



#### **Microcalorimetry**

Isothermal titration calorimetry (ITC) and differential scanning calorimetry (DSC) are powerful analytical techniques for in-depth characterization of molecular binding events and structural stability. Thermodynamic binding signatures not only reveal the strength of a binding event, but the specific or nonspecific driving forces involved. Structural stability profiles from DSC reveal strengths and weaknesses in higher order structure and define the behavior of individual domains and their interactions. The TA Instruments Affinity ITC, Nano ITC and Nano DSC provide the performance, reliability and ease-of-use required for the most demanding applications in drug discovery, protein-protein interactions, structure-function characterization and more.



The Nano DSC has the versatility and precision for characterizing molecular stability, determining high affinity ligand binding and deconvoluting multi-domain structures. There is no other DSC with the proprietary technologies, high performance or the sample throughput of the Nano DSC and Nano DSC Auto.

#### **Features:**

- Highest sensitivity, lowest cell volume for unmatched performance
- Capillary cell design for analysis of samples that tend to aggregate or precipitate
- Built-in precision pressurizing system maintains accurate, constant pressure in the cells
- Solid-state thermoelectric elements for accurate temperature control during heating and cooling scans
- Upgradeable with industry proven HPLC grade autosampler for reliable high sample throughput





### Nano DSC TECHNOLOGY

The Nano DSC is designed for ultra-sensitive measure of heat absorbed or released by dilute in-solution bio-molecules as they are heated or cooled. The capillary cell design, solid-state thermoelectric temperature control and easy cleaning ensure the highest sensitivity and data reproducibility for a wide variety of applications.

#### **Features:**

- 300 µL active volume capillary cells for analyzing hydrophobic samples
- Easy, accurate sample loading with laboratory pipetteman
- Built-in, user-programmable pressurization system (up to 6 atm)
- Flexible data acquisition interface for easy experiment setup
- NanoAnalyze software for accurate model fitting and multi-file batch processing



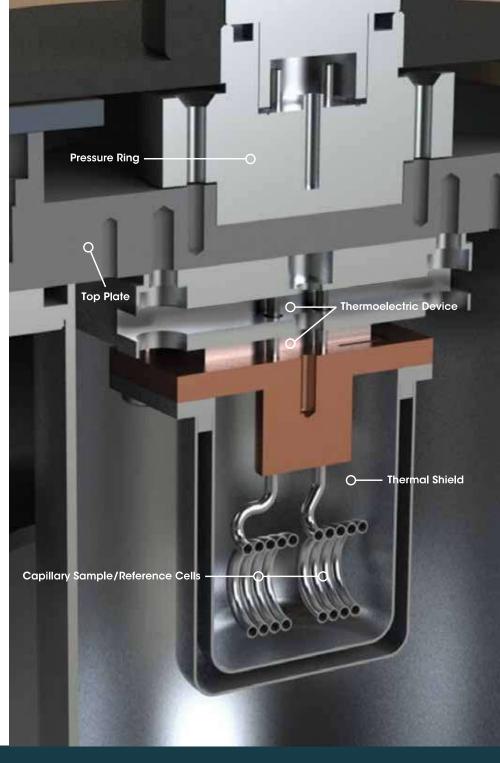
The NanoDSC is a powerful thermal scanning instrument that utilizes a 300 µL capillary cell design and solid-state thermoelectric temperature control to provide unmatched performance.

#### Nano DSC Capillary Platinum Cells

- Fixed-in-place capillary cells attenuate aggregation and precipitation
- Platinum cells are inert and compatible with strong acids, bases and protein cleaning enzymes
- 300 µL active cell volume minimizes sample consumption
- · Sample cell loading with laboratory pipetteman is easy and ensures no trapped air bubbles

#### Nano DSC Solid-State Thermoelectric Temperature Control:

- Accurate, reproducible temperature control for highest sensitivity in both heating and cooling scans and unmatched baseline reproducibility
- Innovative, user-programmable built-in pressure system for complex analysis of water characteristics and molecule structure
- User-programmable scan rates for scan flexibility and highest confidence in data analysis



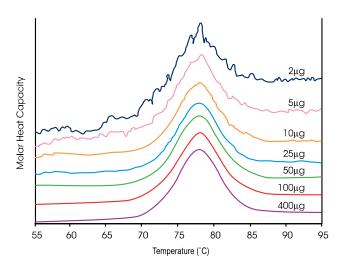


### Nano DSC

**APPLICATIONS** 

#### How much Protein is Required for a DSC Scan?

Determining the thermodynamic parameters of a protein by differential scanning calorimetry (DSC) using the Nano DSC requires about the same amount of protein as surface plasmon resonance or fluorescence studies. Because of the Nano DSC's extreme sensitivity and baseline reproducibility, and the sample cell's small volume (300  $\mu$ L), a complete, interpretable, accurate scan can be obtained on essentially any protein of interest. The sensitivity and accuracy of the Nano DSC is demonstrated by this data. Hen egg white lysozyme (in pH 4.0 glycine buffer) was prepared at various concentrations. As little as 2 µg of lysozyme in the capillary cell is sufficient to provide quality data yielding accurate values of all four thermodynamic parameters!



Lysozyme	Calorimetric		Van't Hoff	
in cell (50µg)	ΔH (kJ mol <sup>-1</sup> )	ΔS (kJ K <sup>-1</sup> mol <sup>-1</sup> )	T <sub>m</sub> (°C)	ΔH (kJ mol <sup>-1</sup> )
400	512	1.46	78.0	515
100	512	1.46	78.0	509
50	517	1.47	77.9	513
25	513	1.46	77.8	513
10	515	1.47	78.0	515
5	490	1.40	78.0	510
2	503	1.43	77.8	499

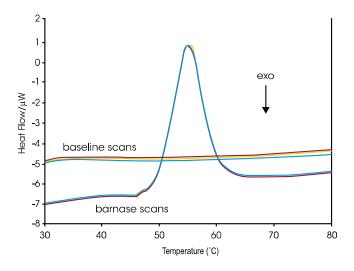
## Nano DSC APPLICATIONS

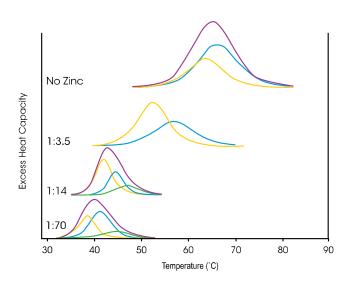
#### **Characterization of Protein Stability**

Analyzing the stability of a protein in dilute solution involves determining changes in the partial molar heat capacity of the protein at constant pressure ( $\Delta$ Cp). The contribution of the protein to the calorimetrically measured heat capacity (its partial Cp) is determined by subtracting a scan of a buffer blank from the sample data prior to analysis. Heating the protein sample initially produces a slightly increasing baseline but as heating progresses, heat is absorbed by the protein and causes it to thermally unfold over a temperature range characteristic for that protein, giving rise to an endothermic peak. Once unfolding is complete, heat absorption decreases and a new baseline is established. After blank subtraction, the data can be analyzed to provide a complete thermodynamic characterization of the unfolding process.

#### **Characterization of Protein Structure**

DSC can be used to characterize both the specific binding of a ligand (for example, a drug to a receptor binding site), or nonspecific binding (for example, detergents binding to hydrophobic patches on a protein surface). In some instances ligand binding, even if to a specific receptor site, results in long-range protein structural rearrangements that destabilize the entire complex. The figure shows DSC scans of Ca²+ saturated bovine a-lactalbumin at various protein:Zn²+ ratios scanned at 1 °C/min. The midpoint of the thermal unfolding of the protein decreases from 65 °C in the absence of Zn²+ to 35 °C at a protein:Zn²+ ratio of 1:70. The enthalpy of unfolding is also decreased substantially by high Zn²+ concentrations.



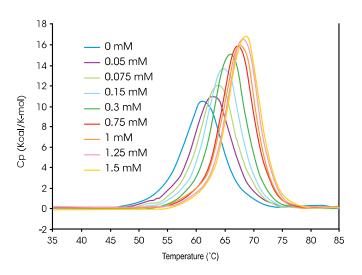


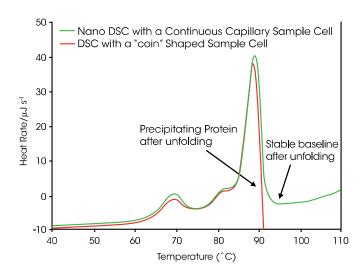
#### **Investigation of Protein-Ligand Binding**

DSC is a valuable tool for studying binding between a biological macromolecule and a ligand such as another biopolymer or a drug. Unlike ITC, DSC allows the thermodynamics that drive binding to be correlated with conformational changes in the macromolecule caused by the binding reaction. DSC is particularly useful for characterizing very tight or slow binding interactions. DSC also allows characterization of binding reactions that are incompatible with the organic solvent requirements of some ITC experiments (i.e., where ligand solubility for an ITC experiment requires concentrations of organic solvent not tolerated by the protein). The data shows DSC scans of RNase A bound with increasing concentrations of 2'-CMP, showing that the protein is stabilized by higher concentrations of the inhibitor, Essentially identical data was obtained in the presence of 5% DMSO, verifying that organic solvents are compatible with the DSC technique.

#### Nano DSC Capillary Cell Advantages

This figure shows two DSC scans of matched samples of human IgG1 at 0.5 mg/ml in physiological buffer. The data from the DSC with a "coin" shaped sample cell shows the easily recognizable exothermic aggregation/precipitation event at approx 89-90 °C, while the data collected on the Nano DSC with a capillary sample cell shows a stable posttransition baseline that will enable complete and accurate determinations of transition temperatures (Tm) and enthalpy ( $\Delta H$ ).





# ITC & DSC SOFTWARE

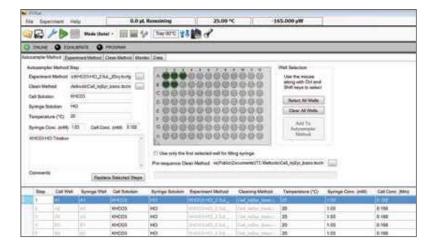
#### **Instrument Control & Data Acquisition Software**

The Affinity and Nano instruments control and data acquisition functions are executed within a Windows-compatible software interface, ITCRun or DSCRun. All experimental parameters and sample information are easily entered into an intuitive graphical user interface and can be saved as an experimental template for future use.

Real-time monitoring of the raw data as the experiment progresses allows rapid assessment of the data quality and instrument performance in individual tabs. Unique icon-controlled functions, such as immediate baseline subtraction, are always available on the display.

#### **ITCRun & DSCRun features:**

- Automatic configuration of user interface for automated or non-automated instruments
- Individual viewing tabs for real-time monitoring of instrument performance characteristics and raw data acquisition
- Easy experiment setup
- Direct autosampler programming and control for automated instruments
- Software passes all experimental parameters to NanoAnalyze™



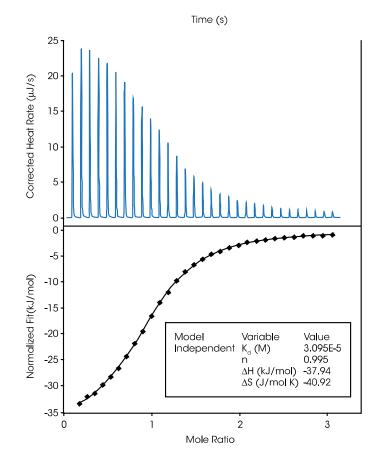
#### Data Analysis with NanoAnalyze™

All ITC and DSC raw data files are easily and quickly analyzed with a powerful ITC/DSC data analysis software, NanoAnalyze. Individual window tabs for each processing step guide the user through the analysis of Individual raw data files or the batch processing of multiple files.

#### NanoAnalyze™ features:

- Easy import of all ITC and DSC raw data files
- User selectable fitting models for ITC and DSC
- Easy set up of new fitting models
- Drag & Drop subtraction of baseline blank files
- · Powerful experiment design and optimization tool
- Flexible overlay graphs for quick data comparisons
- Generates thermodynamic profile bar graph
- Easy export of all data to delimited text files
- Full-featured editing tools for preparation of publication quality images

All instrument control, data acquisition and data analysis software required for ITC and DSC data are provided with all Affinity and Nano instruments. All software updates and feature improvements are available on the TA Instruments website.



#### Nano DSC and DSC Auto

Short-term Noise	0.015 µWatts
Baseline Stability	±0.028 µWatts
Response Time	7 seconds
Operating Temperature	-10 °C to 130 °C or 160 °C
Temperature Scan Rate	up to 2°C/minute
Pressurization Perturbation	Built-in up to 6 atmospheres
Pressurization Perturbation Cell Volume	Built-in up to 6 atmospheres  300 mL
	<u> </u>
Cell Volume	300 mL

#### **Automation**

Sample Capacity	2 standard plates x 96 wells x 1000 µL/well
Sample Tray Temperature Control Range	4 °C to Ambient
Available Wash/Rinse Buffer Ports	4 for Sample/Reference Cells; 2 for Sample Handling Syringe

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