Validation of Cyclosporin on LC-QTOF

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| 1 nave | reviewed | tms data | a and approv | ve tms tes | st for clinic | ar use. | |
|--------|------------|----------|--------------|------------|---------------|---------|--|
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| | | | | | | | |
| | | | | | | | |
| Labora | atory Dire | ector | | | | | |

Note: all CV and bias are in percent

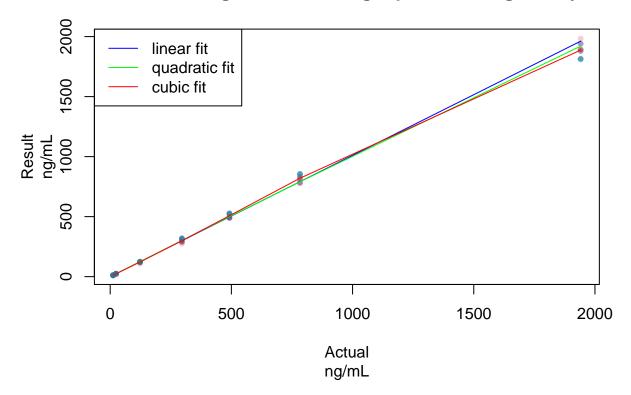
Reportable Range Study

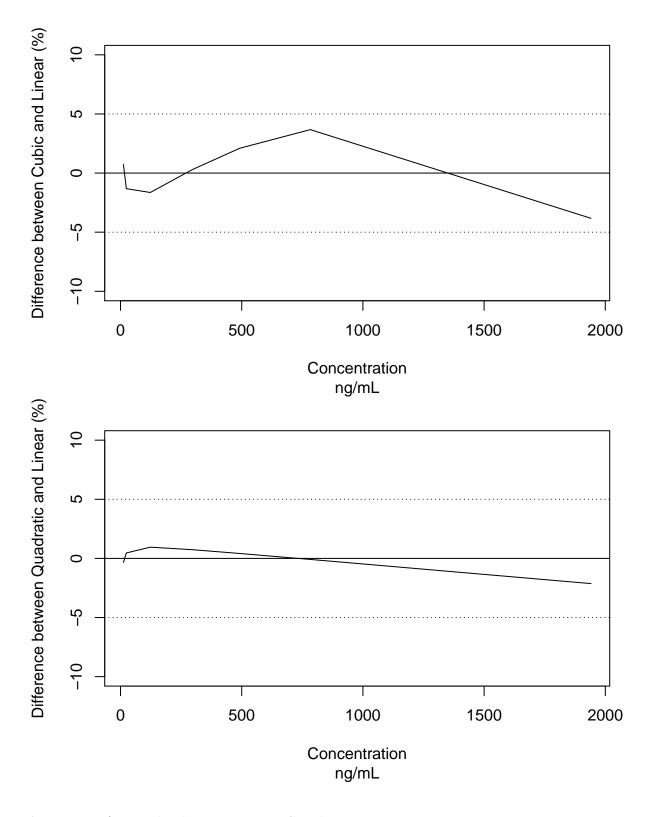
COM.40600

The reportable range study was performed by running standards over 6 days. Analysis was performed by both calculating the %CV at each level and ensuring that it is less than 20% and by using the method of Emancipator and Kroll to evaluate for linearity over the calibration range. A tolerance of 10% for non-linearity was used as the criteria and comparisons were made with both quadratic vs linear and cubic vs quadratic. The assay is validated for the range of 12 - 1941 ng/mL.

| Actual.cals | Mean.cals | SD.cals | CV.cals |
|-------------|-----------|---------|---------|
| 12 | 11.6 | 1.4 | 11.9 |
| 24 | 23.9 | 0.9 | 3.8 |
| 123 | 119.7 | 6.2 | 5.2 |
| 296 | 302.0 | 17.8 | 5.9 |
| 492 | 507.3 | 18.9 | 3.7 |
| 783 | 819.2 | 34.1 | 4.2 |
| 1941 | 1887.5 | 68.0 | 3.6 |

Plot of data generated during reportable range study





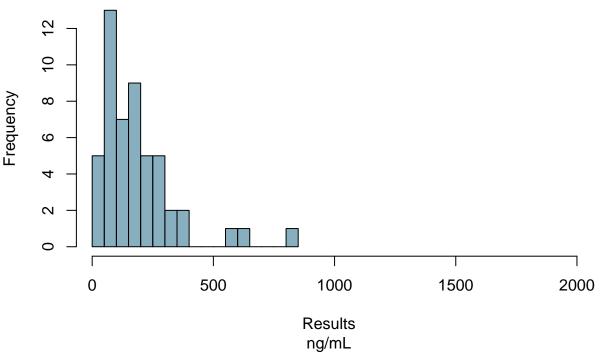
Accuracy & Method comparison Study

 $COM.40300,\ COM.40350,\ CHM.14200\ \&\ CHM.18800$

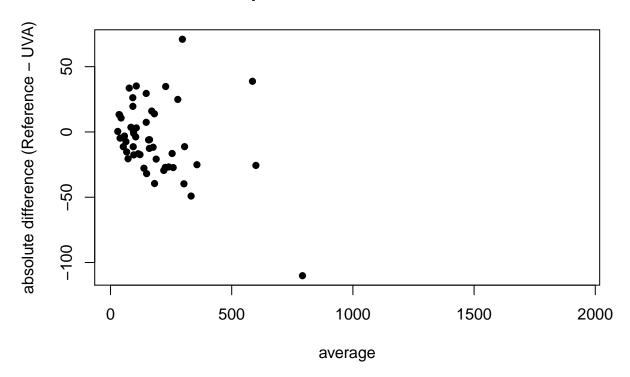
This study consisted of analyzing 51 quantitatively positive patient samples. The UVA method utilizes

LC-MS/MS while the comparator method is LC-MS/MS performed by previous UVA method. The sections that fall within this section include qualitative comparison and accuracy spike.

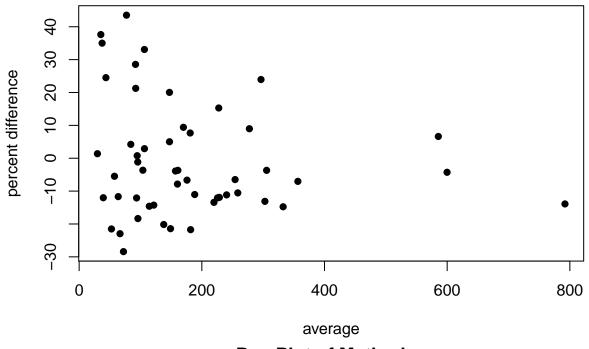
Distribution of Results



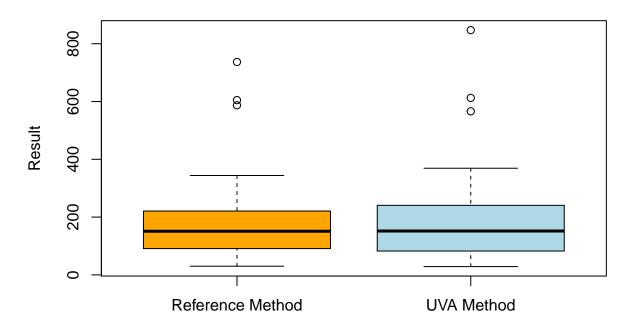
Method Comparison – Absolute Difference Plot



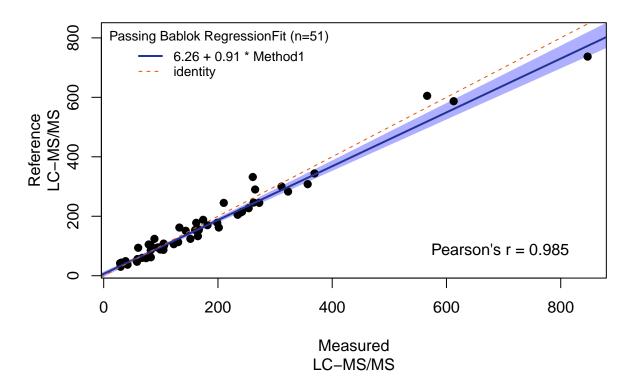
Method Comparison – Percent Difference Plot



Box Plot of Methods



Passing Bablock



Qualitative Comparison

This data compares values where one of the methods reports a value outside of the AMR, since quantitative comparison is not possible. The goal is to evaluate if the interpretation of the samples are in agreement. At least 82 samples were analyzed and were in qualitative agreement by both the reference and UVA method. The chart below shows any discrepancies.

| method.comp.dis.Reference | method.comp.dis.Measured |
|---------------------------|--------------------------|
| | |

Accuracy Spike

This experiment evaluates the accuracy of the assay by spiking a known amount of analyte into at least 10 different patient samples.

| Specimen | Bias | Calculated | Measured |
|------------------|-------|------------|----------|
| Spike Patient 1 | -12.8 | 40 | 45.1 |
| Spike Patient 2 | -5.5 | 40 | 42.2 |
| Spike Patient 3 | -8.0 | 200 | 216.1 |
| Spike Patient 4 | -4.5 | 200 | 209 |
| Spike Patient 5 | 1.8 | 500 | 490.8 |
| Spike Patient 6 | -2.0 | 500 | 510.2 |
| Spike Patient 7 | -10.2 | 1000 | 1102.5 |
| Spike Patient 8 | -3.8 | 1000 | 1038.5 |
| Spike Patient 9 | -9.4 | 1600 | 1750 |
| Spike Patient 10 | -6.2 | 1600 | 1698.4 |
| Average | -6.7 | | |

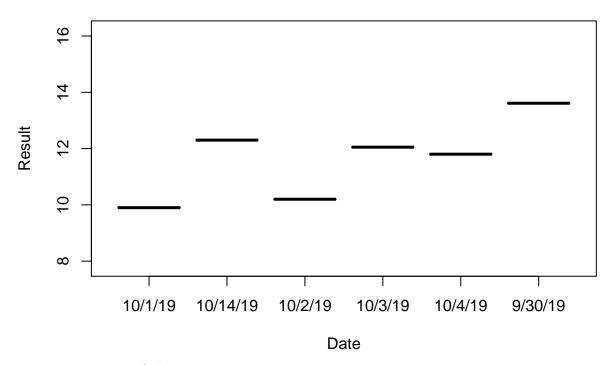
Imprecision Study

COM.40300

Intra-day and inter-day precision was determined at multiple concentrations, and is expressed as the coefficient of variation. The acceptability criteria is an imprecision < 15%.

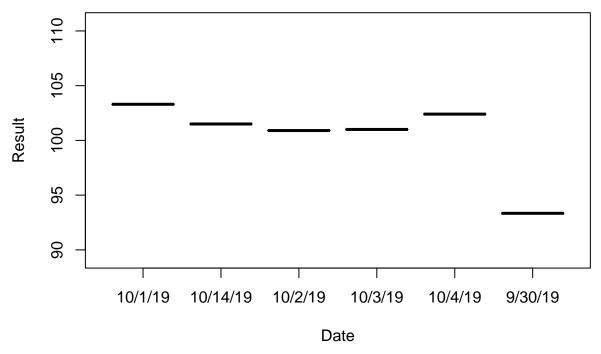
QC Data over multiple days:

Levey-Jennings Chart for Lower Limit of Quantitation



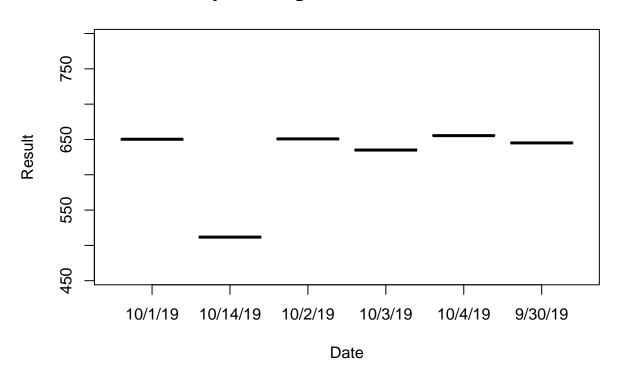
target mean = 12 ng/mL calculated mean = 12 ng/mL SD = 1.4 ng/mL CV = 11.9%Bias = 0 %

Levey-Jennings Chart for Low QC Level



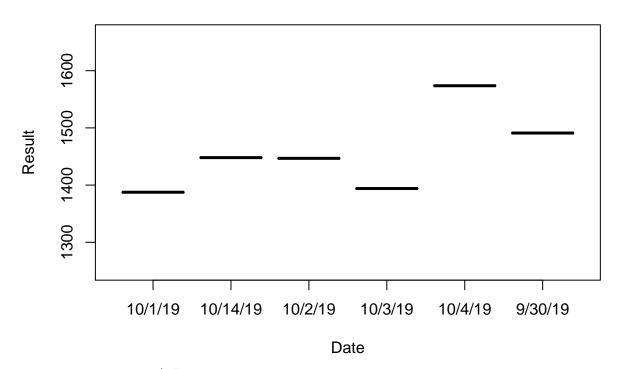
target mean = $105~\rm ng/mL$ calculated mean = $100~\rm ng/mL$ SD = $3.6~\rm ng/mL$ CV = 3.6%Bias = 4.8~%

Levey-Jennings Chart for Mid QC Level



target mean = 550 ng/mLmean = 625 ng/mLSD = 55.8 ng/mLCV = 8.9%Bias = -13.6 %

Levey-Jennings Chart for High QC Level



target mean = 1400 ng/mLcalculated mean = 1457 ng/mLSD = 68.9 ng/mLCV = 4.7%Bias = -13.6 %

QC Data within-run imprecision:

| Actual | Mean | CV | Bias |
|--------|--------|------|--------|
| 12 | 10.0 | 7.00 | 16.70 |
| 105 | 103.0 | 1.19 | 1.90 |
| 550 | 640.0 | 1.35 | -16.36 |
| 1400 | 1368.6 | 2.03 | 2.24 |
| | | | |

Matrix Effect Assessment

CHM.18825

This was performed by spiking the same concentration of drug into 10 separate patient samples. We compare either the peak area or the relative response (RR) for each sample back to a sample where the concentration of

drug was spiked into solvent and generate a ratio. Acceptability includes matrix suppression or enhancement of <25% OR a CV of the effect <15% across all samples.

| Specimen | Response | ratio |
|-----------------|----------|-------|
| Solvent spike | 1876698 | 0.95 |
| Patient #1 | 1858085 | 0.94 |
| Patient #2 | 1900218 | 0.96 |
| Patient #3 | 1828758 | 0.93 |
| Patient #4 | 1881936 | 0.95 |
| Patient #5 | 1939030 | 0.98 |
| Patient #6 | 1927748 | 0.98 |
| Patient #7 | 2000672 | 1.01 |
| Patient #8 | 1863361 | 0.94 |
| Patient #9 | 2056677 | 1.04 |
| Patient #10 | 1976174 | |
| Average | 1913318 | 0.97 |
| % Matrix Effect | | 3.2 |
| %CV | 3.66 | |

Autosampler Stability

Acceptability was determined by preparing and running samples, allowing them to sit on the autosampler for 72 hours, and then repeating the injections. Data is acceptable.

| Specimen. Type | First.run | Repeat.run | Bias |
|------------------------|-----------|------------|-------|
| std 1 | 13.6 | 14.3 | 4.8 |
| std 2 | 25.5 | 24.5 | -3.9 |
| std 3 | 107.9 | 110.4 | 2.3 |
| $\operatorname{std} 4$ | 275.5 | 278.3 | 1.0 |
| std 5 | 487.7 | 476.2 | -2.4 |
| std 6 | 777.4 | 738.8 | -5.2 |
| std 7 | 1983.6 | 2028.6 | 2.2 |
| QC 1 | 93.3 | 93.6 | 0.3 |
| QC 2 | 645.1 | 599.0 | -7.7 |
| QC 3 | 1490.6 | 1496.0 | 0.4 |
| A | 78.8 | 75.3 | -4.6 |
| В | 272.3 | 267.5 | -1.8 |
| $^{\mathrm{C}}$ | 143.6 | 139.8 | -2.7 |
| D | 242.2 | 244.1 | 0.8 |
| \mathbf{E} | 58.3 | 55.4 | -5.3 |
| \mathbf{F} | 612.6 | 618.1 | 0.9 |
| G | 82.5 | 76.2 | -8.3 |
| Н | 104.6 | 100.6 | -4.0 |
| I | 181.7 | 174.0 | -4.4 |
| J | 30.9 | 27.7 | -11.5 |
| K | 96.1 | 84.6 | -13.6 |
| K | 311.3 | 310.6 | -0.2 |
| M | 163.9 | 159.7 | -2.6 |
| N | 99.3 | 94.0 | -5.7 |
| O | 41.7 | 40.3 | -3.5 |
| P | 130.4 | 127.4 | -2.3 |
| Q | 164.9 | 166.3 | 0.8 |
| R | 82.4 | 86.8 | 5.0 |
| S | 322.7 | 310.6 | -3.9 |
| ${ m T}$ | 105.8 | 98.3 | -7.6 |
| U | 67.4 | 64.9 | -3.9 |
| V | 94.3 | 94.6 | 0.3 |
| W | 29.6 | 29.3 | -1.0 |
| X | 59.2 | 56.7 | -4.3 |

Dilutional Linearity

 ${
m QC3}$ was diluted 2, 5 and 10-fold to evaluate quantitation once diluted. Data is acceptable.

| Specimen.Type | Dilution.Factor | Measured.Results | Bias |
|---------------|-----------------|------------------|-------|
| blood | 2 | 1454.6 | -0.5 |
| blood | 2 | 1443.2 | 0.3 |
| blood | 2 | 1453.8 | -0.5 |
| blood | 5 | 1433.0 | 1.0 |
| blood | 5 | 1470.0 | -1.6 |
| blood | 5 | 1494.5 | -3.3 |
| blood | 10 | 1607.0 | -11.1 |
| blood | 10 | 1642.0 | -13.5 |
| blood | 10 | 1634.0 | -12.9 |

Analytical Specificity & Analytical Interferences

COM.40450, COM.40500, GEN.42030

We determined analytical specificity through by comparing our assay with an assay that is clinically validated, as well as comparing patient samples with spiked standard material. We ran specimens by both methods and showed that when they identified the analyte, we also identified and accurately quantitated the same analyte by our method. To further minimize the potential of interferences, we monitor multiple ions for each drug and use ion ratios in our criteria for identifying the presence of a drug.

Carryover

A urine blank is included after the calibration curve to assess for any carryover up until the highest point of the AMR and is repeated over multiple days. The acceptability criteria is a relative response <50% of the relative response of the LLOQ. No carryover was detected up to 1941 ng/mL.