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1 EARLY LABORATORY DIAGNOSIS OF COVID-19 BY ANTIGEN DETECTION IN BLOOD SAMPLES OF THE

2 SARS-COV-2 NUCLEOCAPSID PROTEIN

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- 19
- 20 Running title: Diagnosis of COVID-19 by antigen detection in blood

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21	ABSTRACT
22	The purpose of this study was to characterize the diagnostic performance of a newly developed
23	enzyme-linked immunosorbent assay (ELISA) for detection of SARS-CoV-2 nucleocapsid protein (NP) in
24	blood. Blood samples were collected during hospitalization of 165 inpatients with PCR-confirmed
25	SARS-CoV-2 infection, and from 505 outpatients with relevant symptoms of COVID-19 simultaneously
26	with PCR-testing. For the 143 inpatients who had their first blood sample collected within 2 weeks
27	after PCR-confirmed infection, the diagnostic sensitivity of the ELISA was 91.6%. The mean NP
28	concentration of the 131 ELISA-positive blood samples was 1,734 pg/ml (range: [10-3,840] pg/ml). An
29	exponential decline in NP concentration was observed for 368 blood samples collected over the first 4
30	weeks after PCR-confirmed SARS-CoV-2 infection, and all blood samples taken later had an NP
31	concentration below the 10 pg/ml diagnostic cut-off.
32	The diagnostic sensitivity of the ELISA was 81.4% for the 43 blood samples collected from outpatients
33	with a simultaneous positive PCR-test, and the mean NP concentration of the 35 ELISA-positive
34	samples was 157 pg/ml (range: [10-1,377] pg/ml). For the 462 outpatients with a simultaneous
35	negative PCR-test, the diagnostic specificity of the ELISA was 99.8%.
36	In conclusion, the SARS-CoV-2 NP ELISA is a suitable laboratory diagnostic test for COVID-19.
37	Particularly, for hospitals, where blood samples are readily available, screening of serum or plasma
38	samples by ELISA can facilitate prevention of nosocomial infections and reduce the requirement for

laborious swab sampling and subsequent PCR-analysis to confirmatory tests. 39

41 **INTRODUCTION**

The pandemic corona virus disease 2019 (COVID-19) caused by the severe acute respiratory 42 syndrome–related coronavirus 2 (SARS-CoV-2) virus has led to the rapid development and widespread 43 application of many laboratory diagnostic tests (1). According to the World Health Organization 44 45 (WHO), the standard confirmation of acute infections with SARS-CoV-2 is based on a nucleic acid 46 amplification test, such as real-time reverse-transcription PCR for the presence of unique sequences of SARS-CoV-2 RNA (2). Testing for genomic RNA by PCR is widely supplemented by two other major 47 diagnostic test principles: testing for specific virus antigens and humoral immune response to the 48 infection. Like PCR, analyses for SARS-CoV-2 antigens are typically employed before the onset of 49 50 clinical symptoms of COVID-19 or during the anticipated acute phase of infection. In contrast, 51 immunoassays for humoral antibodies directed against components of SARS-CoV-2 should not be 52 applied until about 10 days after symptom onset, when the expected humoral immune response has matured sufficiently to reach a detectable level (1). 53 54 These three fundamental *in vitro* diagnostic test principles have their individual advantages and limitations, which partially are associated with their respective sampling techniques for appropriate 55 56 test material. For almost all PCR analyses for SARS-CoV-2 RNA and immunoassays for SARS-CoV-2 antigen, the hitherto preferred test material is extracted from swabs collected from the upper 57 58 respiratory tract (URT). In contrast to this heterogeneously composed, individually fluctuating, and 59 somewhat ill-defined test material, immunoassays for antibodies to SARS-CoV-2 rely on a blood sample. In general, blood samples are by far the most used biological material in laboratory diagnostic 60 61 procedures, and consistencies and variations of this sample material are very well characterized.

62	Shortly after the severe acute respiratory syndrome (SARS) epidemic in 2002-2004, it was reported,
63	that the nucleocapsid protein (NP) of the original SARS corona virus (SARS-CoV) could be detected by
64	enzyme-linked immunosorbent assay (ELISA) in serum samples collected from 95% of infected
65	patients three days after symptom onset (3). The SARS-CoV-2 NP is highly conserved and 90.5%
66	identical to the primary structure of SARS-CoV NP, whereas the full proteome identity of these two
67	viruses is 77.1% (4). Inspired by these observations, a new ELISA has been developed and tested for
68	detection of SARS-CoV-2 NP antigen in blood samples collected from COVID-19 patients during the
69	early stages of SARS-CoV-2 infection (5)(6)(7). By using PCR analysis of URT swabs as reference, the
70	present clinical study reports the laboratory diagnostic characteristics and performance of this NP
71	ELISA, when used for SARS-CoV-2 antigen quantification in serum and plasma samples.

72 MATERIALS AND METHODS

73 Patients and blood samples

Venous blood was collected from patients at two Danish university hospitals and prepared as either 74 75 serum or EDTA plasma according to the Standard Operating Procedures in Bio- and GenomeBank, 76 Denmark (8). All blood samples were collected from patients, who had not been COVID-19 vaccinated. 77 Serum samples were obtained from two different patient groups: Inpatients with symptoms of COVID-19 and a confirmatory PCR-positive test result admitted to a COVID-19 specific department at 78 79 University Hospital Rigshospitalet, Copenhagen, and outpatients referred for testing for SARS-CoV-2 infection at an outpatient testing facility at University Hospital Rigshospitalet. Only outpatients with 80 81 symptoms of upper respiratory tract infection (e.g., fever, sore throat, cough) were included in the 82 study. 83 Plasma samples were obtained from two different patient groups: Inpatients with symptoms of COVID-19 and a confirmatory PCR-positive test result admitted to COVID-19 specific departments at 84

Aalborg University Hospital, and outpatients referred for testing for SARS-CoV-2 infection at an

86 outpatient testing facility at Aalborg University Hospital including both persons with and without

87 symptoms of SARS-CoV-2 and persons who had been exposed.

88 For each inpatient, 1 to 10 blood samples collected within the interval from the day of their first PCR-

positive URT swab (Day 0) until Day 201 were all included in the study. For outpatients, only the blood

sample collected simultaneously with their URT swab (Day 0) was included in the study.

91 The serum and plasma samples were stored at -20°C or -80°C until testing by ELISA.

All patients provided written statement being part of Bio- and Genome Bank, Denmark. For
participants under the age of 18 years, a parent or legal guardian provided the consent. The present
methodology study was approved by the Central Denmark Region Committees on Biomedical
Research Ethics (IORG-number: 0005129).

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97 PCR analysis

For all in- and outpatients included in this study, the reference laboratory diagnosis of COVID-19 was 98 performed at the involved hospitals by PCR analysis of URT swabs for the presence of unique 99 sequences of SARS-CoV-2 RNA. All URT swabs were collected according to Danish national guidelines 100 101 (9) by health professionals and taken as oropharyngeal samples. The swabs were then processed as 102 routine samples and analyzed using a real-time reverse-transcription PCR assay. Two different PCR test kits were used, either the "Cobas® SARS-CoV-2 Test" on a Cobas 6800 system (Roche, Basel, 103 104 Switzerland) or the "RealStar® SARS-CoV-2 RT-PCR Kit" (Altona, Hamburg, Germany). The result of PCR 105 analysis was reported as positive or negative for SARS-CoV-2 genomic RNA.

106

107 Quantification of NP in blood samples

The quantification of NP concentration [NP] in serum or plasma was accomplished in approximately 2
hours by the "Solsten SARS-CoV-2 Antigen ELISA Kit" from Solsten Diagnostics International (Aarhus,
Denmark) according to the manufacturer's guidelines. Up to 12 strips of each 8 wells precoated with
antibody to SARS-CoV-2 NP were mounted in each 96-well frame. First, 50 µl of biotin-conjugated
antibody was added to each well and then directly supplemented with 50 µl/well of either internal NP

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113 calibrator, serum, or plasma. Hereafter, the wells were incubated for 1 hour at 37°C, washed, incubated with 100 µl/well of peroxidase-conjugated streptavidin for 30 minutes at 37°C, washed, and 114 then incubated with the provided substrate for 15 minutes at 37°C before stopping the chromogenic 115 116 enzyme reaction and measuring the absorbance photometrically. Standard curves based on ELISA 117 results of the 5 internal calibrators were used for quantification of [NP] between 0 and 160 pg/ml, as 118 defined by the manufacturer. All samples were analyzed twice by the ELISA on different days. The first NP-quantification was done 119 120 blinded for the characteristics of the individual sample except for being serum or plasma. Similarly, the second ELISA analysis was done blinded for COVID-19 status, but with insight into the [NP] 121

determined by the first ELISA run. This allowed the appropriate dilution of samples with previously
determined [NP] higher than 100 pg/ml. Serum and plasma samples, which after a 24-fold dilution,

124 produced an ELISA absorbance value higher than the highest NP calibrator (160 pg/ml) were not

125 further diluted for precise quantification but registered as having an [NP] of 3,840 pg/ml.

126

127 Statistical analysis

The statistical uncertainty of the estimates of diagnostic accuracy for the SARS-CoV-2 NP ELISA,
including sensitivity, specificity, and predictive values of positive and negative results were reported
as 95% confidence intervals (95% Cl). The mean [NP] ± standard deviation (SD) of serum and plasma
were compared by two-tailed t-tests. A p-value less than 0.05 was considered statistically significant.

132	RESULTS

Patients and blood samples 133

The 670 individuals included in this study comprised 414 females aged 14 to 102 years (mean ± SD: 52 134 135 \pm 19 years) and 256 males aged 20 to 100 years (mean \pm SD: 62 \pm 20 years). According to PCR-analysis 136 of URT swabs, 208 of these individuals were infected with SARS-CoV-2. The COVID-19 patients 137 comprised 97 females aged 22 to 96 years (mean \pm SD: 62 \pm 19 years) and 111 males aged 28 to 100 138 years (mean \pm SD: 70 \pm 16 years).

A total of 914 human blood samples were collected from the 670 individuals between March 3, 2020, 139 and February 2, 2021, and prepared as either serum (n=439) or plasma (n=475). Of these, 447 (49%) 140 blood samples were from 165 COVID-19 inpatients and 43 COVID-19 outpatients, including 173 serum 141 142 samples and 231 plasma samples collected from 38 and 127 inpatients, respectively, between 0 and 201 days after their first confirmatory PCR-positive URT swab. Furthermore, 15 serum samples and 28 143 plasma samples were collected from outpatients simultaneously with their first confirmatory PCR-144 145 positive URT swab. Most of the blood samples from COVID-19 patients (n=324, 72%) were collected within 14 days of the first PCR-positive URT swab confirming the patient's infection with SARS-CoV-2, 146 147 and the remaining blood samples (n=123, 28%) were collected from COVID-19 inpatients more than 2 weeks after their first PCR-positive test. All other blood samples (n=467, 51%) were collected from 148 149 462 outpatients without COVID-19 according to a simultaneously collected PCR-negative URT swab 150 and comprised 251 serum samples from 251 outpatients, and 216 plasma samples from 211 151 outpatients.

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152 A schematic overview of patients and samples of the study is provided in Supplemental Material

153 (Figure S1).

154

155 Diagnostic performance of the SARS-CoV-2 NP ELISA

156 Based on the Receiver Operating Characteristic (ROC) curve in Figure 1, and prioritization of a low 157 false positive rate, the manufacturer's recommended diagnostic cut-off value of 10 pg/ml NP was confirmed. When using this cut-off value, the specificity of the SARS-CoV-2 NP ELISA was 99.8% (95% 158 CI: 99.4% - 100%), as 1 of 462 outpatients without COVID-19 had a false positive blood sample with an 159 [NP] of 12 pg/ml. All 462 blood samples were collected simultaneously with a PCR-negative URT swab. 160 161 The diagnostic sensitivity of the SARS-CoV-2 NP ELISA was determined at patient level by using only 162 the [NP] measured for the first blood sample after collection of the confirmatory PCR-positive URT 163 swab. According to results for 160 COVID-19 inpatients, the ELISA sensitivity varied with the time gap from confirmatory PCR-positive URT swab to blood sampling (Figure 2). 164 165 When the first blood sample was collected from COVID-19 inpatients within 1 and 2 weeks from PCRconfirmed infection, the diagnostic sensitivity of the SARS CoV-2 NP ELISA was 92.9% (n=99; 95% CI: 166 167 87.9% - 98.0%) and 91.6% (n=143; 95% CI: 87.1% - 96.2%), respectively (Table 1). The average [NP] (± SD) of the true ELISA-positive blood samples collected within the first 2 weeks from 131 COVID-19 168 169 inpatients was 1,734 ± 1,560 pg/ml (range: [10 - 3,840] pg/ml, median: 1,184 pg/ml). The average [NP] (± SD) of the true ELISA-positive blood samples collected at Day 0 from 35 COVID-19 outpatients 170 was 157 ± 294 pg/ml (range: [10 - 1,377] pg/ml, median: 52 pg/ml). 171

172 According to all patients (n=520) with a blood sample collected simultaneously with the URT swab i.e., at Day 0, the PCR-defined point prevalence of COVID-19 was 11.2%: 15 PCR-positive inpatients and 173 505 outpatients, including 43 PCR-positive. For the SARS-CoV-2 NP ELISA, at this timepoint and 174 175 prevalence, the specificity and sensitivity were 99.8% (95% CI: 99.4-100%) and 82.8% (95% CI: 73.0-176 92.5%), respectively, whereas the positive predictive value (PPV) and negative predictive value (NPV) 177 were 98.0% (95% CI: 94.0-100%) and 97.9% (95% CI: 96.6-99.2%), respectively. 178 For COVID-19 outpatients, who had their blood sample collected at Day 0 i.e., simultaneously with the confirmatory PCR-positive URT swab, the diagnostic sensitivity of the SARS-CoV-2 NP ELISA was 81.4% 179 (n=43; 95% CI: 69.8% - 93.0%). For the present study, where the point prevalence of COVID-19 for the 180 181 analyzed group of 505 outpatients according to PCR analysis was 8.5%, the probability of having 182 COVID-19 was 97.2% for ELISA-positive outpatients (n=36; 95% CI: 91.9% - 100.0%), and the 183 probability of not being infected with SARS-CoV-2 was 98.3% for ELISA-negative outpatients (n=469; 95% CI: 97.1% - 99.5%). 184 185 Serum and plasma analysis by the SARS-CoV-2 NP ELISA 186 187 According to 368 blood samples from 160 COVID-19 inpatients, the correlation between SARS-CoV-2 ELISA and PCR confirmed SARS-CoV-2 infection varied with the time gap from confirmatory PCR-188 189 positive URT swab to blood sampling (Figure 3).

- 190 For 131 serum and 150 plasma samples collected from COVID-19 inpatients within 2 weeks after their
- 191 confirmatory PCR-positive URT swab, the correlation was 89.3% (95% CI: 84.0% 94.6%) and 86.7%
- 192 (95% CI: 81.2% 92.1%), respectively (Table 2). For collection within the first week only, the

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196	inpatients within the first week from their confirmatory PCR-positive URT swab was 1,041 \pm 1,332
197	pg/ml (range: [3 - 3,840] pg/ml, median: 337 pg/ml) and 1,631 ± 1,553 pg/ml (range: [3 - 3,840] pg/ml,
198	median: 1,036 pg/ml), respectively. The median [NP] decreased exponentially with the time gap from
199	collection of the PCR-positive URT swab to blood sampling (Figure 4). No systematic difference was
200	observed in the [NP] levels between serum and plasma samples collected within the first week from
201	COVID-19 inpatients (p=0.0577).
202	For outpatients with SARS-CoV-2 infection according to PCR, the average apparent [NP] (\pm SD) of 15
203	serum samples and 28 plasma samples were 54 ± 69 pg/ml (range: [2 - 274] pg/ml, median: 28 pg/ml)
204	and 169 ± 328 pg/ml (range: [3 - 1,377] pg/ml, median: 31 pg/ml), respectively. No systematic
205	difference was observed in the [NP] levels between serum and plasma samples of COVID-19
206	outpatients (p=0.418).
207	For outpatients without SARS-CoV-2 infection according to PCR, the average apparent [NP] (\pm SD) of
208	251 serum samples and 216 plasma samples were 2.1 \pm 1.3 pg/ml (range: [0 - 12] pg/ml, median: 2.4
209	pg/ml) and 2.5 \pm 1.3 pg/ml (range: [0 - 7.4] pg/ml, median: 2.4 pg/ml), respectively.

corresponding correlations were higher: 92.3% (n=65; 95% CI: 85.8% - 98.8%) for serum and 91.7%

The average [NP] (± SD) of 65 serum samples and 84 plasma samples collected from COVID-19

210

211 Individual dynamics in [NP] levels of blood samples

(n=84; 95% CI: 85.8% - 97.6%) for plasma.

The individual progression in [NP] during the first month after PCR-based diagnosis was observed for 212

40 COVID-19 inpatients, who had at least 3 blood samples collected within 30 days from their first 213

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214	PCR-positive URT swab (Figure 5). For 4 of these inpatients (10%), none of their blood samples (total
215	n=20) reached an [NP] above the diagnostic cut-off value of 10 pg/ml. In two of these cases, the
216	earliest blood samples were collected more than 2½ weeks after their PCR-based diagnosis, when a
217	substantial humoral immune response to infection was measured (Table S1 and Figure S2 in
218	Supplemental Material). For all the remaining 36 COVID-19 inpatients (90%), at least their first blood
219	sample had an [NP] above the diagnostic cut-off value. Despite the clear individual tendency of
220	decline in [NP] over time, all blood samples (total n=122) collected from 23 of the 40 COVID-19
221	inpatients (58%) were positive according to the NP ELISA.
222	None of the 34 plasma samples collected from 10 COVID-19 inpatients 27-201 days after their first
223	PCR-positive URT swab had an [NP] above the diagnostic cut-off value of 10 pg/ml (Figure 6), even
224	though 1 to 3 plasma samples collected earlier from each of these patients were clearly positive for
225	NP (mean ± SD: 1,444 ± 1,448 pg/ml; range: [34 – 3,840] pg/ml).

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227 DISCUSSION

The outbreak of COVID-19 has caused an unparalleled worldwide requirement for laboratory 228 diagnostic tests for virus infection, and PCR analysis for genomic RNA of SARS-CoV-2 in extracts of 229 230 swabs collected from the upper respiratory tract has proven very suitable for early detection of 231 infection, even in patients with mild or no clinical symptoms. Still, the characterization of PCR as the 232 gold standard laboratory diagnostic test for COVID-19 (1), and its wide application as a reference test in performance evaluation of other laboratory diagnostic methods is debated (10). This is particularly 233 due to concerns of false-negative PCR results caused by a low viral load at the chosen time and site of 234 URT sample collection, inadequate URT swabbing technique of some operators, failing storage 235 236 conditions during specimen transportation, laboratory error, and/or mutation of the viral target RNA 237 (11). These concerns have intensified the search for improved and less resource-demanding 238 laboratory test procedures for COVID-19 and has led to the development of complementing and supplementing screening methods, which will contribute to diagnostic triage procedures relying on a 239 240 final confirmation of positive results by PCR-analysis. In the present study, we have characterized the first ELISA test kit for quantification of the SARS-CoV-241 242 2 NP antigen in serum and plasma samples. When used for blood samples collected from COVID-19 inpatients within 2 weeks after PCR-confirmed SARS-CoV-2 infection, the diagnostic sensitivity of the 243 244 ELISA was 91.6% (95% CI: 85.6% - 95.2%). The group of 505 outpatients in this study had an 8.5% point prevalence of SARS-CoV-2 infection, and 245 for outpatients with an ELISA-positive blood sample, the probability of having COVID-19 was 97.2% 246

247 (n=36; 95% CI: 91.9% - 100.0%), whereas those with an ELISA-negative blood sample had a 98.3%

249	stage of infection, the SARS-CoV-2 NP ELISA thereby prove to be a very reliable predictor of COVID-19.
250	The individual NP concentrations of COVID-19 patients varied considerably even for blood samples
251	collected within the first week of PCR-confirmed SARS-CoV-2 infection, and probably reflected the
252	disease severity. The [NP] in blood samples collected at Day 0 from 15 COVID-19 inpatients (median:
253	1,237 pg/ml, mean \pm SD: 1,792 \pm 1,687 pg/ml, range: [3-3,840] pg/ml) was at a substantially higher
254	level than the [NP] in blood samples collected from 43 COVID-19 outpatients simultaneously with
255	their PCR-positive URT swab (median: 29 pg/ml, mean ± SD: 129 ± 271 pg/ml, range: [2-1,377] pg/ml).
256	Despite the variability in [NP] between patients and over time, the individual progressions in [NP]
257	were systematic and declining for almost all the 40 inpatients in the present study, who had at least 3
258	blood samples collected within the first month. Also, for the total of 368 blood samples collected from
259	160 COVID-19 inpatients during the first month, the median [NP] declined exponentially with time,
260	and then consistently remained below the diagnostic cut-off value of 10 pg/ml for all samples
261	collected during the succeeding 6 months after infection.
262	When verifying the diagnostic performance of an antigen test by using PCR analysis of an URT swab as
263	the reference, all misclassifications (false negatives and false positives) by definition, will be ascribed
264	to the antigen test, no matter whether the test material is matching or different. Almost all rapid
265	antigen tests for COVID-19 are lateral flow immunoassays for the qualitative detection of SARS-CoV-2
266	NP in extracts of URT swabs. Possibly, more rightfully characterized as tests of individual
267	infectiousness (12)(13), their diagnostic performance is typically evaluated by comparison to the
268	outcome of PCR analysis of the same or a simultaneously collected URT swab and thereby affected by

probability of not being infected with SARS-CoV-2 (n=469; 95% CI: 97.1% - 99.5%). During this early

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the same risks of a sampling-associated false negative result as PCR. The diagnostic performance of the quantitative ELISA for SARS-CoV-2 NP investigated in the present study, also relied on using PCR analysis of a URT swab as reference. Despite the distinct sampling techniques and test materials of these two laboratory diagnostic procedures, the analysis of blood samples by the NP ELISA highly confirmed the laboratory diagnosis of SARS-CoV-2 infection based on PCR.

274 According to the observed performance data, we conclude that the SARS-CoV-2 NP ELISA is suitable for laboratory diagnosis of COVID-19 when used for testing serum or plasma early after infection. 275 Towards the end of the outbreak of SARS in 2002-2004, an ELISA was developed with an analytical 276 detection limit of approximately 50 pg/ml SARS-CoV NP (14). Using a diagnostic cut-off at 100 pg/ml, 277 278 its diagnostic sensitivity increased from 65% 1-2 days after onset of SARS symptoms to over 95% at 3-279 5 days after first symptoms (3). In comparison, the present ELISA for SARS-CoV-2 NP (5) has a 280 substantially improved analytical sensitivity with a detection limit of around 2 pg/ml, which in combination with the assay's high resistance to irregular hemolytic reactions and potentially 281 282 interfering blood substances, such as rheumatoid factors (5), allows the recommended low and robust diagnostic cut-off at 10 pg/ml SARS-CoV-2 NP. Though differences in shedding of NP into 283 284 circulation and time of symptom onset after infection may vary between SARS and COVID-19, the 10fold reduction in diagnostic cut-off contributes decisively to the very early detection of SARS-CoV-2 285 286 infection achieved by the novel NP ELISA investigated in the present study. 287 WHO has concluded that early laboratory diagnosis of SARS-CoV-2 infection can aid clinical management and outbreak control of COVID-19, and that the standard confirmation of acute 288 289 infection should be based on a nucleic acid amplification test (2). However, URT swab collection

followed by PCR analysis is a tedious and expensive method for COVID-19 screening. In contrast, in
settings such as hospitals and blood banks, where blood samples are collected anyway, the SARS-CoV2 NP ELISA provides a simple and economical screening tool for COVID-19. For example, serum and
plasma samples prepared at hospitals for biochemical and other clinical laboratory analyses may also
be systematically examined by ELISA for the presence of SARS-CoV-2 NP, and thereby contribute
importantly to reduce the risk of nosocomial COVID-19 infection (15).

296 Our study has strengths and limitations. First, only a subset of the included participants had blood samples collected within 0-1 days after their first PCR-positive URT swab, although this period is the 297 most clinically relevant for early detection of SARS-CoV-2. However, the wide range of collection of 298 299 blood samples after the first PCR-positive URT swab allowed us to investigate the individual progress 300 of NP concentration in blood for participants with numerous samples available. Second, we did not 301 have information on onset or duration of symptoms for in- and outpatients. Instead, the first PCR-302 positive URT swab was used as the confirmatory test for COVID-19, although the infection with SARS-303 CoV-2 may have started days before the PCR test was performed. Theoretically, in a setting where this information was available, the false negative rate would be even lower, as participants with longer 304 305 duration of symptoms before blood sampling could be excluded from the main analyses. Third, oropharyngeal swabs were utilized for URT sampling according to Danish national guidelines (9), and 306 307 though also stated by the FDA as appropriate for clinical testing, they may be less sensitive than 308 nasopharyngeal swabs (16). The strengths of the study include a large sample size and a standardized and highly reproducible method for quantification of SARS-CoV-2 NP. 309

5		
	310	Though being a recommended subject for further investigations, we propose that automated routine
5	311	screening of blood samples by the NP ELISA will be a suitable procedure for early identification of
5	312	inpatients, who bring or acquire SARS-CoV-2 infection, while hospitalized. As indicated by the
<u>)</u>))	313	projected confusion matrix (Table S2 in Supplemental Material), even for hospitals with a low
	314	prevalence of COVID-19 among inpatients treated for other diseases, and a high number of routinely
	315	analyzed blood samples, the observed 99.8% specificity of the SARS-CoV-2 NP ELISA will ensure a low
	316	number of false positives and an acceptable PPV, and thereby lead to substantial reductions in the
	317	requirement for laborious swab sampling and subsequent confirmatory PCR analysis.

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- 326 for quantification of the SARS-CoV-2 NP antigen and for measurement of titers of neutralizing
- 327 antibodies to the ACE-2 receptor binding domain of the S1 subunit of SARS-CoV-2 Spike Protein,
- 328 respectively.

329

330 CONFLICTS OF INTEREST DISCLOSURES

- AP and NTF are employees of Solsten Diagnostics International, Aarhus, Denmark, which is the
- 332 company providing the "Solsten SARS-CoV-2 Antigen ELISA Kit".

333

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TABLES

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TABLE 1. Diagnostic performance and clinical relevance of the SARS-CoV-2 NP ELISA, when testing the 401

first blood sample collected from each of 648 patients within the first week (Day 0-6) and the second 402

week (Day 7-13) of their PCR analyzed URT swab. 403

		Blood	Posit	ive	Negative	Total	Relevance:	
_		collected:	Inpatients	Outpatients	Outpatients	Patients	Prediction	
	Docitivo	Day 0-6	92	35	1	128	99.2%	
	POSITIVE	Day 7-13	39			39		
SARS-Co	oV-2 Nogativ	Day 0-6	7	8	461	476	96.8%	
NP ELIS	A	Day 7-13	5			5		
	Total	Day 0-6	99	43	462	604		
	TOLAI	Day 7-13	44			44		
			Sensitivity	Sensitivity	Specificity	648		
	Performance	Day 0-6	92.9%	81.4%	99.8%		-	
	of NP ELISA:	Day 7-13	88.6%					

404

408 samples and the result of PCR analysis for SARS-CoV-2 infection according to of a URT swab collected

409 up to 2 weeks earlier.

Table 2A.						
Number of ser	rum samples: 397	Posit	tive	Negative	Total	Relevance:
Blood samplin	g within: Day 0-13	Inpatients	Outpatients	Outpatients	Patients	Prediction
SARS-CoV-2	Positive	117	13	1	131	99.2%
	Negative	14	2	250	266	94.0%
INF LLIJA	Total	131	15	251	397	
Performance of NP ELISA:		Correlation	Correlation	Correlation		
Сог	rrelation	89.3%	86.7%	99.6%		

410

Table 2B.						
Number of pl	asma samples: 394	Positive Negative				Relevance:
Blood samplir	ng within: Day 0-13	Inpatients	Outpatients	Outpatients	Patients	Prediction
	Positive	130	22	0	152	100%
SARS-CoV-2	Negative	20	6	216	242	89.3%
NF LLISA	Total	150	28	216	394	
	Performance	Correlation	Correlation	Correlation		
	of NP ELISA:	86.7%	78.6%	100%		

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FIGURE LEGENDS

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415	FIGURE 1. ROC curves for the SARS-CoV-2 NP ELISA according to the time gap from the first PCR-
416	positive URT swab to first blood sampling. A zoom of the upper left corner of the curves was inserted.
417	The area under the curve was 0.986 for the 604 blood samples collected within a time gap of 1 week
418	(Day 0-6), 0.982 within 2 weeks (648 blood samples collected Day 0-13), and 0.975 within 3 weeks
419	(662 blood samples collected Day 0-20). In compliance with the recommendations by the
420	manufacturer, a diagnostic cut-off at 10 pg/ml secured a combination of very low false positive rate
421	and high sensitivity (red point with yellow halo).
422	
423	FIGURE 2. Variation in the diagnostic sensitivity of the SARS-CoV-2 NP ELISA according to the time gap
424	from the first PCR-positive URT swab to first blood sampling. The blood sample was collected from
425	each of 160 COVID-19 inpatients within 5 weeks after their confirmatory PCR-positive URT swab ($ullet$),
426	and from each of 43 COVID-19 outpatients simultaneously with their PCR-positive URT swab ($^{ u}$). The
427	data point area is proportional to the number of inpatients contributing to the data point.
428	

430

	431	368 blood samples were collected from 160 COVID-19 inpatients within 5 weeks after their
	432	confirmatory PCR-positive URT swab ($ullet$), and 43 blood samples were collected from COVID-19
	433	outpatients simultaneously with their PCR-positive URT swab ([■]). Each inpatient data point area is
	434	proportional to the number of blood samples contributing to the data point.
	435	
	436	FIGURE 4. The median [NP] declined exponentially over time for 368 blood samples collected from
	437	160 COVID-19 inpatients within 5 weeks after their confirmatory PCR-positive URT swab ($ullet$). The
A/R	438	median [NP] of blood samples collected at Day 0-1 for 32 COVID-19 inpatients (1,045 pg/ml) was 36
	439	times higher than the median [NP] (29 pg/ml) of blood samples collected from 43 COVID-19
	440	outpatients simultaneously with their PCR-positive URT swab ($^{\blacksquare}$). For each inpatient data point, the
	441	time of blood sample collection is illustrated as mean \pm SD number of days after the first PCR-positive
	442	URT swab. Each inpatient data point area is proportional to the number of blood samples contributing
	443	to the data point.
	444	
	445	FIGURE 5. Individual dynamics in [NP] of 40 COVID-19 inpatients within 1 month from first PCR-
_		

positive URT swab (total number of blood samples, n=200). A: 3-10 serum samples collected from 446

FIGURE 3. Correlation between the results of SARS-CoV-2 NP ELISA and PCR analysis according to the

time gap from the confirmatory PCR-positive URT swab to collection of serum (A) or plasma (B). The

each of 23 inpatients; B: 3-4 plasma samples collected from each of 17 inpatients. 447

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FIGURE 6. Plasma [NP] above the diagnostic cut-off value (10 pg/ml) was only observed for COVID-19

50 inpatients within the first 26 days of detection of SARS-CoV-2 infection by PCR.



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SUPPLEMENTAL MATERIAL

Detailed characteristics of patients and blood samples included in the study.

The 439 serum samples in this study were collected from 304 individuals, including 24 female COVID-19 patients (mean age \pm SD: 55 \pm 17 years), 29 male COVID-19 patients (62 \pm 16 years), 199 non-infected females (44 \pm 14 years) and 52 non-infected males (42 \pm 13 years).

The 475 plasma samples in this study were collected from 366 individuals, including 73 female COVID-19 patients (mean age \pm SD: 64 \pm 19 years), 82 male COVID-19 patients (68 \pm 18 years), 118 non-infected females (58 \pm 21 years) and 93 non-infected males (62 \pm 20 years).

Figure S1 shows the categorization of patients and blood samples according to 4 differentiators including PCR-based diagnosis, hospitalization, blood fractionation method, and time of collection of blood sample.

No. of Patients in study 670								No	. of S	amp 91	les in stu 4	ıdy			
	COVID-19 Non-infe 208 462		nfected 62				COVI 44	D-19 17		Non-i 4	nfect 67	ed			
	Inp 10	ots. 65	Out 4	pts. 3		Outp 46	ots. 52		Inp 4(ots. 04	Out 4	pts. 3		Out 46	pts. 57
S=Serum	S	Р	S	Ρ		S	Ρ		S	Р	S	Ρ		S	Ρ
P= Plasma	38	127	15	28		251	211		173	231	15	28		251	216
Inpts. first	sample All outpts. sampled Day 0				Inpts. all	samp	les	All	outpl	ts. samp	led D	ay O			
Day 0-34	38	122						Day 0-34	173	195					
Day 0-13	33	110						Day 0-13	131	150					
Day 0-6	26	73						Day 0-6	65	84					

FIGURE S1. Schematic overview of patients and samples included in the study. Beyond the categorization of patients and blood samples according to PCR-based diagnosis, hospitalization, and blood fractionation method, the time of collection of either the inpatient's first blood sample (Patients) or any blood sample (Samples) is reported for the intervals 1 week (Day 0-6), 2 weeks (Day 0-13) and 5 weeks (Day 0-34) after the first PCR-positive URT swab.

Inverse relationship between [NP] and humoral immune response

The rapid decline in serum and plasma [NP] observed for most COVID-19 patients from approximately Day 10 after the first PCR-positive URT swab is consistent with the expected development of a humoral immune response to the SARS-CoV-2 infection.

Selected blood samples of this study were therefore also analyzed by a new prototype "SARS-CoV-2 Neutralization Antibody ELISA Kit". Very briefly, titrated serum or plasma was added to ELISA wells coated with the ACE-2 receptor binding domain of the S1 subunit of SARS-CoV-2 Spike Protein (RBD). After 15 minutes of preincubation at 37°C, ACE-2 receptor conjugated with peroxidase was added to the wells, and incubation of the mixture continued for 45 minutes at 37°C. Blood samples containing neutralizing antibodies to RBD dose-dependently reduced the absorbance of the final colorimetric reaction of the ELISA. The neutralization antibody titer was calculated at 50% of the maximal color formation and compared to the [NP] of the same blood sample (**Table S1** and **Figure S2**).

TABLE S1 . Inverse relationship between the [NP] and the SARS-CoV-2 neutralizing antibody titer in the
initial blood sample collected from 40 COVID-19 inpatients after confirmation by PCR of infection.

Individual [NP]	Patients	Samples	Initial sar	nple: [NP] pĮ	g/ml	Initial sample: Neutralizing Ab titer			
dynamics			Mean ± SD	[range]	Median	Mean ± SD	[range]	Median	
All samples: Over 10 pg/ml	23	122	2,287±1,714	[26-3,840]	3840	34 ± 104	[0 - 454]	0	
Change with time: Over to under 10 pg/ml	13	58	911±1,344	[18-3,840]	197	43 ± 60	[0 - 195]	21	
All samples: Under 10 pg/ml	4	20	2±2	[1-5]	2	298 ± 293	[13 - 700]	240	



FIGURE S2. Indication of the expected inverse relationship between the [NP] and the neutralizing antibody titer of the initial serum (n=23) or plasma sample (n=17) collected from 40 COVID-19 inpatients up to 3 weeks after collection of their confirmatory PCR-positive URT swab. Downloaded from https://journals.asm.org/journal/jcm on 03 September 2021 by 130.225.247.50.

Projected confusion matrix, if using the NP ELISA for nosocomial screening at a low prevalence.

Based on the estimated specificity (99.78%) and sensitivity (82.76%) of the NP ELISA for the 520 patients in the present study with a blood sample collected simultaneously with the URT swab for PCR-analysis, and on an expected rather low prevalence of 0.2% SARS-CoV-2 infection among inpatients hospitalized for more than 3 days for any other disease than COVID-19, the projected confusion matrix for analysis of 100,000 blood samples by the NP ELISA is presented in Table S2.

Screening for no	socomial infection	SARS-Co						
Projected per 10 Expected prevale	0,000 samples ence: 0.2%	Infected inpatients	Non-infected inpatients	Total inpatients	Clinical relevant Predictive value			
	Calculated Positive	166	220	386	PPV	43.01%		
SARS-COV-2	Calculated Negative	34	99,580	99,614	NPV	99.97%		
NF ELISA BIOOU	Total	200	99,800	100,000				
Performance of I	NP ELISA:	Sensitivity	Specificity					
Given sensitivity	and specificity	82.76%	99.78%					

Table S2. Projected confusion matrix if using the NP for screening at low prevalence.

In conclusion, a preceding screening by the NP-ELISA of blood samples from 100,000 inpatients will reduce the number of required URT swabs and associated PCR-analyses from 100,000 primary analyses to 386 confirmatory tests. The screening will identify 166 of the 200 nosocomial SARS-CoV-2 infections.