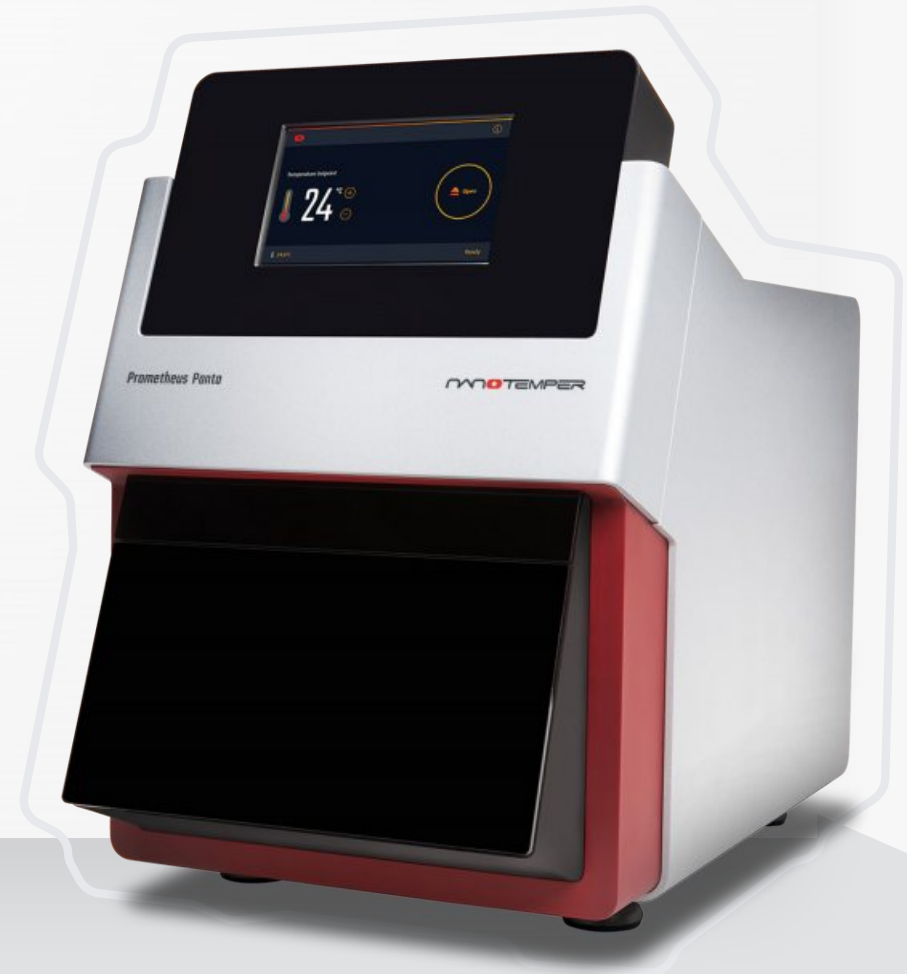


# Prometheus Panta

The new gold standard for challenging stability characterizations brings DLS and SLS to your biologics workflow





Whether you're working in biologics formulation optimization, developability, or comparability assessments, count on Prometheus Panta to provide multi-parameter stability characterization and trustworthy results for your candidate molecules.

**Collect data on thermal unfolding, particle sizing, and aggregation simultaneously throughout the entire thermal ramp for high-resolution, domain-specific stability characterization of biologics.**



## Prometheus Panta combines thermal unfolding, particle sizing, and aggregation for superior characterization of candidate molecules

### **Get high-resolution data that reveals liabilities in drug candidates missed by other technologies**

Having the right technology when it really matters — to detect small differences or subtleties in unfolding events — makes a huge difference. Be confident you're passing high-integrity results on to the next team which they can count on when making critical decisions.

### **Get domain-specific information when you measure thermal unfolding, particle sizing, and aggregation attributes simultaneously throughout the entire thermal ramp**

For the first time correlate particle sizing, thermal unfolding, and aggregation results collected throughout an entire thermal ramp — and get a completely new perspective about stability attributes at the domain level of your candidate molecule.

### **Get an instrument that has flexible throughput and a few choices for sample handling**

Handle any project that comes your way no matter if you need to characterize just a few or a few dozen candidate molecules.

# Monitor essential attributes throughout your biologics workflow

Your candidate molecules go through a long and complicated journey on their way to becoming a final product. Using the same instrument throughout this process ensures there's consistency when you need to compare conformational and colloidal stability data across teams and sites.

Start multi-parameter characterization early on when candidate molecules are being considered, engineered, or modified, and then continue comparing throughout formulation, production, and validation with Prometheus Panta.



## Developability

Determine aggregation propensity

Get self and non-specific interactions

Characterize conformational (thermal) stability



## Antibody engineering & stability enhancement

Determine aggregation propensity

Get self and non-specific interactions

Characterize conformational (thermal) stability



## Downstream process development

Characterize conformational (thermal) stability and determine aggregation propensity and size distribution during scale-up and optimization of processes



## Pre-formulation & formulation

Characterize conformational (thermal) stability

Determine aggregation propensity

Perform buffer and excipient screening and compatibility with melting scans with particle sizing, size distribution, and aggregation propensity



## Investigational new drug (IND) & New drug application (NDA)

Determine thermal stability and particle sizing on reconstitution/dilution/admixing at initial and final time points

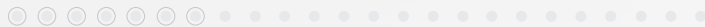
Get thermal stability and particle sizing from forced degradation and photostability studies



## Comparability assessment

Characterize conformational (thermal) stability including accelerated stress conditions spanning a broad concentration range

Determine particle size distribution

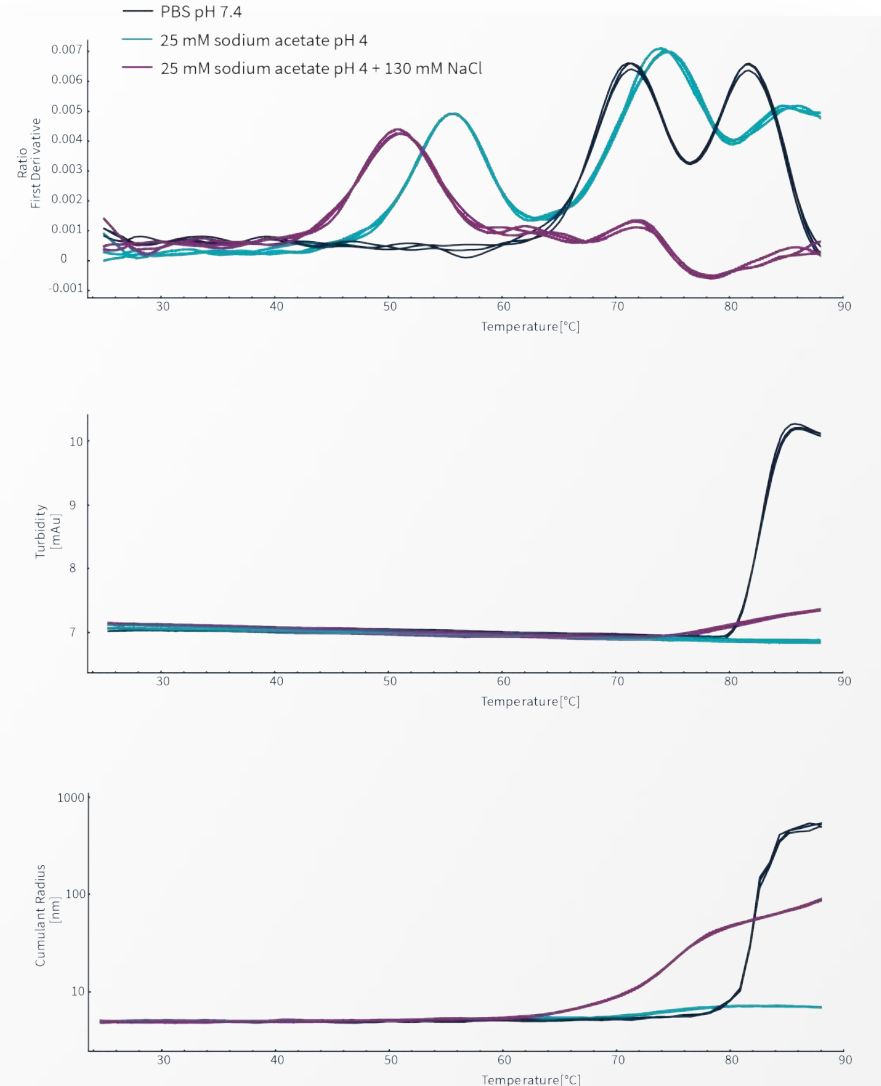




# Become more efficient with simultaneous measurements. Share clear and actionable results sooner.

Providing clear and actionable results gets more complicated when you need to measure multiple parameters. And, when you're up against deadlines and your colleagues are relying on you for reliable results, it's difficult not to feel under pressure. With Prometheus Panta, become more efficient by measuring multiple parameters simultaneously —  $T_m$ ,  $T_{\text{turbidity}}$ , and  $r_H$  — throughout a single thermal ramp. And, provide your colleagues with clear and actionable stability results sooner.

Identify and differentiate stability behavior with simultaneous acquisition of nanoDSF, backreflection, SLS, and DLS measurements collected throughout the entire thermal ramp for Herceptin in three different buffer conditions.

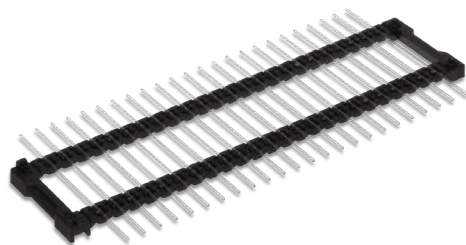


# High performance consumables for consistent results

You don't have to worry about a long list of consumables to run your stability assays. All you need are capillaries.

They're manufactured using the same stringent protocols used for diagnostic-grade capillaries, so you get consistent results.

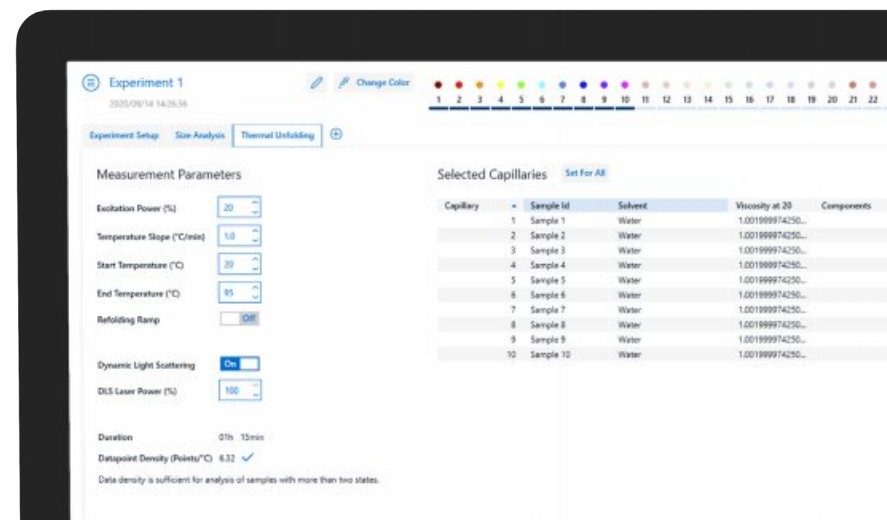
Choose to load up to 48 individual samples, or use chips of 24 capillaries that make it easy to load your samples from standard 384-well plates.



# Software designed to make you more efficient

Panta Control software lets you design experiments that address important questions about your vector's stability. Get the flexibility to queue up experiments, or adapt your work on-the-fly.

With Panta Analysis software displays your results with graphs and charts that help you visually interpret your data. Rank the stability parameters that matter most for your vectors, and quickly compare data within experiments.



# Prometheus Panta specifications, because you need to make decisions



## General Specifications

<b>Sample handling format</b>	Capillaries or capillary chip
<b>Throughput in one run</b>	Up to 48 capillaries or 24 in capillary chip
<b>Sample volume</b>	10 $\mu$ L
<b>Temperature range</b>	15 - 95 °C (up to 110 °C with High Temperature Upgrade)
<b>Heating rate range</b>	Up to 95 °C: 0.1 - 7 °C/min Above 95 °C: 0.1 - 7 °C/min
<b>Precision of 1 °C/min thermal ramp</b>	$\pm$ 0.1 °C
<b>Dimensions</b>	35 cm W x 51 cm H x 52 cm D
<b>Weight</b>	35 kg



# Details for the techie in your lab



## Technology Specifications

### nanoDSF

<b>Measurement parameters</b>	Ratio: $T_{\text{onset}}$ , $T_m$ , $E_a$ , reversibility of unfolding 330 nm, 350 nm: $T_m$ Excitation: 280 nm
<b>Concentration range</b>	5 $\mu\text{g/mL}$ - 250 $\text{mg/mL}$
<b>Inflection point precision @ 75 °C</b>	$\pm 0.1$ °C
<b>Ratio precision/reproducibility</b>	0.008

### DLS

<b>Measurement parameters</b>	$T_{\text{size}}$ , $r_H$ , PDI, $k_D$ , $D_0$ , reversibility of unfolding
<b>Laser wavelength</b>	405 nm $\pm$ 5 nm
<b>Concentration range</b>	0.5 $\text{mg/mL}$ for a 15 kDa protein, up to 40% w/v
<b>Size resolution</b>	Down to 0.5 nm

### Backreflection

<b>Measurement parameters</b>	$T_{\text{turbidity}}$ , reversibility of unfolding
<b>Size resolution</b>	Larger than 12.5 nm radius

### SLS

<b>Measurement parameters</b>	Molecular weight, $B_{22}$ , $T_{\text{scattering}}$ , average scattering intensity
<b>Measurement accuracy</b>	$\leq 10\%$ molecular weight

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