

Biological Variation in haemostasis variables

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Outline presentation

- **Introduction**
- **Aim**
- **Setup study**
- **Results**
- **Conclusion**



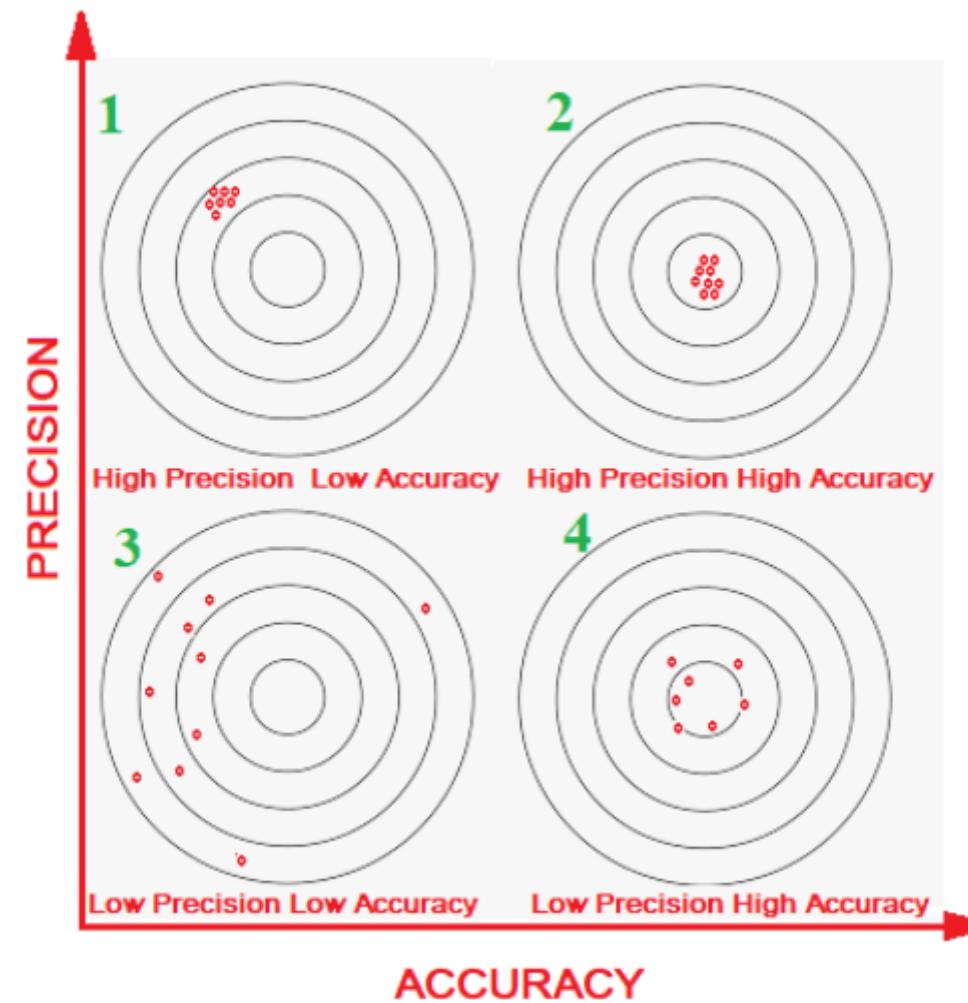


What are the Analytical Performance Characteristics?

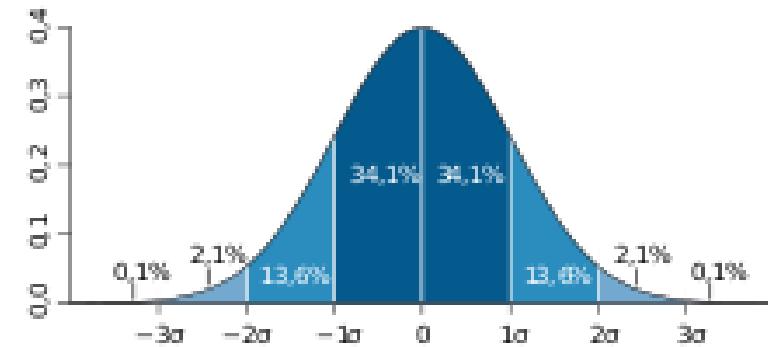
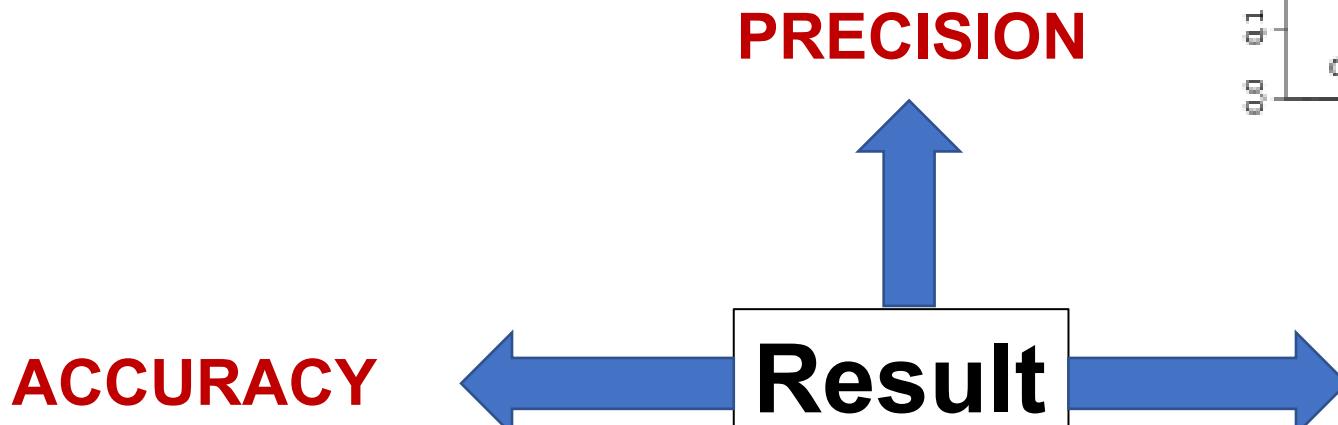
- **Correctness of measurements is defined by analytical performance specifications**



Precision and Accuracy



Correct test result



Every test result is subject to systematic (bias) and random errors (imprecision)



Three models to set analytical performance specification

1. Based on effect of analytical performance on clinical outcomes
2. Based on components of biological variation of the variable
3. Based on state-of-the-art: relates to the highest level of technically available performance of an assay

➤ **Biological variation model is most widely used**



Definitions



Within-subject biological variation (CV_I)

Random fluctuation around a homeostatic set point in an individual



Between-subject biological variation (CV_G)

Difference between the homeostatic set points of different individuals

The equations for desirable performance goals based on the biological variation

$$CV_A < 0.5 CV_I$$

$$B_A < 0.25 (CV_I^2 + CV_G^2)^{1/2}$$

$$TE_A < 0.25 (CV_I^2 + CV_G^2)^{1/2} + 1.65 (0.5 CV_I)$$



Historical online BV database last updated in 2014

	Analyte	Number of papers	Biological Variation		Desirable specification		
			CV _I	CV _g	I(%)	B(%)	TE(%)
P-	Antithrombin III	4	5.2	15.3	2.6	4.0	8.3
S-	D-Dimer (MoM)	1	23.3	26.5	11.65	8.82	28.04
P-	Factor V coagulation	1	3.6	--	1.8	--	--
P-	Factor VII coagulation	2	6.8	19.4	3.4	5.1	10.7
P-	Factor VIII coagulation	2	4.8	19.1	2.4	4.9	8.9
P-	Factor X coagulation	1	5.9	--	3.0	--	--
P-	Fibrinogen	5	10.7	15.8	5.4	4.8	13.6

Reference: <https://www.westgard.com/biodatabase1.htm>



Biological variation (BV) data haemostasis variables

	CV _I			CV _G		
	minimum	median	maximum	minimum	median	maximum
PT	2.3	2.6	5.8	4.0	4.9	6.8
APTT	1.7	3.3	6.8	7.1	7.8	8.9
Fibrinogen	6.8	11.5	18.6	14.7	16.4	20.2
AT	1.1	3.1	5.7	2.6	7.8	10.4

Median and range (minimum and maximum value) based on the results of the combined studies.
CV_I: within-subject coefficient of variation, CV_G: between-subject coefficient of variation

➤ Large heterogeneity in BV data due to heterogeneity in study setup



Sigma Score

$$\text{sigma} = \frac{(TE_A - \text{bias}_A)}{CV_A}$$

- The higher the sigma score, the better the analytical performance of the test



Sigma scores based on CV_A of internal QC data and applying the minimum, median and maximum BV estimates

	CV_A	Sigma		
		Min.	Median	Max.
PT	1.6	1.9	2.2	4.3
APTT	1.9	1.7	2.5	4.3
Fibrinogen	2.7	3.6	5.4	8.3
AT	5.6	0.3	0.8	1.4

CV_A: Analytical variation of an internal QC sample.
Yellow cells represent a sigma score above 3.

- The application of the six-sigma method to BV-derived APS is hampered by the large variation in published BV data



Aim

To generate up-to-date within-subject (CV_I) and between-subject (CV_G) BV data for coagulation and fibrinolytic variables by systematically appraising published BV data and subsequently combine the data in meta-analysis



Setup study

- Literature search including BV variables in healthy adults, study period at least 1 week and three or more samples collected per person
- Appraising publications by the Biological Variation Data Critical Appraisal Checklist (BIVAC):
 - 4 independent assessors reviewed the papers
 - 14 quality items graded A to D
- D-graded studies were excluded from the meta-analysis, data was considered not fit for use
- Meta-analysis was performed with the BIVAC grades given weights [A=4, B=2, C=1]



The Biological Variation Data Critical Appraisal Checklist

Table 1. BIVAC with criteria for achieving A, B, C, and D scores for the different quality items and their rationale. (Continued from page 503)

QI	Quality question	Quality scoring				Rationale
		A	B	C	D ^a	
3: Samples	Are the following documented? (a) Number of samples collected (b) Type of sample material used (c) Timing of sample collections (d) Length of study period	Yes	(a), (c), and (d) are documented. (b) Sample material; insufficient detail is given, but this is not of importance for the measurand in question.	(a), (c), and (d) are documented, but no information or insufficient detail on (b) sample material is given, with consequences for the measurand in question.	(a) Number of samples, (c) timing of sample collections, and/or (d) length of study period are not presented or deducible.	Necessary to characterize how and in what material the study has been performed and to evaluate if variation in timing gives different estimates.
4: Measurand	Are the measurand and the measurement procedure documented?	Yes A detailed method description is presented Or A reference to article where the method is described in detail is provided, Or An identifiable method has been applied and is described with sufficient detail. ^b	Insufficient detail on the method is given, but it is not of importance for the measurand in question.	Insufficient detail on the method is given or an outdated method has been used, which may be of importance for the measurand in question.	The method is obsolete and no longer valid, i.e., methods in use today estimate another measurand, or the method is not fit for the purpose of estimating biological variation in the chosen population.	Adequate description of the measurand and measurement procedure is necessary to ensure transferability of data.

Reference: Aarsand et al., CC (2018) 64: 501-14



Results

- 26 papers were included, representing BV data for 35 variables
- Most publication on BV were found for Fibrinogen (17), Antithrombin (9) and APTT (8).
- Majority of studies were graded as a C => 74%
- 20% were graded as a A, which is the highest score achievable



Reason grading C

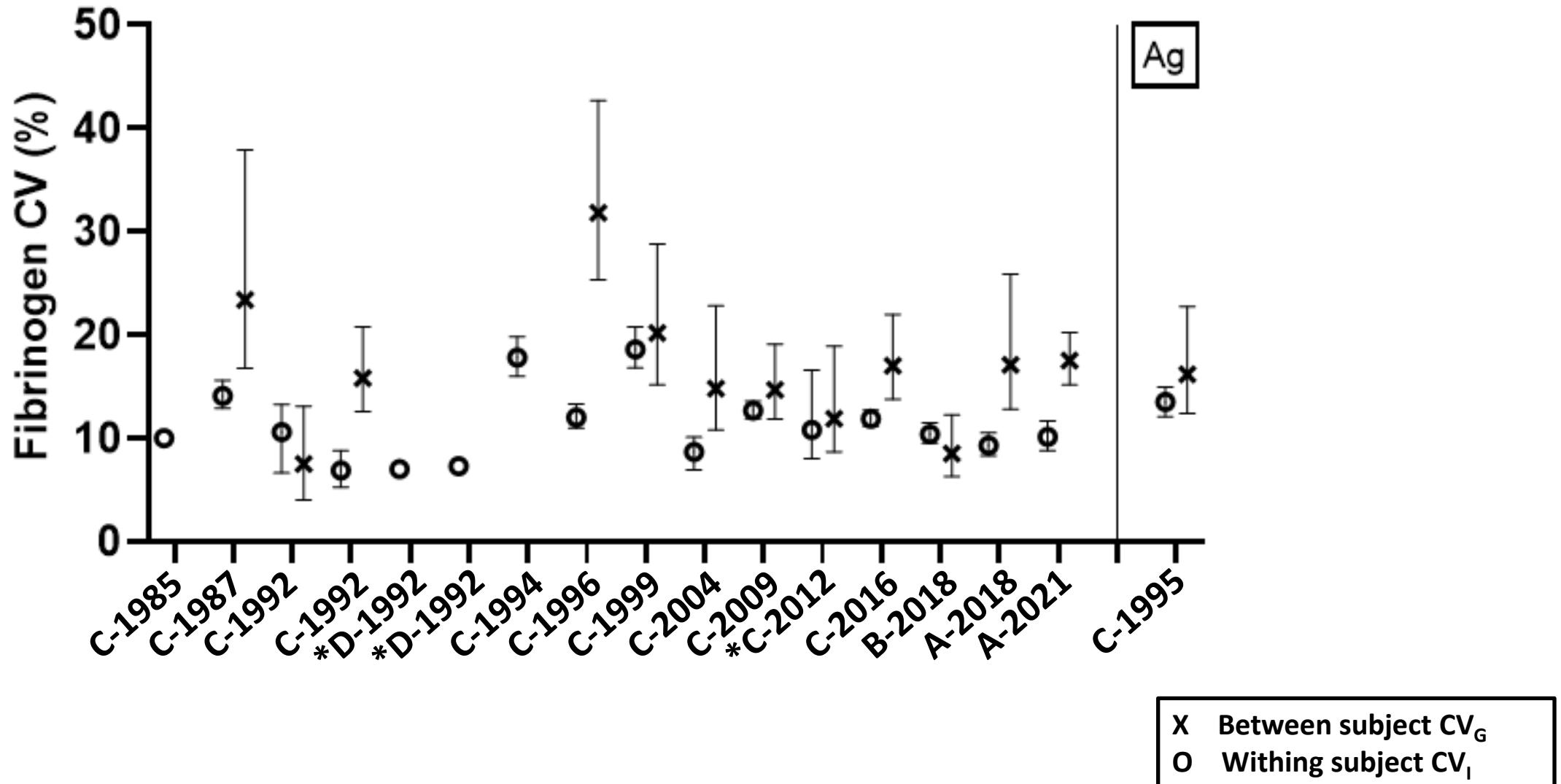
- **Mostly related to statistical issues; related to outlier analysis and testing for homogeneity**

Reason grading D

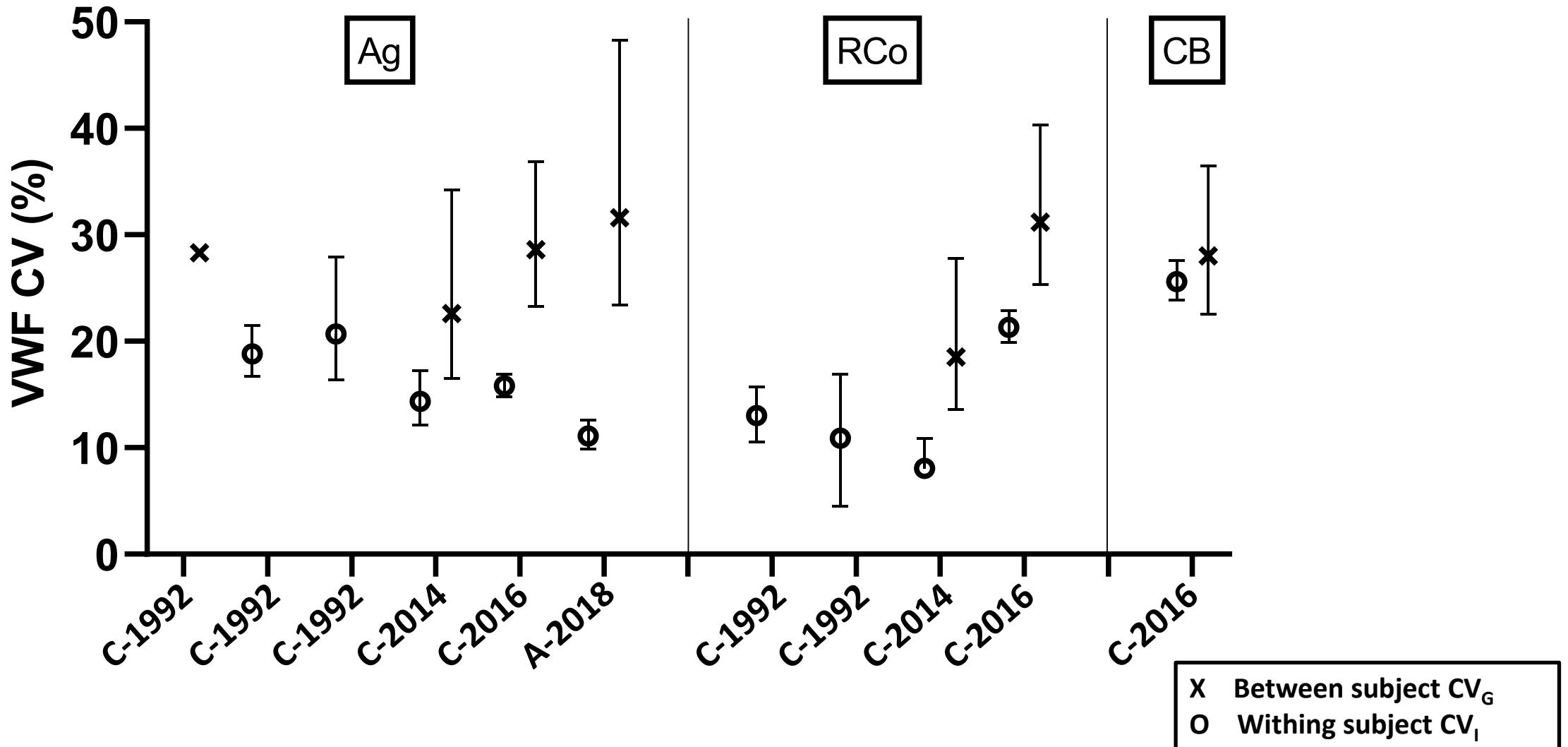
- **Due to obsolete methods being applied**



Fibrinogen



VWF



Comparison of data of different studies

- **No trend observed in BV estimates of different grading**
- **No trend observed in BV estimates of different years**
- **Since robust statistics were used for the meta-analysis, no changes were observed after exclusion of the most extreme BV data**



Meta-analysis derived within-subject (CV_I) and between-subject (CV_G) estimates with 95% CIs of coagulation and fibrinolytic variables

	CV_I	CV_G
APTT	2.8 (1.7-6.8)	7.2 (4.9-8.9)
APCR ratio	1.5 (1.3-6.7)	4.5 (3.8-5.4)
ADAMTS13 Act	12.7 (9.7-15.8)	9.6 (5.6-16.5)
ADAMT13 Ag	9.8 (0.0-13.4)	6.3 (1.9-11.6)
Antithrombin Ag	7.2 (6.3-8.2)	5.0 (3.4-8.3)
Antithrombin Act	3.4 (1.1-7.0)	7.8 (2.6-25.2)
D-Dimer	25.2(17.4-56.4)	35.4 (26.5-89.5)
Factor II	5.8 (5.7-5.9)	9.7 (7.0-15.4)
Factor V	5.3 (3.6-6.6)	18.7 (14.1-27.5)
Factor VII	8.2 (6.9-14.2)	17.8 (16.7-19.4)
Factor VIII	8.7 (4.9-16.0)	22.5 (15.5-31.4)
Factor IX	6.9 (5.8-9.1)	16.3 (15.7-18.2)
Factor X	5.9 (4.6-8.5)	11.4 (8.2-18.2)
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Factor XII	4.0 (3.0-5.1)	23.3 (17.6-34.5)
Fibrinogen Clauss	10.2 (9.3-11.9)	17.1 (8.5-17.3)
Fibringen Ag	13.5 (12.1-14.9)	16.2 (12.4-22.7)

	CV_I	CV_G
Plasmin Inhibitor	5.8 (4.8-5.8)	7.1 (5.2-10.8)
Plasminogen	5.7 (4.2-7.7)	10.5 (7.8-15.8)
Protein C Act	5.5 (5.3-7.9)	16.9 (9.1-55.2)
Protein C Ag	2.2 (0.0-6.2)	13.3 (10.5-17.5)
Protein S Act	7.3 (7.1-8.1)	20.3 (18.8-23.8)
Protein S Total	6.7 (2.9-7.3)	13.3 (8.9-63.4)
Protein S Free	4.2 (4.0-8.7)	16.9 (16.2-25.0)
Prothrombin time	2.6 (2.4-5.8)	5.1 (2.8-5.7)
TAT	19.0 (11.0-26.0)	33.3 (20.0-60.5)
Thrombomodulin	11.4 (9.1-13.2)	16.5 (12.1-25.1)
t-PA Ag	13.3 (11.0-30.9)	38.1 (23.9-191.1)
t-PA Act	32.0 (27.6-37.4)	NA
PAI-1 Ag	48.6 (35.6-55.0)	59.8 (26.0-90.0)
PAI-1 Act	34.9 (30.3-49.0)	90.2 (62.0-181.8)
VWF:RCo	17.0 (8.1-21.3)	24.6 (18.5-31.2)
VWF:Ag	12.7 (11.1-19.4)	29.9 (22.6-31.6)
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Reasons for high BV estimates PAI-1

- PAI-1 is an acute phase reactant
- Influence should be minimized since only healthy adults were included
- PAI-1 levels are influenced by a strong diurnal variation
- Since most samples were taken early in the morning this should be minimized too

→ High BV estimates means wider APS



Green marked variable represent the lowest estimates

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→ Low BV estimates values means
strict APS

Protein C Ag	2.2 (0.0-6.2)	13.3 (10.5-17.5)
Protein S Act	7.3 (7.1-8.1)	20.3 (18.8-23.8)
Protein S Total	6.7 (2.9-7.3)	13.3 (8.9-63.4)
Protein S Free	4.2 (4.0-8.7)	16.9 (16.2-25.0)
Prothrombin time	2.6 (2.4-5.8)	5.1 (2.8-5.7)
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Prothrombin Time, INR: In healthy adults versus patients

Healthy adults

- $CV_I = 2.5\% \text{ (2.3-3.0)}$
- $CV_G = 4.6\% \text{ (2.9-6.8)}$

Patients using anticoagulation treatment (data derived from three studies)

- $CV_I = 9 \text{ to } 23.8\%$
- $CV_G = 12.8 \text{ to } 18.5\%$

- Much higher estimates observed in patients compared to healthy adults
- APS criteria based on patients is less strict compared to healthy adults
- These results show the need for more BV studies in clinical patients to adapt quality criteria for specific clinical questions



Conclusion

- Systematic review and updated estimates of CV_I and CV_G (with 95% confidence interval) for an expanding number of haemostasis variables
- These estimates can form the basis for APS for haemostasis tests used in the diagnostic work-up in bleeding- and thrombosis events and for risk estimation
- Further expansion of high-quality BV studies is necessary to gain more knowledge of BV estimates for different population groups and states of health



Haemostasis BV estimates will be included in the new database

<https://biologicalvariation.eu/>



EFLM Biological Variation Database



BV in haemostasis variables

www.ecat.nl