RESEARCH NOTE

Evaluation of electrochemiluminescence immunoassays for immunosuppressive drugs on the Roche cobas e411 analyzer [version 1; peer review: 2 approved]

Angela W.S. Fung1,2*, Michael J. Knauer2,3*, Ivan M. Blasutig2,4, David A. Colantonio2,5, Vathany Kulasingam2,6

1Department of Pathology and Laboratory Medicine, St. Paul’s Hospital, Vancouver, BC, V6Z 1Y6, Canada
2Department of Laboratory Medicine and Pathobiology, University of Toronto, Toronto, ON, M5S 1A8, Canada
3Lifelabs Medical Laboratories, Toronto, ON, M9W 6J6, Canada
4Division of Biochemistry, Children’s Hospital of Eastern Ontario, Ottawa, ON, K1H 8L1, Canada
5Department of Pediatric Laboratory Medicine, The Hospital for Sick Children, Toronto, ON, M5G 1X8, Canada
6Department of Clinical Biochemistry, University Health Network, Toronto, ON, M5G 2C4, Canada

* Equal contributors

Abstract

Background: Therapeutic drug monitoring of immunosuppressant drugs are used to monitor drug efficacy and toxicity and to prevent organ transplantation rejection. This study evaluates the analytical performance of semi-automated electrochemiluminescence immunoassays (ECLIA) for cyclosporine (CSA), tacrolimus (TAC) and sirolimus (SRL) on the Roche cobas e 411 analyzer at a major transplant hospital to identify method suitability and limitations.

Methods: Residual whole blood samples from patients undergoing immunosuppressant therapy were used for evaluation. Experiments included imprecision, linearity, functional sensitivity, method comparisons and lot-to-lot assessments.

Results: Total imprecision ranged from 3.3 to 7.1% for CSA, 3.9 to 9.4% for TAC, and 4.6 to 8.2% for SRL. Linearity was verified from 30.0 to 960.9 μg/L for CSA, from 1.1 to 27.1 μg/L for TAC, and from 0.5 to 32.3 µg/L for SRL. The functional sensitivity met the manufacturer’s claims and was determined to be <6.5 μg/L for CSA, 1.1 μg/L for TAC, and <0.1 μg/L for SRL (CV≤20%). Deming regression analysis of method comparisons with the ARCHITECT immunoassay yielded slopes of 0.917 (95%CI: 0.885-0.949) and r of 0.981 for CSA, 0.938 (95%CI: 0.895-0.981) and r of 0.974 for TAC, and 0.842 (0.810-1.110) and r of 0.982 for SRL. Deming regression analysis of comparisons with the LC–MS/MS method yielded slopes of 1.331 (95%CI: 1.167-1.496) and r of 0.969 for CSA, 0.924 (95%CI: 0.843-1.005) and r of 0.984 for TAC, and 0.971 (95%CI: 0.913-1.030) and r of 0.993 for SRL.

Conclusions: The cobas e 411 ECLIA for CSA, TAC, and SRL have acceptable precision, linearity, and functional sensitivity. The method...
comparisons correlated well with the ARCHITECT immunoassay and LC–MS/MS and is fit for therapeutic drug monitoring.

**Keywords**
Cyclosporine, tacrolimus, sirolimus, immunoassay, ECLIA
Introduction

Immunosuppressive drugs (ISD), such as the calcineurin inhibitors (cyclosporine (CSA) and tacrolimus (TAC)) and mammalian target of rapamycin (mTOR) inhibitors (sirolimus (SRL) and everolimus), are critical to the maintenance of solid organ transplantation. CSA is a cyclic decapeptide and TAC (also known as FK-506) is a macroline lactone. CSA binds to cyclophilin A/B and inhibits calcineurin. TAC binds to FK506-binding protein 12 (FKBP-12) to form the calcineurin inhibitory complex. Inhibition of calcineurin, a serine/threonine phosphatase, leads to altered calcium-dependent signal transduction, and decreases T-cell activation and downregulates anti-inflammatory response-related genes. SRL (also known as rapamycin) is a 31-membered macroline antibiotic that binds to FKBP-12 and allosterically targets the mTOR pathway, inhibiting cell cycle progression, T-cell proliferation and differentiation. SRL has structural similarities to TAC and competes with TAC for FKBP-12 binding.

All three ISDs are characterized by having variable absorption, similarities to TAC and competes with TAC for FKBP-12 binding. CSA, TAC, and SRL on the Roche cobas e 411 analyzer and calibrated five times during the evaluation by a 2-point calibration method on Abbott ARCHITECT i2000 analyzer. For TAC, 200 µL of sample was mixed with 200 µL of Cyclosporin Solubilizing Reagent (4% saponin) and 400 µL of Cyclosporine Precipitation Reagent (zinc sulfate in methanol and ethylene glycol) was added to 200 µL of sample. For TAC, 200 µL of sample was mixed with 200 µL of Tacrolimus Precipitation Reagent (zinc sulfate in methanol) and vortexed. For SRL, 150 µL of sample was mixed with 300 µL of Sirolimus Precipitation Reagent (zinc sulfate in >50% v/v DMSO and ethylene glycol), vortexed and heated at 42°C for 10 minutes. All ISD samples were then vortexed for 10 seconds and centrifuged for 4 minutes at 15,000 x g, and the supernatant was decanted for analysis. Analysis was performed within 30 minutes of preparation to prevent evaporation of the extracted samples. The ECLA assays were calibrated five times during the evaluation by a 2-point calibration using calibrators traceable to pure standard materials reconstituted in whole blood matrix by gravimetric methods.

Chemiluminescent microparticle immunoassay (CMIA) method on Abbott ARCHITECT i2000

The ARCHITECT CMIA (Abbott Laboratories, Abbott Park, IL, USA) for CSA, TAC, and SRL are based on the competition of analyte in sample with a ruthenium-labeled analogue. A voltage is applied and electrochemiluminescence signal is detected. Testing was performed according to the manufacturer’s instructions. Briefly, the samples (calibrators, QC, whole blood samples) were manually pretreated by combining 300 µL of whole blood with 300 µL of Universal ISD Sample Pretreatment Reagent (containing zinc sulfate and methanol) and vortexed for 10 seconds to lyse the red blood cells, precipitate proteins and extract the analyte. The samples were centrifuged for 4 minutes at 15,000 x g, and the supernatant was decanted for analysis. Analysis was performed within 30 minutes of preparation to prevent evaporation of the extracted samples. The ECLA assays were calibrated five times during the evaluation by a 2-point calibration using calibrators traceable to pure standard materials reconstituted in whole blood matrix by gravimetric methods.

Methods

Specimen source and handling

Ethical approval for this study was waived by the Research Ethics Board at the University Health Network in Toronto, Ontario, Canada (16-6312) for use of routine collected specimens for the evaluation of method performance. Residual EDTA whole blood specimens from 300 patients undergoing immunosuppressant therapy (either cyclosporine, tacrolimus, or sirolimus) at the University Health Network, and CAP proficiency testing samples were used in this evaluation. Samples were collected and either analyzed within the same day or stored as per manufacturer recommendations. Samples were thawed and equilibrated to room temperature for 30 minutes and mixed well prior to analysis.

In accordance with stability studies on whole blood ISD specimens, samples were analyzed within three months of collection and did not undergo more than two freeze-thaw cycles.

Electrochemiluminescence immunoassay (ECLIA) method on Roche cobas e 411

The cobas ECLIAs (Roche Diagnostics GmbH, Mannheim, Germany) for CSA, TAC, and SRL are based on the competition of analyte in sample with a ruthenium-labeled analogue. A voltage is applied and electrochemiluminescence signal is detected. Testing was performed according to the manufacturer’s instructions. Briefly, the samples (calibrators, QC, whole blood samples) were manually pretreated by combining 300 µL of whole blood with 300 µL of Universal ISD Sample Pretreatment Reagent (containing zinc sulfate and methanol) and vortexed for 10 seconds to lyse the red blood cells, precipitate proteins and extract the analyte. The samples were centrifuged for 4 minutes at 15,000 x g, and the supernatant was decanted for analysis. Analysis was performed within 30 minutes of preparation to prevent evaporation of the extracted samples. The ECLA assays were calibrated five times during the evaluation by a 2-point calibration using calibrators traceable to pure standard materials reconstituted in whole blood matrix by gravimetric methods.

Chemiluminescent microparticle immunoassay (CMIA) method on Abbott ARCHITECT i2000

The ARCHITECT CMIA (Abbott Laboratories, Abbott Park, IL, USA) for CSA, TAC, and SRL are based on the competition of analyte in sample with acridinium-labeled analogue. The samples were manually pretreated according to the manufacturer’s instructions and site-specific standard operating procedures. For CSA, 100 µL of Cyclosporine Solubilizing Reagent (4% saponin) and 400 µL of Cyclosporine Precipitation Reagent (zinc sulfate in methanol and ethylene glycol) was added to 200 µL of sample. For TAC, 200 µL of sample was mixed with 200 µL of Tacrolimus Precipitation Reagent (zinc sulfate in methanol) and vortexed. For SRL, 150 µL of sample was mixed with 300 µL of Sirolimus Precipitation Reagent (zinc sulfate in >50% v/v DMSO and ethylene glycol), vortexed and heated at 42°C for 10 minutes.

All ISD samples were then vortexed for 10 seconds and centrifuged for 4 minutes at 15,000 x g. The supernatants were decanted into labelled tubes and assayed within 30 minutes of sample preparation. The ARCHITECT CMIA are calibrated according to the site-specific standard operation procedures, with a 6-point 4-parameter logistic curve fit (4PLC, y-weighted) that is traceable to pure standard materials in a whole blood matrix by gravimetric methods. Internal QC was evaluated with Bio-Rad Lyphochek Whole Blood ISD Controls levels 1, 3, and 4.
Imprecision
Three levels of manufacturer multi-analyte QC materials (Roche Diagnostics PreciControl ISD levels 1, 2, and 3) and third-party multi-analyte QC materials (Bio-Rad Lyphochek Whole Blood ISD Controls levels 1, 3, and 4) were analyzed. QC samples were prepared and measured in duplicate, one run per day over 10 days. The acceptance criterion for total imprecision was based on the recommendation of the International Association of Therapeutic Drug Monitoring and Clinical Toxicology (IATDMCT) expert consensus group of ≤10%.

Functional sensitivity
Residual patient sample with levels 2–3 times the claimed limit of quantification (LoQ) was used to generate a series of dilutions with blank whole blood. The neat sample and dilutions were measured in triplicates within one day. The precision profile curve was used to calculate the LoQ concentration corresponding to a CV of 20% with the upper 95% confidence limit.

Linearity
Since there is a lack of elevated CSA and TAC patient specimen, CSA and TAC linearity were assessed using CAP EQA linearity materials (6 concentrations measured in duplicate). SRL linearity was assessed using a patient sample above the upper measuring range diluted with blank whole blood to 6 concentrations and measured in duplicate. The acceptance criterion was defined as slope of 1.00 ± 0.05 and deviation <10%.

Method comparison
Method comparison experiments were assessed where anonymized residual patient samples spanning the analytical measuring range for each analyte were measured one replicate per method. CSA samples concentrations ranged from 41.0 to 1808.0 µg/L, TAC ranged from 2.1 to 30.0 µg/L, and SRL ranged from 1.8 to 34.6 µg/L, as determined by ARCHITECT CMIA. Roche ECLIA measurements were compared to ARCHITECT CMIA (n=100). To further elucidate the accuracy between immunoassays, a subset of samples was also analyzed by LC-MS/MS (n=20) (Supplementary File 1). Lot-to-lot assessment was also performed between two lots of reagents for each ISD (n=20). The slope, intercept, correlation coefficient r were calculated by Deming regression analysis. The acceptance criteria for method comparison were defined as a slope of 1.00 ± 0.15 and deviation <10%.

Statistical analysis
Microsoft Excel (version 1708, Microsoft Office) and/or EP Evaluator (version 7.0.0.307, Data Innovations) were used for statistical analysis.

Results and discussion
To assess imprecision, three levels of manufacturer (Roche PreciControl) and third-party (Bio-Rad Lyphochek) multi-analyte QC materials were analyzed using one lot of reagents in duplicate, one run per day over 10 days (Table 1). For the PreciControl, the total imprecision was <7.1% for CSA, <9.4% for TAC, and <5.6% for SRL. Imprecision for CSA and TAC were comparable to other studies. Our study additionally evaluated third-party QC performance on ECLIA ISD assays, a total imprecision of <4.7% for CSA, <6.3% for TAC, and <8.2% for SRL were determined. The imprecision goal of ≤10%, based on the recommendation of the IATDMCT expert consensus group, was achieved for all QC samples. Overall, the ECLIA methods demonstrate acceptable precision.

The ECLIA methods offer a wider linear analytical measuring range for CSA and TAC than CMIA methods. ECLIA CSA, TAC, and SRL were linear up to 960.9 µg/L, 27.1 µg/L, and 32.3 µg/L, respectively. The higher upper limit allows TDM and pharmacokinetic analysis of ISD at different time points and peak concentrations offering additional flexibility.

The claimed functional sensitivity of the ECLIA ISD methods are improved for TAC and SRL compared to CMIA ISD methods. The functional sensitivities were assessed and the precision profile was used to calculate the LoQ corresponding to a CV of 20% with upper 95% confidence limit. The functional sensitivities were determined to be <6.5 µg/L for CSA, 1.1 µg/L for TAC, and <0.1 µg/L for SRL, which meets the 2007 European consensus guideline and IATDMCT expert consensus group recommended LoQ of 20.0 µg/L for CSA and a LoQ of 1.0 µg/L for both TAC and SRL.

### Table 1. Total imprecision for cyclosporine, tacrolimus, and sirolimus determined by the ECLIA method (duplicate per run, 1 run per day for 10 days).

<table>
<thead>
<tr>
<th>Samples</th>
<th>Mean Conc. (µg/L)*</th>
<th>Total CV (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Roche ISD L1</td>
<td>63.1</td>
<td>7.1</td>
</tr>
<tr>
<td>Roche ISD L2</td>
<td>271.1</td>
<td>5.1</td>
</tr>
<tr>
<td>Roche ISD L3</td>
<td>976.9</td>
<td>3.8</td>
</tr>
<tr>
<td>Bio-Rad WB ISD L1</td>
<td>67.3</td>
<td>4.7</td>
</tr>
<tr>
<td>Bio-Rad WB ISD L3</td>
<td>346.5</td>
<td>4.5</td>
</tr>
<tr>
<td>Bio-Rad WB ISD L4</td>
<td>740.1</td>
<td>3.3</td>
</tr>
<tr>
<td>Roche ISD L1</td>
<td>1.2</td>
<td>6.3</td>
</tr>
<tr>
<td>Roche ISD L2</td>
<td>9.2</td>
<td>6.9</td>
</tr>
<tr>
<td>Roche ISD L3</td>
<td>16.9</td>
<td>4.1</td>
</tr>
<tr>
<td>Bio-Rad WB ISD L1</td>
<td>4.1</td>
<td>6.3</td>
</tr>
<tr>
<td>Bio-Rad WB ISD L3</td>
<td>7.8</td>
<td>5.3</td>
</tr>
<tr>
<td>Bio-Rad WB ISD L4</td>
<td>15.2</td>
<td>3.9</td>
</tr>
<tr>
<td>Roche ISD L1</td>
<td>3.4</td>
<td>4.6</td>
</tr>
<tr>
<td>Roche ISD L2</td>
<td>8.7</td>
<td>5.6</td>
</tr>
<tr>
<td>Roche ISD L3</td>
<td>15.4</td>
<td>4.7</td>
</tr>
<tr>
<td>Bio-Rad WB ISD L1</td>
<td>5.9</td>
<td>4.7</td>
</tr>
<tr>
<td>Bio-Rad WB ISD L3</td>
<td>9.5</td>
<td>5.7</td>
</tr>
<tr>
<td>Bio-Rad WB ISD L4</td>
<td>14.2</td>
<td>8.2</td>
</tr>
</tbody>
</table>

*Conventional unit: 1 µg/L = 1 ng/mL.
For method comparison, anonymized residual patient samples spanning the analytical measuring range for each analyte were measured. CSA samples concentrations ranged from 41.0 to 1808.0 µg/L, TAC ranged from 2.1 to 30.0 µg/L, and SRL ranged from 1.8 to 34.6 µg/L as determined by CMIA. ECLIA ISDs measurements were compared to CMIA ISDs (n=100). The acceptance criteria were defined as a slope of 1.00 ± 0.15 and \( r \) of ≥ 0.95. Figure 1 shows the Deming regression and Bland-Altman analysis for CSA, TAC, and SRL. ECLIA and CMIA CSA (Figure 1A) showed good agreement with a slope of 0.917 (95% CI: 0.885-0.949), intercept of -15.2 (95% CI: -39.4-9.0), and \( r \) of 0.985. For TAC, the ECLIA TAC also showed good agreement with CMIA TAC (Figure 1B) with a slope of 0.938 (95% CI: 0.895-0.981), intercept of 0.2 (95% CI -0.4-0.8), and \( r \) of 0.974. Similar trends were observed by others (slopes of 0.87 for CSA, and 0.96-0.98 for TAC)\(^{14,29}\). Reported for the first time, method comparison of ECLIA and CMIA SRL (Figure 1C) showed a slope of 0.842 (95% CI: 0.810-1.110), intercept of 0.9 (95% CI: 0.4-1.4), and \( r \) of 0.982. Overall, all three ECLIA ISDs met our acceptance criteria, with SRL slightly exceeding the limit for the slope.

To further examine ECLIA performance with CMIA, a subset of samples was also analyzed by LC-MS/MS (n=20). For both CSA immunoassays, a positive bias was observed when compared with LC-MS/MS, similarly observed by others\(^{13,29,30}\) (Figure S1). For TAC, both the ECLIA and CMIA TAC had good agreement with LC-MS/MS, also similarly observed by others for different cohorts of solid organ transplant\(^{14,21,29}\) (Figure S2). Reported for the first time, the ECLIA SRL compared to LC-MS/MS showed a slope of 0.971 (95% CI: 0.913-1.030), intercept of 2.4 (95% CI: 1.6-3.3), and \( r \) of 0.993 (Figure 2). Meanwhile, CMIA SRL compared to

**Figure 1.** Method comparisons and Bland-Altman plots for (A) cyclosporine (CSA), (B) tacrolimus (TAC), and (C) sirolimus (SRL) between cobas e411 ECLIA and ARCHITECT i2000 CMIA.
LC-MS/MS showed a slope 1.119 (95% CI: 1.051-1.187), intercept of 1.4 (95% CI: 0.5-2.4), and \( r \) of 0.993. Based on our small sample size, both ECLIA and CMIA SRL generally showed good correlation with LC-MS/MS, with ECLIA SRL with better agreement to LC-MS/MS.

For lot-to-lot comparisons, Deming regression analysis of 20 residual patient samples tested by 2 lots of reagent for CSA shows a slope of 0.998 (95% CI: 0.967-1.029), intercept of 8.5 (95% CI: -10.2-27.2), and \( r \) of 0.998. For TAC, a slope of 0.972 (95% CI: 0.936-1.008), intercept of -0.4 (95% CI: -0.8-0.0), and \( r \) of 0.997. And for SRL, a slope of 0.913 (95% CI: 0.841-0.985), intercept of 0.1 (95% CI: -0.9-1.2), and \( r \) of 0.988. All three ISDs had good correlation between 2 different lots of reagents.

Evaluation on practical considerations included ease-of-use, throughput, and workflow of the method. The sample pretreatment for the ECLIA method is faster, simpler, and more convenient than CMIA method due to the use of a single universal sample pretreatment reagent and protocol for all three ISDs. Additionally, there is no heating step for the SRL ECLIA method, which leads to a simpler workflow. The ECLIA universal sample pretreatment reagent and protocol would enable better workflow, simpler sample handling and inventory control.

Dataset 1. File containing the raw data and a table of contents for the data file
http://dx.doi.org/10.5256/f1000research.12775.d18003

Conclusion
In conclusion, the overall analytical evaluation of the ECLIA method for CSA, TAC, and SRL met acceptable performance. ECLIA CSA showed better precision than our current CMIA
CSA. ECLI A CSA and TAC showed better linearity range, and ECLI A TAC and SRL showed good functional sensitivity than CMIA methods. Method comparisons showed good correlations and agreement between ECLI A ISDs and CMIA ISDs.

Data availability
Dataset 1: File containing the raw data and a table of contents for the data file. doi. 10.5256/f1000research.12775.d18003

Competing interests
No competing interests were declared.

Supplementary material
Supplementary File 1: Supplementary Methods.

Click here to access the data.

Figure S1: Method comparisons and Bland-Altman plots for cyclosporine (CSA) between (A) cobas e 411 ECLI A and LC-MS/MS, and (B) ARCHITECT i2000 CMIA and LC-MS/MS.

Click here to access the data.

Figure S2: Method comparisons and Bland-Altman plots for tacrolimus (TAC) between (A) cobas e 411 ECLI A and LC-MS/MS, and (B) ARCHITECT i2000 CMIA and LC-MS/MS.

Click here to access the data.

References

Grant information
The author(s) declared that no grants were involved in supporting this work.

Acknowledgements
We thank the technologists and support staff at University Health Network and the Hospital for Sick Children in Toronto, Ontario, Canada for their contributions to this study. The cobas ECLI A ISD reagents used in the method evaluation were provided by Roche Diagnostics.

Roche Diagnostics had no role in the design, analysis, or interpretation of this study.


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Version 1

Reviewer Report 30 November 2017

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Julie Shaw
Department of Pathology and Laboratory Medicine, The Ottawa Hospital, Eastern Ontario Regional Laboratories Association, University of Ottawa, Ottawa, ON, Canada

Here the authors describe evaluation of three immunosuppressant drug immunoassays on the Roche Cobas e411 analyzer. The authors evaluated the performance of assays for cyclosporine, tacrolimus and sirolimus on the system, which included imprecision, linearity and functional sensitivity studies as well as method comparisons and lot-to-lot comparisons. They compared the assay performance to the performance of the same assays on the Abbott immunoassay system and to LC MS/MS assays for the same analytes.

I have a few comments that, if the authors could address, I feel would strengthen the manuscript.

1. In the methods section, it’s not clear to me whether the measurements were performed using the Roche, Abbott and LC-MS/MS methods on the same day. It is mentioned that some specimens were stored prior to analysis. Could the authors clarify this?

2. What are the immunoassays and LC-MS/MS methods traceable to? Is there a standard? This information would be helpful.

3. The authors mention that the Roche assays were calibrated five times over the course of the study. Is this in fitting with the manufacturer’s recommendations? How does this calibration frequency compare to the Abbott system?

4. The authors should discuss the limitation of performing imprecision studies using only one lot number of reagent.

5. The authors should include a description of the LC MS/MS method that was used. There is currently no mention of this methodology until the results section.

6. There are several grammatical errors throughout the manuscript and I would recommend that the authors read it through carefully to correct these.
Is the work clearly and accurately presented and does it cite the current literature?  
Yes

Is the study design appropriate and is the work technically sound?  
Yes

Are sufficient details of methods and analysis provided to allow replication by others?  
Yes

If applicable, is the statistical analysis and its interpretation appropriate?  
Yes

Are all the source data underlying the results available to ensure full reproducibility?  
Yes

Are the conclusions drawn adequately supported by the results?  
Yes

**Competing Interests:** No competing interests were disclosed.

I have read this submission. I believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

Reviewer Report 20 October 2017

https://doi.org/10.5256/f1000research.13840.r26931

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Jennifer Shea  
Laboratory Medicine, Saint John Regional Hospital, Horizon Health Network, Fredericton, NB, Canada

Fung et al. present data evaluating three new ECLIA assays for the immunosuppressant drugs cyclosporine, tacrolimus, and sirolimus on a Roche Cobas e411 analyzer. Overall the study design is sound generally following CLSI guidelines for method validation. The introduction provides a thorough background on the drugs/assays being evaluated and the methods section clearly describes each portion of the method validation that would easily allow another researcher to replicate the study if desired. The results/discussion section succinctly presents their data in a cohesive and systematic manner and the authors do a great job at highlighting the novel aspects of their study. Their conclusion that these three new assays are fit for use for therapeutic drug monitoring is supported by the data presented.

Is the work clearly and accurately presented and does it cite the current literature?  
Yes

Is the study design appropriate and is the work technically sound?  
Yes
Are sufficient details of methods and analysis provided to allow replication by others?
Yes

If applicable, is the statistical analysis and its interpretation appropriate?
Yes

Are all the source data underlying the results available to ensure full reproducibility?
Yes

Are the conclusions drawn adequately supported by the results?
Yes

**Competing Interests:** No competing interests were disclosed.

I have read this submission. I believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

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