Rapid Centrifugation in the Routine Hemostasis Laboratory

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► centrifugation
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Abstract

Background The use of short and uniform centrifugation schemes contributes significantly to the successful automation of laboratory procedures. It is however unclear if this is applicable to the hemostasis laboratory.

Objectives This article assesses the accuracy of measurements obtained with a rapid, high-speed centrifugation scheme in a large set of hemostasis tests, covering the full spectrum of values obtained in clinical practice, and using meaningful statistical measures.

Methods Two citrated plasma samples were obtained from consecutive patients of a tertiary hospital with suspected abnormal hemostasis tests and processed with two centrifugation schemes in parallel: 1,500 × g for 10 minutes and 3,137 × g for 7 minutes. The following tests were conducted: prothrombin time (n = 125), international normalized ratio (n = 146), activated partial thromboplastin time (n = 119), thrombin time (n = 105), fibrinogen (n = 125), factor (F)II (n = 69), FV (n = 64), FVII (n = 64), FX (n = 67), FVIII (n = 55), FIX (n = 37), FXI (n = 35), and FXIII (n = 20), D-dimer (n = 34), antithrombin (n = 31), anti-Xa activity (n = 30), von Willebrand antigen (n = 25), and von Willebrand activity (VWF:GPIbM; n = 27).

Results A wide range of results were obtained in all tests. Spearman’s rank correlation coefficient was at least 0.95 for all tests except FV, FIX, and FXI. The coverage probability π at a given deviation index κ of 15% was above 0.9 for all tests except FV, FVII, FX, FVIII, FIX, FXI, and VWF:GPIbM, suggesting a lack of agreement.

Conclusion Our results suggest that high-speed centrifugation is applicable to the majority of routine hemostasis parameters. The coverage probability was more sensitive than Spearman’s rank correlation to detect disagreement among centrifugation schemes.

These authors contributed equally to this study.


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**Introduction**

Total laboratory automation (TLA) is seen as an important strategy to tackle today’s challenges in laboratory medicine and related health care. Integrating diverse analyzers of multiple specialities as well as preanalytic and sorting modules to one track can replace many manual tasks and save labor. TLA can save a relevant amount of long-term health care costs despite high initial investments. The efficiency can be improved by reducing the turnaround time (TAT) and increasing the throughput. Risky and laborious manual sample management can be avoided by algorithms identifying samples and triggering rerun or reflex tests, thus improving accuracy of results. Controlling all phases of the analytical process using software programs run centrally enhances standardization, supports accreditation procedures, and improves quality in general. One important requirement for the integration of analyzers and specialities into TLA is uniform and ideally rapid centrifugation schemes. This might prevent delay of samples on the track and improve total TAT.

It is however uncertain whether rapid centrifugation schemes—used in clinical chemistry as well as immunochemistry—can be applied in the hemostasis laboratory. Coagulation-based assays are considered to be influenced by many preanalytic factors. The residual platelet count after centrifugation is regarded as the most important influencing factor and the platelet count is subject to the centrifugal forces applied. Mechanical stimulation is discussed as a contributing factor as well. Some authors recommend avoiding centrifugal forces above $1,500 \times g$ but others do not. Several scientific societies in the United States (Clinical and Laboratory Standards Institute [CLSI]) and the United Kingdom (British Committee for Standards in Haematology) as well as other authors have released guidance documents and recommend centrifugation schemes that aim to achieve a platelet count below $10 \times 10^9/L$ using $1,500 \times g$ for 10 to 15 minutes. Whether this can be accelerated using higher centrifugal forces remains unclear. Limitations of previous investigations include (1) a limited number of patients or samples, (2) a paucity of hemostasis tests studied, (3) a very limited spectrum of test results covered (normal only), (4) nonstandard centrifugation schemes as comparators, and (5) use of insensitive statistical measures.

With the present study, we aimed to comprehensively assess the accuracy of a rapid, high-speed centrifugation scheme compared with an established and recommended centrifugation scheme in a comprehensive set of hemostasis tests, covering a large spectrum of values, and using meaningful statistical measures.

**Methods**

**Study Design, Setting, and Population**

This prospective cohort study was conducted in a university hospital (Inselspital, Bern University Hospital, Bern, Switzerland) between October 2015 and March 2016. We included consecutive patients from a variety of departments (cardiovascular surgery, internal medicine, intensive care unit, and hemophilia outpatient unit). The inclusion criterion was “suspected abnormal hemostasis laboratory test result” to obtain a sample set covering a broad spectrum of results. Exclusion criteria were an age below 18 years and refused general consent. Patients were screened by treating physicians in the respective departments. The protocol was approved by the local ethics committee (Kantonale Ethikkommission Bern; March 18, 2014) and all patients signed the general consent of the hospital.

**Collection and Handling of Samples**

An in-house protocol for blood drawing was implemented at our institution to ensure adequate preanalytic conditions. Plasma samples were collected after ethylenediaminetetraacetic acid (EDTA) and serum samples to avoid contamination with residual heparin or tissue factor if blood was accidentally obtained from a vascular access device or using a butterfly needle. Venous blood was drawn into two plastic syringes of 4.5 mL containing 0.106 mol/L trisodium citrate (Monovette, Sarstedt, Nümbrecht, Germany). Samples were transported immediately to the central laboratory and the handling of both samples was identical.

**Centrifugation Schemes**

Both plasma samples were centrifuged within 15 minutes after arrival in the laboratory. The established standard centrifugation scheme in our institution was $1,500 \times g$ for 10 minutes (established after determining residual platelet count in 30 samples). The high-speed centrifugation scheme was defined as $3,137 \times g$ for 7 minutes ($3,137 \times g$ is the established speed in clinical chemistry at our institution and the time of 7 minutes was chosen after conducting a pilot study analyzing residual platelet concentration). The centrifuges were used (HERMLE Labortecnik GmbH, Wehingen, Germany), implemented in a GLP laboratory track (GLP Systems GmbH, Hamburg, Germany). The centrifuges work at $20°C$ using a brake. After centrifugation, samples were transferred into aliquots and snap-frozen at $-80°C$. Samples were transferred to $-80°C$ if not analyzed within 1 to 2 weeks. All hemostasis parameters were determined within 2 months following centrifugation.

**Determination of Laboratory Assays**

Platelet count was analyzed using an automated hematology analyzer (Coulter Counter LH750, Beckman-Coulter Inc., Nyon, Switzerland). Prothrombin time was measured using Dade Innovin reagent (Siemens Healthcare Diagnostics Products GmbH, Marburg, Germany) on a Sysmex CS 5100 analyzer (distributed by Siemens Healthcare Diagnostics Products GmbH) and expressed as prothrombin ratio (PR, %). Activated partial thromboplastin time (aPTT) was determined using Pathrombin SL (Siemens Healthcare Diagnostics Products GmbH) on a Sysmex CS 5100. Thrombin time was determined using Thromboclotin reagent (Siemens Healthcare Diagnostics Products GmbH) on a Sysmex CS 5100. D-dimers were measured immunoturbidimetrically by Innovaone D-Dimer.
(Siemens Healthcare Diagnostics Products GmbH) on a Sysmex CS 5100. Functional antithrombin activity was determined using INNOVANCE Antithrombin (Siemens Healthcare Diagnostics Products GmbH) on a Sysmex CS 5100. Fibrinogen was assessed in plasma by the Clauss method using Dade Thrombin Reagent (Siemens Healthcare Diagnostics Products GmbH) on a Sysmex CS 5100. Factor (F)II, FV, FVII, and FX were measured by a one-stage coagulometric assay using Dade Innovin and factor-deficient plasma (Siemens Healthcare Diagnostics Products GmbH) on a Sysmex CS 5100. FVIII, FIX, and FXI were determined using a one-stage coagulometric assay using Pathrombin SL and factor-deficient plasma on a Siemens BCS XP analyzer (Siemens Healthcare Diagnostics Products GmbH). FXIII was additionally created. Coefficients of variation were calculated for within-run and day-to-day imprecision. Analyses were repeated 10 times each.

**Statistical Analysis**

Median (interquartile range, IQR) and numbers (percentages) were used for descriptive purposes. To analyze concordance between the two centrifugation schemes, we determined the Spearman's rank correlation coefficient as well as the coverage probability (CP π). The CP π outlines agreement with several advantages compared with more traditional measures: the computation is simple, discordant measurements can be identified rapidly, the interpretation is intuitive, and it can be applied to continuous as well as categorical data in a broad spectrum of observers and populations. The CP π can be interpreted as the proportion of observations falling within the paired absolute difference k, a priori defined as acceptable (the tolerable total deviation index, TDI). In our study, we defined a TDI of 0.15 as acceptable. The tolerable CP π was assumed at 0.9. Bland–Altman difference plots for determination of bias as well as lower and higher limits of agreement were additionally created. Coefficients of variation were calculated for within-run and day-to-day imprecision. Analyses were restricted to observations in a meaningful range: FII, 0 to 110%; international normalized ratio (INR), 0.9 to 5.0; thrombin time, 10 to 70 seconds; aPTT, 15 to 100 seconds; fibrinogen, 0 to 6.0 g/L; D-dimer, 0 to 2,000 µg/L; antithrombin, 0 to 120%; FII, FV, FVII, FX: 0 to 120%; FVIII, FIX, FXI, FXIII: 0 to 200%; and VWF: GPIbM and VWF:Ag: 0 to 200%. The CP π was observed for fibrinogen, D-dimer, and antithrombin activity measurements: rπ was above 0.95, CP π above 0.9 (see Table 1). Bias and limits of agreement of the Bland–Altman difference plot were close to zero. According to the routine parameters PT, INR, thrombin time, and aPTT, correlation (rπ) was clearly above 0.95 and CP π above 0.9 (Fig. 1). Bias and limits of agreement of the Bland–Altman difference plot were close to zero. According to the normal range for within-run and day-to-day imprecision of fibrinogen and D-dimer: 6.7%.

**Concordance between Centrifugation Schemes**

All results are reported in detail in Table 1. Scattergrams of PR, aPTT, fibrinogen, and D-dimer are shown as an example in Fig. 2. Scattergrams and Bland–Altman difference plots for all tests are shown in Supplementary Figs. S1–S3 (available in the online version). Fig. 3 summarizes the accuracy of all tests and contrasts rπ to CP estimates.

With regard to the routine parameters PT, INR, thrombin time, and aPTT, correlation (rπ) was clearly above 0.95 and CP π above 0.9 (Fig. 1). Bias and limits of agreement of the Bland–Altman difference plot were close to zero. Accordingly, coefficient of variation (CV) was below 5% with regard to within-run and day-to-day imprecision (day-to-day imprecision of the abnormal sample: 6.7%).

**Platelet count**

Platelet counts in plasma after centrifugation with 3,137 × g for 7 minutes was 8 × 10^9/L (IQR: 4–12 × 10^9/L; n = 30). Platelet count was 6 × 10^9/L (IQR: 3–9; n = 30) in samples centrifuged at 1,500 × g for 10 minutes.

**Results**

**Patients**

Samples from 198 patients were analyzed, the flow is illustrated in Fig. 1. Seventy-four patients were recruited from hematology outpatient unit (37%), 51 patients from intensive care unit (26%), 48 patients from general internal medicine (24%), and 25 patients from the surgery department (13%). The median age of the patients was 56.0 years (range: 19.7–94.4) and 47.5% were female (n = 94). The number of observations per test is shown in Table 1. The range of values is reported in Supplementary Figs. S1–S3, available in the online version.

**Concordance between Centrifugation Schemes**

All results are reported in detail in Table 1. Scattergrams of PR, aPTT, fibrinogen, and D-dimer are shown as an example in Fig. 2. Scattergrams and Bland–Altman difference plots for all tests are shown in Supplementary Figs. S1–S3 (available in the online version). Fig. 3 summarizes the accuracy of all tests and contrasts rπ to CP estimates.

With regard to the routine parameters PT, INR, thrombin time, and aPTT, correlation (rπ) was clearly above 0.95 and CP π above 0.9 (Fig. 1). Bias and limits of agreement of the Bland–Altman difference plot were close to zero. Accordingly, coefficient of variation (CV) was below 5% with regard to within-run and day-to-day imprecision (day-to-day imprecision of the abnormal sample: 6.7%).

**Platelet count**

The median platelet count in plasma after centrifugation with 3,137 × g for 7 minutes was 8 × 10^9/L (IQR: 4–12 × 10^9/L; n = 30). Platelet count was 6 × 10^9/L (IQR: 3–9; n = 30) in samples centrifuged at 1,500 × g for 10 minutes.
Accordingly, the agreement interval of the Bland–Altman plot was wide. With regard to FII, $r_s$ was 0.97 and CP $\pi$ 0.95. Within-run as well as day-to-day imprecision was low (CV < 5% for FVII, and < 10% for FII, FV, and FX).

In the case of FVIII, $r_s$ was 0.96 but CP $\pi$ was 0.30; $r_s$ was below 0.95 and CP $\pi$ below 0.9 in the case of FIX. The agreement interval of the Bland–Altman plot was wide (~Table 1). Within-run as well as day-to-day imprecision was below 10%.

With regard to FXIII, $r_s$ was 0.96 and CP $\pi$ 0.96; CV was below 10%.

Discussion

In the present study, we tested the accuracy of measurements obtained with rapid, high-speed centrifugation scheme studying a large spectrum of hemostasis tests and covering the full range of values obtained in clinical practice.
### Table 1
Concordance results of routine hemostasis measurements obtained with a rapid, high-speed centrifugation scheme in comparison to a traditional scheme.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Obs.</th>
<th>Results</th>
<th>Correlation coefficient</th>
<th>Coverage probability na</th>
<th>Bland–Altman difference plot</th>
<th>Analytical variability b</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>Numbers</td>
<td>Median (range)</td>
<td>(95% CI)</td>
<td>(95% CI)</td>
<td>(95% CI)</td>
</tr>
<tr>
<td>PR (%)</td>
<td>125</td>
<td>78.5 (6.6, 107.8)</td>
<td>0.99 (0.98, 0.99)</td>
<td>0.99 (0.84, 1.00)</td>
<td>0.2 (–0.3, 0.7)</td>
<td>−5.6</td>
</tr>
<tr>
<td>INR</td>
<td>146</td>
<td>1.04 (0.9, 3.5)</td>
<td>0.99 (0.99, 0.99)</td>
<td>1.00 (0.96, 1.00)</td>
<td>0.0 (–0.0, 0.0)</td>
<td>−0.2</td>
</tr>
<tr>
<td>Thrombin time (s)</td>
<td>105</td>
<td>16.5 (13.2, 57.3)</td>
<td>0.97 (0.95, 0.98)</td>
<td>1.00 (1.00, 1.00)</td>
<td>−0.1 (–0.3, 0)</td>
<td>−1.6</td>
</tr>
<tr>
<td>aPTT (s)</td>
<td>119</td>
<td>34.9 (22.0, 96.0)</td>
<td>0.99 (0.99, 0.99)</td>
<td>1.00 (1.00, 1.00)</td>
<td>0.0 (–0.2, 0.2)</td>
<td>−2.1</td>
</tr>
<tr>
<td>Fibrinogen (g/L)</td>
<td>125</td>
<td>2.3 (0.0, 5.8)</td>
<td>0.98 (0.97, 0.99)</td>
<td>0.96 (0.91, 0.99)</td>
<td>−0.0 (–0.0, 0.0)</td>
<td>−0.3</td>
</tr>
<tr>
<td>D-dimer (μg/L)</td>
<td>34</td>
<td>395 (17.1970)</td>
<td>0.99 (0.99, 1.00)</td>
<td>1.00 (0.99, 1.00)</td>
<td>13.4 (3.5, 23.4)</td>
<td>−43.4</td>
</tr>
<tr>
<td>Antithrombin (%)</td>
<td>31</td>
<td>76.6 (40.4, 108.5)</td>
<td>0.99 (0.99, 1.00)</td>
<td>1.00 (0.99, 1.00)</td>
<td>0.2 (–0.7, 1.1)</td>
<td>−4.7</td>
</tr>
<tr>
<td>Anti-Xa activity (UFH; U/mL)</td>
<td>30</td>
<td>0.10 (0.01, 0.92)</td>
<td>0.99 (0.98, 1.00)</td>
<td>0.94 (0.86, 0.99)</td>
<td>0.00 (–0.01, 0.01)</td>
<td>−0.03</td>
</tr>
<tr>
<td>Anti-Xa activity (LMWH; U/mL)</td>
<td>27</td>
<td>0.06 (0.00, 0.51)</td>
<td>0.99 (0.97, 0.99)</td>
<td>0.93 (0.77, 1.00)</td>
<td>0.00 (–0.01, 0.01)</td>
<td>−0.03</td>
</tr>
<tr>
<td>Factor II (%)</td>
<td>69</td>
<td>84.0 (16.0, 119.0)</td>
<td>0.97 (0.96, 0.98)</td>
<td>0.95 (0.81, 1.00)</td>
<td>0.6 (–0.6, 1.8)</td>
<td>−9.2</td>
</tr>
<tr>
<td>Factor V (%)</td>
<td>64</td>
<td>85.0 (0.0, 120.0)</td>
<td>0.92 (0.86, 0.95)</td>
<td>0.41 (0.20, 0.65)</td>
<td>−1.6 (–4.3, 1.1)</td>
<td>−23.3</td>
</tr>
<tr>
<td>Factor VII (%)</td>
<td>64</td>
<td>65.3 (15.0, 120.0)</td>
<td>0.95 (0.92, 0.97)</td>
<td>0.57 (0.36, 0.84)</td>
<td>−1.1 (–1.2, 0.9)</td>
<td>−17.5</td>
</tr>
<tr>
<td>Factor X (%)</td>
<td>67</td>
<td>75.1 (5.0, 118.8)</td>
<td>0.95 (0.91, 0.97)</td>
<td>0.60 (0.46, 0.74)</td>
<td>−0.4 (–2.4, 1.7)</td>
<td>−17.2</td>
</tr>
<tr>
<td>Factor VIII (%)</td>
<td>55</td>
<td>79.5 (0.3, 198.0)</td>
<td>0.96 (0.93, 0.98)</td>
<td>0.30 (0.21, 0.38)</td>
<td>−17.5 (–22.3, −12.7)</td>
<td>−53.0</td>
</tr>
<tr>
<td>Factor IX (%)</td>
<td>37</td>
<td>104.0 (15.0, 138.0)</td>
<td>0.84 (0.70, 0.91)</td>
<td>0.79 (0.61, 0.90)</td>
<td>−0.9 (–4.5, 2.7)</td>
<td>−22.5</td>
</tr>
<tr>
<td>Factor XI (%)</td>
<td>35</td>
<td>96.0 (47.0, 163.0)</td>
<td>0.79 (0.61, 0.89)</td>
<td>0.80 (0.68, 0.90)</td>
<td>−1.3 (–4.8, 2.2)</td>
<td>−21.6</td>
</tr>
<tr>
<td>Factor XIII (%)</td>
<td>20</td>
<td>110.5 (15.1, 162.0)</td>
<td>0.96 (0.89, 0.98)</td>
<td>0.96 (0.82, 0.99)</td>
<td>−2.5 (–5.9, 1.0)</td>
<td>−17.0</td>
</tr>
<tr>
<td>VWF:Ag (%)</td>
<td>25</td>
<td>87.0 (12.0, 191.3)</td>
<td>0.99 (0.97, 0.99)</td>
<td>0.96 (0.83, 0.99)</td>
<td>−1.2 (–2.9, 0.5)</td>
<td>−9.4</td>
</tr>
<tr>
<td>VWF:GPIbM (%)</td>
<td>27</td>
<td>84.0 (7.0, 178.9)</td>
<td>0.98 (0.95, 0.99)</td>
<td>0.76 (0.59, 0.89)</td>
<td>−3.8 (–6.6, −1.0)</td>
<td>−18.1</td>
</tr>
</tbody>
</table>

Abbreviations: aPTT, activated partial thromboplastin time; CV, coefficient of variation; HLA, higher limit of agreement; INR, international normalized ratio; LLA, lower limit of agreement; LMWH, low molecular weight heparin; Obs., observations; PR, prothrombin ratio; r_s, Spearman’s rank correlation coefficient; UFH, unfractionated heparin; VWF:Ag, von Willebrand factor antigen; VWF:GPIbM, von Willebrand factor functional activity as measured using the spontaneous binding of VWF to a gain-of-function mutant GPIb fragment.

*Based on an accepted total deviation index of 0.15.

Obtained in two samples, one with results in the normal range [n] and one with results in the abnormal range [p], 10 times each.

CP π identified more assays with nonconcordant results than Spearman’s correlation coefficient. Whereas an appropriate concordance between measurements were observed for all routine parameters, relevant discrepancies were observed with regard to most single-factor determinations as well as VWF:GPIbM.

Our observations and data reported here are in line with the limited data published to date on this topic. Lippi et al compared samples obtained in 15 healthy volunteers and centrifuged with 3,000 × g versus 1,500 × g. Significant differences were found with regard to FVIII and IX, but not for PT, aPTT, or fibrinogen. Several routine tests (fibrinogen, Thrombosis and Haemostasis Vol. 119 No. 12/2019
INR, aPTT, D-dimer) as well as single factors have also been studied (1,500 × g for 5 minutes vs. 3,000 × g for 7 minutes) in a limited number of patients with mostly normal test results and a correlation coefficient of all routine tests as well as most factors studies was above 0.9. Suchsland et al have compared PT, aPTT, FVIII, and protein S in one versus two runs of 3,280 × g for 5 minutes. The outcome of that work reported a correlation coefficient above 0.9 for all but protein S. Boissier et al compared a centrifugal spin protocol of 4,500 × g for 2 minutes to a 2,200 × g for 15 minutes protocol in 200 patients in the following assays PT, INR, aPTT, fibrinogen, thrombin time, FII, FV, antithrombin, and anti-Xa activity. These authors concluded that the agreement was adequate. We have also essentially repeated the Boissier centrifugation protocols in samples of patients treated with direct oral anticoagulations and measurements of anti-Xa activity, and we have arrived at similar conclusions. In another study, PT, aPTT, and INR measurements were compared in platelet-poor plasma and platelet-rich plasma of 100 samples, and again found to be similar.

**Strengths and Limitations/Implications**

In contrast to previous studies, we have studied a broad range of hemostasis tests covering the full spectrum of test results in a large number of patients recruited in clinical practice. In addition, an established centrifugation scheme was used, and sensitive statistical measures were applied. Thus, our results are meaningful and are applicable to routine clinical practice. Our study nevertheless has several limitations. First, lupus anticoagulant and other tests used in clinical practice were not studied, and our results can therefore not be extrapolated to these tests. Second, even though our routine centrifugation scheme (1,500 × g for 10 minutes), which was used as comparator, was tested according to the CLSI guidelines, most institutions are known to use a 15-minute scheme instead. We cannot fully exclude that this might have affected our results. Third, only one centrifuge was tested, and we have not excluded that different results might have been obtained using alternative centrifuges. Thus, future investigations will seek to confirm our results in various settings using different centrifuges as well. Forth, citrated samples were...
collected after EDTA samples and we cannot fully exclude that residual EDTA might have influenced the results. We believe that the risk of a bias is low because the tubes were applied to the centrifugation schemes randomly and previous studies did not demonstrate a relevant difference of standard coagulation test results. Fifth, for reasons of practicability not all pairs of samples were analyzed within the 2-week interval recommended by current guidelines but were transferred to a −80°C freezer after several days and analyzed within 2 months. However, all pairs were handled identically, and we do not think that this might have introduced any bias. In addition, it might be argued that double centrifugation decreases preanalytical influences in case of freezing. However, aiming to compare two different (single) centrifugation schemes we did not apply double centrifugation for the purpose of this study. We do not believe that this might have introduced any bias. Seventh, plasma pathologically enriched with soluble fibrin might entrap VWF/FVIII by its complexes and technically diminish these levels under low temperature conditions. The risk of a systematic bias is low however, because both pairs of samples were treated identically.

We believe that our results add sufficient data on previous studies to justify rapid centrifugation for routine hemostasis tests (PT, thrombin time, aPTT, fibrinogen, D-dimer, antithrombin, and anti-Xa activity). Traditional centrifugation schemes shall be used for all other tests. The problem of sample identification can be solved by means of different adhesive labels (“routine” vs. “specialized hemostasis”) or by means of laboratory automation procedures as done in our institution (samples with certain tests requested will be put out of the laboratory track for “manual centrifugation”).

Fig. 3 Accuracy of routine hemostasis measurements obtained with a rapid, high-speed centrifugation scheme. Spearman’s rank correlation coefficient ($r_s$) is contrasted to the coverage probability (CP $\pi$) at a given acceptable total deviation index $\kappa$ of 0.15. The acceptable $r_s$ was set at 0.95, and the acceptable CP $\pi$ at 0.9. aPTT, activated partial thromboplastin time; INR, international normalized ratio; LMWH, low molecular weight heparin; PR, prothrombin ratio; UFH, unfractionated heparin; VWF:Ag, von Willebrand factor antigen; VWF:GPIbM, von Willebrand factor activity.
Conclusion

We tested the accuracy of measurements obtained with a rapid, high-speed centrifugation scheme studying a large spectrum of hemostasis tests and covering the full spectrum of values obtained in clinical practice. The new agreement measure CP identified more assays with nonconcordant results than Spearman’s correlation coefficient. A clinically appropriate concordance between measurements was observed for all routine parameters, but relevant discrepancies were observed with regard to most single-factor determinations as well as von Willebrand activity. We believe that our results add sufficient data on previous studies to justify rapid centrifugation for routine hemostasis tests (PT, thrombin time, aPTT, fibrinogen, D-dimer, antithrombin, and anti-Xa activity). A traditional centrifugation scheme shall be used for all other tests.

What is known about this topic?

- Short and uniform centrifugation schemes contribute significantly to successful laboratory automation.
- To date, it is unclear if this is applicable to the hemostasis laboratory.

What does this paper add?

- We assessed a rapid, high-speed centrifugation scheme in a large set of hemostasis tests, covering the full spectrum of values obtained in clinical practice, and using meaningful statistical measures.
- We demonstrated that high-speed centrifugation is applicable to the majority of routine hemostasis parameters.
- The coverage probability was more sensitive than Spearman’s rank correlation to detect disagreement among centrifugation schemes.

Acknowledgments

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References

18. CLSI. Collection, Transport, and Processing of Blood Specimens for Testing Plasma-Based Coagulation Assays and Molecular Hemostasis.