

The Development of a High Throughput Solid Phase Extraction Method for PFAS in Human Serum Using an Automated Janus® Workstation

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Introduction

Per- and Polyfluoroalkyl Substances (PFAS) are a large group of organic persistent pollutants that are of interest in human health studies. PFAS can be quantitated in human serum with ultra high-performance liquid chromatography-tandem mass spectrometry (UHPLC-MS/MS). Samples typically undergo solid phase extraction (SPE) before instrument analysis. The PFAS method used at the New Jersey Public Health Laboratories (NJPHL) employs an online automated sample micro-SPE prior to injection and chromatographic separation. Since these events occur sequentially, sample throughput is a limitation of this approach. We aim to develop and optimize this SPE process and automate the pre-sample preparation steps. This method will be validated for precision and accuracy. This new SPE method reduces the duration to prepare 48 samples from eight to two hours. Initial chromatographic peaks shapes were poor but was resolved by dissolving the sample in initial mobile phase conditions. Low analyte peak intensities were first observed because lower sample volumes were injected but increasing the injection volume from 40 µL to 100 µL fixed this problem. The new method produces acceptable calibration curves, the R² are between 0.991 to 0.998 for all analytes. The accuracy of matrix spiked samples is within ±30% of the nominal (accepted) value.

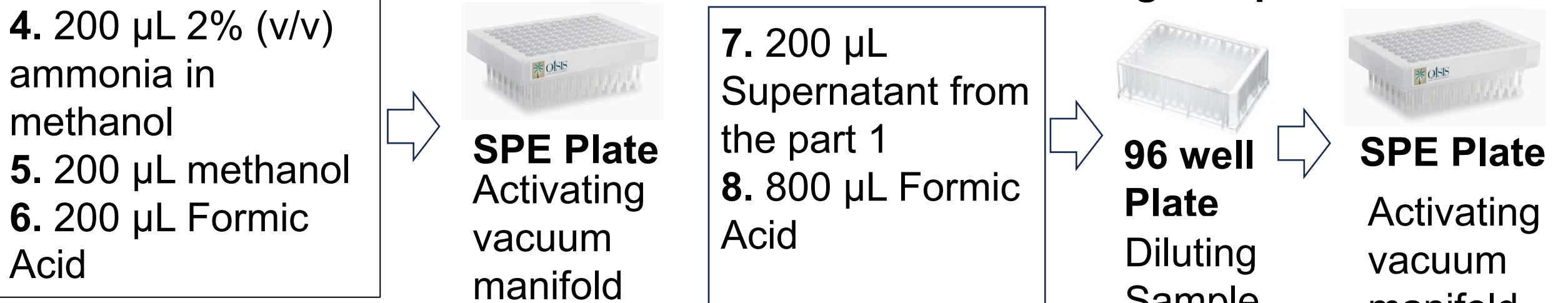
Method

A SPE protocol was created using “WinPREP® for Janus®” software on a Janus® workstation. The protocol is built by combining pipetting orders on the list provided by the software and run on the Janus® application assistant program. The protocol combines 50 µL human serum sample, 50 µL internal standard and 200 µL acetonitrile. This mixture is centrifuged at 4,000 RPM for 20 minutes. 200 µL of the supernatant is loaded on the SPE plate (Oasis® WAX) and washed with 200 µL 1% formic acid and 1:3 acetonitrile: water solution. A conditioning step is then performed with 200 µL 2% ammonia in methanol, 200 µL methanol, and 200 µL 1% formic acid. The analytes are eluted with 50 µL methanol and 100 µL 2% ammonia in methanol. The samples are dried at 70 °C and reconstituted with 120 µL an initial mobile phase solution. The sample is injected into Sciex 7500 UHPLC-MS/MS instrument with UHPLC Guard 3PK column/InfinityLab® PFC Delay column for LC-MS/MS analysis.

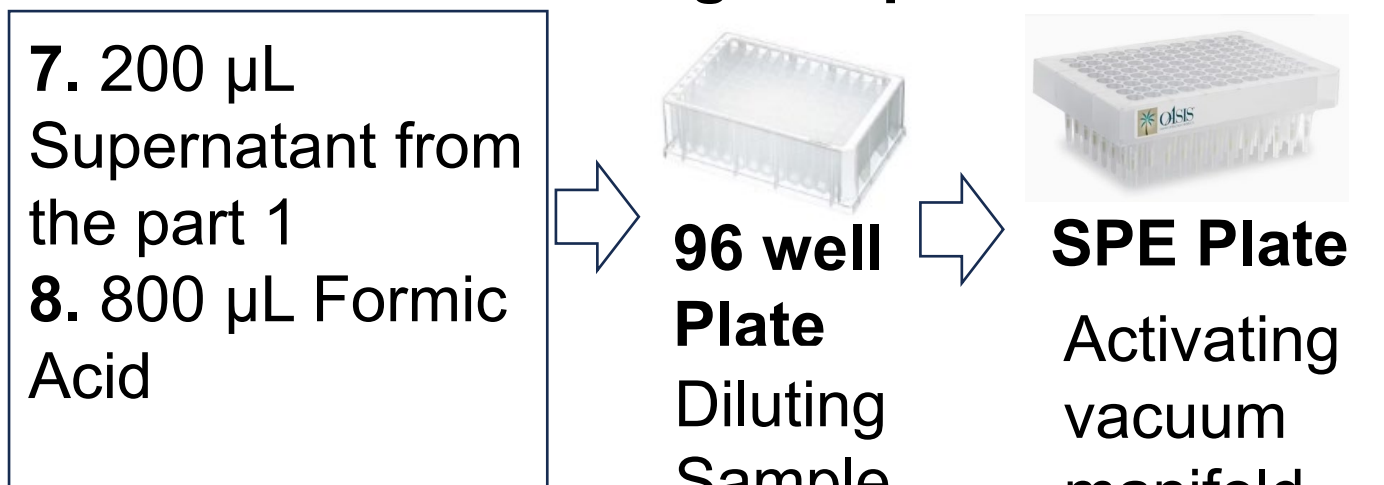
Protocol Part 1: Sample Pre-preparation



Protocol Part 2: Condition



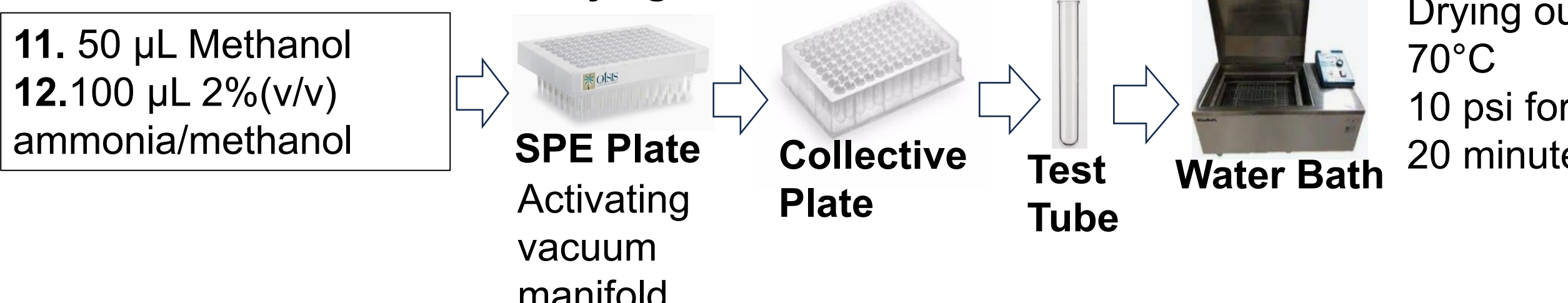
Protocol Part 3 Loading Sample



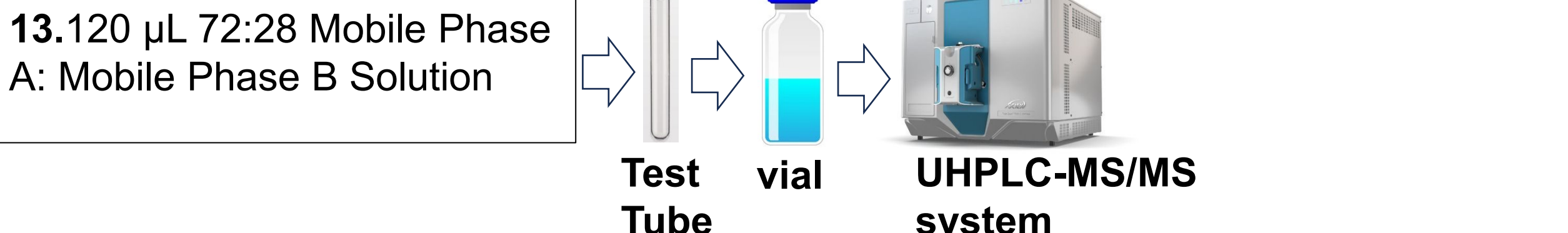
Protocol Part 4: Wash Plate



Protocol Part 5: Elution and Drying out



Protocol Part 6: Reconstitution



Results

Peak Shapes

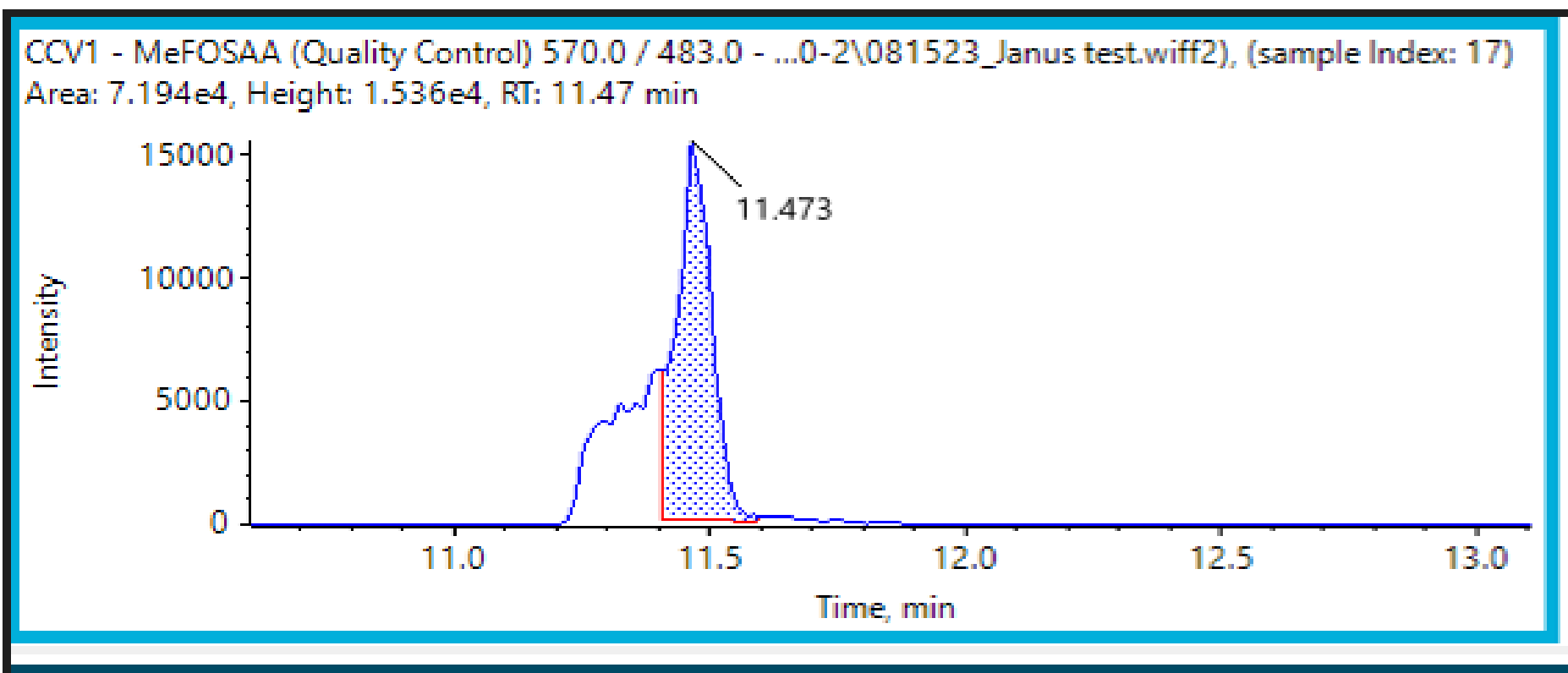


Figure 1: Representative peak: MeFOSAA without drying down and reconstitution procedure

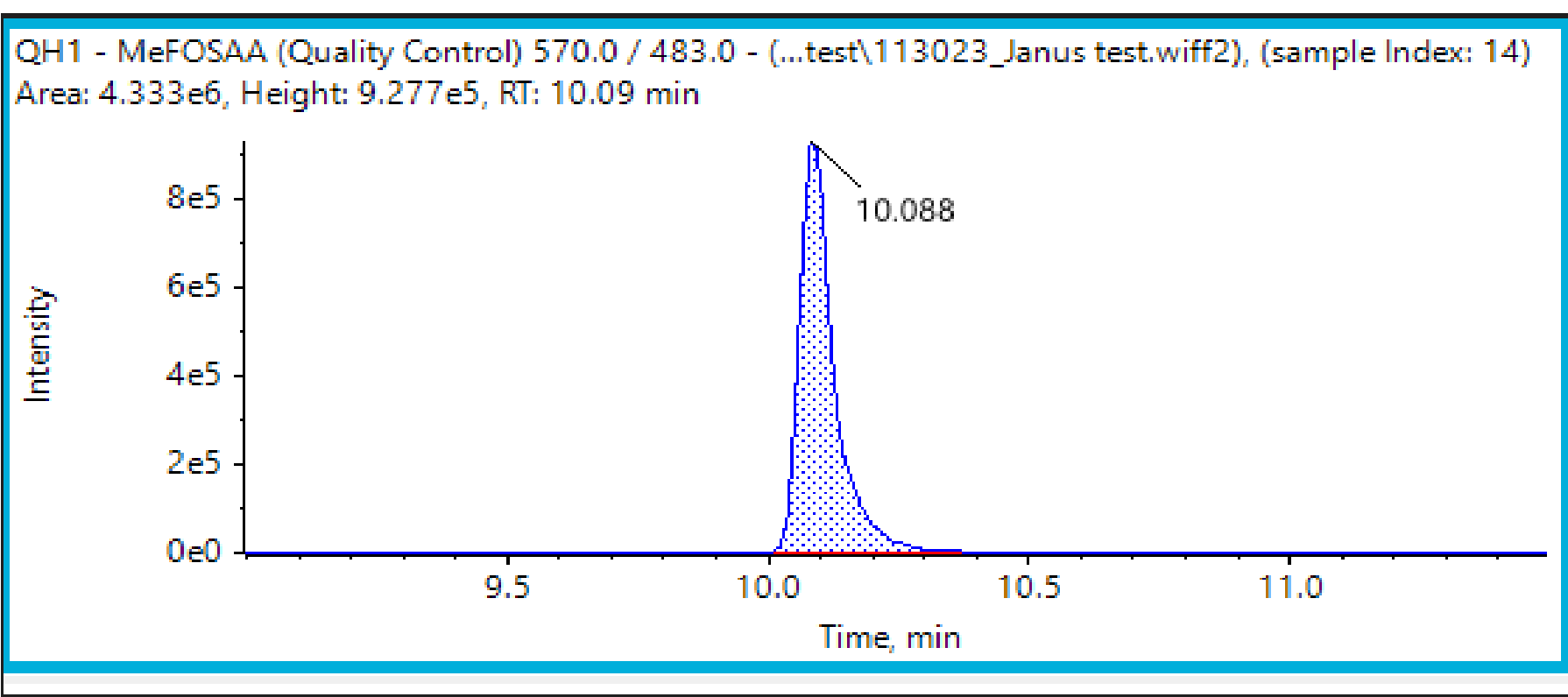


Figure 2: Representative peak: MeFOSAA with drying down and reconstitution procedure

Calibration Curve and Correlation Coefficient

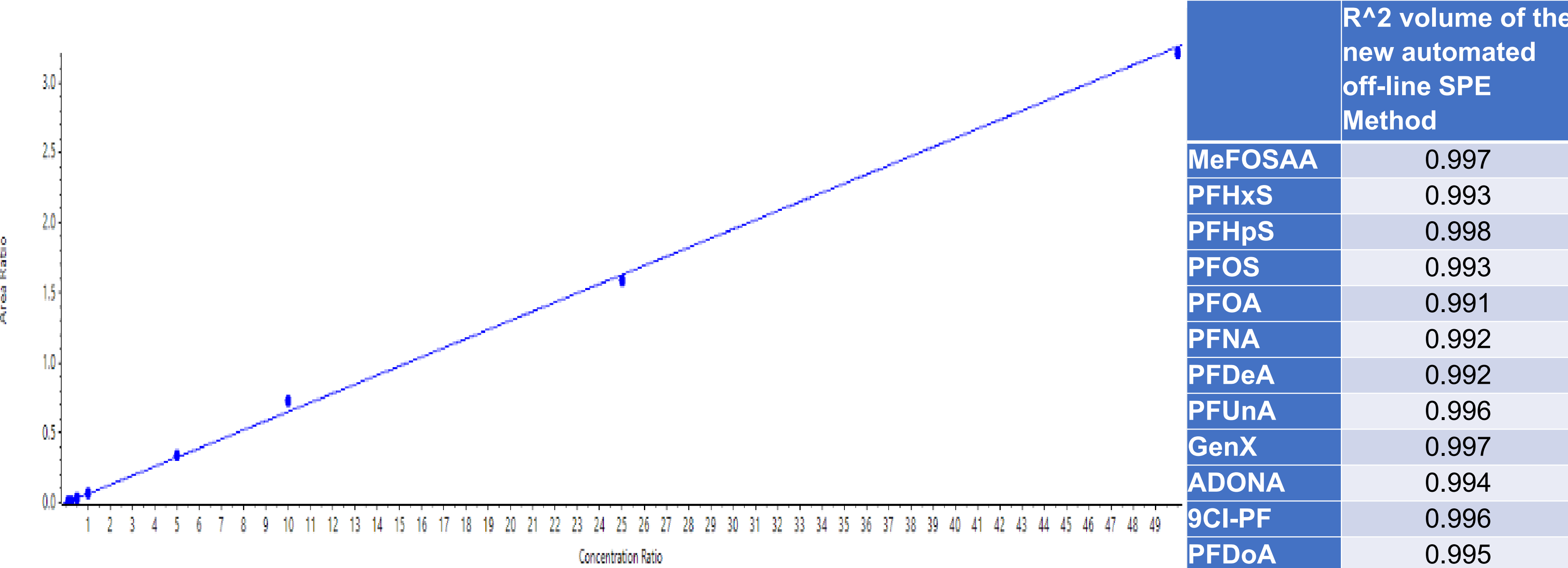


Figure 3: MeFOSAA as a representative 10 points calibration curve (Automated off-line SPE method)

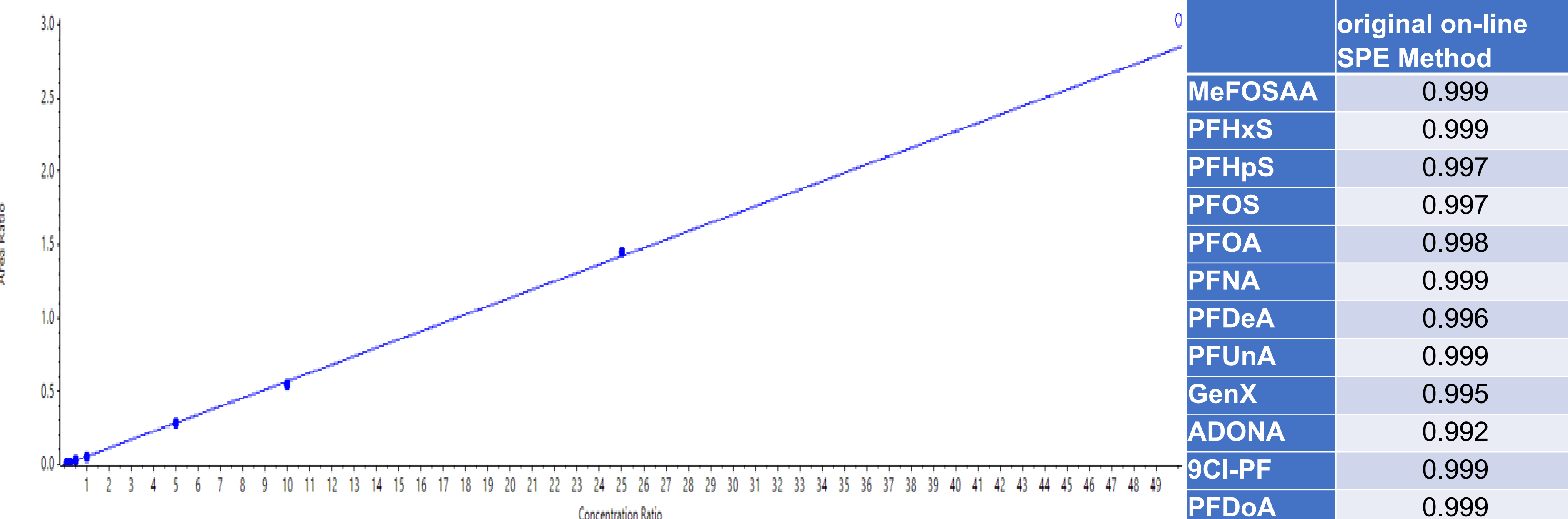


Figure 4: MeFOSAA as a representative 10 points calibration curve (on-line SPE method)

Recovery Rate

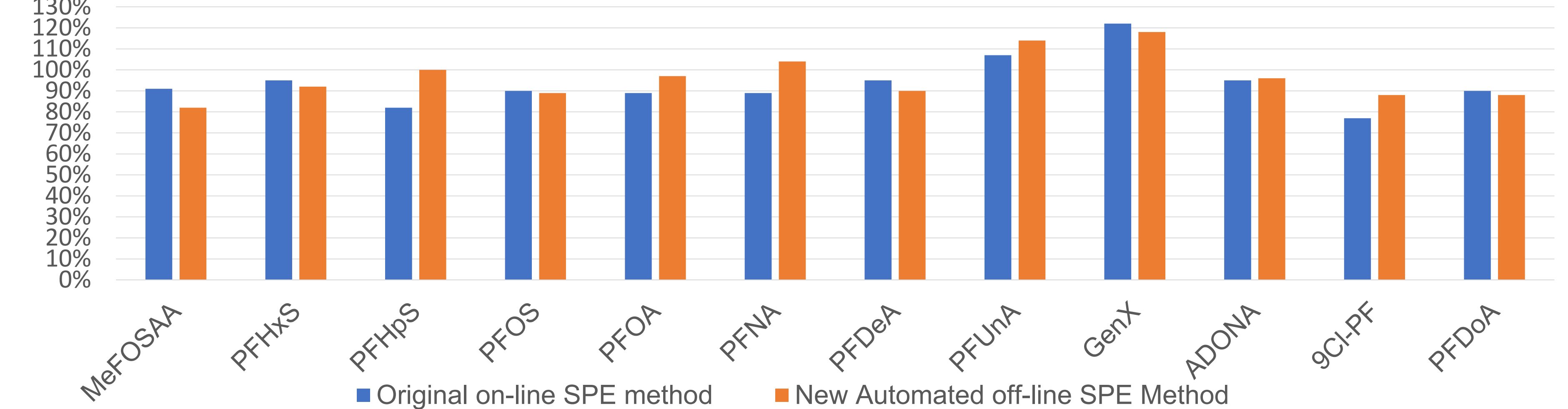


Figure 5. Recovery Rate of 14 analytes with the original online SPE method and the new automated off-line SPE method

Conclusions

1. A new Janus protocol has been created for the new automated off-line SPE method.
2. The poor peak shape issue in the UHPLC-MS/MS analysis is a major challenge of the new method development. Drying out and reconstituting the analytes before the injection of the UHPLC-MS/MS system can improve the peak shape significantly (See Figure 1 and Figure 2.)
3. The new protocol has already been tested on the Janus. This new SPE method reduces the time it takes to prepare 48 samples from eight to two hours.
4. The test results (See Figure 3) showed that the new off-line SPE method produces acceptable calibration curves, the R² are between 0.991 to 0.998 for all analytes as the original on-line SPE method does (See Figure 4.)
5. According to the test result (See Figure 5), the new automated SPE method has the same recovery rate as the original on-line SPE method had. Like the original online SPE method, the accuracy of the matrix spiked samples is within ±30% of the nominal value. This means the usage of the automated off-line method will not decrease the accuracy and precision of the sample preparation.
6. We will continue to improve and test the protocol for the further method validation.
7. This method will be used in our biomonitoring projects for PFAS after method validation.

References

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