



Essential tools
for the characterization
of proteins, polymers
and nanoparticles

It's Personal

For over forty years we have brought family commitment to the analytical industry business, and that's not going away.

From the beginning, Wyatt Technology has been dedicated to our customers by creating strong, personal connections, a philosophy instilled by our founder, Dr. Philip Wyatt and his sons, Geoffrey and Clifford. We take the work of our customers personally, and when they succeed, we couldn't be prouder.

In May 2023, Wyatt Technology joined Waters Corporation, another organization that sits at the intersection where curiosity meets inspiration. We remain committed to our mission of delighting our customers by providing outstanding analytical tools and unparalleled levels of personal service. We personalize customer interactions through direct contact with the entire technical staff.

We invite you to join us and experience our customer-first philosophy, to unlock together the potential of science by solving problems that matter.



Clifford D. Wyatt, former President, Wyatt Technology
Dr. Philip J. Wyatt, Founder and former Chairman of the Board, Wyatt Technology
Dr. Udit Batra, President & CEO, Waters Corporation
Geoffrey K. Wyatt, former CEO, Wyatt Technology

Light Scattering University

Wyatt Technology's founders established one of the crown jewels of Wyatt Technology – a course we call Light Scattering University (LSU).
LSU is the starting point of successful, life-long relationships with our customers.

Wyatt Technology

It's Personal	2
Wyatt's Rich History	4
What Can I Measure and Analyze?	6

SEC-MALS Products for HPLC & UHPLC

Characterize molar mass, size and conformation	8
------------------------------------------------------	---

RT-MALS Products for Process Analytics

Monitor molar mass, size and particle concentration	18
-----------------------------------------------------------	----

Dynamic & Electrophoretic Light Scattering Products

Characterize size, zeta potential and stability	22
-------------------------------------------------------	----

Field-Flow Fractionation

Advanced separation technology	32
--------------------------------------	----

CG-MALS Products

Analyze biomolecular interactions	38
-----------------------------------------	----

Training, Service & Support

Service and Support	42
Light Scattering University	43
World Wide Support	44

Wyatt Technology's Rich History

2020

- 2024 Integration of Waters LCs with SEC-MALS and FFF-MALS with launch of HPLC CONNECT™ 4 and VISION™ 4 softwares
- 2023 DynaPro™ ZetaStar™ is the first DLS/SLS/ELS instrument offering both walk-up and automated measurement modes
- 2022 DynaPro NanoStar™ II instrument introduces walk-up measurements with DYNAMICS Touch™ software interface
- 2020 Next-generation Eclipse™ FFF system adds dilution control, EAF4 and advanced ease of use
- 2019 ultraDAWN™ instrument for real-time process monitoring and control is introduced
- 2019 DAWN™, miniDAWN™, microDAWN™, Optilab™ and ViscoStar™ instruments are introduced
- 2018 Integration of field-flow fractionation technology with acquisition of Superon GmbH

2010

- 2017 DynaPro Plate Reader III instrument, with true molar mass capability, launched
- 2017 miniDAWN TREOS™ II instrument, with field-serviceability and upgradeability to μ DAWN™ instrument, introduced
- 2017 DAWN HELEOS™ II instrument wins Scientist's Choice Award® from SelectScience for Instrument of the Year
- 2017 Wyatt Technology™ expands headquarters by 50% to 45,000+ square feet
- 2016 Completely re-engineered ViscoStar III detector revealed
- 2014 First MALS detector for UHPLC, the μ DAWN, featured

2000

- 2011 Tibbetts Award for exemplifying notable lifetime achievements in innovation
- 2010 Mobius™ zeta potential instrument, first with flow-through and pressurization capabilities
- 2008 DynaPro NanoStar DLS detector introduced with front panel graphic display
- 2008 Scientist Magazine Award: Best Places to Work in Industry—also awarded in 2009, 2010 and 2012
- 2007 miniDAWN TREOS instrument introduced with front panel graphic display
- 2007 Calypso™ composition-gradient system introduced for reversible and irreversible interactions
- 2005 R&D 100 Award for Optilab rEX™ RI detector
- 2005 DAWN HELEOS (18-angle) instrument introduced with front panel display, integrated QELS™ and COMET™ modules
- 2005 First DynaPro Plate Reader instrument for automated DLS measurements introduced
- 2004 Optilab rEX (Extended Range) diode array dRI detector arrives
- 2004 Wyatt Technology China office formed

1990

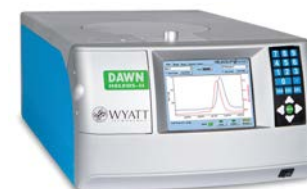
- 2004 ViscoStar differential viscometer enters the market
- 2004 ASTRA GPC-MALS software with 21 CFR Part 11 compliance released
- 1999 DAWN EOS (18-angle Enhanced Optical System) instrument introduced with solid state laser
- 1995 Optilab DSP (Digital Signal Processing) RI detector comes to market
- 1994 Major sensitivity improvements arrive with the DAWN DSP (Digital Signal Processing)
- 1993 Wyatt Technology Europe formed in Germany
- 1992 miniDAWN (3-angle) MALS detector introduced with solid state laser
- 1989 ASTRA™ 1.0 GPC-MALS software released

1980

- 1988 Optilab differential refractive index detector line acquired from Perstorp Analytical, Sweden
- 1986 First high-temperature (150 °C) DAWN F instrument placed
- 1985 DAWN B (Batch-mode) instrument introduced
- 1984 AMOCO Production Company orders 1st DAWN 16-angle GPC-MALS detector
- 1983 SC Johnson & Son orders 1st DAWN F MALS instrument with 7-angle flow-through detector
- 1982 Wyatt Technology formed with \$50,000 contract to detect toxicants in drinking water



DAWN NEON
Multi-touch screen,
health indicators and
full field serviceability,
2019



DAWN HELEOS
Front panel graphic
interface, integrated
QELS and COMET,
2005



DAWN EOS
18-angle Enhanced
Optical System with
solid-state laser, 1999



DAWN B
Batch mode, 1985



DAWN F
First ever DAWN
instrument created for
SC Johnson & Son, 1983



In 1970, Wyatt Technology's founder, Philip Wyatt, and some of his colleagues, formed a company that developed the world's very first multi-angle light scattering instruments using a laser as the light source. Since those days, Dr. Wyatt spearheaded the definition and redefinition of state-of-the-art analytical instrumentation at Wyatt Technology.

Science and discovery are deeply-rooted passions we share with our customers. We take pride that Wyatt instrumentation has been cited more than 40,000 times in patents and peer-reviewed publications.

What can I measure?



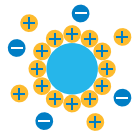
Molar Mass

Absolute molecular weight
from 200 to 1,000,000,000 g/mol



Size

RMS radius from 10 to 500 nm
and hydrodynamic radius
from 0.2 to 1,000 nm



Charge

Zeta potential and net molecular
charge for particles from 2 nm
to 100 μm



Turbidity

Turbidity/opalescence from
1 to 100 NTU



Particle
Concentration

Physical titer and quantitative
nanoparticle size distributions



Conformation

Shape, structure and
branching parameters



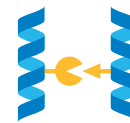
Payload

Content of gene vectors
or nanoparticles



Conjugation

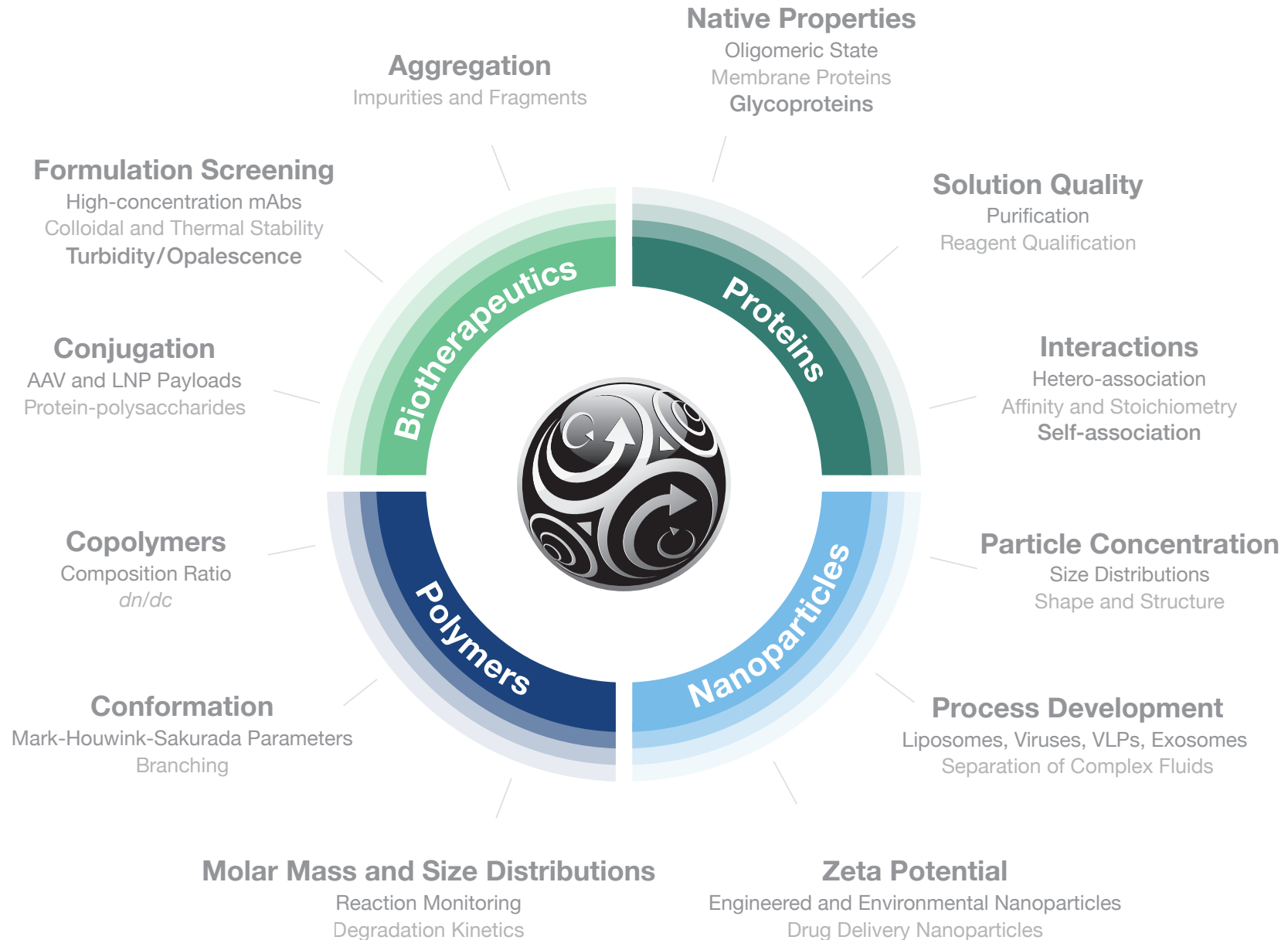
Molecular weight and fraction
of each constituent in a conjugated
biomolecule



Interactions

Binding affinity from pM to mM
and absolute stoichiometry
of complex interactions

What can I analyze?

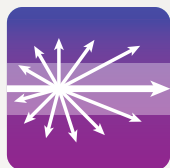




SEC-MALS Products

For HPLC & UHPLC

Characterize molar mass, size and conformation



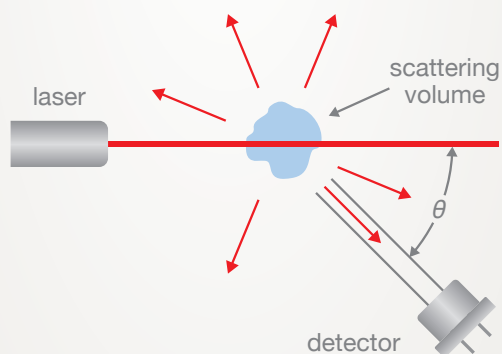
MALS

multi-angle light scattering

Based on first principles, MALS determines the molar mass and size of macromolecules and nanoparticles in solution.

Characterize:

- Peptides and proteins
- Conjugated proteins
- Polymers and copolymers
- Nanoparticles
- Viral vectors and VLPs
- Liposomes, LNPs and exosomes



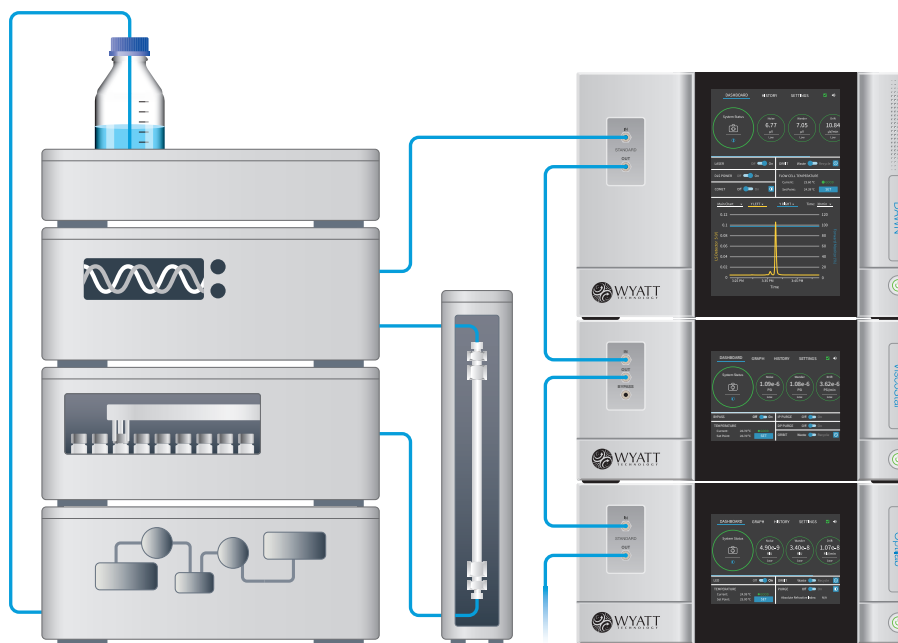
Multi-angle light scattering determines molar mass from the scattered intensity and the molecular radius from the angular scattering pattern.



DAWN

Premier family of MALS detectors

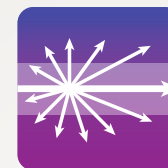
Choose between the DAWN instrument for the highest sensitivity and widest measurement range or the miniDAWN instrument for fundamental analysis of proteins and small polymers. Also available is the microDAWN instrument, uniquely suited for UHPLC systems.



Wyatt's MALS detectors interface to most industry-standard HPLC, GPC and FPLC systems.

	DAWN	miniDAWN	microDAWN
Description	The premier SEC-MALS detector for absolute molar mass and size, offering the highest sensitivity	The best in fundamental multi-angle light scattering	The only MALS detector uniquely designed for UHPLC with superb sensitivity
Applications	Peptides, proteins and polymers; plus viruses, vesicles and nanoparticles up to 500 nm in radius	Peptides, proteins small polymers, small viruses, VLPs and nanoparticles	Peptides, proteins and small polymers compatible with UHPLC
Molar Mass Range	200 Da to 1 GDa	200 Da to 10 MDa (proteins) or 1 MDa (polymers)	200 Da to 10 MDa (proteins) or 1 MDa (polymers)
Molecular Size Range (MALS — R_g)	10 to 500 nm	10 to 50 nm	10 to 50 nm
Molecular Size Range (DLS — R_h)*	Flow: 0.5 to 300 nm Batch: 0.5 nm to 1 μ m	Flow: 0.5 to 50 nm Batch: 0.5 nm to 1 μ m	Flow: 0.5 to 30 nm 0.5 nm to 1 μ m
Compatibility	HPLC	HPLC	UHPLC/APC
Flow Cell	Standard and high-temperature flow cells, COMET cell cleaning module included	Standard flow cell, COMET cell cleaning module included	Micro flow cell, COMET cell cleaning module included
Detectors	18 angles	3 angles	3 angles
MALS Sensitivity: BSA in Aqueous Buffer	0.2 μ g typical, 30 cm SEC column	0.5 μ g typical, 30 cm SEC column	70 ng typical, 15 cm UHPLC-SEC column
MALS Sensitivity: 100 kDa Polystyrene in THF	10 ng typical, 30 cm GPC column	25 ng typical, 30 cm GPC column	3.5 ng typical, 15 cm UHPLC-SEC column
Temperature Control	Ambient; Heated/cooled from -15 °C to 150 °C; Ultra-high: 20 °C to 210 °C	Ambient only	Ambient only
Options	Temperature control, Fluorescent polymer configuration, WyattQELS embedded DLS	WyattQELS embedded DLS	WyattQELS embedded DLS

* Size range will depend on flow rate, application and instrument configuration.



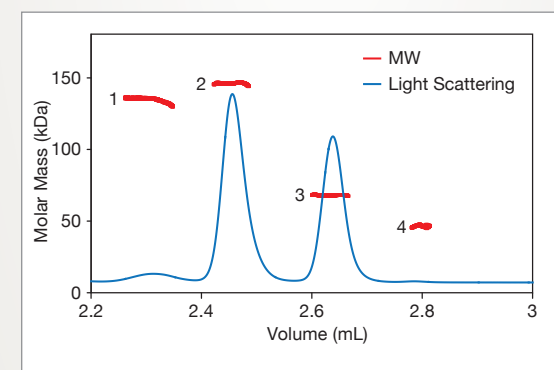
SEC-MALS

size exclusion chromatography
combined with multi-angle
light scattering

SEC-MALS is an absolute method that does not rely on column calibration for analyzing:

- Molar mass
- Size distributions
- Oligomeric state
- Conformation
- Conjugation ratio
- Polymer branching

SEC-MALS combines MALS, intrinsic viscosity (IV) and differential refractive index (dRI) instruments with SEC separation.



Even though Peak 1 elutes earliest, MALS shows that it does not have the largest molar mass for this example of protein aggregates and fragments.



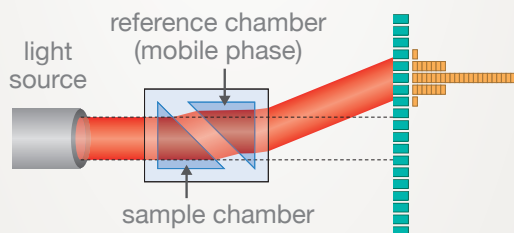
dRI

differential refractive index

dRI is a universal concentration measurement technique that does not depend on chromophores or fluorophores.

Optilab online dRI instruments are used in:

- MALS analysis of molar mass
- Intrinsic viscosity determination for polymer conformation and branching
- Triple-detection characterization of copolymers and protein conjugates
- Basic quantitation of chromatographic peaks
- Measurement of dn/dc in different mobile phases
- Determination of solvent absolute refractive index



Optilab's 512-detector array means it can reliably quantify a tiny peak at the nanogram level superimposed on a milligram-level peak!



Optilab

Extended dRI measurement range

The only RI detector designed to operate at the same wavelength as the MALS detector for dn/dc measurements, Optilab is available in a variety of configurations depending on your application. It can also measure the absolute refractive index (aRI) of the solvent.

	Optilab	Optilab HC	microOptilab
Description	dRI detector for standard HPLC, offering the highest sensitivity and dynamic range	dRI detector for CG-MALS, protein purification and other high-concentration analyses	dRI detector for UHPLC, offering the highest sensitivity and dynamic range
Application	Quantify a few ng/mL up to 25 mg/mL	Measure proteins up to 180 mg/mL	UHPLC/APC
dRI Range	-4.7×10^{-3} RIU to $+4.7 \times 10^{-3}$ RIU (refractive index unit)	-2.6×10^{-3} RIU to $+3.4 \times 10^{-2}$ RIU	-4.7×10^{-3} RIU to $+4.7 \times 10^{-3}$ RIU
Dynamic Range	12,000,000:1	23,000,000:1	6,000,000:1
dRI Sensitivity	0.75×10^{-9} RIU	1.5×10^{-9} RIU	1.5×10^{-9} RIU
aRI Range	1.2 to 1.8	1.2 to 1.8	1.2 to 1.8
aRI Sensitivity	± 0.002	± 0.002	± 0.002
Temperature Control	4 °C to 65 °C	4 °C to 65 °C	4 °C to 65 °C

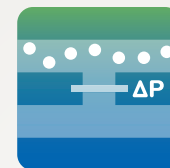


ViscoStar

Unsurpassed differential viscometer

Incorporating patented thermal bridge balancing, as well as proprietary technology to suppress pressure pulse noise and temperature gradients, the ViscoStar detector offers the best performance in differential viscosity measurements.

	ViscoStar	microViscoStar
Description	The ultimate differential viscometer for GPC	Differential viscometer for UHPLC/APC
Applications	Polymers below ~ 1 MDa for conformational analysis; all polymers for Mark-Houwink-Sakurada parameters	Polymers suitable for UHPLC-SEC separation
Sensitivity	0.1 µg of 100 kDa polystyrene in THF	5 ng of 100 kDa polystyrene in THF
Dynamic Range	135,000:1	135,000:1
Drift	2.5 Pa/hr	1.25 Pa/hr
Temperature Control	4 °C to 70 °C	4 °C to 70 °C
Capillary Bridge Tuning	Automated thermal tuning	Automated thermal tuning
Pump Pulse Suppression	Full impedance matching of the capillary bridge and proprietary software algorithms	Full impedance matching of the capillary bridge and proprietary software algorithms
Delay Column Options	16.2, 10.8, or 5.4 mL; 32.4 mL optional	5.4 mL



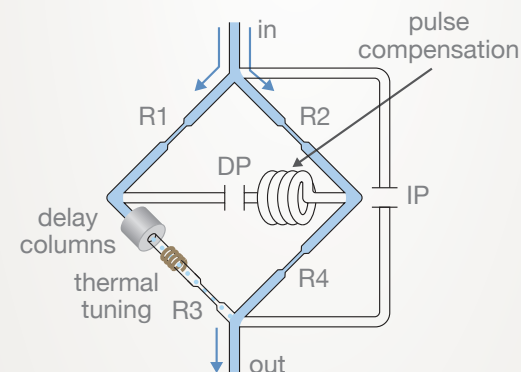
IV

intrinsic viscosity

Differential viscometers are used in conjunction with SEC to measure the specific and intrinsic viscosities of polymer solutions.

Combined with a MALS instrument, SEC-MALS-IV determines:

- Intrinsic viscosity
- Conformation
- Branching analysis
- Hydrodynamic radius
- Mark-Houwink-Sakurada parameters

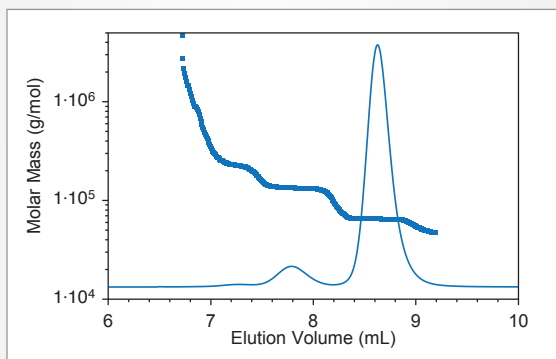


Without delay columns, the impedance of the capillary bridge would be fully balanced. The pulse compensation element matches the additional impedance of the delay columns, eliminating the effect of pump pulses on the DP transducer.



ASTRA

advanced software
for macromolecular and
nanoparticle characterization



Absolute molar mass analysis

ASTRA software's Band Broadening Correction accounts for interdetector dispersion to match signals from each detector in the chromatographic elution series.

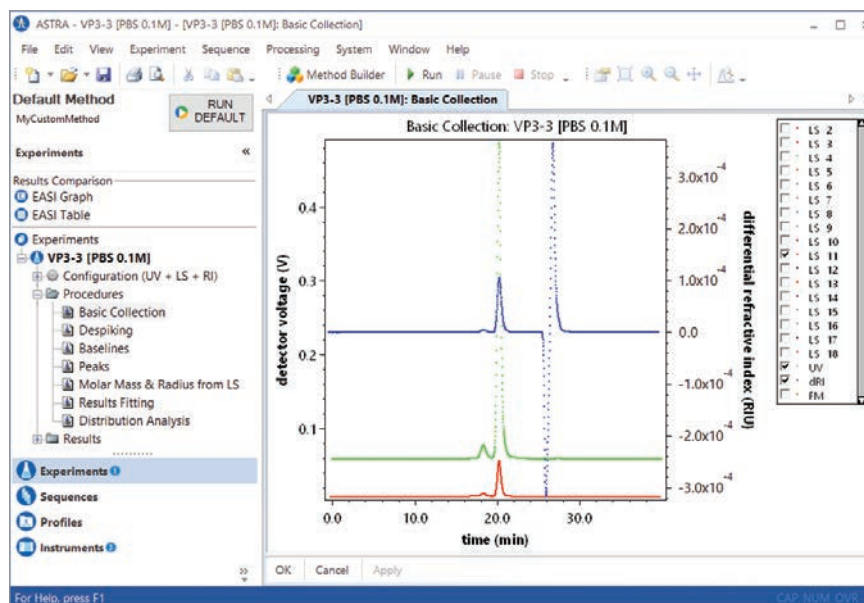
This algorithm is responsible for proving uniform molecular weights across the BSA monomer, dimer and trimer peaks.



ASTRA

The premier software for analyzing macromolecules and nanoparticles by multi-angle light scattering

ASTRA software integrates MALS, UV, refractive index, dynamic light scattering and intrinsic viscosity data for comprehensive characterization of the physical properties of materials in solution/suspension.



ASTRA software provides absolute determination of:

- Molar mass and size
- Conformation, shape and conjugation ratio
- Differential and cumulative distributions; moments of the distribution and polydispersity
- Intrinsic viscosity and Mark-Houwink-Sakurada parameters
- Nanoparticle concentration, total viral titer; viral and drug nanocarrier payload

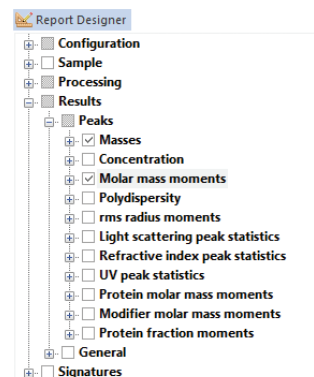
Compile key results:

ASTRA software gives you a quick and easy overview of your most important results in one compact table.

EASI Table			
Experiments:	All	Peaks:	All
Abscissa:	min		
	Peak 1		
	Mn (kDa)	Mw (kDa)	Polydispersity (Mw/Mn)
sample01	66.8 (±0.2%)	66.8 (±0.2%)	1.00 (±0.23%)
sample02	65.9 (±0.0%)	65.9 (±0.0%)	1.00 (±0.05%)
sample03	65.7 (±0.4%)	65.7 (±0.4%)	1.00 (±0.54%)
sample04	66.6 (±0.0%)	66.6 (±0.0%)	1.00 (±0.07%)
sample05	66.8 (±0.2%)	66.8 (±0.2%)	1.00 (±0.26%)
sample06	65.8 (±0.2%)	65.8 (±0.2%)	1.00 (±0.29%)
Average	66.3	66.3	1.00
Standard deviation	0.5	0.5	0.00
% Standard deviation	0.8	0.8	0.00
Minimum	65.7	65.7	1.00
Maximum	66.8	66.8	1.00

Customized reports:

ASTRA software provides customized reporting options so you can export the exact information you need. It even allows you to customize the report with your company's logo and descriptive text.



Regulatory Compliance

Following industry standards, ASTRA software offers an optional 21 CFR Part 11 compliance package, including IQ/OQ documents and procedures.

ASTRA software's Security Pack includes:

- Administrator, researcher, technician and guest access levels
- Full audit trails
- Electronic signatures
- Sign-in/sign-out during a run
- Secure SQL server database
- Local or remote database connectivity
- Data integrity validation
- Full IQ/OQ procedures and documentation validation

Molar mass in a single click

1. Select experiment type
2. Input parameters
3. Click 'Run'

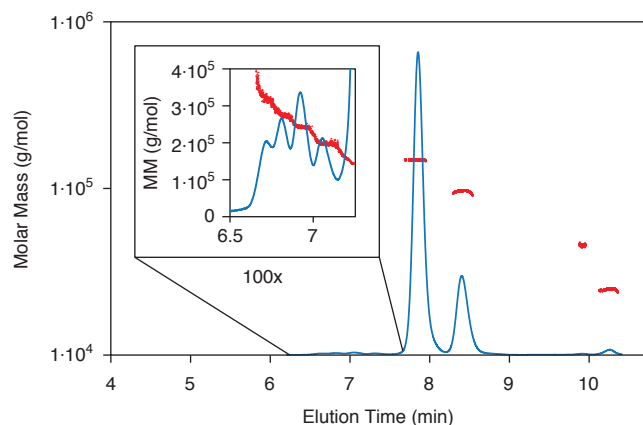


HPLC CONNECT Software

- ASTRA software performs all SEC-MALS control, data acquisition and analysis
- Full digital synchronization of the HPLC pump, autosampler and detectors
- Direct digital acquisition of multiple wavelengths from the HPLC UV/Vis

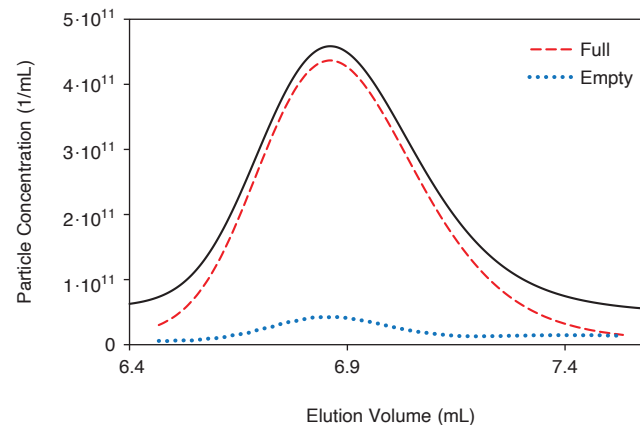
SEC-MALS Applications

Aggregates and Fragments



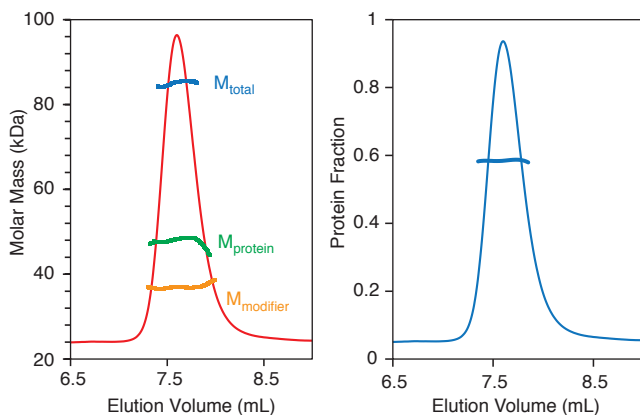
The power of UHPLC for separating aggregates and fragments combines with MALS to unequivocally identify small quantities of impurities in an IgG sample. Each of the aggregate peaks shown in the 100x inset represent a fraction of one percent of the monomer total mass yet is well-quantified by MALS.

Viral Vector Particle Concentrations



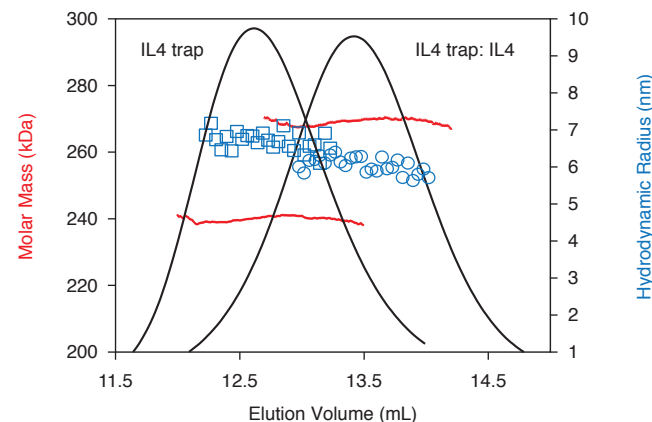
The *Viral Vector Analysis* method determines multiple critical quality attributes. This graph shows an overlay of the size-exclusion chromatogram of an adeno-associated virus (black solid line) with particle concentrations determined at each data slice for sub-populations of full capsids (red, long dash) and empty capsids (blue, dotted).

Protein Conjugate and Copolymer Analysis



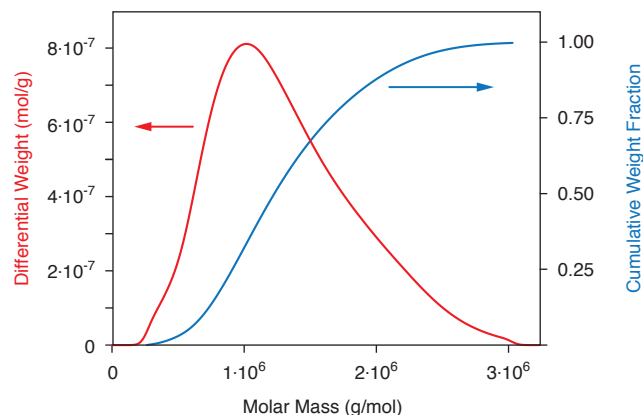
ASTRA software's Protein Conjugate algorithm makes use of data from MALS, UV and RI detectors to characterize conjugated proteins and copolymers. This analysis determines the molecular weights of the protein, modifier and complete conjugate as well as average extinction coefficient and dn/dc .

Protein Complexes and Conformations



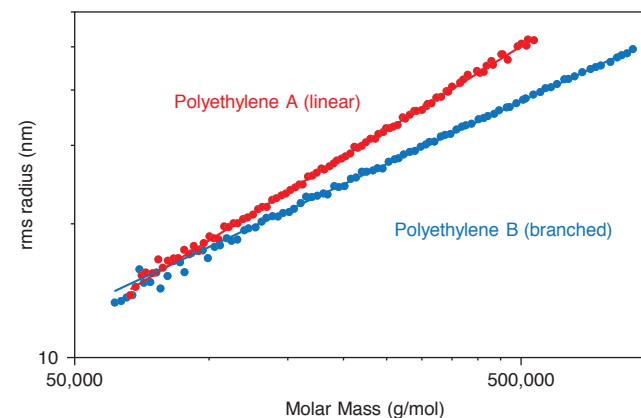
Pure interleukin 4 trap (IL4-trap) elutes earlier than the IL4 : IL4-trap complex, despite its lower molecular weight. MALS MW analysis (small red symbols) indicates the expected MW values. Online DLS R_h data (open blue symbols) show the reason for the late elution: IL4 stabilizes the trap to form a compact IL4 : IL4-trap complex.

Molar Mass and Size Distributions



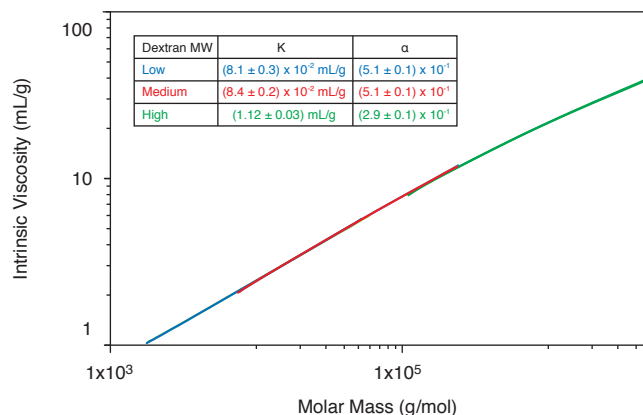
In addition to plotting the molar mass and size determined by multi-angle light scattering over a chromatogram or fractogram, ASTRA software can convert the data into distributions. These graphs show differential and cumulative distributions of molar mass as measured for hyaluronic acid.

Polymer Branching



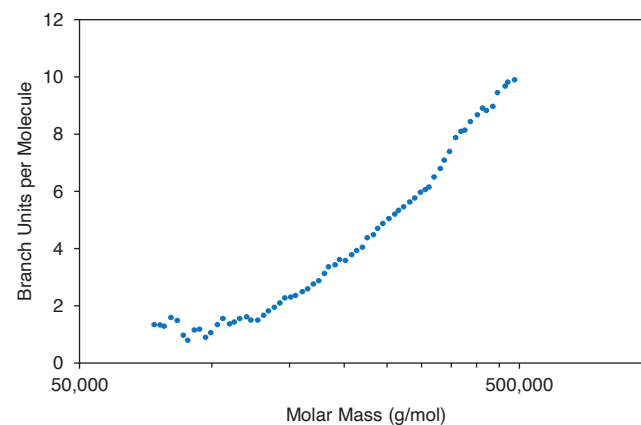
A MALS instrument measures rms radius vs. molar mass to reveal a polymer's branching properties. Here, the branching of Polyethylene B is apparent by its significantly lower slope in relation to Polyethylene A, which is known to be linear.

Conformational Change with MW

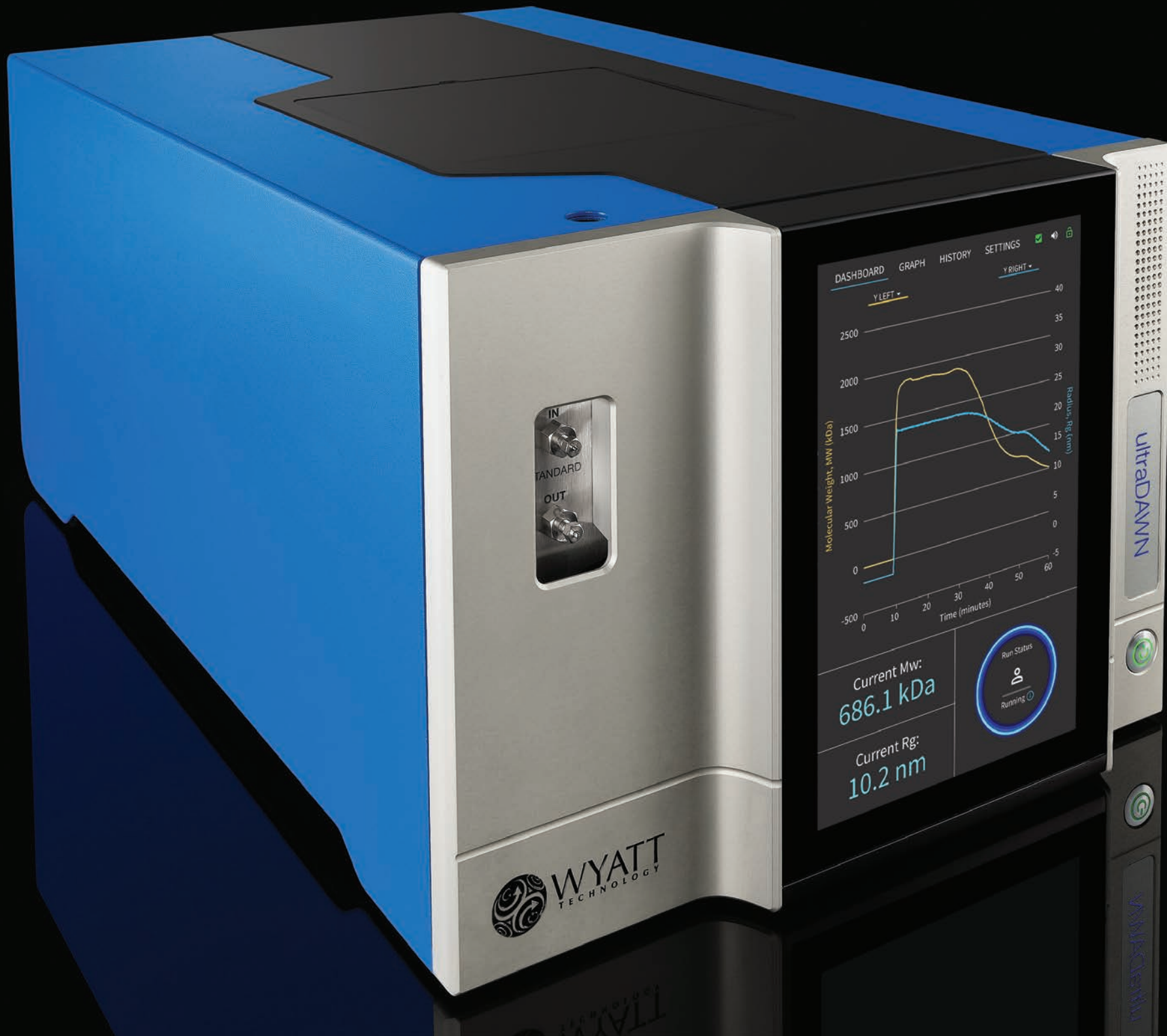


A Mark-Houwink-Sakurada (MHS) plot shows intrinsic viscosity as a function of molar mass—revealing the polymer conformation. The MHS plots of low, medium and high MW dextrans, shown here, indicate conformational change with increasing molar mass of the molecules.

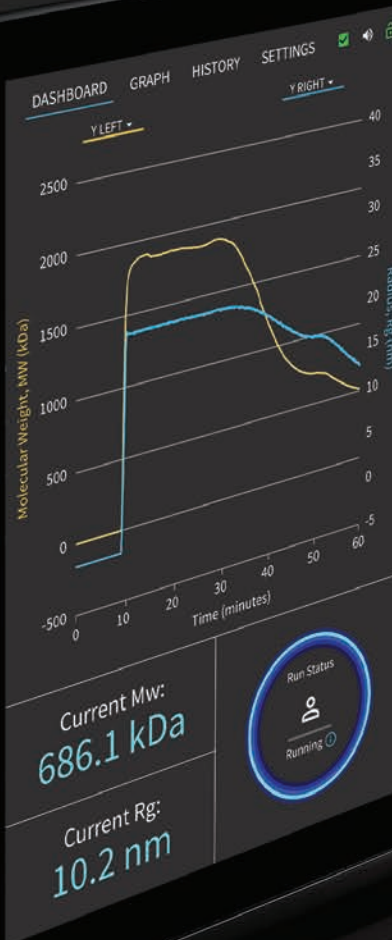
Branching Calculations



ASTRA software compares linear and branched polymers in order to determine branching ratio. The data in the top chart (Polymer Branching) were analyzed to yield the average number of branching units per molecule and its dependence on molar mass. Branching begins above a molar mass of ~100,000 g/mol.



IN
STANDARD
OUT



WYATT
TECHNOLOGY

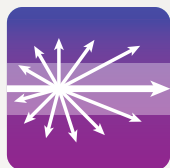
ultraDAMN

WYATT
TECHNOLOGY

RT-MALS Products

For Process Analytics

Monitor biophysical product attributes in real time



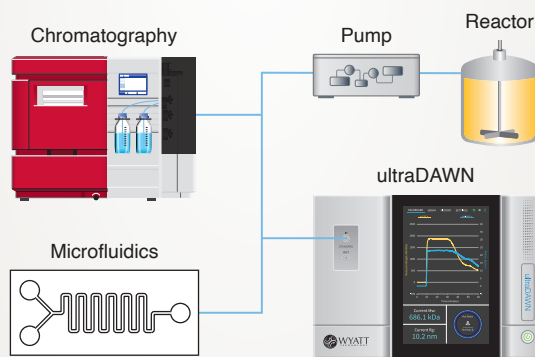
RT-MALS

real-time
multi-angle light scattering

RT-MALS monitors molar mass, size and particle concentration for quality assurance and control of production processes.

Optimize and control:

- Viral vector downstream processing
- Nano-pharmaceutical formulation
- Protein or nucleic acid purification
- Non-viral vector production
- De/polymerization and conjugation



The ultraDAWN instrument supports inline measurements with flow rates up to 150 mL/min. Online operation utilizes a pump to divert samples from a vessel or high-flow-rate process to the ultraDAWN instrument.



ultraDAWN

*Measure the product,
not the process*

ultraDAWN is a breakthrough instrument in process analytical technology for nanomedicines, gene vectors, vaccines and biotherapeutics. It measures key product attributes in real time for process development, scale-up and production.

Measurement capabilities

- **Macromolecules:** Weight-average molar mass from 10^3 to 10^9 g/mol and rms radius from 10 to 250 nm
- **Nanoparticles:** z-average radius from 10 to 250 nm, and corresponding particle concentration
- **Viral Vectors:** Vg/Cp (full:total ratio); full, empty and total viral concentrations; capsid, genome and total molar masses. For viruses and VLPs smaller than 20 nm in radius

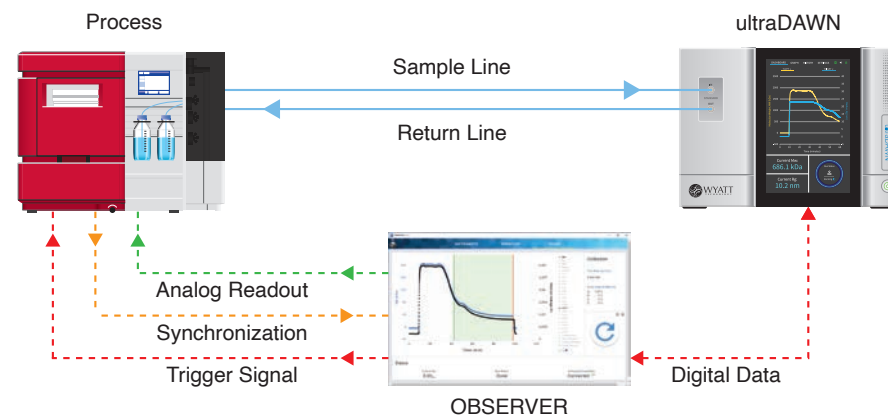


OBSERVER Software

*Built-in workflows for flexible
process integration*

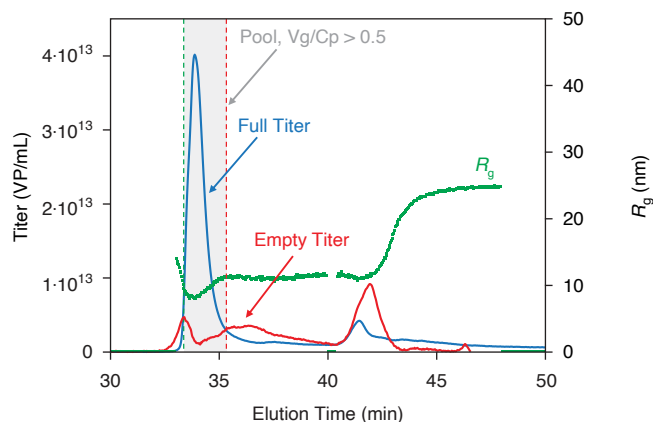
OBSERVER software monitors MALS data from the ultraDAWN instrument, calculating MW, radius and/or particle concentration up to 300 times per minute. PAT workflows let users specify product attribute criteria for trigger activation.

OBSERVER software communicates digitally via OPC-UA with process-control software such as SIPAT or DeltaV, or via analog signals with lab-scale equipment such as FPLC, TFF or microfluidic nanoparticle production.



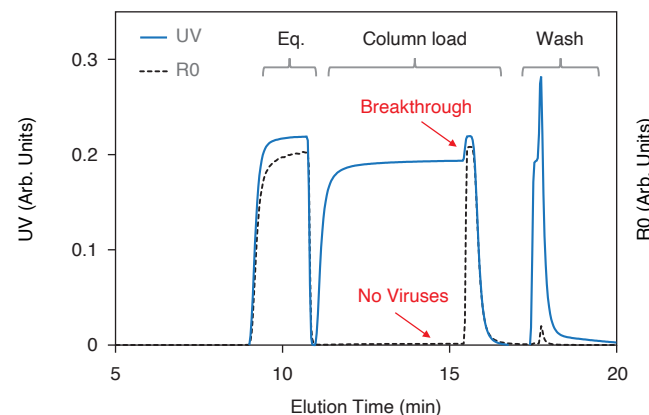
RT-MALS Applications

AAV Vg/Cp, capsid titer and aggregates



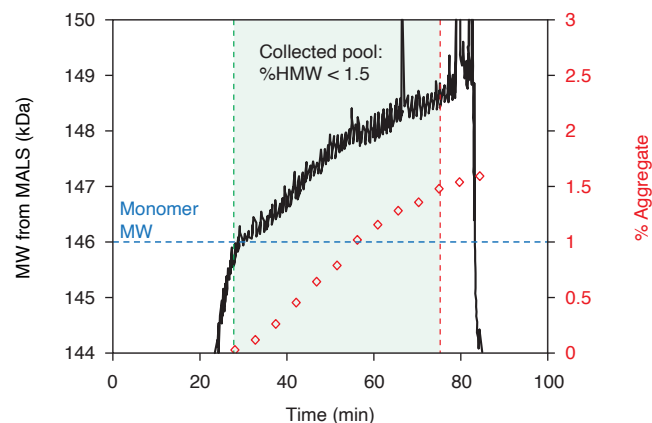
During AAV enrichment by IEX chromatography, RT-MALS determines the empty and full titers and more. A trigger was set to indicate pool collection for Vg/Cp > 0.5 (shaded region). Increasing particle radius (green) identifies aggregates, starting at the 42nd minute of elution time onward.

Optimize adenovirus purification



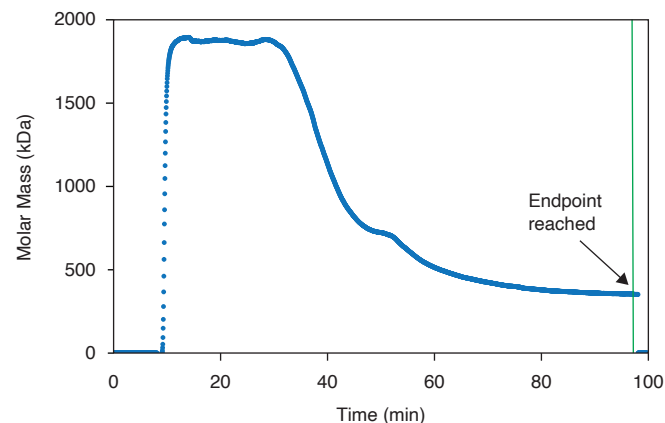
During sample application, MALS and UV are combined to differentiate small impurities flowing through the column from viral breakthrough toward the end. The signals confirm no viral loss during the wash step. Courtesy Janssen Vaccines & Prevention.

Protein purification – flow through hydrophobic interaction chromatography



RT-MALS improves mAb yield when purification is controlled by inline, real-time MW. As seen here for flow-through HIC, MW by RT-MALS correlates well to aggregate content determined by offline SEC on fractions. The shaded region is where RT-MALS set the trigger to ensure collection of low-aggregate product. Data from Patel et al., mAb 2018.

End-point determination of a polysaccharide depolymerization process



A critical depolymerization step must reduce the polysaccharide's initial molar mass (M_w) from over 1800 kDa down to less than 350 kDa. RT-MALS tracked reduction of M_w and triggered reaction shutdown once it fell below 350 kDa.



DLS & ELS Products

Measure in Cuvettes and Well Plates

Characterize size, zeta potential, turbidity and stability



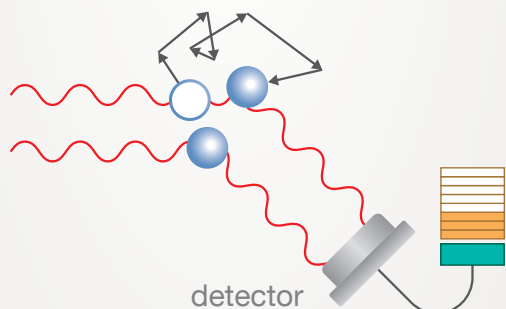
DLS

dynamic light scattering

DLS determines the diffusion coefficients, size and size distributions of particles in a fluid by measuring the light intensity fluctuations arising from their Brownian motion.

In addition to basic sizing applications for sub-micrometer macromolecules and nanoparticles, DLS measures:

- Quality
- Aggregation
- Stability
- Propensity for aggregation
- Particle concentration
- Turbidity



Brownian motion of sub-micrometer particles gives rise to intensity fluctuations in the scattered light. The rate of fluctuation is analyzed to determine the diffusion coefficient.



DynaPro ZetaStar and NanoStar

Versatility and simplicity for particle analysis

The DynaPro NanoStar and ZetaStar are intuitive walk-up instruments with advanced analytical capabilities. The NanoStar instrument combines dynamic and static light scattering (DLS/SLS) to analyze size distributions, particle concentrations, turbidity and molar mass. The ZetaStar instrument adds ELS, enabling zeta potential and surface charge measurements.

	ZetaStar	NanoStar
Description	DLS/SLS/ELS instrument for batch and online measurements	DLS/SLS instrument for batch and online measurements
Application	Size, particle concentration, molar mass, turbidity, and zeta potential. Optional MALS integration.	Size, particle concentration, molar mass, and turbidity. Optional MALS integration.
Hydrodynamic Radius Range (R_h)	0.2 nm to 400 nm (flow cell); 0.2 nm to 1000 nm (cuvette)	0.2 nm to 1000 nm
DLS Sensitivity	90° (DLS/SLS): 0.1 mg/mL lysozyme 163.5° (DLS): 1 mg/mL lysozyme	90° (DLS/SLS): 0.1 mg/mL lysozyme
Molar Mass Range	300 g/mol to 10 ⁷ g/mol*	300 g/mol to 10 ⁷ g/mol*
Particle Concentration	10 ⁸ to 10 ¹⁵ mL ⁻¹ (dependent on size)	10 ⁵ to 10 ¹⁵ mL ⁻¹ (dependent on size)
Turbidity/Opalescence	0 to 100 NTU	1 to 80 NTU
Mobility Sensitivity	1 mg/mL lysozyme	n/a
ELS Conductivity Range	0 to 7 mS/cm; 0 to 100 mS/cm [†]	n/a
Minimum Sample Volume	2 µL (DLS/SLS, quartz cuvette); 65 µL (ELS, quartz cuvette); 500 µL (flow cell)	2 µL (DLS/SLS, quartz cuvette)
Temperature Control	-10 °C to 120 °C	-10 °C to 120 °C
Automation	Autosampler compatible	n/a
Fluorescence Suppression	785 nm laser avoids fluorescence	n/a

*Upper limit depends on conformation: It is limited to a maximum R_h of 50 nm. [†]PEEK flow cell and either Atlas™ or Arc™ autosampler.



Automation

The ZetaStar instruments' flow cell enables automated sample handling with an Arc pump and autosampler. The integrated system is controlled directly by DYNAMICS software to run an entire sequence of dozens of samples, measuring size and zeta potential.

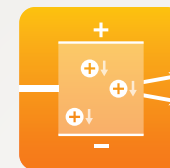
Zeta Potential in Formulation Buffer

Make measurements that are truly relevant by obtaining size and zeta potential in native or physiological buffer. With the aid of an Arc autosampler and pump, or the Atlas module, the ZetaStar instrument's pressurized flow cell suppress bubbles that otherwise interfere with zeta potential determination in high-salt buffers.



DLS for Separations

The ZetaStar and NanoStar instruments' DLS detector can be fiberoptically coupled to the DAWN MALS instrument line, positioned downstream from a liquid chromatography or Eclipse field-flow fractionation system, resulting in high-resolution size distributions that cannot be achieved with batch DLS. Alternating between online and offline measurements is simple in DYNAMICS or DYNAMICS Touch software.



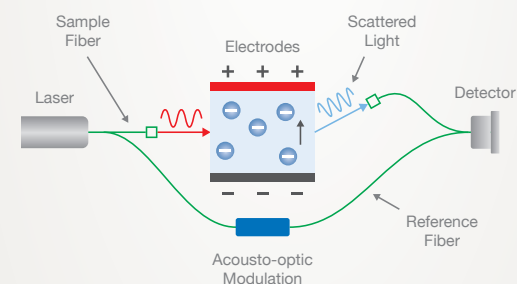
ELS

electrophoretic light scattering

ELS determines the electrophoretic mobility of particles in a fluid by measuring their velocity under an applied electric field. With additional measurement of R_h by DLS, zeta potential and the net charge are calculated.

ELS is used to study:

- Stability against flocculation of colloids and emulsions
- The isoelectric point of protein formulations in native formulation buffer or under physiological conditions
- Surface charges that impact gene and drug delivery



Wyatt's advanced FIDELIS (fiber-Interferometric Doppler electrophoretic light scattering) technology employs fiber-coupled elements, including high-frequency acousto-optic phase modulators, for a robust ELS optical system providing excellent sensitivity and speed.



HT-DLS

high-throughput
dynamic light scattering

The DynaPro Plate Reader high-throughput DLS/SLS instrument is compatible with industry-standard plates, allowing for seamless integration with automated liquid handling systems. This flexibility ensures adaptability to changing needs and configurations.



To increase throughput, the DYNAMICS Automation API enables integration with a robot for well plate transfer and liquid handling. The API interfaces to both hardware control and data analysis functions.

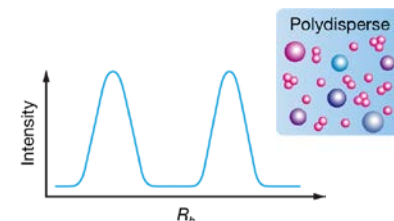
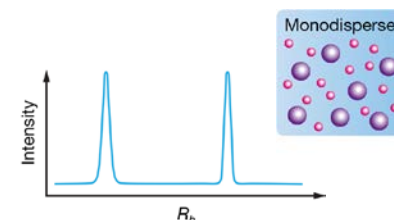


DynaPro Plate Reader

High-throughput size and stability analysis

The DynaPro Plate Reader instrument combines dynamic and static light scattering (DLS/SLS) in industry-standard microwell plates. Perform high-throughput experiments and acquire, in one day, data that would otherwise take weeks to collect.

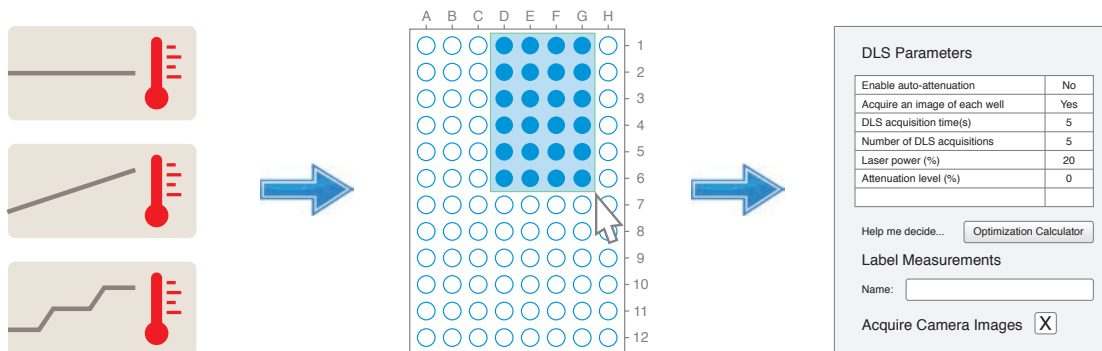
	Plate Reader
Description	Automated DLS/SLS measured directly in standard microwell plates
Application	High-throughput screening and other automated measurements of multiple samples
Plate Scan Time	As little as 1.5 hours for a 384 well plate
Hydrodynamic Radius Range (R_h)	0.5 nm to 1000 nm
Sensitivity	0.125 mg/mL lysozyme
Molar Mass Range*	1000 g/mol to 10^6 g/mol
Particle Concentration	10^8 to 10^{11} mL ⁻¹ (dependent on size)
Minimum Sample Volume	4 μ L (1536 well plate), 10 μ L (384 well plate), 60 μ L (96 well plate)
Temperature Control	4 °C to 85 °C



The DynaPro DLS instrument line determines size distribution without fractionation, providing polydispersity estimates as well as hydrodynamic radii.

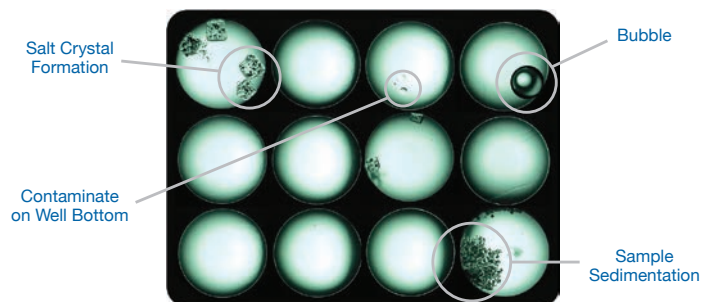
*Upper limit depends on conformation: It is limited to a maximum R_h of 12 nm.

Design High-throughput Experiments in 3 steps



Tiny Drop, Massive Data

Obtain up to 5 critical parameters with just 4 μL of sample. With a sample volume this small, valuable metrics like stability and aggregation can be obtained during earlier development stages.



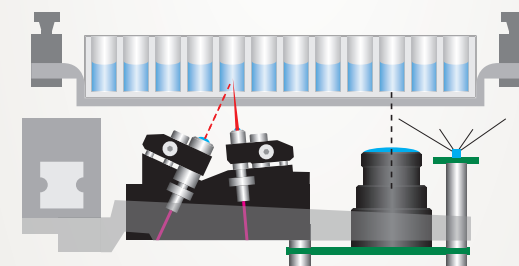
Advanced Imaging

An onboard camera views each well from below, aiding in diagnostics by revealing sample behavior beyond nanoparticle size, showing clean wells, bubbles, crystals, and precipitates for confident data interpretation.

Standard plates, exceptional data



The DynaPro Plate Reader is compatible with a variety of standard clear-bottom microwell plates in 96-, 384-, or 1536-well formats. High measurement sensitivity can be achieved with sample volumes as low as 4 μL per well.

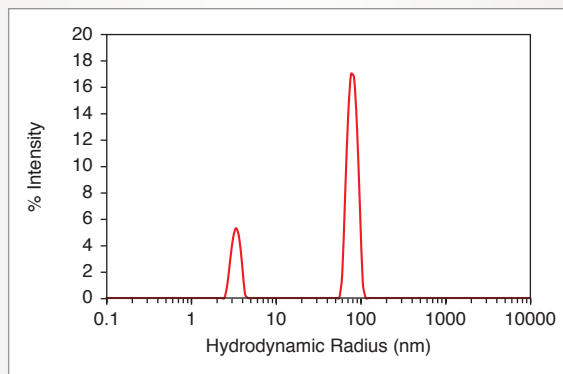


For both static and dynamic light scattering in a well plate, laser illumination and detection take place from below.



DYNAMICS

comprehensive software for
dynamic and electrophoretic
light scattering



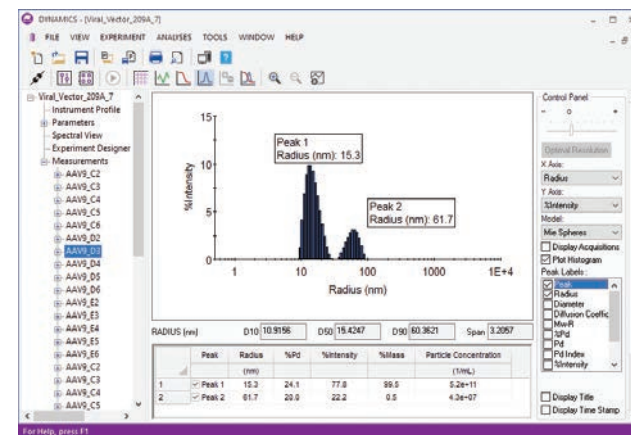
Size distributions from sub-nanometers
to micrometers

Dynamic light scattering determines size
distributions without any separation. This
regularization graph shows the presence of
an 80 nm nanoparticle in a protein solution.
The particle concentration of each peak
may be determined and displayed as well.

Essential Size and Zeta Potential

DYNAMICS software is a full-featured package
for collecting, analyzing, and reporting data
acquired with one of Wyatt's batch dynamic
or electrophoretic light scattering instruments:
DynaPro Plate Reader, NanoStar and ZetaStar.
It combines flexibility and intelligence with
rigorous calculations that can be as seamless
or fine-tuned as you choose.

Intuitive yet powerful, DYNAMICS software
gives you access to all the information needed
to ensure correct and thorough analysis
of dynamic light scattering (DLS), static light
scattering (SLS), and electrophoretic light
scattering (ELS) data.



Regulatory Compliance

Following industry standards, DYNAMICS
software offers an optional 21 CFR Part 11
compliance package, including IQ/OQ
documents and procedures.

DYNAMICS software's Security Pack includes:

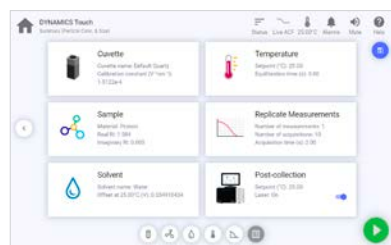
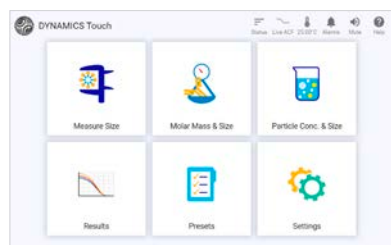
- Administrator, researcher, technician,
and guest access levels
- Full audit trails
- Electronic signatures
- Sign-in/sign-out during a run
- Secure SQL server database
- Local or remote database connectivity
- Data integrity validation
- Full IQ/OQ procedures and
documentation validation

Data Quality with Every Measurement

DYNAMICS and DYNAMICS Touch software packages include a data quality report for every measurement. The automated data quality assessment allows even first-time users to understand their results and troubleshoot any issues that may have led to poor measurements.



Results with Just a Tap



DYNAMICS Touch software's graphically rich interface enables almost anyone to immediately perform measurements and see the results. It guides you every step of the way.



Effortless Result Documentation

For easy viewing and reporting, DYNAMICS and DYNAMICS Touch software offer compact, preset reports containing only the essential results. In addition, you can fully customize extended reports to document exactly the information you need.



DYNAMICS Touch

particle size, concentration and zeta potential measurements

DYNAMICS Touch software, installed on the DynaPro NanoStar and ZetaStar instruments, intuitively guides users to top-quality light scattering measurements with virtually no training.

Working from a touchscreen display, you can:

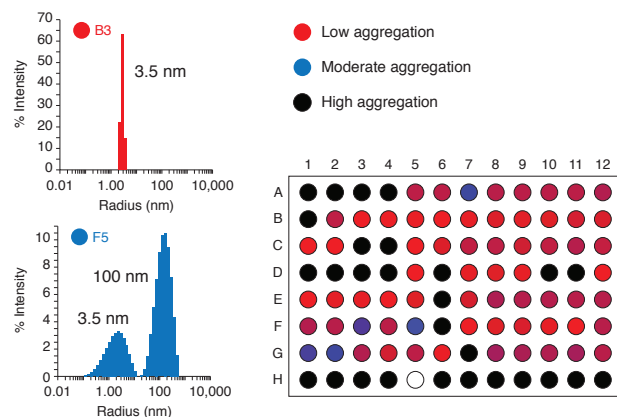
- Set up methods
- Run samples
- Assess data quality
- Analyze data
- Produce reports



Data files may be sent to a network location or downloaded onto a USB device for storage and extended analysis in Wyatt's DYNAMICS desktop application.

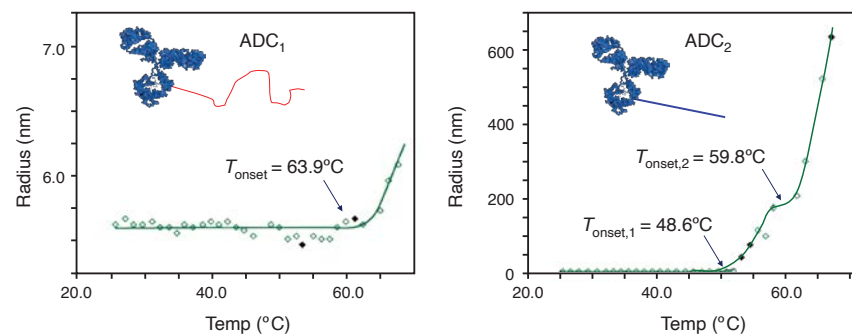
DLS | SLS | ELS Applications – Biologics

Aggregation in a 96 Well Plate



The SpectralView feature in DYNAMICS supports color-coded visualization of the results of a plate scan, which might include hundreds of samples. Here the visualization represents the degree of aggregation for a rapid, intuitive assessment of the optimal formulation.

Conformational Stability



Conjugating the same monoclonal antibody and drug via different linkers can have significant impact on stability. Here, ADC₂ exhibits two thermal transitions, one at 60 °C, similar to ADC₁, while the other is near 50 °C. DLS highlights the degree of thermally-induced aggregation, negligible in ADC₁ yet rapid and extensive in ADC₂.

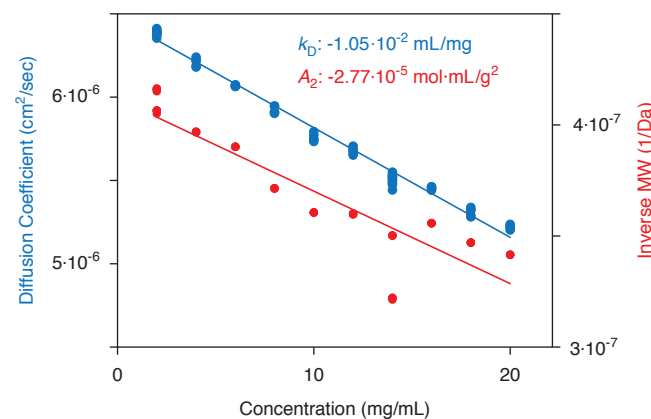
Biosimilarity

Sample	Radius (nm)	%PD	Mobility ($\mu\text{m cm/s V}$)	Zeta Potential (mV)
IgG 1	5.5 ± 0.1	4 ± 1	1.26 ± 0.17	22 ± 3
IgG 2	5.5 ± 0.1	5 ± 1	1.13 ± 0.27	20 ± 5

Average and standard deviation from 10 measurements

For biosimilars, matching the reference product is desirable. The DynaPro ZetaStar instrument's simultaneous DLS/ELS measurement provides orthogonal data in less than 1 minute. Size, polydispersity (PD), electrophoretic mobility and zeta potential, shown in the table, can be used to help establish biosimilarity.

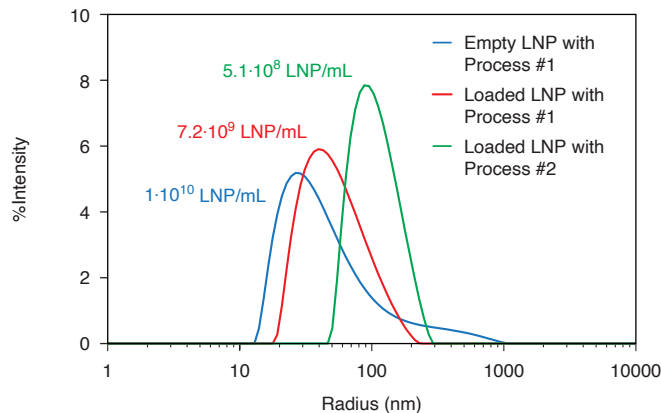
Aggregation Propensity



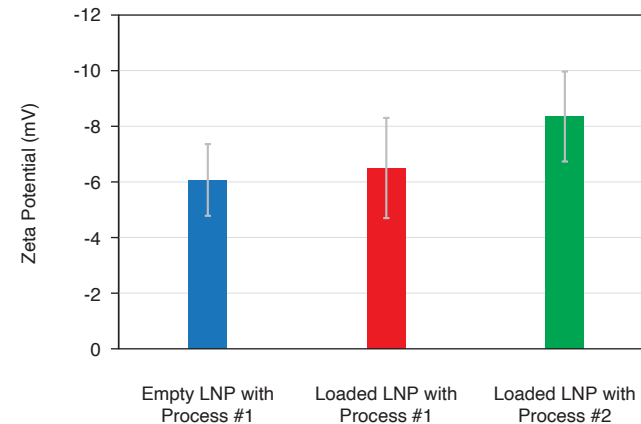
Non-specific protein-protein interactions, important for selecting and optimizing biotherapeutic candidates and formulations such as IgG, are characterized by means of a concentration series. Both static light scattering (A_2) and dynamic light scattering (k_D) may be used.

DLS | SLS | ELS Applications – Nanoparticles

Lipid Nanoparticle Formulations

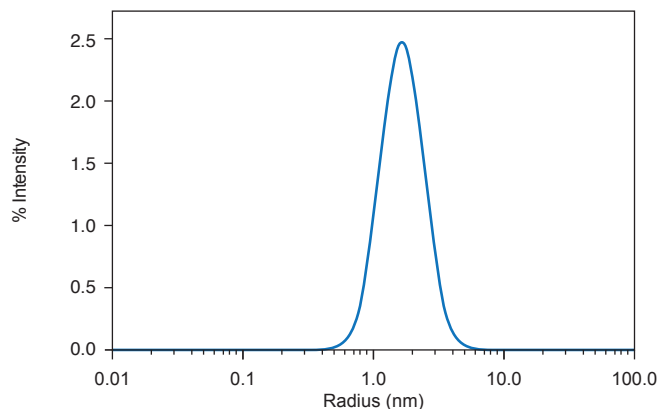


Particle size, crucial for biodistribution and cellular uptake, is an important property of LNPs. In the figure above, LNPs prepared using manual mixing (green) yielded larger particles with a 100 nm average radius, whereas microfluidic mixing (red) resulted in the desired 45 nm average radius.



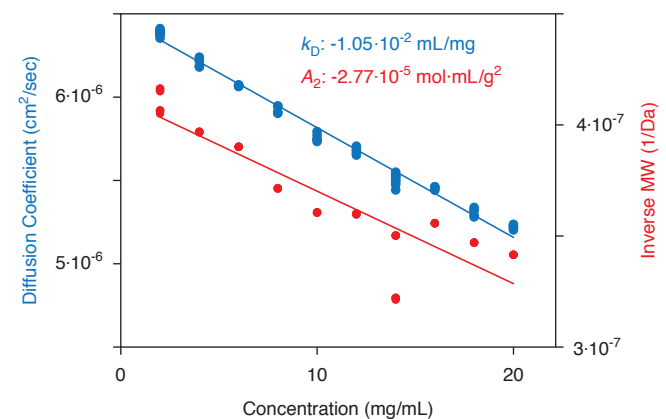
The zeta potential analysis reveals that all three LNPs exhibit a consistent negative charge, regardless of their preparation method or whether they are loaded or empty, with no statistically significant differences observed.

Quantum Dots Characterization



Quantum dots' properties depend greatly on size and zeta potential. However, they are tricky to analyze because they absorb and emit light in the visible range. This isn't an issue with the ZetaStar instrument, as its near-IR laser source suppresses emissions, ensuring precise analysis of these challenging nanoparticles.

In-process AAV Quantitation



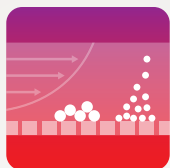
Direct measurement of AAV attributes during upstream and downstream processing (USP and DSP) can be accomplished at-line with a NanoStar or ZetaStar instrument. Sample purity progresses from crude USP product (blue) to the first DSP purification (red) and the final DSP filtrate (green), where a single size population is present with concentration of $2 \cdot 10^{12}$ particles/mL.



FFF Products

Advanced Separation Technology

Separate complex samples for extended characterization



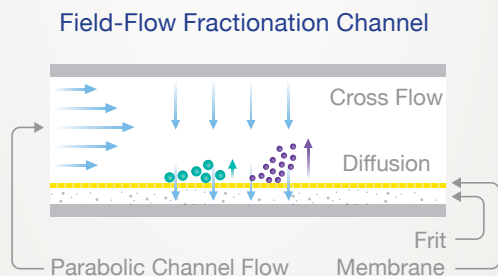
FFF

field-flow fractionation

FFF is a powerful separation technique covering a size range of 1 to 1000 nm and beyond. Having very low surface area and no stationary phase, FFF generates very little shear and is an excellent choice when non-ideal sample-surface interactions are a concern. MALS, DLS and dRI detectors are placed downstream of the separation channel for complete characterization.

FFF fractionates and characterizes:

- Colloids and nanoparticles
- Macromolecules and assemblies
- Complex samples



FFF separation power can be tuned by changing the ratio of cross flow to channel flow.



Eclipse

Advanced FFF technology

The Eclipse FFF is a sophisticated system for performing analytical and semi-preparative separations over a wide range of analytes.

The Eclipse system combines an industry-standard autosampler and pump for maximum reliability, convenience, repeatability and intelligence. With multiple online detectors, FFF-MALS provides extended characterization.

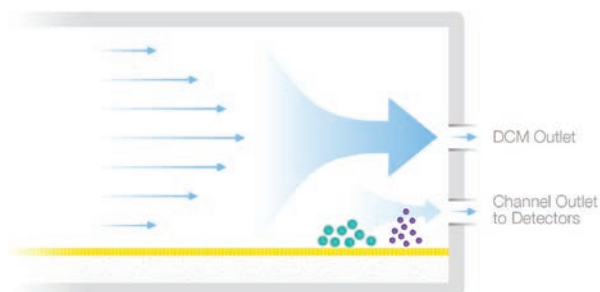
	Feature	Benefit
Single-pump technology	Standard	High system reliability
System Ready Monitor and Health Indicators	Standard	High productivity: eliminate bad runs and shorten troubleshooting
Injection method	Tip or Focus-zone	Supports different FFF separation techniques and SEC
FFF-SEC switching	Optional	Share FFF system with SEC
Dilution Control Module	Optional	Higher sensitivity, fraction concentration and repeatability
Mobility	Optional	Measure zeta potential of each fraction



Dilution Control Module

The Dilution Control Module (DCM) increases the concentration of sample eluting from the channel by a factor up to 5x or more. Benefits of the DCM:

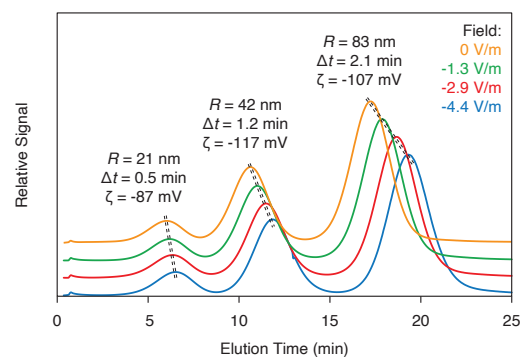
- Higher sensitivity at the detector
- Higher concentration in collected fractions
- Extended dynamic light scattering size range
- Highly reproducible retention time



Mobility

Mobility combines an innovative EAF4 channel design with precise current control, and the pH and conductivity measurements essential for zeta potential interpretation.

The Mobility channel is rigorously engineered for long life and high reliability. It incorporates a DCM port to increase retention accuracy, sensitivity and DLS measurement range.



Fixed-Height Channel



Key Features and Benefits

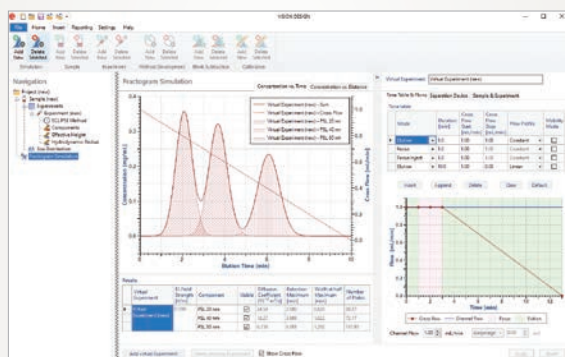
- Top plates with integrated channel heights eliminate the need for a spacer
- Robust channel sealing and fast assembly
- Ideal for routine and QC applications
- All Eclipse channels come with temperature regulation control from ambient up to 50 °C
- Semi-preparative fixed-height channel option available

Channel Type	Benefits	Applications
Analytical Short	Rapid nano/microgram separations	Versatile all-purpose
Analytical Long	Nano/microgram separations	Polymers, less prone to overloading
Semi-Prep	Milligram separations	Extracellular vesicle, virus and LNP isolation
Dispersion Inlet	For aggregation-prone samples	Monoclonal antibodies, liposomes
Mobility EAF4	Apply electric field to channel	Zeta potential distributions



VISION

intelligent design,
operation and analysis for
field-flow fractionation



In silico method development

VISION DESIGN software eliminates long cycles of trial-and-error method development. Its FFF simulation engine predicts the fractogram based on the specified separation conditions, leading to method optimization with a single sample run.

Regulatory Compliance

VISION software offers an optional 21 CFR Part 11 compliance package, including IQ/OQ documents and procedures.

VISION

The brains behind FFF

VISION software is the intelligent interface to FFF-MALS. It streamlines complex procedures and provides critical diagnostics to ensure simplicity and productivity. The software turns FFF-MALS into a routine analytical tool for scientists and technicians alike.

VISION DESIGN software helps users design optimal FFF separation methods from their desks. It also calculates diffusion coefficients and zeta potential from FFF and EAF4 measurements.

VISION RUN software is a sophisticated control center designed to run FFF methods. It seamlessly coordinates the pump, autosampler, Eclipse FFF controller, detectors and ASTRA software, and records FFF and UV signals for diagnostics and analysis in VISION DESIGN software.



VISION DESIGN

DESIGN METHOD

- Estimate particle size in sample
- Select spacers and flow profiles
- Use FFF theory to simulate and predict separation



Transfer optimized method



VISION RUN

RUN EXPERIMENTS

- Acquire flow, electrical and UV data
- Collect fractions
- Monitor and diagnose system



Launch and synchronize ASTRA software

ANALYZE AND REPORT



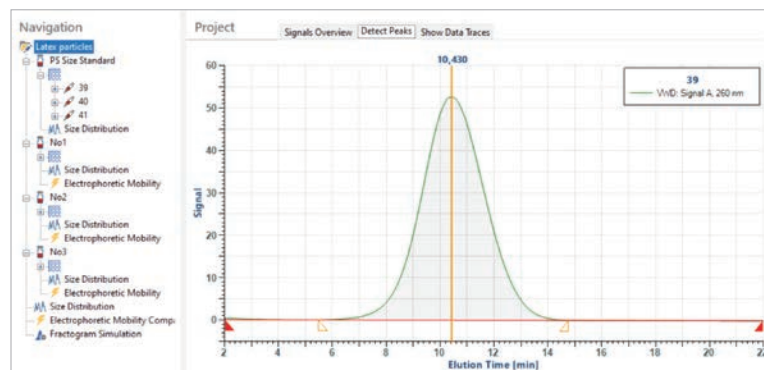
VISION DESIGN

- Collate project files
- Estimate particle size from retention time
- Refine method
- Determine zeta potential



ASTRA

- Basic and advanced MALS-DLS-UV-RI characterization
- EASI Graph overlays
- EASI Table consolidated results
- Customized reports

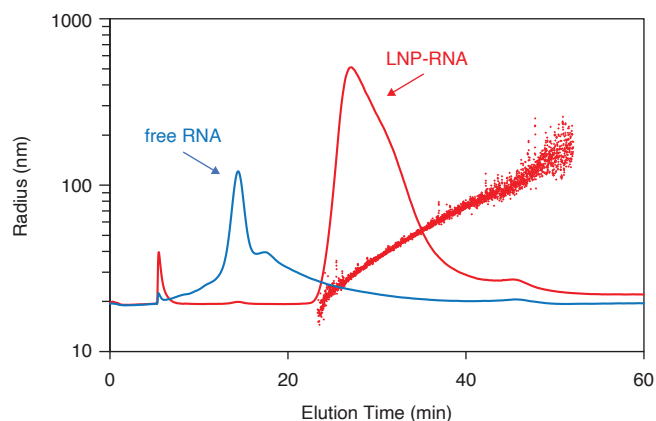


Smart project administration

VISION software organizes all FFF and MALS data as projects, for convenient review and replication, as well as comparison and reporting. Projects can be merged and experiments added or deleted at will, making this a powerful and flexible way to handle large sets of FFF experiments.

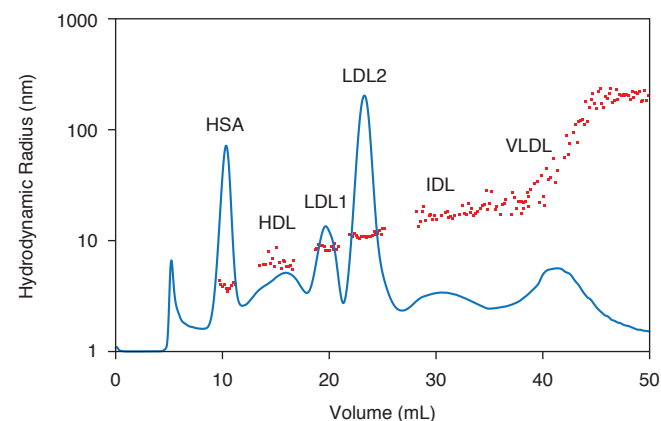
FFF Applications

DNA/RNA Lipoplex



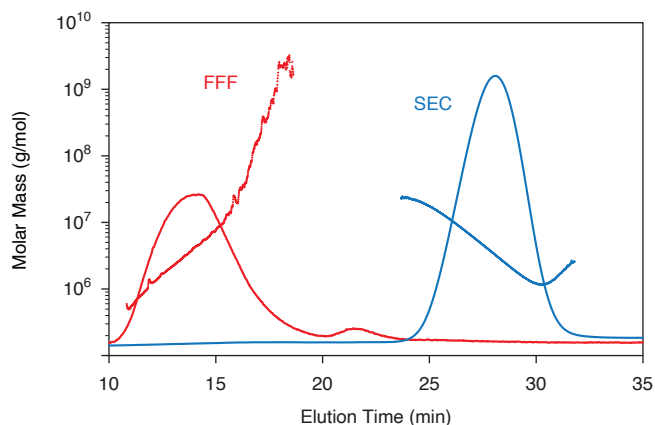
mRNA, siRNA, and plasmid DNA are often formulated into and delivered by non-viral vectors such as lipid nanoparticles (LNP). For such lipoplexes it is important to know the amount of DNA/RNA encapsulated by the particle. In this example, free RNA is well resolved from the mRNA-LNP complex by FFF separation and therefore the amount of free RNA is readily quantified.

Blood Serum Components



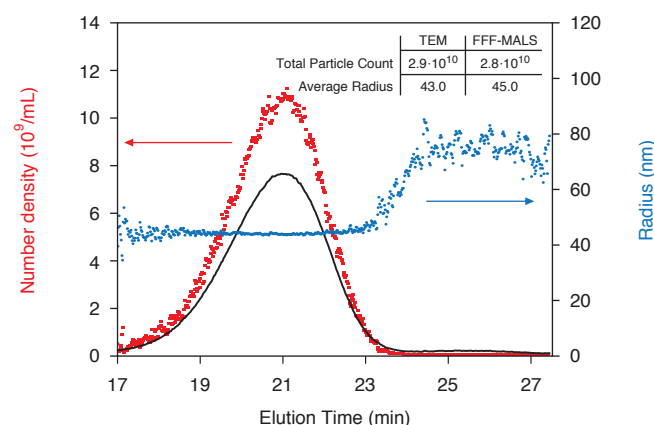
AF4-MALS-DLS separation of whole serum, with distinct peaks for serum albumin, IgG, and various types of lipoproteins. Hydrodynamic radii (R_h) were determined by online dynamic light scattering embedded in the DAWN MALS detector. Not shown, MALS determines molar masses of each peak; it also determines rms radius R_g for species larger than ~ 10 nm.

HMW complexes and conjugates



A high-molecular-weight protein-polysaccharide conjugate (PPC), spanning four orders of magnitude in MW, was characterized by both FFF-MALS (red) and SEC-MALS (blue). The SEC elution was non-ideal and HMW fractions were removed by the column, while FFF provides near-ideal fractionation of this large conjugate and is conducive to accurate MALS analysis.

Viruses and Viral Vectors



FFF-MALS provides quantitative, high-resolution size distributions based on large particle ensembles. This adenovirus analysis indicates the concentration (number density) in billion/mL at each elution time along with the radius. The LS fractogram is overlaid in black. The results compare well with TEM analysis.



CG-MALS Products

Analyze biomolecular interactions

Label-free, in solution, from pM to mM



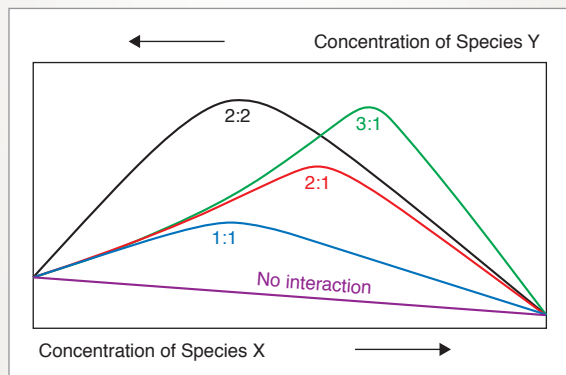
CG-MALS

composition-gradient
multi-angle light scattering

CG-MALS is a label-free, immobilization-free technique for characterizing:

- Protein-protein interactions
- Protein-DNA complexes
- Batch polymer interactions

CG-MALS characterizes biomolecular interactions from first principles by measuring the change in the weight-average molar mass (M_w) of a solution as a function of concentration and composition.



CG-MALS analyzes the light scattering signals from composition gradients to calculate K_d and absolute stoichiometry. It can differentiate between complexes with the same stoichiometric ratio but different overall number of bound monomers.



Calypso

Composition-gradient stop-flow system for biomolecular interactions and reaction kinetics

- K_d from pM to mM
- Reaction times from seconds to hours
- Self- and hetero-associations
- Interfaces with DAWN, miniDAWN and Optilab instruments for automated MALS and concentration measurements.



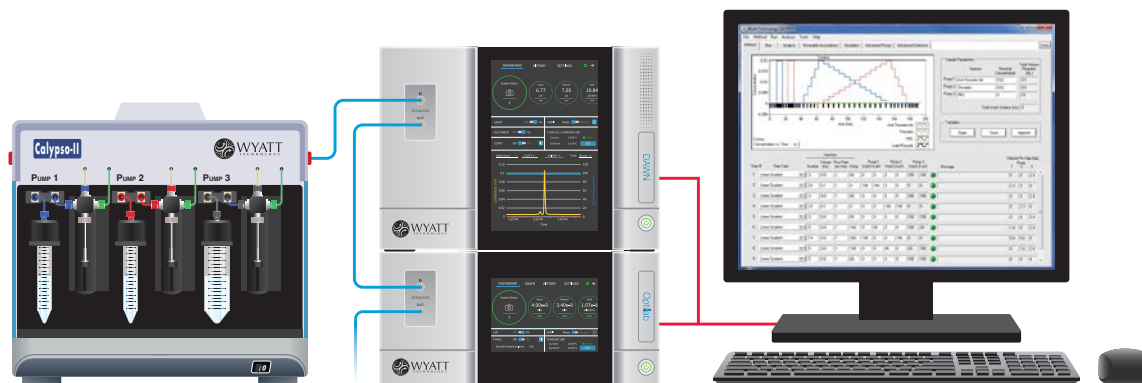
CALYPSO Software

Comprehensive set of association models covering simple to complex interactions

- Versatile, easy-to-use method programming for multiple gradient types, system preparation and post-experiment cleanup
- Simulation capabilities for experiment design and interpretation

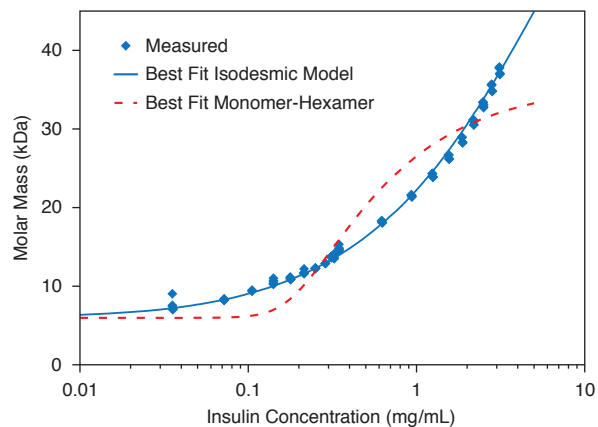
Versatile association model design for:

- Standard homodimer, heterodimer and progressive self-association
- Multivalent interactions and multiple oligomers in equilibrium
- Simultaneous self- and hetero-association
- High-concentration proteins
- Non-specific interactions of cosolutes



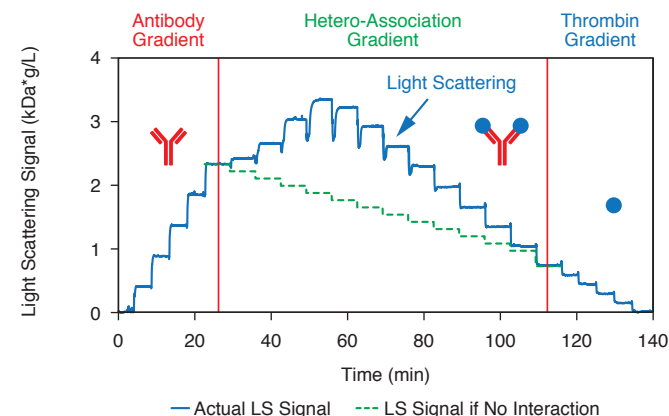
CG-MALS Applications

Insulin Self-Association



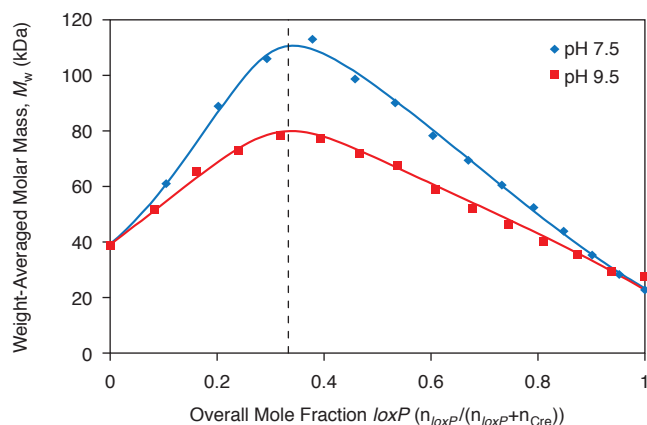
CG-MALS analyzes self-association by measuring the weight-average molar mass over a concentration series. In the absence of zinc, insulin is found to self-associate isodesmically (progressively) with a K_d of 52 μM . A monomer-hexamer model fits poorly and can be ruled out.

Antibody-Antigen Binding



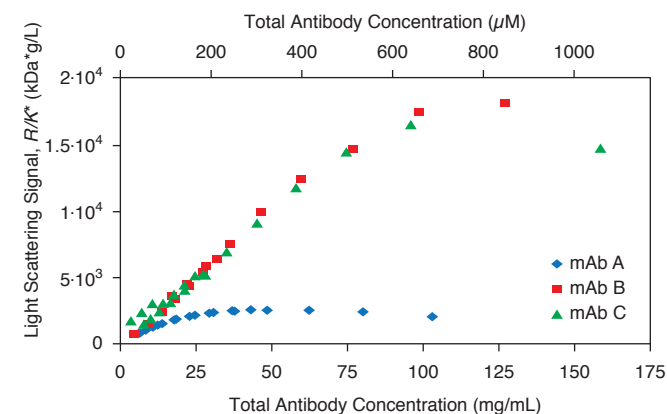
A Calypso stop-flow measurement of antibody-antigen interactions. Here the CALYPSO software found that thrombin binds to an anti-thrombin monoclonal antibody with $K_d=9$ nM at two equivalent, non-cooperative binding sites on the mAb and no self-association.

Cooperative Binding vs. pH



Cre recombinase binds to the *loxP* DNA segment in a pH-dependent manner. CG-MALS determines that at pH 7.5, each *loxP* binds two Cre molecules with positive cooperativity, and the 2:1 complex dimerizes to form a synapse tetramer; while at pH 9.5, cooperativity and synapsis are lost.

High-Concentration IgG



mAbs A, B and C exhibit widely varying viscosities at high protein concentration, a consequence of differing degrees of self-attraction. CG-MALS is one of very few techniques capable of analyzing protein self-interaction at high concentrations. For these mAbs, self-interaction correlates well with viscosity.

Service & Support Plans

continued instrument service
and software support



Service & Support Plan

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Service & Support

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We look forward to meeting you at Light Scattering University!



Michelle Radeke, Ph.D.
Head of Customer Service
Joined Wyatt Technology 2015

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Michelle Chen, Ph.D.
Senior Director, Research and Development
Joined Wyatt Technology 1996

Light Scattering University



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**Dr. InKwan Han,
Merck & Co. Inc.**

Highlights of LSU

Many trainees come away from LSU inspired with new ideas for how light scattering can solve some of their analytical challenges. One of the most popular aspects of LSU is the opportunity to meet and work with the scientists and engineers behind the products, as well as get acquainted with support staff that they usually only contact over the phone.

Another not-to-be missed session (available only in Santa Barbara) is the Light Scattering Museum tour, a fascinating journey through the history of light scattering technology.



LSU

light scattering university

Often described by participants as the best instrument user training they have ever attended, Light Scattering University (LSU) is an intensive experience that combines hard work, good food and a friendly atmosphere.

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- How to interpret your data
- Instrument best practices
- History of light scattering



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