

Bauer Core Standard Protocol		
Title: Using the Spectramax Plus 384 UV-Visible Plate Reader		
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Author(s): Claire Reardon	Reviewers: Michelle Li	
Contact: claire@cgr.harvard.edu	Comment:	

1. Purpose

This protocol provides instructions for using the Spectramax Plus 384 UV-Visible Plate Reader to read absorbance between 190 – 1000 nm. The protocol is designed as a reference and is not a substitute for training. Users must complete a training session before using any of the Bauer Core’s instrumentation.

2. Materials

- 2.1. UV Transparent Plates (e.g. Corning plate #3635) are needed at wavelengths below 340nm. Above 340nm, any 96 or 384 well plate can be used.
- 2.2. Samples (minimum 100µl volume)

3. Instrumentation

Molecular Dynamics Spectramax Plus 384 UV-Visible Plate Reader

4. Reagent preparation

none

5. Procedure

- 5.1. Open the SoftMax Pro software using the Spf shortcut on the desktop.
- 5.2. Click on the Instrument Icon.
 - 5.2.1. Choose COM 4 and Spectramax Plus 384.
- 5.3. Click on Plate 1 – Setup.
 - 5.3.1. Assay type.
 - 5.3.1.1. Endpoint: most common assay, 1 scan/well, up to 4 wavelengths. Use autocutoff to ensure excitation wavelength isn’t detected.
 - 5.3.1.2. Kinetic: will scan several times over selected run time. Choose run time and interval.
 - 5.3.1.3. Spectrum: will scan over a range of wavelengths.
 - 5.3.2. Wavelengths – choose the wavelength(s) you would like to read.
 - 5.3.3. Pathcheck: Used to normalize the results to a 1cm pathlength.
 - 5.3.3.1. Choose the “water constant” to use a pre-programmed constant.
 - 5.3.3.2. Choose cuvette reference determine the constant from a water sample in the cuvette holder.
 - 5.3.3.3. Background constant: If you don’t want to run a pre-read blank

- enter your pre-determined background constant for each wavelength.
- 5.3.4. Automix and Blanking:
 - 5.3.3.1. Automix will shake the plate for 5s (not very powerful).
 - 5.3.3.2. Use “pre-read” to read a blank water plate before the samples
this is the most accurate way to blank.
 - 5.3.5. AutoCalibrate: on. The instrument self zeroes against an internal reference.
 - 5.3.6. Assay Plate Type: choose from the list of well types.
 - 5.3.7. Strips: choose “read entire plate”.
 - 5.3.8. Speed Read: Choose no/off.
 - 5.3.9. Column Priority: choose column priority
 - 5.3.10. Autoread: off.
 - 5.4. Click on Template to choose blanks, standards, and unknowns or just leave blank.
The average of the Blanks will be subtracted from the data wells
 - 5.5. Click on Reduction.
 - 5.5.1. Click on absorbance and use both pre-read and path check.
 - 5.5.2. Input any formulas you would like to use.
 - 5.6. Click on Display to determine what data is shown (raw only or reduced).
 - 5.7. Click on Temperature to set the assay temperature from 4°C above ambient to 45°C.
 - 5.8. Place your blank plate in the reader.
 - 5.8.1. Click the “drawer” button on the instrument to open and close the drawer.
 - 5.8.2. Well A1 should be in the back left corner.
 - 5.9. Click Read.
 - 5.9.1. If a pre-read is being done, insert the blank plate and click Pre-Read.
 - 5.9.1.1. The results for the pre-read will be “Path?”– this is normal.
 - 5.9.2. Click Read again and choose “Normal” to read the real samples.
 - 5.10. Your data will now appear in the plate display
 - 5.10.1. Note that the linear range of absorbance is from 0-2.35.
 - 5.11. You can copy and paste the results into excel for further analysis.
 - 5.11.1. If you pre-read a plate, the results will appear in two sections:
The upper section is the raw result and the bottom section is the pre-read value.
Subtract the pre-read value from the raw result to get the corrected data.
 - 5.12. To enter another plate click Experiment – New Plate .
 - 5.13. To save the settings, click Save As.
this will save a .pda file (readable only in softmax).
 - 5.14. Remove your plate, close the drawer, and log off from the computer when finished.