# LC-MS/MS ANALYSIS OF IMMUNOSUPPRESSANT DRUGS IN WHOLE BLOOD USING THE XEVO TQ ABSOLUTE WITH CAPITAINER® B 50 DEVICES FOR CLINICAL RESEARCH

Analyte

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### **INTRODUCTION**

Traditional laboratory analysis of the immunosuppressant drugs cyclosporine, everolimus, sirolimus and tacrolimus is wellestablished in clinical research. However there remains a need for individuals to undergo an invasive, time-consuming and disruptive process under the supervision of trained staff in order to collect a sufficient volume of whole blood for laboratory analysis.

A reliable, remote sampling method may find utility in a clinical research setting. Here we describe the use of Capitainer® B 50 Devices to obtain analytically sensitive, precise and accurate data for cyclosporine, everolimus, sirolimus and tacrolimus analysis using small sample volumes for clinical research studies.

The Waters<sup>™</sup> ACQUITY<sup>™</sup> UPLC<sup>™</sup> I-Class FL with Xevo<sup>™</sup> TQ Absolute mass spectrometer was used to analyze these samples.

### **METHODS**

#### **Materials and Sample Preparation**

- MassTrak Immunosuppressant Calibrator and Control Sets were used.
- 30 μL of whole blood was pipetted onto the inlet of the Capitainer® B 50 device, which resulted in a 10 μL dried blood spot.
- Following overnight drying, the dried blood spot was removed and placed in a 2mL microcentrifuge tube.
- 200  $\mu$ L of internal standard (12.5 ng/mL <sup>2</sup>H<sub>12</sub>-cyclosporine, 1 ng/mL ascomycin, <sup>13</sup>C<sub>2</sub><sup>2</sup>H<sub>4</sub>-everolimus and <sup>2</sup>H<sub>3</sub>-sirolimus in 10% methanol) was added, and the tube underwent mixing and sonication steps.
- Add 10 μL of 0.05M hydrochloric acid and 1 mL *tert*-methyl butyl ether, vortex mix and centirfuge.
- 850 μL of the top layer was transferred to a clean, TruView Total Recovery vial (p/n: 186005669CV), and dried under nitrogen at 40°C.
- Samples were reconstituted in 200 μL mobile phase A:mobile phase B 50:50 (v:v).

	(m/z)	(m/z)	(s)	(V)	Energy (V)
Cyclosporine	1219.8	1202.8 (1184.8)	0.02	50	18 (34)
<sup>2</sup> H <sub>12</sub> -Cyclosporine	1231.8	1214.8	0.02	50	18
Everolimus	975.6	908.6 (926.6)	0.02	50	16 (12)
<sup>13</sup> C <sub>2</sub> <sup>2</sup> H <sub>4</sub> -Everolimus	981.6	914.6	0.02	50	16
Sirolimus	931.6	864.5 (882.5)	0.02	50	16 (12)
<sup>2</sup> H <sub>3</sub> -Sirolimus	934.6	864.5	0.02	50	16
Tacrolimus	821.5	768.5 (786.5)	8.5 (786.5)0.0250756.50.0250		20 (18)
Ascomycin	809.5	756.5			22

Daughter

Parent

Cone

Collision

Dwell

**Table 2.** MRM transitions and parameters of the immunosuppressants and internal standards(qualifier parameters in parentheses)



### RESULTS

Five analytical runs were performed using this method.

#### Chromatography and Analytical Sensitivity

• The analytical sensitivity of the developed method for the lowest calibrator (1 ng/mL for everolimus, sirolimus and tacrolimus; 25 ng/

#### Linearity and Precision

• Linearity of the calibration ranges (1-30 ng/mL for everolimus, sirolimus and tacrolimus; 25-1500 ng/mL for cyclosporine) was demonstrated with mean r<sup>2</sup> values for the calibration lines >0.99 over five analytical runs.

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• Total reproducibility and repeatability across the immunosuppressants at the QC three concentrations (2, 8 and 22 ng/mL for all analytes except cyclosporine, which were 150, 400 and 900 ng/mL), with five\* replicates over five analytical runs (n = 25\*) was ≤7.6%CV (Figure 3).

\*except cyclosporine, four runs and n=20



#### Accuracy

- LGC (Bury, UK) whole blood External Quality Assurance samples were sourced and analyzed to assess method accuracy. A summary is presented in Table 3.
- All individual samples met the criteria of the scheme, with mean bias ≤8.7%.

Analyte	Number of samples analyzed (n)	Range (ng/mL)	Mean %bias from Scheme LC-MS ALTM
Cyclosporine	20	31.0-1814.1	+6.0
Everolimus	20	0-23.2	+2.9
Sirolimus	25	1.9-22.9	-8.7
Tacrolimus	25	1.6-27.0	+5.6

 Table 3. EQA accuracy summary (note: ALTM is the all-laboratory trimmed mean)

CONCLUSION

#### **LC-MS/MS** Parameters

- Using an ACQUITY UPLC I-Class FL System, samples were injected onto an ACQUITY UPLC HSS C<sub>18</sub> SB Column, 1.8µm 2.1x30mm (p/n: 186004117), using a water/acetonitrile/ammonium fluoride gradient (Table 1) and analyzed with a Xevo TQ Absolute Mass Spectrometer (Figure 1) in ESI+, using MRM parameters described in Table 2.
- The run time is 1.5 minutes (approximately 2.2 minutes injection-to-injection).

Time [min]	Flow rate [mL/min]	A [%]	B [%]	Curve	Kagy
Initial	0.8	50	50	Initial	
0.2	0.8	50	50	1	
0.6	0.8	0	100	6	
1.2	0.8	50	50	11	

**Table 1.** LC gradient for analysis of the immunosuppressants using the ACQUITYUPLC I-Class FL System

mL for cyclosporine) is shown in Figure 2, with S/N (PtP) > 10 for the four analytes over the five runs.



*Figure 2.* Representative chromatograms of the lowest immunosuppressant calibrator, demonstrating analytical sensitivity

- Capitainer B 50 devices prepared dried blood spots from a low initial whole blood volume, enabling simultaneous analysis of the immunosuppressant drugs cyclosporine, everolimus, sirolimus and tacrolimus in 2.2 minutes (injection-to-injection).
- The performance characteristics of the method indicate good analytical sensitivity, total precision and repeatability (≤7.6%CV) across all analytes and concentrations tested.
- Finally, good agreement was obtained when analyzing External Quality Assurance samples, providing confidence in the accuracy of the method and sample collection devices.

## ACKNOWLEDGEMENT

• Capitainer are thanked for the provision of Capitainer B 50 devices for this study.

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