

1 **Are Hemoglobin A<sub>1c</sub> Point-of-Care analyzers fit for purpose? The**  
2 **story continues.**

3 **HbA<sub>1c</sub> POCT: the story continues**

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30 **Keywords:** : HbA<sub>1c</sub>, Hemoglobin A<sub>1c</sub>, POCT, quality, diabetes, Hb-variant

31 **Abbreviations:** Point-of-Care (POC); Clinical and Laboratory Standards Institute  
32 (CLSI); International Federation of Clinical Chemistry and Laboratory Medicine  
33 (IFCC); National Glycohemoglobin Standardization Program (NGSP), Secondary  
34 Reference Measurement Procedures (SRMP); External Quality Assessment (EQA);  
35 European Reference laboratory for Glycohemoglobin (ERL); System Usability Scale  
36 (SUS).

37 1

1 **Abstract**

2 **Background** Point of care (POC) analyzers are playing an increasingly important role  
3 in diabetes management but it is essential that we know the performance of these  
4 analyzers in order to make appropriate clinical decisions. Whilst there is a growing  
5 body of evidence around the more well-known analyzers, there are many 'new kids on  
6 the block' with new features, such as displaying the presence of potential Hb-variants,  
7 which do not yet have a proven track record.

8 **Methods** The study is a comprehensive analytical and usability study of six POC  
9 analyzers for HbA<sub>1c</sub> using Clinical and Laboratory Standards Institute (CLSI) protocols,  
10 international quality targets and certified International Federation of Clinical Chemistry  
11 and Laboratory Medicine (IFCC) and National Glycohemoglobin Standardization  
12 Program (NGSP) Secondary reference Measurement Procedures (SRMP). The study  
13 includes precision (EP-5 and EP-15), trueness (EP-9), linearity (EP-6), sample  
14 commutability (fresh, frozen and lyophilised), interference of Hb-variants (fresh and  
15 frozen samples).

16 **Results** Only two of the 6 analyzers performed to acceptable levels over the range of  
17 performance criteria. Hb-variant interference, imprecision or variability between lot  
18 numbers are still poor in 4 of the analyzers.

19 **Conclusions** This unique and comprehensive study shows that out of 6 POC  
20 analyzers studied only 2 (The Lab 001 and Cobas B101) met international quality  
21 criteria (IFCC and NGSP), 2 (A1Care and Innovastar) were borderline and 2  
22 (QuikReadgo and Allegro) were unacceptable. It is essential that the scientific and  
23 clinical community are equipped with this knowledge in order to make sound decisions  
24 on the use of these analyzers.

25

## 1 **Introduction**

2 Diabetes is a global health burden and a leading cause of morbidity and mortality  
3 worldwide. It is estimated that up to 50% of people with diabetes are currently  
4 undiagnosed, and there is an urgent need for rapid, accurate and timely diagnostic  
5 testing to identify those both with and at risk of the disease [1].

6 Point of care analyzers play an increasingly important role in wide range of clinical  
7 settings and there is increasing desire from clinicians to have access to more POC  
8 tests and a wider range of tests [2]. There is belief that POC enables faster clinical  
9 decision making, increased rapport with patients and reduced referrals to secondary  
10 care and subsequent healthcare costs. Over 50% of primary care physicians surveyed  
11 by Horwick et al (2014) wanted increased access to HbA<sub>1c</sub> POC testing, bespeaking  
12 a clear demand for HbA<sub>1c</sub> POC [3]. However, Jones et al (2013) also highlighted an  
13 apparent nervousness amongst primary care physicians around the accuracy of POC  
14 testing [4]. Coupling the desire for increased HbA<sub>1c</sub> POC availability and the prudent  
15 concerns on quality it is essential that we understand how well POC HbA<sub>1c</sub> analyzers  
16 perform.

17 Understanding the quality of POC testing has been a topic of key interest for over a  
18 decade with the stark message of 6 out of 8 analyzers not meeting the accepted quality  
19 criteria in 2010 [5]. Since this seminal study, there have been numerous evaluations  
20 of POC HbA<sub>1c</sub> performance with a focus around the more common analyzers [6].  
21 External quality assessment (EQA) provides a snap shot of 'real world' data on the  
22 performance of POC analyzers, although only a fraction of analyzers in use are  
23 currently enrolled in EQA schemes [7, 8].

24 Whilst there is a growing body of evidence around the more well-known analyzers,  
25 there are many 'new kids on the block' with new features, such as displaying the

1 presence of potential Hb-variants, which do not have a proven track record. Clinicians  
2 and laboratory scientists need robust and rigorous evaluation data on performance,  
3 acceptability and usability of POC analyzers to support informed decision making  
4 around use of POC testing analyzers.

5 The evaluation of POC analyzers is not without issue. Whilst many HbA<sub>1c</sub> POC  
6 analyzers are scaled down versions of laboratory analyzers they have their own  
7 unique differences which require adaptations in order to complete a comprehensive  
8 evaluation. One key issue is that several POC analyzers are not compatible with  
9 frozen or lyophilized blood samples, meaning conventional evaluation protocols  
10 cannot be directly applied.

11 Whilst there are numerous method comparisons published, it is important to note that  
12 these are often single comparisons to routine laboratory methods (which will have their  
13 own imprecision and bias to consider), which provide insight into local performance  
14 but do not provide a robust picture of performance against internationally accepted  
15 secondary reference measurement procedures (SRMPs) or international quality  
16 criteria [9,10].

17 This study aims to understand the performance of a range of POC HbA<sub>1c</sub> analyzers  
18 using a rigorous evaluation protocols which examines issues such as; interference  
19 from Hb-variants with fresh and frozen samples, sample compatibility (fresh, frozen  
20 and lyophilised) and system usability whilst comparing analytical performance to IFCC  
21 and NGSP SRMPs.

22

## 23 **Materials and Methods**

24 ***Analyzers evaluated*** The six POC analyzers included in this study and their key  
25 characteristics, are summarized in Table 1. The choice of analyzer was both

1 manufacturer led (Allegro, QuikReadgo and The lab 001) and investigator led  
2 (Cobas B101, InnovaStar and A1Care). The manufacturers of the last three  
3 analysers were approached by the authors as previous evaluations had highlighted  
4 some performance issues. An initial familiarization protocol was undertaken with  
5 each instrument and the results were shared with the manufacturers to enable them  
6 to decide if they wished to continue to a full evaluation. This supports a collaborative  
7 approach to working with manufacturers with the aim to improve quality. In some  
8 cases when a product is new in development, feedback at an early stage will enable  
9 further development before a product is brought to full evaluation, saving time and  
10 resources [11]. Six analyzers were fully evaluated, however two new analyzers were  
11 not ready yet for a full evaluation.

12

13 ***Imprecision study (EP-5 and EP-15)*** The CLSI EP-5 protocol was used to investigate  
14 assay imprecision. It is known that some POC methods have a bias with frozen  
15 material and as it is not known if frozen samples have an impact on the imprecision,  
16 also EP-15 was performed with two fresh patient samples (HbA<sub>1c</sub> values of 48 and 75  
17 mmol/mol). Both samples were analyzed five-fold for 5 days. CVs were also calculated  
18 on the basis of the duplicates of the fresh patient samples in the EP-9 protocol.

19

20 ***Method comparison (trueness; EP-9)*** The CLSI EP-9 protocol was performed with  
21 40 fresh patient samples and the data was used to investigate the bias between each  
22 instrument and 4 SRMPs (n=40, 8 samples per day for 5 days, duplicate  
23 measurements). Values were assigned with 4 IFCC and NGSP Certified SRMPs  
24 [12,13]:

- 1 • Roche Tina-quant Gen.3 HbA<sub>1c</sub> on Cobas c513, immunoassay, IFCC and  
2 NGSP certified (Roche Diagnostics);
- 3 • Premier Hb9210, affinity chromatography HPLC, IFCC and NGSP certified  
4 (Trinity Biotech);
- 5 • Tosoh G8, cation-exchange HPLC, IFCC certified (Tosoh Bioscience);
- 6 • Abbott Enzymatic method on Alinity, IFCC and NGSP certified (Abbott  
7 Diagnostics).

8

9 **Linearity (EP-6)** Linearity was assessed using the CLSI EP-6 protocol. After  
10 adjustment for Hb concentration, patient samples with a low HbA<sub>1c</sub> value and a high  
11 HbA<sub>1c</sub> value were mixed in incremental amounts to generate a series of equally  
12 spaced samples over a broad HbA<sub>1c</sub> concentration range. Eleven samples were  
13 analyzed in duplicate in one day. The samples were made fresh and then frozen at -  
14 80 °C degrees until analysis. Whilst some analyzers display a bias with frozen samples  
15 this is generally a consistent bias and therefore these can still be used to assess  
16 linearity.

17 The difference between the fitted values of the best polynomial line and the regression  
18 line for the 11 samples were compared. CLSI states for EP-6 that goals for linearity  
19 should be derived from goals for bias, and should be less than or equal to these goals  
20 [14]. The IFCC Task Force on Implementation of HbA<sub>1c</sub> Standardization has set an  
21 TAE of 10% at an HbA<sub>1c</sub> concentration of 50 mmol/mol (19). Taking into account the  
22 whole clinical relevant range, we have set a TAE of 6 mmol/mol with a nonlinearity  
23 budget of 50% (=3 mmol/mol). If the deviation exceeds allowable nonlinearity (3  
24 mmol/mol) the data was considered nonlinear.

25

1 ***Hemoglobin variants AS, AC, AE, AD, elevated A2 and HbF***

2 Twenty patient samples of each heterozygous Hb variant, from our frozen whole blood  
3 biobank, were measured on each of the different POC analyzers. Values were  
4 assigned using an IFCC calibrated boronate affinity HPLC (Premier Hb9210). For  
5 samples with increased HbF, HbA<sub>1c</sub> values were assigned using an IFCC calibrated  
6 cation-exchange HPLC (Menarini HA8180V, Diabetes Mode, (frozen) and Tosoh G8  
7 (fresh)). Percentage HbF (3.5 – 42.0%) was determined using the Sebia Capillarys 2  
8 Flex Piercing Hemoglobin program.

9 In addition to the frozen samples, 16 HbAS, 7 HbAC, 5 HbAD, 9 HbAE and 4 HbF (9.1,  
10 20.5, 20.8 and 27.5%) fresh Hb-variant samples were also analyzed on each analyser  
11 as two of the analyzers (InnovaStar and QuikReadgo) showed a bias with frozen  
12 samples.

13 Any bias observed due to the presence of variants is a compound of both the bias in  
14 normal samples (identified by the EP-9 protocol) and the bias associated with the  
15 variant. In order to account for this, the results were adjusted for the bias found during  
16 EP-9 (Premier Hb9210 for fresh and frozen Hb-variant), thus any residual bias would  
17 be due to the Hb-variant. For the two analyzers that also display a bias with frozen  
18 samples, the bias correction was done using the 24 frozen EQA samples rather than  
19 the EP-9 data. Whilst this is not a perfect solution it avoids a two-step correction.

20 For an Hb variant to be considered as not causing a clinically relevant interference,  
21 the results of the Hb variant should fall within a defined scatter line of  $\pm 10\%$  (SI units)  
22 of the regression line derived from the comparison of the test instrument and the  
23 Premier Hb9210 with the nonvariant samples (HbAA).

24

25 ***Schiff Base, Icteric samples, different Hemoglobin concentrations***

1 In order to create labile samples (Schiff Base) 12, 16 and 20 mg/ml glucose was added  
2 to aliquots of high, medium and low HbA<sub>1c</sub>, EDTA samples. Icteric samples were  
3 generated by removing the plasma of a non-icteric sample and replacing with plasma  
4 with 219, 236 and 258 µmol/L bilirubin (icteric sample), again at three different HbA<sub>1c</sub>  
5 levels. Similarly addition or removal of plasma was used to create a range of samples  
6 with varying hemoglobin levels. The samples were stored frozen at –80 °C until  
7 analysis. A mean relative difference of ± 10% (in SI units) pre and post treatment of  
8 the samples, was considered a significant interference.

9

10 ***EQA Programs (assessing sample commutability)*** In order to assess sample  
11 commutability, samples from both the IFCC Certification Program for manufacturers  
12 [15] and the European Reference Laboratory for Glycohemoglobin (ERL) EQA  
13 Program [16] were used to provide data on frozen and lyophilized samples  
14 respectively.

15

### 16 ***System Usability Scale (SUS)***

17 This study included a SUS score generated by the two technicians who performed the  
18 evaluation study. SUS is a simple technology diagnostic tool consisting of ten  
19 questions which gives a global view of subjective assessment of the usability of the  
20 device tested [17].

21 A SUS score >81 can be considered as excellent, between 71 and 80 as good,  
22 between 52 and 70 okay and <51 poor [18].

23

### 24 **Defining the quality criteria**



1 **International Quality Standards** This study used the previously published global  
2 guidance on acceptable quality and performance criteria for HbA<sub>1c</sub> testing from the  
3 IFCC Task Force on Implementation of HbA<sub>1c</sub> Standardization [19].

4

5 **NGSP Manufacturer Certification Criteria** Thirty six out of 40 results must be within  
6 5% (relative) of an individual NGSP SRMP to pass certification [20].

7

8 **Statistical analysis** Calculations were performed using Microsoft® Excel 2016  
9 (Microsoft Corporation). Statistical analyses were performed using Analyse-It®,  
10 version 5.40 (Analyse-It Software) and EP Evaluator Release 12 (Data Innovations).

11

## 12 **Results**

13 **Imprecision (EP-5 and EP-15)** Table 2 displays the CVs derived from both EP-5 and  
14 EP-15 and the duplicates from the EP-9 protocol. Only the Lab 001 and the InnovaStar  
15 achieved the performance criteria of <3% CV (SI units) (<2% in NGSP units) across  
16 all protocols and both high and low HbA<sub>1c</sub> levels [21, 22]. The Lab 001 actually  
17 achieved <2 % CV (SI units) showing very low levels of imprecision. However the  
18 Allegro failed to achieve <3% CV (SI units) in any protocol at either level, with CVs as  
19 high as 4.2%, showing unacceptable levels of imprecision. The A1Care had mixed  
20 results, performing better with fresh samples and at higher HbA<sub>1c</sub> values. The B101  
21 also had mixed results, with better performance at higher HbA<sub>1c</sub> values. The  
22 QuikReadgo met the criteria in both EP-5 and EP-15 with little difference between  
23 fresh and frozen samples. However, the performance with the duplicates from EP-9  
24 was mixed with one lot failing and one lot passing.

25

1 **Method comparison (trueness; EP-9)** Table 3 details the results of the method  
2 comparison study and also NGSP pass/fail rate. From these data it is clear that all  
3 analyzers suffered some degree of bias, with some such as the Allegro showing this  
4 across multiple levels and between lot numbers. The data for the individual POC  
5 analyzers versus the individual SRMPs in NGSP units is available in supplemental  
6 table 1. From this table and table 3 it can be seen that only the Cobas B101 passed  
7 the NGSP criteria with two lot numbers compared to all four individual SRMPs and  
8 that the QuikReadgo and the Allegro failed the NGSP criteria with both lot numbers  
9 for all four individual SRMPs. Figure 1 shows the regression lines for each POC device  
10 versus the mean of the SRMPs. All analyzers suffered some degree of bias. The Lab  
11 001 and the Cobas B101, had the least bias across the lots and the HbA<sub>1c</sub> range and  
12 all other POC analyzers had a statistically significant difference either between the lot  
13 numbers or at different HbA<sub>1c</sub> levels or both and showed a large dispersion around the  
14 deming regression line compared to the mean of the SRMPs.

15  
16 **Linearity (EP-6)** Supplemental table 2 details the results of the linearity study. The  
17 maximum deviation is shown between the fitted values of the best polynomial line  
18 and the regression line for the 11 samples. If the deviation exceeds allowable  
19 nonlinearity (3 mmol/mol) the data was considered nonlinear. Based on this criteria  
20 all POC analyzers were linear except for the Cobas B101 and the InnovaStar.  
21 However, the detection limit of the InnovaStar was >30 mmol/mol. Excluding the  
22 lowest sample for the calculations showed that the InnovaStar was linear. The HbA<sub>1c</sub>  
23 result of the highest sample for the Allegro was above the detection limits (> 130  
24 mmol/mol) therefore the linearity was assessed with 10 samples instead of 11.

25

1 **Hemoglobin variants AS, AC, AE, AD, elevated A2 and HbF** Table 4 shows the  
2 mean relative difference of the frozen and fresh Hb-variants samples and  
3 supplemental figure 1 to 6 shows the graphs of the interference of Hb-variants for  
4 frozen and fresh Hb-variants for the different POC analyzers. All methods, except for  
5 the A1Care, had an interference with one or more of the Hb-variants with frozen or  
6 fresh samples (mean relative difference was >10%). All Hb-variants were detected  
7 and correctly identified by the Lab 001 via a S-, C-, D-, E- or F-window.

8

### 9 **Schiff Base, Icteric samples, different Hemoglobin concentrations**

10 None of the POC analyzers showed an interference for schiff base, icteric samples or  
11 different hemoglobin concentrations. Supplemental tables 3 to 5 show the data.

12

### 13 **Sample commutability**

14 To investigate the impact of sample type in relation to different clinical applications,  
15 fresh (EP-15 and EP-9), frozen (EP-5 and IFCC certification program samples) and  
16 lyophilized (ERL EQA scheme) samples were compared. Figure 2 (panel A and B)  
17 show the data from EP15 and EP-9 (red circles) representing all fresh samples, IFCC  
18 certification samples (blue circles) representing all frozen samples and the ERL EQA  
19 (green circles) representing lyophilized samples. In addition panel B shows EP-5 and  
20 EP-9 data (grey circles) in order to assess the impact of using EP-5 versus EP-15 for  
21 routine method evaluations. The EP-5 and EP-15 studies were both used to compare  
22 performance with fresh and frozen samples and for the majority of analyzers the EP-  
23 15 evaluation provided sufficient data to assess performance.

24 The data clearly demonstrates that lyophilized material was only commutable with The  
25 Lab 001, all other POC analyzers showed a large positive bias when analyzing

1 lyophilized material. Frozen material was not commutable with the InnovaStar and the  
2 QuikReadgo. When using fresh patient samples all, except the Allegro, passed the  
3 IFCC criteria of having a sigma  $>2$  at an HbA<sub>1c</sub> concentration of 50 mmol/mol.  
4 Conversely the Allegro actually performed better when using frozen samples instead  
5 of fresh samples.

6

### 7 ***SUS scores***

8 Table 1 shows the SUS scores of the different POCT analyzers. The usability of all the  
9 POCT analyzers was good to excellent (mean SUS score  $> 80$ ) except for the  
10 QuikReadgo (mean SUS score was 60).

11

## 12 **Discussion**

### 13 ***What progress has been made?***

14 In the study of 2014 the InnovaStar showed an interference with fresh patient samples,  
15 which was likely due to the instrument being calibrated using frozen samples [23]. The  
16 previous publication led the manufacturer to switch to fresh patient samples, which are  
17 available from the ERL, to calibrate their cartridges resulting in lower bias in fresh  
18 samples [24].

19 The Lab 001 device is new to the market and the sigma graphs show excellent  
20 performance however there was a small bias at higher HbA<sub>1c</sub> levels. Paradoxically,  
21 had the imprecision of the instrument been higher than the bias would not have been  
22 detected as the confidence intervals would be wider. This device shows that the field  
23 of POCT has moved on and quality improvements are possible.

24

25 ***There are still significant issues with the performance of some analyzers***

1 The key issues we still see are: a) lot to lot variability, b) high imprecision, and c)  
2 significant interference from variants. Four out of the 6 evaluated analyzers still do not  
3 demonstrate acceptable and/or consistent performance.

4 The InnovaStar had poor performance between lot numbers that was not acceptable.  
5 The new to the market QuikReadgo also showed a statistically significant difference  
6 between the lot numbers, and failed to meet the NGSP criteria with either lot number  
7 when compared to any of the 4 SRMPs. The Allegro also showed similarly poor  
8 performance. It should be noted that both the Allegro and A1Care were evaluated in  
9 a previous study (data not presented) in which the results were acceptable, to good.  
10 As the data for certain elements of the current study was acceptable it is likely that  
11 there is an inconsistency in the manufacturing chain that needs to be identified. The  
12 considerable variability in performance across the analyzers, albeit less than in earlier  
13 studies, shows that there is still work to be done [6].

14 A key issue with POC analyzers still appears to be interference from variant  
15 hemoglobins. A complicating factor when evaluating Hb-variants is the fact that some  
16 analyzers (QuikReadgo and Innovastar) are not compatible with frozen samples. This  
17 study addresses this issue with the use of fresh Hb-variant samples. However it was  
18 not possible to obtain as wide a range or number of fresh Hb-variant samples for  
19 investigation as would be desirable. Explaining the interferences seen in these  
20 analyzers, which are nearly all immunoassay, is difficult, why would frozen and fresh  
21 samples perform so differently? Why are they causing an interference at all when the  
22 epitopes that the antibodies bind to do not contain the mutations that cause the Hb  
23 variant? HbE for example is an inherited single base mutation at codon 26 of the beta-  
24 globin gene, leading to substitution of glutamic acid (46 amino acid of the beta chain)  
25 for lysine which should not, theoretically, interfere with the antibodies used in the POC

1 analyzers. A possible explanation could be that the mutation causes folding of the  
2 hemoglobin molecule in such a way that position 46 is then very close to the first 4  
3 amino-acids of the beta chain and interacts with the antibodies used in the  
4 immunoassay [25]. Alternatively this may be due to differences in the immobilization  
5 of the antibodies which lead to differences in the surface chemistry and thus the  
6 binding of the antibody.

7 The Lab 001 suffers from interference with fresh HbAS samples, but less so with  
8 frozen samples which are easier to obtain for method development. This does  
9 potentially pose a problem for patient samples, however this is mitigated by the fact  
10 that Lab 001 as a capillary electrophoresis method is capable of identifying the  
11 presence of a variant – unlike most other POC analyzers.

12 It is important to clarify that some of the manufacturers do clearly state that the  
13 presence of variants may alter the HbA<sub>1c</sub> results however these claims and the findings  
14 of this study do not always correlate.

15

### 16 ***It is not all about analytical performance***

17 Whilst many evaluations focus on analytical performance, it is important to consider  
18 the wider context of the use of POC analyzers. The usability/user-friendliness of each  
19 of the analyzers was assessed and found to be variable.

20 A crucial factor in the practical, clinical usability of an instrument is how long it takes  
21 to generate a result, with the benefit of providing real time results often touted as a key  
22 selling point for POC analyzers. The time from a 'cold start' (turning the power on and  
23 warming reagents if needed) to a result was assessed. The range of time needed was  
24 wide at ~3.0 minutes for The Lab 001 to ~16.5 minutes for the InnovaStar. This is

1 important information for users of the analyzers as they may have little advance  
2 warning that a test may need to be undertaken.

3

#### 4 ***Key messages***

5 This complex and detailed evaluation provides a comprehensive overview of 6 HbA<sub>1c</sub>  
6 POC analyzers. Whilst there are areas of excellence in performance there are still  
7 significant areas for improvement with the performance of some being unacceptable.  
8 It is possible for an analyzer to meet certification criteria for the IFCC and/or NGSP  
9 and perform well in one evaluation and then perform very poorly in subsequent  
10 evaluations. From a clinical and scientific perspective this is alarming. It is essential  
11 that performance of an analyzer is stable, especially with increased use for both  
12 monitoring and diagnosis of people with diabetes. Four of the analyzers in this study  
13 showed highly variable performance which is not acceptable.

14 It is unclear why such discrepant results are seen when fresh or frozen samples are  
15 used, especially as this is not commonly seen with routine laboratory analyzers. Whilst  
16 many evaluations and method development often necessitate the use of frozen  
17 samples, it is rare that a POCT would be used with anything other than fresh samples.  
18 We have shown here that there can be marked differences in performance with each  
19 sample type.

20 One way to identify variability in performance is through the use of EQA schemes. The  
21 authors strongly advocate the use of EQA to identify ongoing performance issues, and  
22 although POC analyzers are often exempt from the need to participate in EQA it is a  
23 valuable and powerful tool for monitoring performance. A caveat to this is that POC  
24 analyzers may not be able to utilize the frozen or lyophilized samples often used in  
25 EQA schemes. None of the manufacturers claim in their information for users that

1 lyophilized samples can be used and this is supported by the data from the ERL-EQA  
2 samples (see figure 2), EQA program leads need to be cognoscente of this issue and  
3 work towards providing commutable samples for POC analyzers.

4 As discussed earlier, the interference from Hb-variants in a number of the analyzers  
5 is perplexing. The disparity in results seen between fresh and frozen samples is of  
6 concern as many manufacturers will likely develop their methods using frozen samples  
7 but in the 'real world' setting where fresh samples are used, the variants pose a  
8 potential unseen problem. Not all manufacturers are accurate in their claims for Hb-  
9 variant performance.

10



1 **Author Contributions** All authors confirmed they have contributed to the intellectual  
2 content of this paper and have met the following three requirements: (a) significant  
3 contributions to the conception and design, acquisition of data, or analysis and  
4 interpretation of data; (b) drafting or revising the article for intellectual content; and (c)  
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13

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19

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1 **Legends**

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3 **Figure 1:** EP-9 data for each of the POC analyzers when compared to the mean of  
4 the SRMPs with two lot numbers. Panel A - the A1Care, B - the Lab 001, C - the  
5 Cobas B101, D - the QuikReadgo, E - the InnovaStar, F - the Allegro

- Linear (X=Y)
- - - Linear (-10%)
- - - Linear (+10%)

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7 **Figure 2:** Sigma metrics graphs showing the impact of different sample types on the  
8 ability to meet the International quality performance criteria. Figure 2A shows EP- 15  
9 and EP- 9 (red) values compared to frozen (blue) and lyophilized samples (green).  
10 Figure 2B shows EP-5 (grey) data compared to frozen and lyophilized samples  
11 showing minimal difference in performance between EP-5 and EP-15 protocols.  
12 Panel A - the A1Care, B - the Lab 001, C - the Cobas B101, D - the QuikReadgo, E -  
13 the InnovaStar, F - the Allegro

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Analyzer	Manufacturer	Sample Volume (µl)	Analysis Time (min)	Time from a cold start to first result (min)	Method Principle	Weight (kg)	Dimensions (W x H x D mm)	Operating temperature	Storage temperature cartridges	Hb-variant visible?	Mean SUS score <sup>(a)</sup> N=2
A1Care	I-Sens	2.5	4.2	≈13.7	Enzymatic	3.8	290 x 250 x 130	10 - 32 °C	1 - 30 °C	No	88
Cobas B101	Roche Diagnostics	2	6.0	≈9.5	Immuno-assay	2.0	135 x 184 x 234	15 - 32 °C	2 - 30 °C	No	95
InnovaStar	DiaSys	10	6.5	≈16.5	Immuno-assay	4	200 x 150 x 170	15 - 35°C.	2 - 8 °C	No	80
The Lab 001	Arkray	1.5	1.5	≈3.0	Capillary electrophoresis	10	220 x 298 x 330	10 - 30 °C	2 - 30 °C	Yes	91
QuikReadgo	Aidian	1	6.0	≈7.4	Immuno-assay	1.7	145 x 155 x 270	15 - 35°C.	2 - 8 °C, 2 months at 18 -25 °C	No	60
Allegro	Nova Biomedical	1.5	6.5	≈9.2	Immuno-assay	9.1	203 x 381 x 381	15 - 32 °C	2 - 8 °C	No	84

<sup>a</sup>= System Usability Scale

**Table 1:** Analyzers included in the study and their key characteristics.

	CV (%) in SI units (sample value)	CV (%) in NGSP units (sample value)
<b>A1Care</b>		
EP-5	<b>3.5</b> (46.6 mmol/mol)	<b>2.4</b> (6.41%)
Frozen samples	2.7 (71.3 mmol/mol)	<b>2.1</b> (8.68%)
EP-15	<b>3.1</b> (46.3 mmol/mol)	1.9 (6.38%)
Fresh samples	1.9 (79.8 mmol/mol)	1.5 (9.44%)
Lot number A <sup>a</sup>	<b>3.2</b>	2.3
Lot number B <sup>a</sup>	2.9	2.0
<b>Cobas B101</b>		
EP-5	<b>3.1</b> (44.7 mmol/mol)	<b>2.2</b> (6.24%)
Frozen samples	1.7 (71.4 mmol/mol)	1.3 (8.69%)
EP-15	<b>3.6</b> (44.2 mmol/mol)	<b>2.3</b> (6.20%)
Fresh samples	1.2 (81.5 mmol/mol)	0.9 (9.61%)
Lot number A <sup>a</sup>	2.3	1.6
Lot number B <sup>a</sup>	1.6	1.1
<b>InnovaStar</b>		
EP-5	2.0 (48.7 mmol/mol)	1.3 (6.61%)
Frozen samples	2.4 (75.5 mmol/mol)	1.8 (9.06%)
EP-15	1.5 (45.1 mmol/mol)	1.0 (6.28%)
Fresh samples	1.4 (84.0 mmol/mol)	1.1 (9.84%)
Lot number A <sup>a</sup>	2.4	1.7
Lot number B <sup>a</sup>	1.5	1.0
<b>The Lab 001</b>		
EP-5	1.4 (46.1 mmol/mol)	0.9 (6.37%)
Frozen samples	1.8 (72.0 mmol/mol)	1.5 (8.74%)
EP-15	1.7 (45.4 mmol/mol)	1.1 (6.30%)
Fresh samples	1.1 (82.4 mmol/mol)	0.8 (9.69%)
Lot number A <sup>a</sup>	1.8	1.2
Lot number B <sup>a</sup>	1.6	1.1
<b>QuikReadgo</b>		
EP-5	2.5 (50.9 mmol/mol)	1.9 (6.79%)
Frozen samples	<b>3.0</b> (82.7 mmol/mol)	<b>2.1</b> (9.71%)
EP-15	2.5 (43.5 mmol/mol)	1.5 (6.14%)
Fresh samples	2.1 (86.8 mmol/mol)	1.8 (10.08%)
Lot number A <sup>a</sup>	<b>3.5</b>	<b>2.5</b>
Lot number B <sup>a</sup>	2.7	1.9
<b>Allegro</b>		
EP-5	<b>4.2</b> (44.3 mmol/mol)	<b>2.8</b> (6.23%)
Frozen samples	<b>4.1</b> (70.0 mmol/mol)	<b>2.8</b> (8.56%)
EP-15	<b>4.2</b> (43.6 mmol/mol)	<b>3.0</b> (6.16%)
Fresh samples	<b>3.8</b> (85.4 mmol/mol)	<b>3.0</b> (9.96%)
Lot number A <sup>a</sup>	<b>3.4</b>	<b>2.4</b>
Lot number B <sup>a</sup>	<b>3.6</b>	<b>2.6</b>

<sup>a</sup>based on the duplicates in EP-9

**Table 2:** Imprecision results based on EP-5, EP-15 and on the duplicates in EP-9. Red: fail performance targets (CV <3% in SI units and <2% in NGSP units).

	Bias at different HbA <sub>1c</sub> levels (mmol/mol)			
	30 mmol/mol (95% CI)	48 mmol/mol (95% CI)	75 mmol/mol (95% CI)	Pass NGSP criteria? X out of 4 SRMP*
<b>A1Care</b>				
Lot A vs mean SRMP	<b>31.7</b> (30.62 to 32.74) <sup>a</sup>	<b>47.9</b> (47.29 to 48.57)	<b>72.3</b> (70.92 to 73.69) <sup>a</sup>	
Lot B vs mean SRMP	<b>31.6</b> (30.42 to 32.73) <sup>a</sup>	<b>48.9</b> (48.15 to 49.60) <sup>a</sup>	<b>74.8</b> (73.67 to 75.98)	
Lot A vs Lot B	<b>30</b> (29.5 to 31.2)	<b>49</b> (48.3 to 49.4) <sup>*</sup>	<b>77</b> (75.5 to 77.5) <sup>a</sup>	
Lot A				2/4
Lot B				1/4
<b>Cobas B101</b>				
Lot A vs mean SRMP	<b>30.1</b> (28.81 to 31.47)	<b>47.6</b> (46.82 to 48.34)	<b>73.7</b> (72.95 to 74.55) <sup>a</sup>	
Lot B vs mean SRMP	<b>29.3</b> (28.29 to 30.29)	<b>47.2</b> (46.63 to 47.85) <sup>a</sup>	<b>74.2</b> (73.41 to 74.93) <sup>a</sup>	
Lot A vs lot B	<b>29.0</b> (28.0 to 30.7)	<b>48.0</b> (46.9 to 48.4)	<b>75.0</b> (74.2 to 76.2)	
Lot A				4/4
Lot B				4/4
<b>InnovaStar</b>				
Lot A vs mean SRMP	<b>30.4</b> (29.63 to 31.21)	<b>47.3</b> (46.88 to 47.81) <sup>a</sup>	<b>72.7</b> (71.57 to 73.87) <sup>a</sup>	
Lot B vs mean SRMP	<b>28.6</b> (27.73 to 29.52) <sup>a</sup>	<b>46.2</b> (45.68 to 46.80) <sup>a</sup>	<b>72.7</b> (71.88 to 73.45) <sup>a</sup>	
Lot A vs Lot B	<b>28.1</b> (27.73 to 28.53) <sup>a</sup>	<b>46.7</b> (46.49 to 46.94) <sup>a</sup>	<b>74.6</b> (74.04 to 75.13)	
Lot A				4/4
Lot B				2/4
<b>The Lab 001</b>				
Lot A vs mean SRMP	<b>29.7</b> (28.73 to 30.58)	<b>47.1</b> (46.58 to 47.55) <sup>a</sup>	<b>73.2</b> (72.30 to 74.06) <sup>a</sup>	
Lot B vs mean SRMP	<b>30.0</b> (29.44 to 30.60)	<b>47.4</b> (47.07 to 47.71) <sup>a</sup>	<b>73.4</b> (72.75 to 74.13) <sup>a</sup>	
Lot A vs Lot B	<b>29.9</b> (29.47 to 30.36)	<b>47.8</b> (47.57 to 48.08)	<b>74.7</b> (74.26 to 75.12)	
Lot A				3/4
Lot B				4/4
<b>QuikReadgo</b>				
Lot A vs mean SRMP	<b>28.4</b> (27.17 to 29.69) <sup>a</sup>	<b>46.5</b> (45.84 to 47.21) <sup>a</sup>	<b>73.7</b> (71.65 to 75.66)	
Lot B vs mean SRMP	<b>30.3</b> (28.70 to 31.92)	<b>48.3</b> (47.30 to 49.36)	<b>75.4</b> (73.79 to 76.93)	
Lot A vs Lot B	<b>32.0</b> (30.7 to 33.0) <sup>a</sup>	50.0 (49.1 to 50.5) <sup>a</sup>	<b>77.0</b> (74.8 to 78.6)	
Lot A				0/4
Lot B				0/4
<b>Allegro</b>				
Lot A vs mean SRMP	<b>32.4</b> (30.89 to 34.00) <sup>a</sup>	<b>50.0</b> (49.01 to 51.01) <sup>a</sup>	<b>76.4</b> (74.27 to 78.45)	
Lot B vs mean SRMP	<b>30.3</b> (28.48 to 32.03)	<b>48.4</b> (47.22 to 49.56)	<b>75.6</b> (74.06 to 77.12)	
Lot A vs Lot B	<b>28.0</b> (26.5 to 29.0) <sup>a</sup>	<b>46.0</b> (45.6 to 47.1) <sup>a</sup>	<b>74.0</b> (72.8 to 75.5)	



Lot A				0/4
Lot B				0/4

<sup>a</sup> Statistically significant differences were observed for bias when 30 and/or 48 and/or 75 mmol/mol are not within 95% confidence interval limits.

**Table 3:** Bias at different HbA<sub>1c</sub> levels for each of the different POC analyzers using two different lot numbers, compared with the mean of the SRMPs. Bias between the two lot numbers. Results for NGSP show how many times the criteria were met out of a possible 4 comparisons.

	<b>HbAS</b> (frozen n=20, fresh n=16)	<b>HbAC</b> (frozen n=20, fresh n=7)	<b>HbAD</b> (frozen n=20, fresh n=5)	<b>HbAE</b> (frozen n=20, fresh n=9)	<b>A2</b> (frozen n=15, fresh n=0)	<b>HbF<sup>a</sup></b> (frozen n=15, fresh n=4)	<b>Manufacturer claims for interference if present</b>
<b>A1Care</b>							
Frozen	2.1	1.8	-0.7	5.5	2.9	>9.3%	Hb F >10%
Fresh	-0.4	-1.1	-2.2	4.7			
<b>Cobas B101</b>							
Frozen	-0.1	0.2	<b>9.9</b>	<b>11.1</b>	6.3	>9.5%	None
Fresh	-0.1	1.8	6.2	<b>10.9</b>			
<b>InnovaStar</b>							
Frozen	1.3	<b>9.5</b>	3.3	1.8	2.8	>5.4%	AE and elevated F
Fresh	-2.2	<b>14.0</b>	4.5	7.1			
<b>The Lab 001</b>							
Frozen	-6.6	-6.1	-0.3	-1.1	0	>42%	Hb F >30%
Fresh	<b>-10.1</b>	-0.5	-0.5	-3.1			
<b>QuikReadgo</b>							
Frozen	2.0	<b>18.5</b>	<b>-11.5</b>	-2.2	<b>-10.8</b>	>3.5%	AC and F >7%
Fresh	-1.7	<b>22.1</b>	-6.1	5.7			
<b>Allegro</b>							
Frozen	<b>-9.5</b>	7.0	<b>1.1</b>	6.2	-1.7	>3.5%	None
Fresh	<b>-10.1</b>	8.5	<b>9.2</b>	7.0			

<sup>a</sup>= % of HbF at which a significant negative bias results

**Table 4** Mean relative difference (%) of the common Hb-variants compared to the assigned value after correction for bias in non-variant samples (number of samples for frozen and fresh samples). Red: equals at or near 10% difference. Green: this variant would not be seen to affect the value if only evaluated using frozen samples. The manufacturer's claims for are also listed.