Validation of a point-of-care handheld blood total calcium analyzer in postpartum dairy cows

Rita Couto Serrenho,1* Tony C. Bruinjé,1 Emma I. Morrison,1 David L. Renaud,1 Trevor J. DeVries,2 Todd F. Duffield,1 and Stephen J. LeBlanc1

Graphical Abstract

Summary
Our objective was to validate a point-of-care handheld blood calcium analyzer to estimate circulating calcium concentrations in postpartum dairy cows. An accurate, rapid, and inexpensive tool to screen for hypocalcemia on-site would allow for implementation of selective treatment protocols but the meter assessed was not sufficiently accurate to quantify blood calcium concentration.

Highlights
• It is desirable to identify cows for selective administration of Ca supplements
• The objective was to validate a cow-side meter to estimate blood Ca in dairy cows
• The meter was not sufficiently accurate to quantify blood Ca concentration
• The meter might be useful to classify subclinical hypocalcemia in fresh plasma

1Population Medicine, University of Guelph, Guelph, ON, Canada N1G 2W1, 2Animal Biosciences, University of Guelph, Guelph, ON, Canada N1G 2W1.
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Validation of a point-of-care handheld blood total calcium analyzer in postpartum dairy cows

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Abstract: Our objective was to validate a point-of-care handheld blood total calcium analyzer (Ca meter, CM; TD-5220 Vet Ca®, TaiDoc, New Taipei, Taiwan) to estimate circulating Ca concentrations in postpartum dairy cows. Whole blood was collected from 251 multiparous cows within the first 4 d after parturition (Martinez et al., 2012; McArt and Neves, 2020). The cut-point to define SCH varies among studies, depending on the outcome of interest, and when postpartum and how many samples per cow were assessed. Subclinical hypocalcemia is associated with increased risk of metritis and other postpartum diseases (Chapinal et al., 2011; Martinez et al., 2012; Rodriguez et al., 2017). It may also be associated with reduced milk yield in early lactation, depending on when blood Ca is measured and the duration of SCH (Chapinal et al., 2012; Neves et al., 2018a; McArt and Neves, 2020). Cows with blood total Ca (tCa) ≤2.14 mmol/L in the first 3 DIM tend to have lower pregnancy rates and to take longer to return to cyclicality (Martinez et al., 2012; Caixeta et al., 2017). Because SCH is common and associated with undesirable outcomes, peripartum Ca supplementation is a common practice in dairy herds (USDA-NAHMS, 2018).

However, the effects of postpartum Ca supplementation appear to depend on parity, milk production potential, or pretreatment tCa. In multiparous cows, postpartum Ca supplementation enhanced reproductive performance, and increased milk yield in cows with milk yield greater than their herd average in the previous lactation, but decreased production in cows with below-average production (Martinez et al., 2016b). Calcium supplementation reduced disease risk in multiparous cows with lower blood Ca concentration (Leno et al., 2018) or those with excessive BCS or pre-existing lameness (Oetzel and Miller, 2012; Leno et al., 2018). In primiparous cows, Ca supplementation reduced pregnancy at first insemination and increased the risk of postpartum disease when tCa at parturition was >2.15 mmol/L (Martinez et al., 2016a,b; Leno et al., 2018). Therefore, it would be desirable to selectively treat multiparous cows likely to benefit from Ca supplements. An accurate, rapid, and inexpensive tool to screen for SCH on-site would allow for implementation of treatment protocols based on circulating blood Ca concentration at specific days in milk and to avoid blanket treatments that could incur costs without benefits.

To date, only one portable clinical analyzer has been validated for hypocalcemia testing in cattle (Peiró et al., 2010). However, its purchase cost and the cost of individual cartridges represent a barrier to its routine use on farm. Therefore, the objective of this cross-sectional diagnostic accuracy study was to validate a point-of-care handheld blood total Ca analyzer (TD-5220 Vet Ca®, TaiDoc, New Taipei, Taiwan; calcium meter, CM) to estimate circulating Ca concentration in postpartum, multiparous dairy cows.

This work is reported using the Standards for Reporting of Diagnostic Accuracy Studies (STARD) Guideline (Cohen et al., 2016). Sample collection was approved by the University of Guelph Animal Care Committee (Guelph, ON, Canada; AUP# 3951). Whole blood was collected from the coccygeal vessels from 251 multiparous cows at 1, 2, 3, or 4 DIM (1 sample per cow) between September 2019 and February 2020 from 2 commercial dairy herds.
in Ontario, Canada, milking approximately 450 (farm A) or 400 (farm B) Holstein cows.

Blood was collected with a 22-gauge, 1.5-inch needle into a 10-mL tube without anticoagulant (BD Vacutainer Precision Glide; Becton Dickinson, Franklin Lakes, NJ) and into 4-mL and 6-mL sodium heparinized tubes (Vacutainer, Becton Dickinson). All testing with the CM was performed following the manufacturer’s guidelines. In short, 0.5 mL of air followed by 0.5 mL of the sample were drawn into a 1-mL syringe; a small plastic filter (provided by the manufacturer) was attached to a syringe; a test strip was introduced into the meter; and a drop of filtered blood or fresh or thawed plasma was placed in the strip well. After 180 s, a result was displayed on the screen.

Within 60 min of sampling, a sample of heparinized whole blood (4-mL tube) was tested at the farm with the CM, indoors at approximately 22°C. The 6-mL heparinized sample tube was placed on ice and the 10-mL whole blood sample was maintained at room temperature until processing in the laboratory. Within 4 h of sampling, the 6-mL and 10-mL tubes were centrifuged at 1,500 × g at room temperature for 15 min. A sample of the fresh plasma was used to measure tCa with the CM as previously described, while the serum and remaining plasma were stored in aliquots at −20°C for further analysis. Frozen plasma was thawed at room temperature and tested with the CM, whereas frozen serum was sent to the Animal Health Laboratory (University of Guelph, ON, Canada) for measurement of tCa using a Cobas Calcium Gen 2 kit (Roche Diagnostics, Indianapolis, IN), which was considered the reference test (RT). Although atomic absorption spectrophotometry is considered the gold standard to measure calcium concentration, photometric methods are commonly used in clinical practice (Kimura et al., 1996; Bazydlo et al., 2014). These validated methods are available at a lower cost (Kimura et al., 1996; Bourguignon et al., 2014).

The CM consisted of a colorimetric method to estimate blood tCa concentration between 0.5 and 3.0 mmol/L in increments of 0.1 mmol/L, with outputs of “low” or “high” if values were out of the lower or upper limits of quantification. The manufacturer provided instructions to test tCa in heparinized whole blood and fresh plasma (sample size of 45 μL). Because serum samples were frozen and thawed before being tested at the Animal Health Laboratory, we also tested thawed plasma. The RT is a photometric method with a range of quantification of 0.20 to 5.0 mmol/L, and intra- and interassay coefficients of variation of 1.3 and 2.3%, respectively. Independent of the samples used to assess the CM, we compared tCa measured by the RT method in 20 paired plasma and serum samples from the same cows at the same time (blood collected into tubes with lithium heparin or no anticoagulant, respectively). Regression of these data gave the relationship serum Ca = 0.373 × 0.812(plasma Ca); R² = 0.94, ρ = 0.92, with a mean difference of 0.05 ± 0.06 mmol/L, where ρ is Lin’s concordance correlation coefficient.

Sample size was calculated using the method described by Budriner (1996). A prevalence of SCH (defined as serum concentration ≤2.14 mmol/L) of at least 50% was used based on previous studies in multiparous cows (Martinez et al., 2012; Caixeta et al., 2017). To provide a conservative estimate, we calculated the sample size using expected sensitivity (Sn) and specificity (Sp) of 50%, and a clinically acceptable width of the 95% CI for Sn and Sp of 10%, requiring a minimum of 193 samples.

Assumptions of normality were assessed with the Shapiro-Wilk test and histograms for all continuous variables. Data from the CM results were censored at <0.5 and >3.0 mmol/L, which compromised its normal distribution. Transformations of the data did not improve the distribution (negatively skewed) so the data were not transformed. Lin’s concordance correlation coefficient (ρ; Lin, 1989) was calculated using the macro “CCC V9” (Crawford et al., 2007) in SAS version 9.4 (SAS Institute Inc., Cary, NC) to measure the agreement between the RT (thawed serum) and each of the CM results (whole blood, fresh plasma, or thawed plasma). To evaluate bias between each CM test and the RT, Bland-Altman (B-A) plots (Bland and Altman, 1986) were generated in SAS. For calculation of ρ and the B-A plots, values reported as “high” or “low” by the CM were excluded. Graphical representation of Lin’s regressions and the B-A plots were created using Excel (Windows 10, version 1903; Microsoft Corp., Redmond, WA). Total calcium values from RT were categorized as SCH (≤2.14 mmol/L; Martinez et al., 2012) or normocalcemia (NC; >2.14 mmol/L). Contingency tables were created to calculate Sn, Sp, positive and negative predictive values, and accuracy (percentage of samples correctly classified).

Table 1. Descriptive statistics of blood total calcium concentrations (tCa) on d 1, 2, 3, or 4 after calving from 251 multiparous cows, measured with a point-of-care meter or in a diagnostic laboratory (reference test)

| Variable       | Point-of-care calcium analyzer1,2 | Reference test4 | Reference test4 |
|----------------|----------------------------------|-----------------|-----------------|
| Samples (no.)  | Whole blood                      | Fresh plasma    | Thawed plasma   | Thawed serum    |
| “High”         | 9                                | 5               | 3               | —               |
| “Low”          | 11                               | 1               | 0               | —               |
| Blood tCa (mmol/L)4 | Mean (SD) | 2.3 (0.6) | 2.5 (0.2) | 2.5 (0.3) | 2.03 (0.29) |
|                | Median (IQR)5                      | 2.5 (0.8)       | 2.5 (0.3)       | 2.5 (0.3)       | 2.05 (0.38)   |
|                | Maximum                            | 3.0             | 3.0             | 3.0             | 2.63          |
|                | Minimum                            | 0.5             | 1.9             | 1.6             | 1.05          |

1TD-5220 Vet Ca2+ (TaiDoc, New Taipei, Taiwan).
2All samples were filtered with a device provided by the manufacturer.
3Cobas Calcium Gen 2 kit (Roche Diagnostics, Indianapolis, IN).
4Samples with “high” (>3.0 mmol/L) or “low” (<0.5 mmol/L) readings from the calcium analyzer were not included.
5Interquartile range.
formed using PROC LOGISTIC in SAS to calculate the area under the curve and identify the threshold with highest combined Sn and Sp (the point on ROC curve closest to \( x = 0 \) and \( y = 1 \)); false negatives and false positives were weighted equally. For all categorical analyses, “high” and “low” readings from the CM were replaced by 3 or 0.5 mmol/L, respectively, and included as classifications above or below the cut-point of interest. Because of damage to the CM at the beginning of the study, fewer samples were analyzed as whole blood or fresh plasma than as thawed plasma (Table 1). While we waited for a new meter, blood sample collection continued, providing frozen plasma but no fresh samples. We repeated all the statistical analyses above after applying a correction to the serum data based on the regression of serum versus plasma \( tCa \) measured with the RT.

A total of 251 samples from multiparous cows within the first 4 DIM were included in this study. During the time that the CM was not functional, 153 whole blood and 151 fresh plasma samples were not tested. The calculation of \( \rho \) and analysis of B-A plots did not include 3, 6, and 20 thawed plasma, fresh plasma, and whole blood samples, respectively, due to the lack of a numeric result from the CM. The prevalence of SCH (≤2.14 mmol/L) using the RT was 63.7% [farm A: 73.8% (96/130); farm B: 52.9% (64/121)]. Descriptive statistics are presented in Table 1. Lin’s correlation coefficient demonstrated poor agreement between the CM and

![Figure 1](image_url)

**Figure 1.** Agreement between a point-of-care calcium analyzer (CM; TD-5220 Vet Ca\(^{2+}\), TaiDoc, New Taipei, Taiwan) used with thawed plasma (top row), fresh plasma (middle row), or whole blood (bottom row) and serum total Ca concentration measured in a diagnostic laboratory (reference test, RT; Cobas Calcium Gen 2 kit, Roche Diagnostics, Indianapolis, IN). (a) Regression showing Lin’s concordance correlation coefficient (\( \rho \)) and its 95% CI; the solid line represents perfect concordance, and the dashed line represents the observed concordance regression. (b) Bland-Altman plot of differences between the CM and the mean of the results from the CM and the reference test. For thawed plasma (top row), there were 247 samples (1 sample per cow between 1 and 4 DIM), excluding 3 for which no numeric result was provided by the CM. For fresh plasma (middle row), there were 94 samples (1 sample per cow between 1 and 4 DIM), excluding 6 for which no numeric result was provided by the CM. For whole blood (bottom row), there were 78 samples (1 sample per cow between 1 and 4 DIM), excluding 20 for which no numeric result was provided by the CM.
RT measurement for blood tCa concentration (thawed plasma: \( \rho = 0.16 \); fresh plasma: \( \rho = 0.21 \); and whole blood: \( \rho = 0.23 \); with the plasma vs. serum regression applied to the data, \( \rho = 0.18, 0.24, \) and 0.26, respectively). The inferences were the same for all the following analyses whether or not we adjusted the serum values, so unadjusted data are presented here. Most of the data points are located above the line of perfect agreement, indicating overestimation by the CM (Figure 1a). The same is indicated by the B-A plots, where the negative bias present in all 3 plots (Figure 1b) indicated that the CM values were greater than the RT for the same sample. For thawed plasma, fresh plasma, and whole blood, these plots showed a mean difference (bias) of \(-0.44, -0.53, \) and \(-0.26 \) mmol/L, respectively. This means that, on average, the CM gave a result on thawed plasma that was 0.44 mmol/L ± 0.31 greater than the RT value. The B-A plots also illustrated random variability between the results from the CM and RT. Although most of the data fall within the confidence limits on the B-A plots, that does not indicate that differences within these bounds are biologically or practically acceptable. Moreover, mean errors of 0.26 to 0.53 mmol/L are meaningful differences when measuring tCa postpartum. For context, the allowable error for measurement of tCa in human samples is ±0.25 mmol/L (Verma et al., 2019). The variability between CM and RT appeared to be randomly distributed for both fresh and thawed plasma but not for whole blood. The B-A plots demonstrated a positive bias with low mean tCa concentrations (i.e., the RT result was higher) and a negative bias with a greater mean tCa concentration. It is possible that the fewer samples in the low tCa range contributed to this difference. However, this should be taken into consideration because this meter is recommended for use with whole blood. The large mean bias with substantial variability and low concordance between tests led us to conclude that the CM results, on a continuous scale, were not good estimates of tCa.

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observed in the B-A plots were 5- to 10-fold greater than the mean difference between serum and plasma.

Based on low concordance with the RT, we conclude that the CM was not sufficiently accurate to quantify tCa concentration. However, with an adjusted cut-point, when used with fresh plasma, with sensitivity of 72% and specificity of 86%, it might be useful as a screening tool for SCH. The performance of the CM for on-farm assessment of tCa with whole blood was not adequate to select cows to receive a Ca supplement after calving.

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Notes

Rita Couto Serrenho © https://orcid.org/0000-0002-7956-7039
Tony C. Brunjie © https://orcid.org/0000-0003-4046-8081
Emma I. Morrison © https://orcid.org/0000-0003-2602-0336
David L. Renaud © https://orcid.org/0000-0002-3439-3987
Trevor J. DeVries © https://orcid.org/0000-0001-9364-2456
Todd F. Duffield © https://orcid.org/0000-0001-6035-4669
Stephen J. LeBlanc © https://orcid.org/0000-0003-2027-7704

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