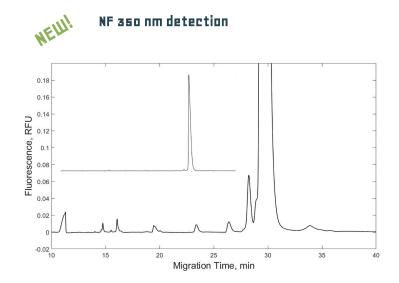


Protein labeling is not needed

Requires high sample concentration

Non-reduced CE-SDS

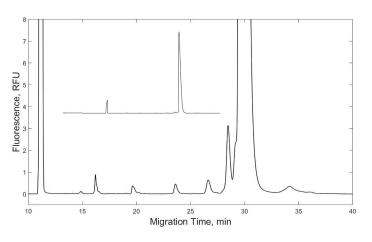
Sample: 1 mg/mL [1 mg/mL] non-reduced NIST Ig6



Protein labeling is not needed

Improved S/N ratio & flat baseline compared to UV detection

LIF 600 nm detection



LIF dye conjugation to protein is required

Improved S/N ratio > 10X & flat baseline compared to UV detection



BioPhase 8800 system with NF: non-reduced purity for mAb

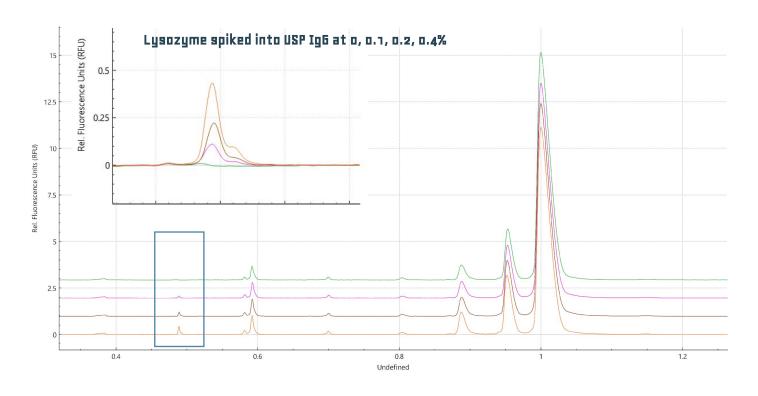
- The measured relative abundance of total fragments was comparable among the three detection approaches
- The measured relative abundance of aggregates with NFD is equivalent to that with LIF
- The aggregate peaks were not detectable with UV due to interference by the baseline anomalies

Comple Let	Detection	Detection Relative Abundance, %					, %		
Sample Lot	Detection	L	Н	HL	НН	HHL	Total	Intact mAb	Aggregates
	UV	0.29	0.13	0.15	0.18	0.78	1.53	98.48	ND*
1	NFD	0.24	0.14	0.15	0.24	1.00	1.77	97.84	0.40
	LIF	0.18	0.15	0.15	0.22	0.92	1.63	97.97	0.40
2	UV	0.29	0.14	0.11	0.17	1.07	1.78	98.22	ND*
2	NFD	0.22	0.17	0.14	0.21	1.26	2.00	97.62	0.38
	LIF	0.24	0.14	0.14	0.25	1.09	1.86	97.77	0.37
	UV	0.28	0.19	0.13	0.18	0.96	1.73	98.27	ND*
3	NFD	0.21	0.20	0.15	0.22	1.11	1.88	97.74	0.38
	LIF	0.24	0.14	0.16	0.26	1.06	1.85	97.80	0.35

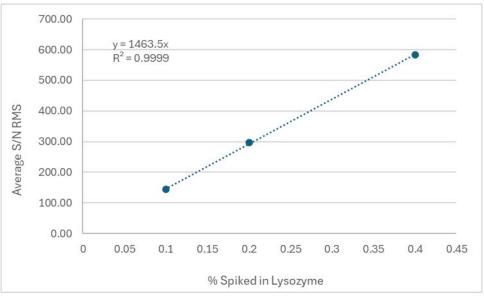
*ND – not detected. With UV absorbance detection, the aggregate peaks were not detected due to interference by the baseline anomalies.



BioPhase 8800 system with NF: Quantitation of 0.01% spiked in lysozyme



NF 350 nm



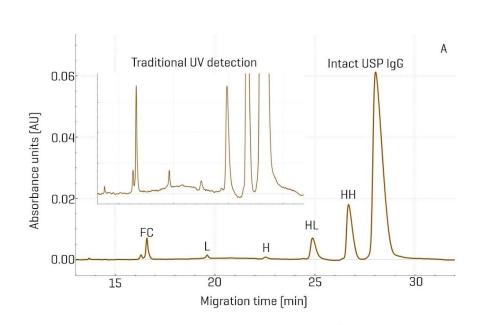
Sample: non-reduced USP Ig6 at 1 mg/mL.

- NFD 350 nm: S/N RMS at 0.1% spike in > 1000, linear extrapolation S/N RMS at 0.01% spike in > 10.
- UV 220 nm: S/N RMS at 0.1% spike in ~ 20



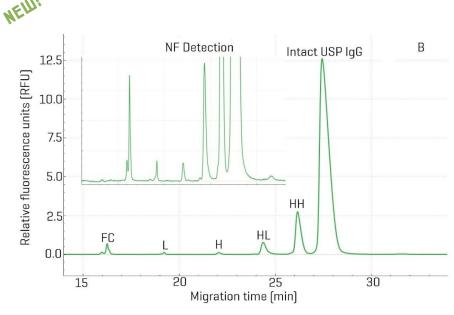
BioPhase 8800 system with NF: Quantification of protein fragments





C

NF detection USP Ig6

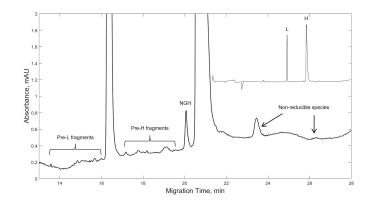


	UV det	ection	NF det	ection
	CPA %	Average S/N RMS	CPA %	Average S/N RMS
USP IgG-HC	0.53	78	0.44	185
NIST mAb- HL	0.25	20	0.19	76
NIST mAb-HH	0.22	19	0.21	83



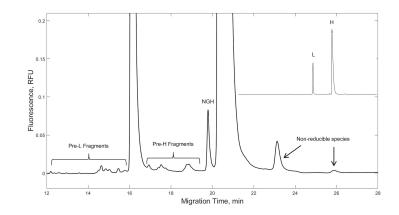
BioPhase 8800 system with NF: reduced CE-SDS

Traditional UV 220 nm detection

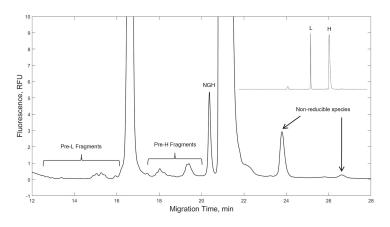


NFD 350 nm

NFD 350 nm detection



LIF-600 nm detection



Protein labeling is not needed

Requires high sample concentration

Protein labeling is not needed and Improved sensitivity & flat baseline compared to UV detection LIF dye conjugation to protein is required

Improved sensitivity >10X & flat baseline compared to UV detection

Reduced CE-SDS

Sample: 1000 µg/mL [1 mg/mL] reduced NIST Ig6



BioPhase 8800 system with NF: glycan occupancy

Comple Let	Glycosylation Site Occupancy on Heavy Chain, %					
Sample Lot	UV Absorbance	NFD	LIF			
1	99.30	99.33	99.33			
2	99.31	99.32	99.32			
3	99.31	99.33	99.33			

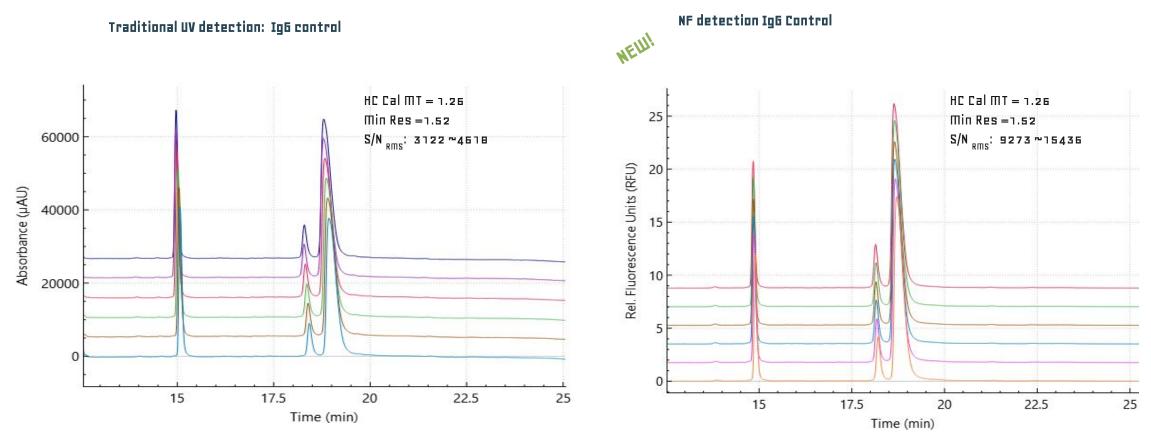
The determined glycosylation site occupancy was consistent, in the range of 99.30—99.33%, using the three detection approaches.

NIST mab

6lycosylation site occupancy = 1- N6H/H %



BioPhase 8800 system with NF: Consistency and repeatability



Comparable profiles observed with the same level of repeatability



BioPhase 8800 system with NF: Consistency and repeatability

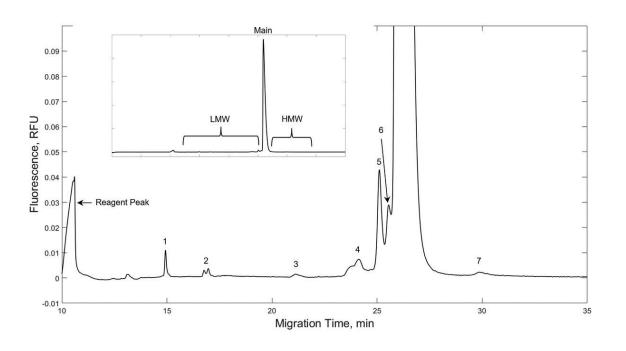
L	(LC+HC	C)%- UV	(LC+HC)%- NFD	HC/(HGH_	HC)% -UV	HC/(HGH_1	HC)% -NFD
Intra-Capillary	Average	% RSD	Average	% RSD	Average	% RSD	Average	% RSD
A	92.55	0.07	91.63	0.04	89.86	0.11	89.77	0.04
В	92.50	0.05	91.63	0.03	89.79	0.07	89.76	0.02
С	92.50	0.06	91.63	0.01	89.80	0.07	89.75	0.02
D	92.52	0.04	91.63	0.02	89.80	0.06	89.76	0.02
E	92.56	0.06	91.62	0.01	89.84	0.10	89.74	0.02
F	92.51	0.04	91.62	0.03	89.79	0.06	89.75	0.03
G	92.53	0.07	91.61	0.03	89.84	0.11	89.73	0.03
Н	92.52	0.07	91.64	0.05	89.81	0.11	89.76	0.06
Inter-capillary N = 48	92.52	0.06	91.63	0.03	89.82	0.09	89.75	0.03

Comparable sample purity with the same level of repeatability



BioPhase 8800 system with NF: Emicizumab analyzed with CE-SDS

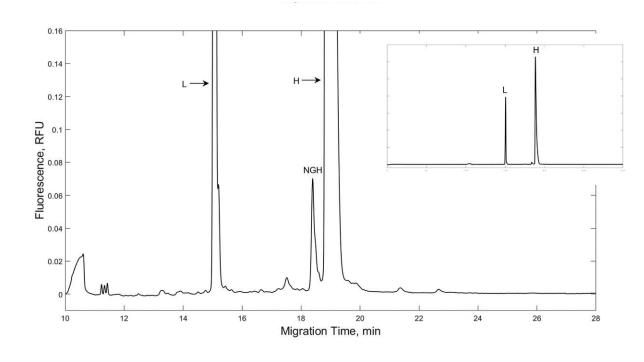
Non-reduced



LMW	Main	HMW
3.02%	96.85%	0.13%

Sample: o.s mg/mL Emicizumab, Injection: 20 s @-s kV

Reduced

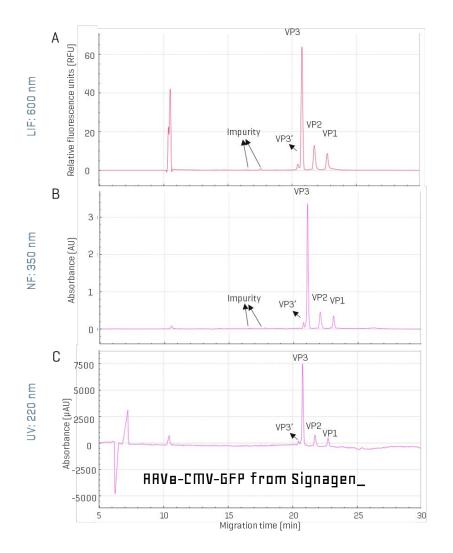


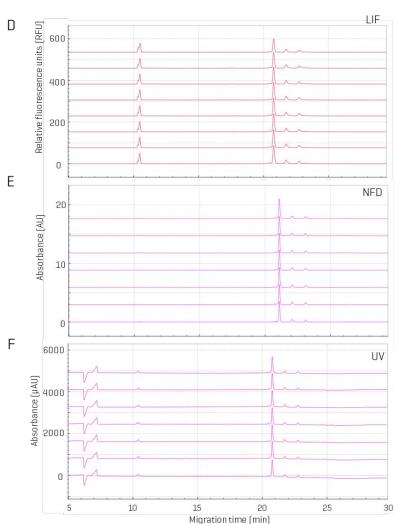
6lycosylation site occupancy: 98.01%





AAVe capsid protein analysis on BioPhase eeoo system

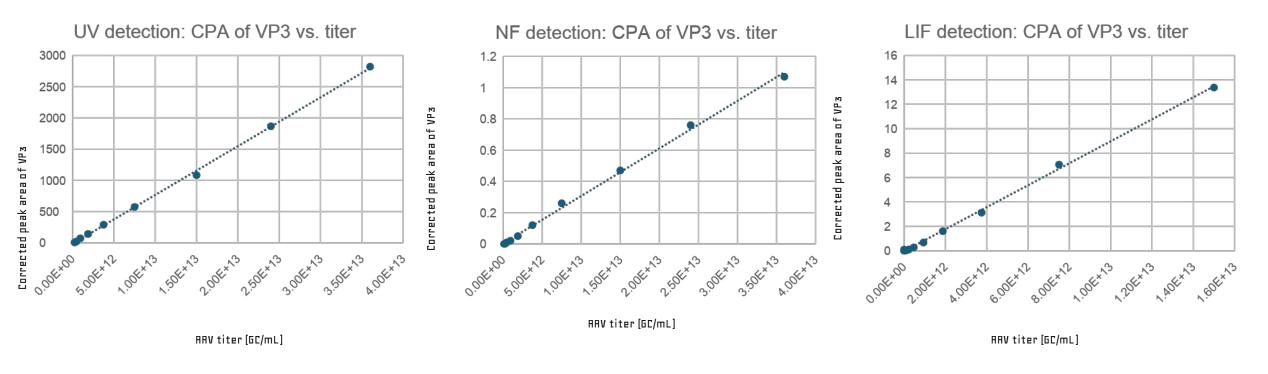




- Comparable capsid protein profiles across three different detections.
- Comparable repeatability across capillary, across injections was observed.



AAVe capsid protein analysis on BioPhase eeoo system: dynamic range



The linearity plot of the CPA of VP3 versus the titer of an AAV8-CMP-GFP sample using the CE-SDS with UV 220nm, NF, LIF detection on the BioPhase 8800 system

	LDR [6C/mL]	L00 (6C/mL)	LOD [6C/mL]	R²
ШV	4.7E+11 - 3.6E+13	9.4E+11	4.69E+11	0.999
NF	1.2 E+11 - 3.6E+13	4.7E+11	1.1 7 E+11	0.999
LIF	9.2E+08 - 1.5E+13	9.16E+08	3.05E+08	0.999



AAVe capsid protein analysis on BioPhase eeoo system: capsid ratio

VP 1

VP2

VP3

Capsid Ratios

Experimental	Experimental value						
	VPı	VP2	VP3				
LIF	1.0	1.2	4.5				
NFD	1.0	1.5	7.8				
ШV	1.0	1.4	7.5				
Amino acid co	intent						
Lys	32	22	17				
Тгр	15	12	12				
88	737	600	534				
Normalized b	y amino acid	content					
LIF	1.0	1.7	8.5				
NFD	1.0	1.9	9.8				
ШV	1.0	1.7	70.4				
Expected	1.0	1.0	10				

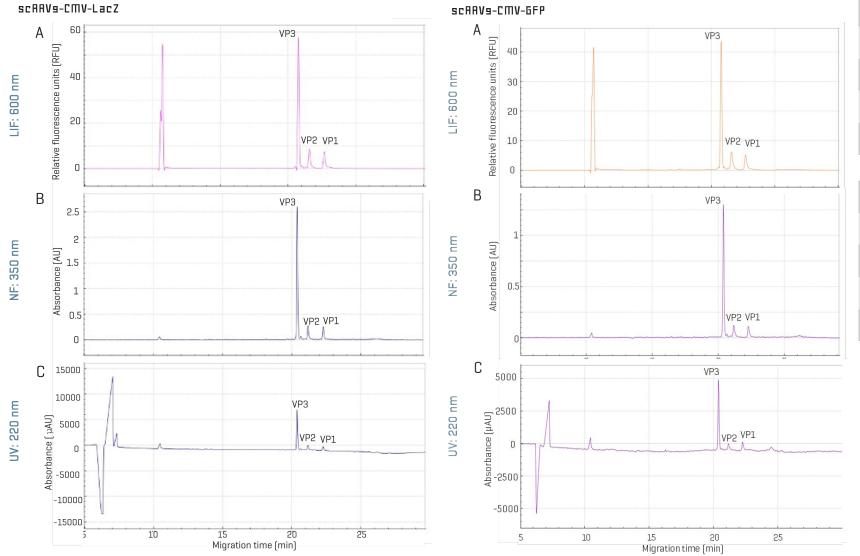
HADGYLPDWLEDNLSEGIREWWALKPGAPKPKANQQKQDDGRGLVLPGYKYLGPFNGLDKGEPVNAADA
HALEHDKAYDQQLQAGDNPYLRYNHADAEFQERLQEDTSFGGNLGRAVFQAKKRVLEPLGLVEEGAKT
HPGKKRPVEPSPQRSPDSSTGIGKKGQQPARKRLNFGQTGDSESVPDPQPLGEPPARPSGVGPNTM
HAGGGAPMADNNEGADGVGSSSGNWHCDSTWLGDRVITTSTRTWALPTYNNHLYKQISNGTSGGATNDN
TYFGYSTPWGYFDFNRFHCHFSPRDWQRLINNNWGFRPKRLSFKLFNIQVKEVTQNEGTKTIANNLTSTI
QVFTDSEYQLPYVLGSAHQGCLPPFPADVFMIPQYGYLTLNNGSQAVGRSSFYCLEYFPSQMLRTGNNFQ
FTYTFEDVPFHSSYAHSQSLDRLMNPLIDQYLYYLSRTQTTGGTANTQTLGFSQGGPNTMANQAKNWLP
GPCYRQQRVSTTTGQNNNSNFAWTAGTKYHLNGRNSLANPGIAMATHKDDEERFFPSNGILIFGKQNAAR
DNADYSDVMLTSEEEIKTTNPVATEEYGIVADNLQQQNTAPQIGTVNSQGALPGMVWQNRDVYLQGPIWA
KIPHTDGNFHPSPLMGGFGLKHPPPQILIKNTPVPADPPTTFNQSKLNSFITQYSTGQVSVEIEWELQKEN

Capsid ratio using NF is similar to the values using UV, and closer to theoretical values

SKRWNPEIQUTSNUUKSTSVDFAVNTEGVUSEPRPIGTRULTRNL



AAVs capsid protein analysis on BioPhase 8800 system



scAAV9-CMV-LacZ capsid ratio							
	VPı	VP2	VP3				
LIF	1.0	1.0	4.5				
NF	1.0	1.0	B.1				
ШV	1.0	1.0	B.2				

scAAV9-CMV-GFP capsid ratio							
	VPı	VP2	VP3				
LIF	1.0	1.2	5.2				
NF	1.0	1.5	8.0				
ЦV	٦.٥	7.4	8.6				

Comparable trends for capsid protein ratios were observed when RRVs is analyzed



ARV Capsid protein analysis using three different detection method

Sionagen AAVa-C	NP3		VP2		VP1			
mv-6FP	MT (min)	RSD% of MT	CPR%	RSD% of CPA%	of CPA% CPA%	RSD% of CPA%	CPR%	RSD% of CPA%
LIF-600 nm	20.94	0.6	65.8	0.3	17.1	0.7	13.4	1.1
NF -350 nm	21.28	0.7	74.7	0.3	13.30	1.2	8.80	2.9
UV -220 nm	20.90	0.5	74.5	0.8	13.00	3.5	9.70	5.4

Signagen_AAV9-		V	P3		VI	P2	VP1	
CMV-lacZ	MT (min)	RSD% of MT	СРЯ%	RSD% of CPA%	СРЯ%	RSD% of CPA%	СРЯ%	RSD% of CPA%
LIF-600 nm	20.50	0.4	70.30	1.2	14.00	1.90	13.40	1.1
NF -350 nm	20.89	0.8	B1.34	0.6	9.48	1.27	9.18	4.0
UV -220 nm	20.38	0.6	80.85	1.1	10.06	5.76	9.09	6.6

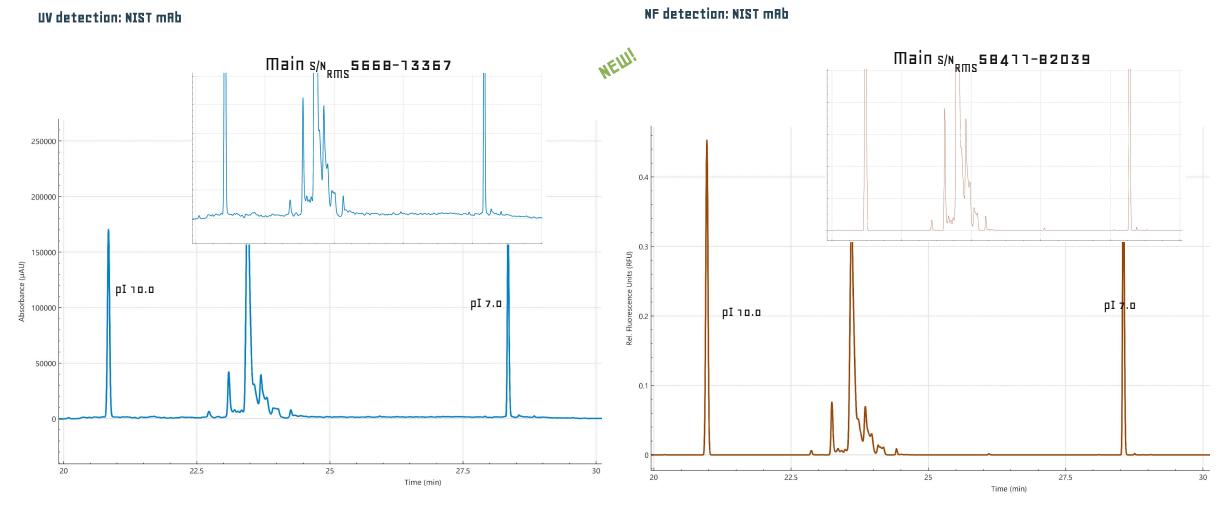
Comparable repeatability was observed between LIF, NF, and UV

Signagen_scAA		VP3				P2	VP1	
V9-CMV-6FP	MT (min)	RSD% of MT	СРЯ%	RSD% of CPA%	CPR%	RSD% of CPA%	СРЯ%	RSD% of CPR%
LIF-600 nm	20.44	0.5	69.02	0.3	14.89	0.5	12.06	1.0
NF -350 nm	21.20	1.1	80.12	0.5	9.56	1.1	9.15	2.8
UV -220 nm	20.56	0.9	87.90	1.2	9.56	7.1	8.55	12.0





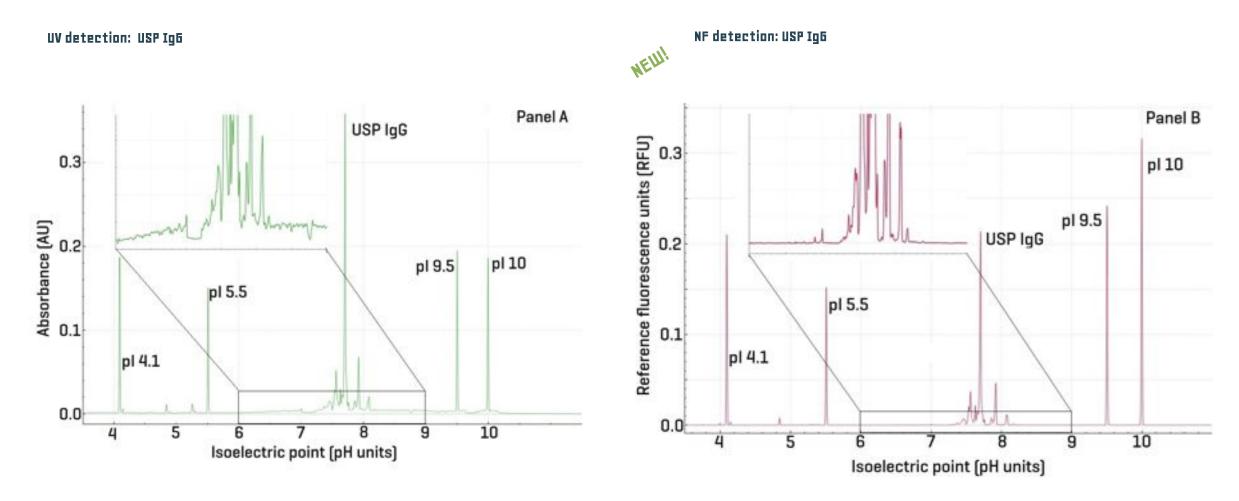
BioPhase 8800 system with NF: cIEF of NIST mAb



NF detection shows similar profiles for all charge variants with improved S/N ratios



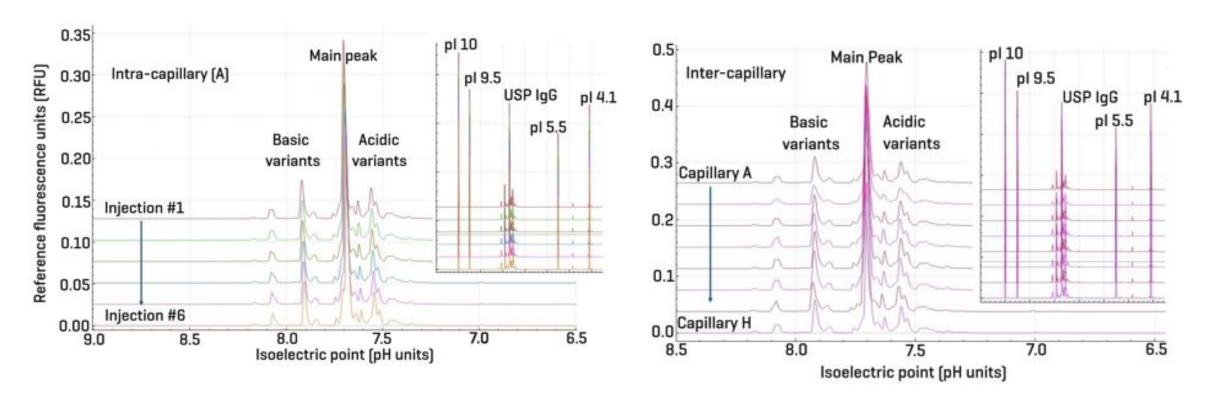
BioPhase 8800 system with NF: cIEF of USP Ig6



NF detection shows similar profiles for all charge variants with improved S/N ratios



BioPhase 8800 system with NF: repeatability



Intra and Inter-capillary reproducibility. The inset shows the zoomed-out electropherograms.



BioPhase 8800 system with NF: Consistency and repeatability

Intra-Capillary	Acidic %		Main %		Basi	ic %	Main pI Values		
	Average	% RSD	Average	% RSD	Average	% RSD	Average	% RSD	
A	15.72	0.4	63.81	0.5	20.46	1.18	7.66	0.1	
В	16.07	1.2	61.80	0.6	22.13	0.95	7.66	0.1	
С	16.25	0.6	61.25	0.5	22.50	0.91	7.66	0.2	
D	16.30	0.6	61.36	0.7	22.35	1.47	7.66	0.1	
E	15.94	0.8	62.43	0.8	21.63	1.82	7.66	0.1	
F	15.93	0.6	62.75	0.8	21.32	1.94	7.65	0.2	
G	15.75	0.4	63.03	0.5	21.22	1.27	7.66	0.1	
Н	15.46	1.1	64.46	0.7	20.08	1.49	7.66	0.1	
Inter-capillary 48 injections	15.93	1.8	62.61	1.8	21.46	2.0	7.66	0.1	

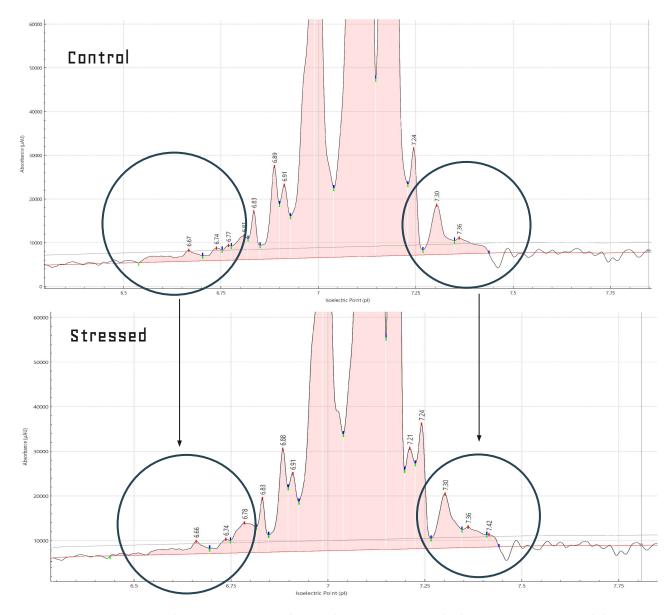
Comparable sample purity with the same level of repeatability



cIEF Profile comparison between control and stressed sample

UV detection

Baseline drifts make integration on the basic side more subjective/uncertain





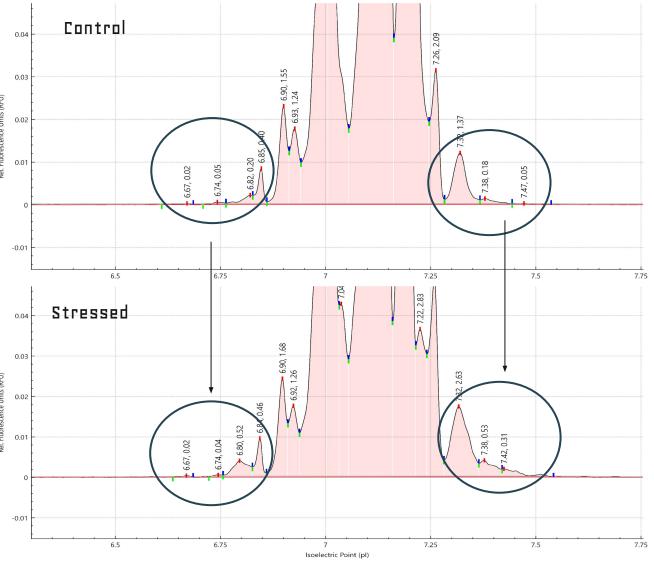
cIEF Profile comparison between control and stressed sample

NEW!

NF detection

More confident to integrate the basic side and reflect ξ the changes

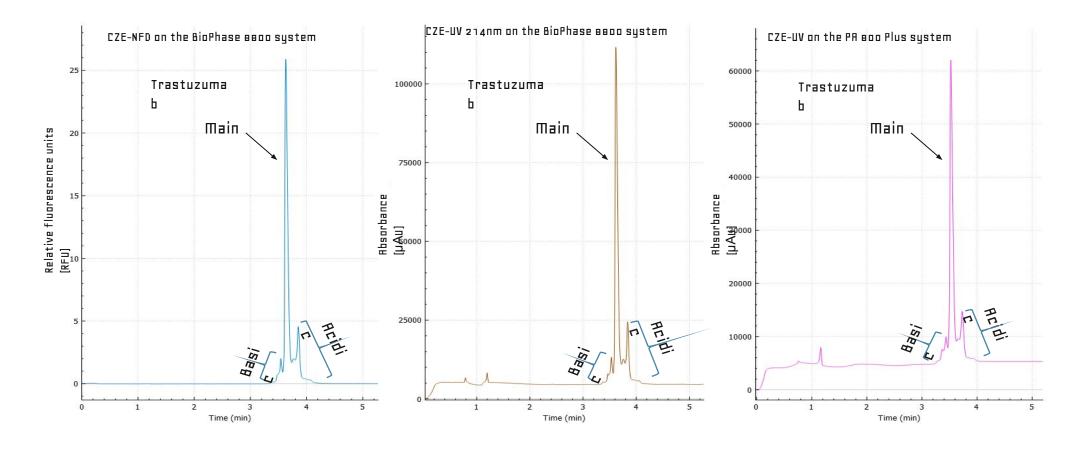
		Basic %	Main%	Acidic%	
Control	UV	3.39 78.20		18.12	
ИЕШ!	NFD	3.39	78.55	18.06	
Stressed	UV	6.28	71.84	21.81	
иеш!	NFD	8.84	69.57	21.55	







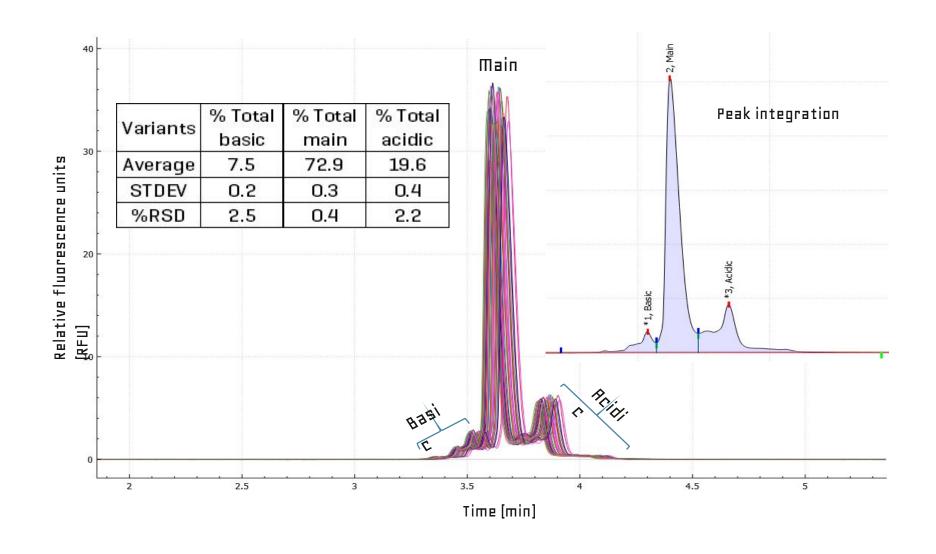
CZE analysis of Trastuzumab on the BioPhase 8800 system and the PR 800 Plus system.



	S/N ratio [RMS] for the main peak					
	Trastuzumab					
Detection	NFD on BP	UV on BP	UV on PA			
# of injections	42	42	6			
Average	10658	8856	5533			



CZE analysis of Trastuzumab on the BioPhase 8800 system with NF: repeatability





CZE analysis of Trastuzumab on the BioPhase 8800 system with NF: repeatability

	% Total basic variants			% Total main peak			% Total acidic variants		
Detection	NFD on BP	UV on BP	UV on PR 800	NFD on BP	UV on BP	UV on PR 800	NFD on BP	UV an BP	UV on PA800
* of injections	42	42	6	42	42	6	42	42	6
Average	7.5	8.3	9.2	72.9	71.3	71.2	19.6	20.4	19.6
STDEV	0.2	0.3	٦. ٦	0.3	0.6	0.5	0.3	0.5	0.4
%RSD	2.5	3.7	1.6	0.4	0.8	0.7	1.7	2.3	2.2



Summary- NF detection for all protein workflows on BioPhase 8800 system

Native: signal relies on the naturally existing aromatic amino acid in the protein, eliminating the protein labeling process

Fluorescence: no direct transmission/absorbance interference from background media in the capillary, providing a clean background

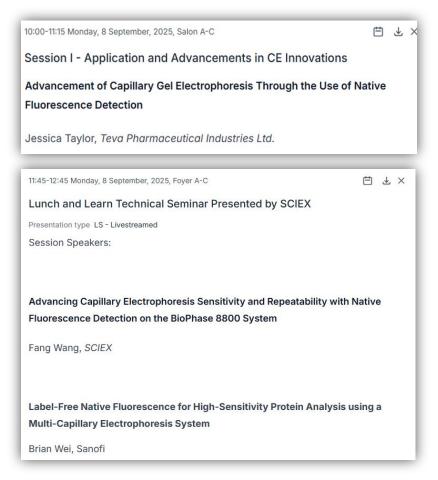
BioPhase 8800 system: 8-capillary parallel processing and easy-to-integrate data, reducing the time to answer with high confidence

BioPhase 8800 system with NF: demonstrated to be suitable for all existing protein-related workflows, providing comparable results, faster and easier data processing.



Place holder for all the Sciex-related materials in CE Pharm.

Monday, Sep eth



Tuesday and Wednesday, Sep 9-10th



Sciex Booth #8





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