

Bioanalytical Method Development with UNIFI: Gradient and Temperature Optimization

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Abstract

The Waters UNIFI Scientific Information System employs a workflow approach with acquisition templates and promotable parameters to enable the rapid simple screening and optimization of gradient and column temperature conditions. Further optimization of the chromatographic parameters can significantly improve assay robustness and analytical throughput.

Benefits

Maximize bioanalytical assay productivity and throughput using promoted parameters and template acquisition with the Waters UNIFI Scientific Information System.

Introduction

Developing and optimizing the chromatographic conditions for a bioanalytical method is a multi-step process that includes screening column chemistries, mobile phase organic modifier, mobile phase pH, and column

After the mobile phase solvents and column chemistry have been selected, the gradient steepness, starting concentration and final composition can be optimized to maximize analyte resolution and minimize the analytical run time to improve sample throughput.

Welcome to UNIFI

Home

Getting Started
A tutorial describing how to get started with UNIFI.

Administration
Configure administrative settings and customise functionality within UNIFI.

My Work
Start new tasks and keep track of your work.

UNIFI™

Show help on...

☒ Show welcome screen at startup

Develop a bioanalytical workflow
Quickly develop sensitive and robust bioanalytical assays. Learn how to set up and configure hardware, run samples, and analyze results.

Analyze Data
View your chromatograms and spectra, perfect the integration, review the results, and compare reference and unknown samples.

Approve Reports
When you create and share reports that contain statistical information about your results, reviewers can sign off on those results using an electronic signature.

Manage Projects
Use folders to manage collections of data such as analysis results, reports, methods, and imported files that reside in UNIFI.

Define Methods
Methods are the specific instructions you want used to configure your system and to acquire, process, report, and export data.

Chadwick, Stuart (Chemist)

Plexis Analytica (Reg Bio 1 TQ System)

Figure 1. UNIFI Scientific Information System.

Experimental

UPLC Conditions

LC system:	Waters ACQUITY UPLC
LC column:	ACQUITY UPLC BEH C ₁₈ 1.7- μ m, 2.1 x 50 mm

MS Conditions

MS system:	Waters Xevo TQ-S
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The separations were performed using a mobile phase of 0.1% (v/v) formic acid and acetonitrile as the organic modifier

The gradient steepness and column temperature were adjusted as a variable during the analytical acquisition process.

The mass spectrometer was operated in electrospray positive ion mode with MRM data collection, RADAR full scan data collection and precursors of $m/z = 184$ to monitor the phospholipid elution profile.

Biological samples were prepared by spiking the target analyte into control human plasma. The spiked sample was then precipitated using a 2:1 ratio of acetonitrile to plasma. The resulting sample was centrifuged at 13,000 RPM and the supernatant removed for analysis by LC-MS.

Results and Discussion

The acquisition of data within the Waters UNIFI Scientific Information System is workflow-driven, allowing scientific managers to develop and administer templates for their scientist to employ for multiple tasks, including routine batch analysis, matrix effects calculation, gradient screening and gradient optimization. This use of a workflow-driven approach ensures consistency and allows the less experienced scientist a structured route to analytical method development and optimization.

Within the UNIFI Scientific Information System, the user is presented with a series of workflows to select from, Figure 2.

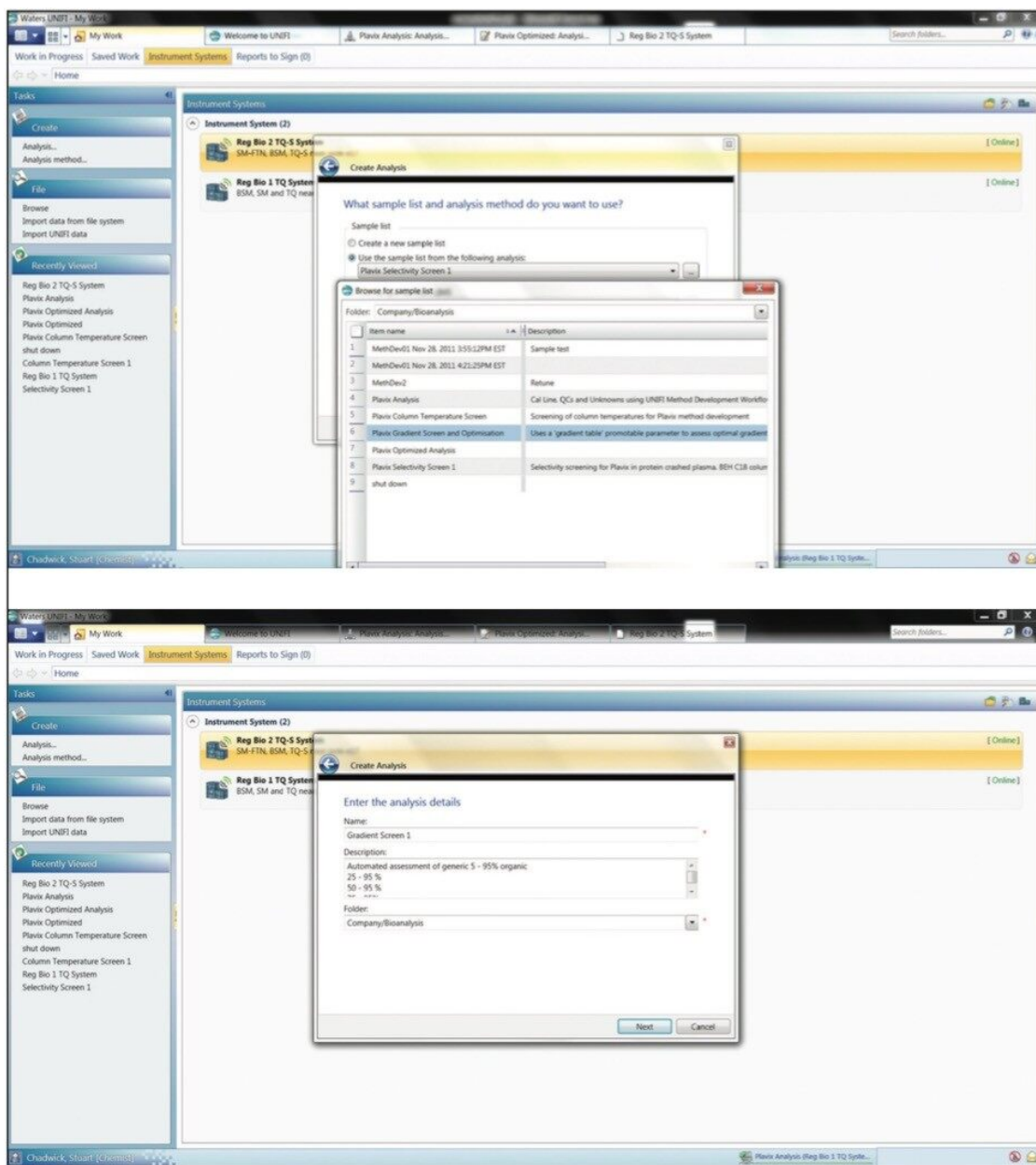


Figure 2. Selecting workflows.

After selection of the analysis method and sample list to be used, the user then saves the analysis to the appropriate folder. The scientist can then select the different gradient steepness and shapes to be evaluated via a drop-down menu, as shown in Figure 3. This prevents the propagation of numerous methods, by different scientists and ensures consistent operation.

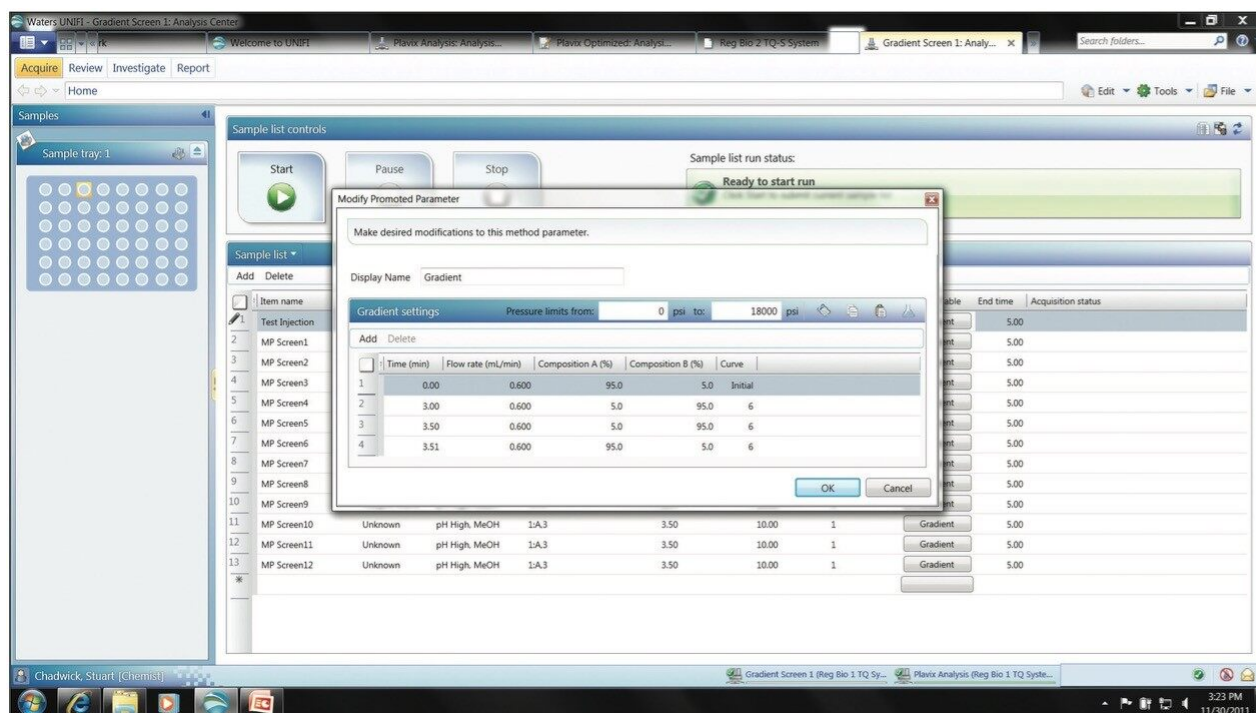


Figure 3. Selecting gradient steepness and shapes.

Once the various gradient slopes to be evaluated have been selected the acquisition can be started and the data acquired. Following data acquisition the data can be reviewed to select not only the system that will allow the greatest throughput, but also that which will deliver the greatest robustness.

In this case four different gradient slopes were evaluated: A 5-95%; B 5-50%; C 25-75%; and D 50-75%. The evaluation of the resolution and effectiveness of the different gradient shapes was performed by comparing the retention time of the target analytes with that of the endogenous components in the sample. The endogenous components were monitored by the simultaneous acquisition of full scan MS data and precursors of $m/z = 184$ to track the elution profile of the phospholipids. The data displayed in Figure 4 shows the elution profile of the target compound and the endogenous components in the sample

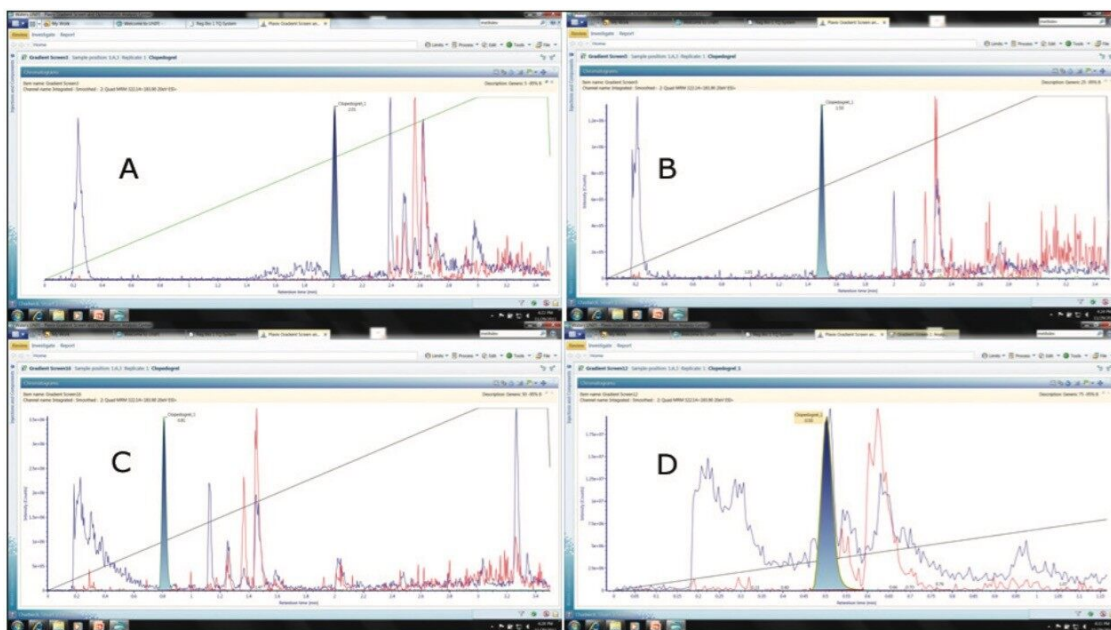


Figure 4. Elution profile.

We can see from this data that composition A gave clear resolution of the analytes from the endogenous components as did gradients B and C, with gradient C delivering the shortest analysis time of the three. Gradient D had the greatest throughput of the four, however the analyte peak eluted in the same region of the chromatogram as the endogenous components. Thus for this assay gradient C would be the best option, delivering the best compromise of throughput and resolution. By employing gradient C for the assay the compound elution time has been reduced from 2.01 minutes to 0.81 minutes allowing the LC condition to be modified to effectively double the throughput.

Chromatographic column temperature can be evaluated in a similar manner to that employed for gradient steepness and slope for method development. Using a similar template approach, the column temperature can be varied, Figure 5. In this case column temperatures of 40 °C, 50 °C, and 60 °C were evaluated. The results generated showed that, in this case, little benefit is gained in terms of peak shape or retention time when the column temperature is varied from 40 to 60 °C, Figure 6.

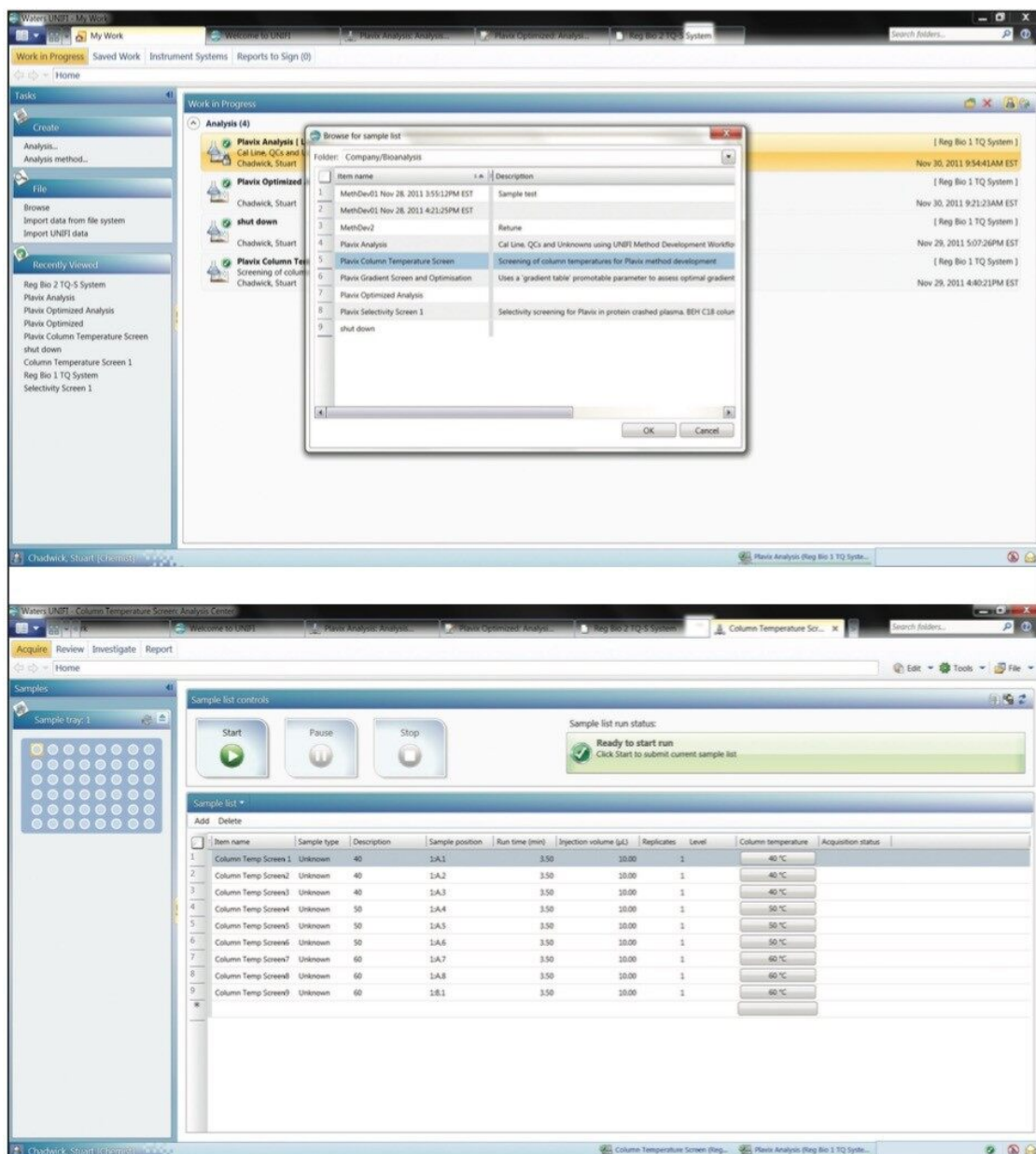


Figure 5. Evaluating varying column temperatures.

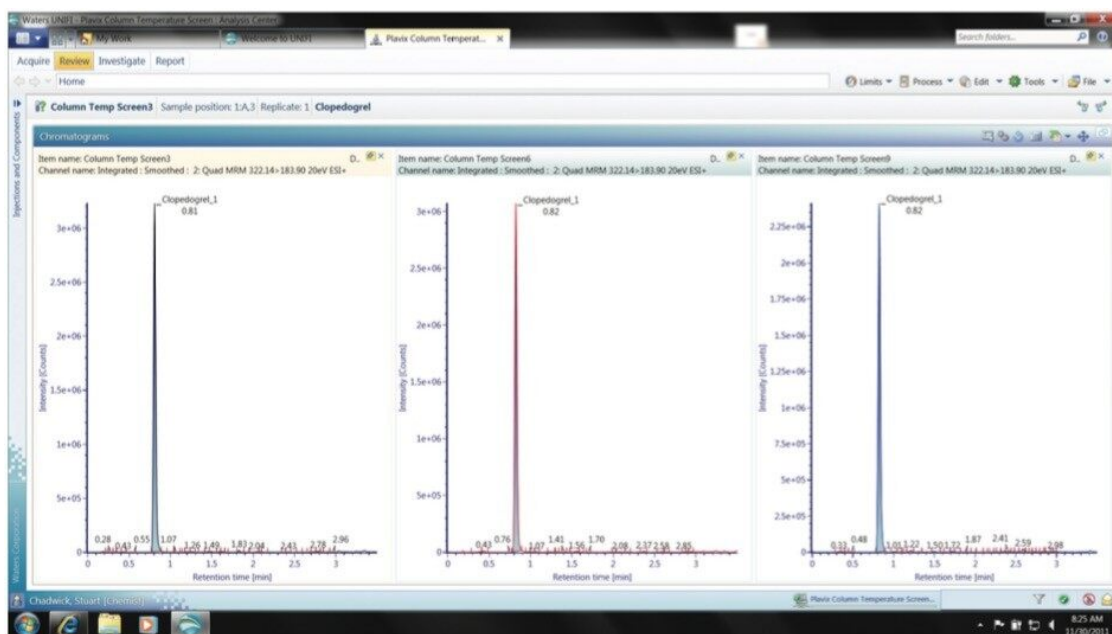


Figure 6. Effect of column temperature on separation of clopidogrel in plasma.

Conclusion

The Waters UNIFI Scientific Information System employs a workflow approach with acquisition templates and promotable parameters to enable the rapid simple screening and optimization of gradient and column temperature conditions. Further optimization of the chromatographic parameters can significantly improve assay robustness and analytical throughput. This workflow approach delivers:

- Streamlines data acquisition
- Promotes consistency of the method development process
- Allows expert users to define templates to be used by less experienced scientists
- Utilizes promotable parameters to allow multiple parameters such as gradient profile and column temperature to be evaluated using a pre-determined template

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UNIFI Scientific Information System <<https://www.waters.com/134801648>>

Regulated Bioanalysis Platform Solution with UNIFI <<https://www.waters.com/134613325>>

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