# **LETTER TO THE EDITOR**

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# Revalidating the prognostic COVID-19 severity assessment (COSA) score for variants of concern



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Dear Editor,

Many prediction models have been published for Coronavirus disease 2019 (COVID-19) to support clinical decision-making for either diagnosis, prediction of mortality risk, or disease progression. An ongoing review concluded that most of these were poorly reported and had a high risk of bias, casting doubt over their real-world predictive value [1]. Additionally, dominant strains are no longer the ancestral type but variants of concern (VOCs) with different pathogenicity, and vaccinations and herd immunity can influence individual outcomes. We aimed to assess the performance of our previously published COSA (COVID-19 Severity Assessment) score on admitted patients in the twelve months following its development, a time during which the score was used clinically for risk stratification.

The COSA score was developed using data from patients who tested positive for severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) by reverse-transcriptase polymerase chain reaction (RT-PCR) between February 1st and November 16th, 2020 (1st and 2nd waves in Switzerland, i.e. the original cohort) [2]. The revalidation cohort consisted of patients who tested positive for SARS-CoV-2 between November 17th, 2020 and

November 16th, 2021. The primary outcome was disease severity, as determined by the worst outcome at any point after diagnosis:

- Non-severe outcome: no intensive care unit (ICU) admission or death of any cause during the observational period
- Severe outcome: ICU admission at any stage and / or death of any cause

All patients were discharged or had died by the time the revalidation was performed.

The COSA score (Table 1) was calculated for each patient using the most extreme laboratory values within 3 days prior to 1 day after the positive SARS-CoV-2 test. The results of the original and the revalidation cohort were then compared.

Data wrangling, analysis, and visualization were performed in GNU R (version 4.0.2, [3]). Statistical significance levels were defined at a p-value of < 0.05, and determined with the Wilcoxon rank sum test for continuous variables, and the Chi-square test for categorical variables using the *stats* package (version 4.0.2). The area under the receiver operating characteristic curve (AUROC) and the 95% confidence intervals (CI) were

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 Table 1
 COVID-19 severity assessment (COSA) score (from Ref. [2])

Parameter	Value	Score points
Sex	Male	1
CRP	≥ 25 mg/L	3
Sodium	≥ 144 mmol/L	2
Hemoglobin	≤100 g/L	1
eGFR according to CKD-EPI	≤75 mL/min	1
Glucose	≥ 8.6 mmol/L	1
Leucocytes	≥ 10 G/L	1

The COSA score was calculated for each patient using the most extreme values within 3 days prior to 1 day after the positive SARS-CoV-2 test, with a score of 6 or higher indicating a high risk (> 50%) for a severe progression

**Table 2** General demographics and laboratory parameters of the original and revalidation cohort

	31 71 3					
	Non-severe (N = 758)			Severe (N = 512)		
	Original (N = 457)	Revalidation $(N=301)$	P value	Original (N = 171)	Revalidation (N = 208)	P value
Demographics						
Age (years)						
Median (IQR)	64.00 (49.00, 76.00)	72.00 (59.00, 82.00)	< 0.002	68.00 (57.00, 78.00)	72.00 (59.75, 82.25)	0.058
Sex						
Female, n (%)	188 (41.05)	125 (41.81)	0.895	45 (26.32)	63 (30.29)	0.460
Hospitalization						
Inpatients, n (%)	319 (69.58)	299 (100.00)	< 0.002	163 (95.27)	208 (100.00)	0.005
Deaths						
Deceased, n (%)	0 (0.00)	0 (0.00)	-	51 (29.82)	120 (57.69)	< 0.002
Weight (kg)						
Median (IQR)	77.90 (66.23, 88.00)	76.30 (65.62, 88.00)	0.526	81.00 (70.40, 93.70)	75.20 (65.20, 90.00)	0.026
Height (cm)						
Median (IQR)	170.00 (165.00, 176.00)	170.00 (163.00, 177.00)	0.241	170.00 (165.00, 176.00)	170.00 (165.00, 176.75)	0.883
Body Mass Index (BN	/I, kg/m2)					
Median (IQR)	25.98 (23.38, 29.74)	26.20 (23.29, 30.32)	0.489	28.07 (25.20, 31.20)	26.87 (23.28, 29.81)	0.018
Laboratory parameters						
Maximum C-reactive	protein (CRP) levels					
Median (IQR)	37.00 (10.00, 78.00)	60.00 (21.00, 107.50)	< 0.002	106.00 (59.00, 175.00)	125.50 (62.00, 202.50)	0.300
Maximum sodium le	vels					
Median (IQR)	139.00 (137.00, 141.00)	139.00 (137.00, 142.00)	0.262	142.00 (139.00, 145.00)	142.00 (138.00, 146.00)	0.901
Minimum hemoglob	oin levels					
Median (IQR)	132.00 (118.00, 144.00)	126.00 (111.00, 142.00)	0.004	113.00 (89.50, 125.00)	107.50 (84.75, 124.25)	0.372
Minimum glomerula	r filtration rate (GFR) values					
Median (IQR)	82.00 (61.00, 97.00)	75.00 (52.00, 93.00)	0.007	64.00 (39.50, 87.00)	56.00 (31.75, 84.25)	0.158
Minimum glucose va	alues					
Median (IQR)	6.40 (5.67, 7.90)	6.90 (6.00, 8.46)	0.002	9.50 (7.46, 12.55)	9.45 (7.18, 11.80)	0.393
Minimum leukocyte	s values					
Median (IQR)	6.32 (4.70, 8.61)	6.41 (4.91, 9.09)	0.235	8.92 (6.65, 13.70)	10.90 (7.03, 15.12)	0.072

Laboratory parameters were considered from three day prior to until 1 day after the first positive SARS-CoV-2 PCR test result *IQR* interquartile range

Bold numbers indicate significant differences (p < 0.05) between severe and non-severe cases.

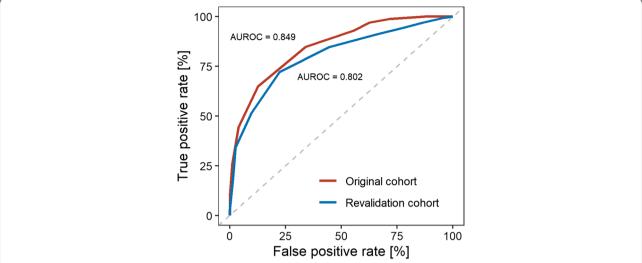


Fig. 1 Area under the receiver operating characteristic (AUROC) of the COVID-19 severity assessment (COSA) score. Original (red) and revalidation cohort (blue)

calculated to assess the discriminatory power of the COSA score to predict a severe outcome. The COSA score was considered validated if the 95% CI of the AUROC from fitting the original and the revalidation data set were overlapping.

The original cohort consisted of 626 patients (457 non-severe and 169 severe outcomes), and 508 patients were included in the revalidation cohort (301 non-severe and 208 severe outcomes). A comparison of the demographics and laboratory parameters of both cohorts is provided in Table 2. The patients in the original non-severe cohort were significantly younger with a lower share of inpatients than in the non-severe revalidation cohort. The latter had significantly higher peak C-reactive protein (CRP) and glucose, and lower minimal hemoglobin and estimated glomerular filtration rate values. For severe patients, we noted a higher proportion of inpatients and deaths, and significantly lower body weights and body mass index in the revalidation cohort. There was no significant difference in laboratory parameters.

The AUROC was 0.85 (95% CI 0.82–0.88) and 0.80 (95% CI 0.76–0.84) for the original and revalidation cohort, respectively (Fig. 1). A closer look at the severity distribution per score value (Fig. 2) revealed a greater share of severe cases with low score values (0–4 points) in the revalidation cohort than in the original cohort. No major differences are noticeable for score points greater than 4.

Patients in the revalidation cohort presented with overall worse laboratory markers and a greater likelihood of severe outcomes compared to the original cohort (69.1% vs. 37.4%). This could be due to more selective laboratory testing for hospitalized patients only and differences in circulating variants. Although no sequencing data is available for either cohort, national surveys indicate that the dominant strain during the 1st and 2nd waves (original cohort) was the ancestral type, whereas the 3rd and 4th waves (revalidation cohort) were driven by the more transmissible and virulent VOCs Alpha (B.1.1.7) and Delta (B.1.617.2) [4]. Vaccination campaigns and an emerging population-level immunity likely mitigated the individual disease severity [5].

While the performance of the score remains robust with strong positive discriminative ability we did note a decrease in specificity towards the lower end of the scale (Fig. 2). This suggests that readjustment of the cut-offs might be beneficial for a better separation of the severity classes as new variants emerge.

Despite changes in the viral landscape and population immunity, the COSA score still delivered good predictions of disease progression one year after its development. We attribute this to the simple set of covariates and the rigorous internal and external validation in the original model-building process. Adaptations to the score could become necessary in the near future as vaccination effects begin to wane and if drastically different VOCs appear.

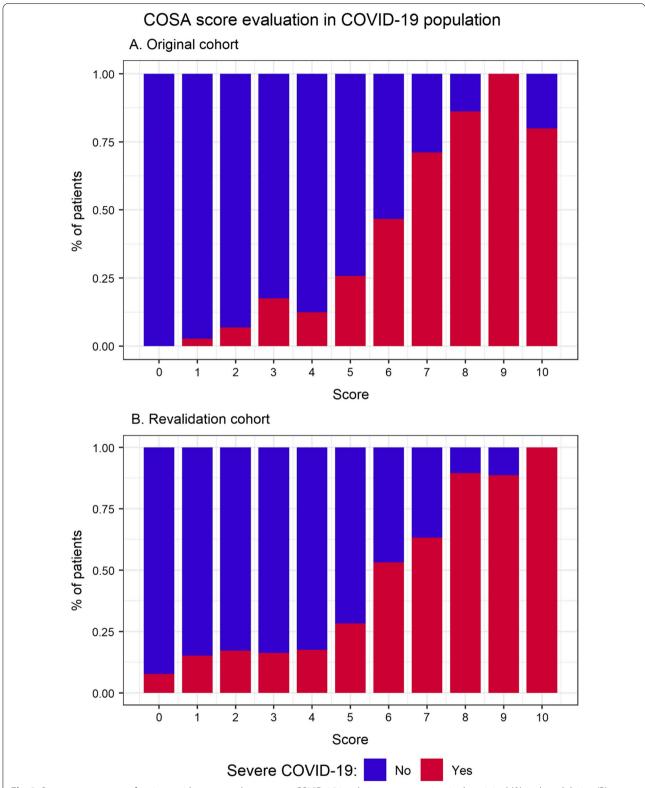


Fig. 2 Percentage amount of patients with severe and non-severe COVID-19 in relation to score points in the original (A) and revalidation (B) cohort

#### Abbreviations

AUROC: Area under the receiver operating characteristic; BMI: Body mass index; COSA: COVID-19 severity assessment; COVID-19: Coronavirus disease 19; CRP: C-reactive protein; eGFR: Estimated glomerular filtration rate; ICU: Intensive Care Unit; RT-PCR: Reverse-transcriptase polymerase chain reaction; SARS-CoV-2: Severe acute respiratory syndrome coronavirus 2; VOC: Variants of concern.

# Acknowledgements

We thank Noel Frey, Myoori Wijayasingham, and the Insel Data Coordination Lab for database and infrastructure support.

#### **Author contributions**

FH and VS conceptualized this study. VS and FH performed the data analysis and drafted the manuscript. AA assessed the machine learning models and statistics, EL, CB, AE, and WH assessed the clinical score. All authors read and approved the final manuscript.

#### **Funding**

None to declare.

#### Availability of data and materials

The source code and corresponding input file are available on GitHub: https://github.com/cptbern/COSAscore. The datasets used and/or analysed during the current study are available from the corresponding author on reasonable request.

#### **Declarations**

## Ethics approval and consent to participate

The study was approved by the Cantonal Ethics Committee of Bern (Project-ID 2020–00973). Participants either agreed to a general research consent or, for participants with no registered general research consent status (neither agreement nor rejection), a waiver of consent was granted by the ethics committee.

# Consent for publication

Not applicable

#### **Competing interests**

The authors declare that they have no competing interests.

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Received: 8 September 2022 Accepted: 10 September 2022 Published online: 23 September 2022

## References

- Wynants L, Van Calster B, Collins GS, Riley RD, Heinze G, Schuit E, Bonten MMJ, Dahly DL, Damen JA, Debray TPA, et al. Prediction models for diagnosis and prognosis of covid-19: systematic review and critical appraisal. BMJ. 2020;369: m1328.
- Schöning V, Liakoni E, Baumgartner C, Exadaktylos AK, Hautz WE, Atkinson A, Hammann F. Development and validation of a prognostic COVID-19 severity assessment (COSA) score and machine learning models for patient triage at a tertiary hospital. J Trans Med. 2021;19:56.
- R Core Team. R: A language and environment for statistical computing. Vienna: R Foundation for Statistical Computing; 2022.
- Covid-19 Schweiz: Epidemiologischer Verlauf, Schweiz und Liechtenstein, Virusvarianten. https://www.covid19.admin.ch/de/epidemiologis/virus-variants?time=total&variants=VariantB11529,VariantB16172,Varia

- ntP1,VariantB1351,VariantB117,VariantC37,VariantB16171,VariantP2,VariantB1525,VariantB1526, Accessed 5 July 2022.
- Zeng B, Gao L, Zhou Q, Yu K, Sun F. Effectiveness of COVID-19 vaccines against SARS-CoV-2 variants of concern: a systematic review and metaanalysis. BMC Med. 2022;20:200.

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