

### ATR FTIR Spectoscopy of Aqueous Cell Cultures V.2

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#### ABSTRACT

ATR-FTIR Spectroscopy of aqueous cell culture samples is explained step-by-step. Infrared spectra acquisition, processing and analysis is included briefly.

### Spectra acquisition

- Arrange the temperature of the working environment to § 25 °C
- Clean the ATR crystal using a cellulose wipe and water-ethanol-water sequence. Scratching the crystal surface must be avoided.



- Leave the crystal to air-dry
- Select "Scan and Instrument Setup" and input sample name, scan range (650-4000 cm<sup>-1</sup>), scan number (128), and spectral resolution (2 cm<sup>-1</sup>). This step needs optimization for specific sample being used.



- Scan the backgroung spectrum to overcome the probable atmospheric interference that can be triggered by H<sub>2</sub>O and CO<sub>2</sub> molecules present in the air, subtracted automatically by the software.
- Additionally, record PBS as sample (subtract this manually from each spectrum) This eliminates interference of PBS and flattens the water band located around 2125 cm<sup>-1</sup>

Put 📮 1 µl cell samples and let to dry with a mild N<sub>2</sub> flux for 🕲 00:05:00 . This allows cells to settle on the crystal.

5m

- 8 Scan the sample. Process takes about **© 00:07:00** to finish.
  - As the scanning process goes on, one must be sure that the quality check sign is everytime green.
- 9 Scan each sample under the same conditions in three independent scans (the average constitutes a replicate n). Replicates (n=5) are used in data analysis.
  - This minimizes intra-sample variability and eliminates possible variation that might arise from experimental conditions.

### Spectral pre-processing

10 The avarage of triplicates is taken manually.



- 11 13-point Savitzky-Golay smoothing
- 12 Baseline correction at 3800-2750-1800-900 cm<sup>-1</sup> points.

# Principal Component Analysis (PCA)

13 Import spectra as data file into



- 14 Define the data sets to 3000-650 cm<sup>-1</sup> region.
- Make the PCA Model.
  Define the regions with high variation using the Loadings plots.
  Check how the replicates are grouping and if there are outliers in Scores plots.

# **Spectral Calculations**

Peak labelling can be done automaticaly. For increased precision we measured manually: Measure the center of weight of each band. Read the frequency. Compare the peaks you obtained, to the literature:



Make an excel file of your peaks and their assignment to the functional groups. For side-peak band assignment, the second derivative spectrum is used.

7 Concentration studies are carried out by integrating the curves under the peaks.

For conformational alteration studies, bandwidth measurements are done manually using the 75% height of the peaks.

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