

# Agilent ChemStation for UV-visible Spectroscopy: Biochemical Analysis Software

### **Specifications**

#### **General Description**

The biochemical analysis software for Agilent ChemStation adds single cell kinetics, multicell kinetics and thermal denaturation (DNA melt) capabilities to the general purpose software.

#### **Kinetics** — **Data Acquisition**

- For time-based measurements the run, start, and cycle time can be defined. For non-equidistant time intervals, a percent increase of the cycle time can be defined.
- Full or partial spectra can be acquired.
- Measurements can be made in parallel using the cell multicell transport.
- The positions of the blank and sample cells for time-based measurements are defined by the
- All traces and the spectra of one selected cell can be monitored online in a trace monitor and a spectra monitor.

#### **Kinetics** — Data Analysis

- Data from single (multiple cell acquisition) or multiple wavelengths (single cell acquisition only) can be extracted from timebased spectra.
- A choice of four rate calculation types is available: initial rate, delta AU, zero and first order.
- Time traces may be interactively processed by the user with mathematical functions.
- Time traces and rate data can be exported as CSV and DIF files which are common ASCII text file formats.

## <u>Thermal Denaturation —</u> <u>Instrument Control and Data</u> <u>Acquisition</u>

- For temperature-based measurements, a ramp with multiple start, step, and end temperatures can be defined.
- The temperature can be taken from the cell holder or the optional dipping probe.
- Full or partial spectra can be acquired.
- The temperature trace can be monitored online.

#### <u>Thermal Denaturation —</u> <u>Temperature Trace Evaluation</u>

- Data from single wavelength can be extracted from temperaturebased spectra.
- The transition temperature can be determined in a user-defined calculation range by absorbance average or first derivative.
- An equation allows the user to enter an equation for the calculation of a result from the Tm value. The default equation is the calculation of %GC.
- Temperature traces can be exported as CSV and DIF files which are common ASCII text file formats.



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