1	A novel in-cell ELISA assay allows rapid and automated quantification of SARS-CoV-2 to		
2	analyse neutralizing antibodies and antiviral compounds		
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11			
12	Keywords: SARS-CoV-2; COVID-19; neutralizing; antibodies; interferon		
13	Running title: Rapid & automated SARS-CoV-2 icNT assay		
14	Brief summary: Knowledge concerning SARS-CoV-2-neutralizing antibodies (NAbs) is		
15	indispensable for COVID-19 convalescent plasma selection, evaluation of vaccines, and immunity		
16	certificates. Our in-cell ELISA (icELISA) test allows rapid (<48h) and high-throughput detection		
17	and quantification of SARS-CoV-2-specific NAbs and antiviral activity of drug candidates.		
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28 Abstract

The coronavirus disease 2019 (COVID-19) caused by the severe acute respiratory syndrome 29 coronavirus 2 (SARS-CoV-2) is currently the most pressing medical and socioeconomic challenge. 30 Constituting important correlates of protection, determination of virus-neutralizing antibodies 31 (NAbs) is indispensable for convalescent plasma selection, vaccine candidate evaluation, and 32 immunity certificates. In contrast to standard serology ELISAs, plaque reduction neutralization 33 tests (PRNTs) are laborious, time-consuming, expensive, and restricted to specialized laboratories. 34 35 To replace microscopic counting-based SARS-CoV-2 PRNTs by a novel assay exempt from genetically modified viruses, which are inapplicable in most diagnostics departments, we 36 37 established a simple, rapid, and automated SARS-CoV-2 neutralization assay employing an in-cell ELISA (icELISA) approach. 38

After optimization of various parameters such as virus-specific antibodies, cell lines, virus doses, 39 40 and duration of infection, SARS-CoV-2-infected cells became amenable as direct antigen source for quantitative icELISA. Using commercially available nucleocapsid protein-specific antibodies, 41 viral infection could easily be quantified in human and highly permissive Vero E6 cells by 42 icELISA. Antiviral agents such as human sera containing NAbs or antiviral interferons dose-43 dependently reduced the SARS-CoV-2-specific signal. Applying increased infectious doses, the 44 icNT was superior to PRNT in discriminating convalescent sera with high from those with 45 intermediate neutralizing capacities. 46

The SARS-CoV-2 icELISA test allows rapid (<48h in total, read-out in seconds) and automated quantification of virus infection in cell culture to evaluate the efficacy of NAbs as well as antiviral drugs, using reagents and equipment present in most routine diagnostics departments. We propose the icELISA and the icNT for COVID-19 research and diagnostics.

51 Introduction

By the time of writing, more than 6.3 million people experienced a laboratory confirmed infection with the severe acute respiratory syndrome coronavirus (SARS-CoV)-2 and more than 378,000 people died while having coronavirus infectious disease 19 (COVID-19). First surveillance studies and calculations of excess mortality rates indicate that the precise number of infections and the true number of fatalities exceed above-mentioned numbers by far.

Coronaviruses (CoVs) are positive strand RNA viruses widespread among various vertebrate hosts 57 including bats and rodents (1). Together with four seasonal human CoVs (hCoVs) as well as the 58 two other emerging hCoVs SARS-CoV-1 and MERS-CoV, SARS-CoV-2 is the seventh hCoV 59 causing widespread human diseases (2). In December 2019, SARS-CoV-2 was first recognized in 60 the Hubei province in China (3) from where it rapidly spread throughout the world. In addition to 61 62 its genetic similarity, SARS-CoV-2 shares some clinical characteristics with SARS-CoV-1 (4), but also exhibits some highly relevant particularities such as an increased spreading efficacy and the 63 length of the course of disease (5). On January 31, the WHO declared the SARS-CoV-2 outbreak 64 a Public Health Emergency of International Concern. On March 11, 2020, WHO started to denote 65 the outbreak a global pandemic. Since its beginning, the centre of the pandemic shifted from China, 66 via Europe and Northern Americas to Central and Southern Americas. This dynamic nature of the 67 68 pandemic poses an inherent danger of repetitive local and temporal reintroduction circles. Thus, even countries which coped relatively well with the first wave must prepare in terms of diagnostics 69 70 capacities for potential future re-emergences.

Most SARS-CoV-2 infections lead to mild or moderate illnesses. However, a considerable fraction
 of cases proceeds to severe pneumonia or life-threatening acute respiratory distress syndrome.

Elderly individuals and people with pre-existing comorbidities such as impaired immunity, chronic
respiratory diseases, cardiovascular diseases, and cancer are more prone to suffer from severe
COVID-19. The case fatality rate (CFR) is difficult to calculate in the midst of the pandemic.
Depending on the age, CFR estimates of up to 18.4% for individuals older than 80 years and 1.38%
(range 1.23 - 1.53) for the general population have been reported (6).

Given the extent, pace, and severity of the COVID-19 pandemic, diagnostics departments even in countries with highly developed medical systems struggle to provide sufficient and timely test capacities. Since nucleic acid-based pathogen detection, usually by quantitative RT-PCR based on naso- or oropharyngeal swabs, has a very short window of opportunity, assays detecting longlasting immune responses such as antibodies are required to monitor virus spread and to estimate potential herd immunity.

84 With some delay, most infected individuals raise a detectable humoral immune response including 85 specific immunoglobulins (Ig) M, IgA, and IgG (7-9). Neutralizing antibodies (NAbs) bind and abrogate the function of viral proteins such as the SARS-CoV-2 Spike (S) protein that are essential 86 for virus entry into host cells, e.g., through recognition of the cognate receptor ACE2 (10). 87 Accordingly, monoclonal NAbs exhibit strong therapeutic and prophylactic efficacies in SARS-88 CoV-2-infected human ACE2-transgenic mice (11). A recent vaccination study conducted in non-89 human primates identified NAbs as correlate of protection (12), indicating that a human SARS-90 CoV-2 vaccine should also be capable to elicit potent NAb responses. Additionally, monoclonal 91 NAbs and NAb-containing hyper-immunoglobulin preparations may be applicable as treatment 92 against COVID-19 (13). NAbs are also the backbone of convalescent plasma (CP) therapies (14-93 16) that are one of 7 clinical recommendations of the IDSA (17). Based on the havoc COVID-19 94 causes to the global economy, immunity passports and vaccination certificates, documenting 95

protection through NAbs, have been discussed (e.g., (18)). Taken together, SARS-CoV-2-specific
NAbs and their quantification appear to be of central importance for the medical and socioeconomic management of the pandemic.

99 Different types of neutralization tests (NT) have been developed for SARS-CoV-2. However, to our knowledge, these assays rely on laborious microscopic counting of virus plaques or antibody-100 stained foci by trained personnel (19-21) or on genetically modified viruses such as transgenic 101 SARS-CoV-2 mutants (22) or pseudo-typed viruses (e.g., Vesicular stomatitis virus [VSV] or 102 Human immunodeficiency virus [HIV]) expressing the S protein of SARS-CoV-2 (23, 24). 103 Genetically modified viruses are generally prohibited in routine diagnostics laboratories and 104 usually not applicable in less developed regions. Due to the central importance of NAbs and the 105 limitations of the currently available methodology, we established a cheap, simple, fast, reliable, 106 and automatable in-cell ELISA (icELISA)-based icNT applicable in most routine diagnostics 107 108 departments.

109

110 Methods

111 Cells, viruses, interferons, and sera

112 Caco-2 (ATCC HTB-37) and Vero E6 (ATCC CRL-1586) were cultivated in Roswell Park 113 Memorial Institute (RPMI) and Dulbecco's minimal essential medium (DMEM), respectively, 114 supplemented with 10% (v/v) FCS, penicillin, and streptomycin at 37°C in an atmosphere of 5% 115 CO. SARS-CoV-2 was isolated from a patient sample using Vero E6 and confirmed by SARS-116 CoV-2 diagnostic qRT-PCR. Viral titers were determined by TCID50 titration. Human IFN α 2 and 117 IFN β were purchased from PBL Assay Science (#11101) and Peprotech (#300-02BC), respectively. The collection of serum samples has been approved by the ethics committee of the medical faculty of the University of Duisburg- Essen (20-9208-BO). Anti-SARS-CoV-2 IgG antibodies were detected using ELISA detecting SARS-CoV-2 Spike protein (Euroimmun Medizinische Labordiagnostika, Germany) according to manufacturer's instructions

122 SARS-CoV-2 icELISA and icNT

123 Defined doses of SARS-CoV-2 were incubated with different serum dilutions for 1 h at 37°C prior 124 to Vero E6 infection. Neutralizing capacities were evaluated by icELISA. A detailed icELISA/ 125 icNT protocol is provided in Suppl. File 1. The α -N mAb1 (ABIN6952435), α -N mAb2 126 (ABIN6952433), α -S Ab (ABIN1031551), and POD-coupled secondary antibodies (Dianova) 127 were used.

128

129 **Results**

130 Quantification of SARS-CoV-2 replication and its inhibition by antiviral compounds using
131 icELISA

We hypothesized that virus-encoded proteins expressed by infected cells should be amenable as source of viral antigens for the detection and quantification by ELISA. We optimized the experimental conditions such as cell type, virus dose, infection period, cell fixation method, blocking reagent, as well as type and concentrations of primary and secondary antibodies (Fig. 1 and data not shown). As described in the Methods section and provided as detailed laboratory protocol in the supplementary information (Suppl. File 1), we compared different commercially available antibodies for the icELISA-based quantification of SARS-CoV-2 proteins. Based on existing data on virus entry (*10*), we infected human Caco-2 cells with graded virus doses (ranging from 0.03125 to 2 PFU/cell). At 3 days post infection (d p .i.), we fixed and stained the cells with different antibodies either recognizing the S or the N protein of SARS-CoV-2. In accordance with high expression level of the N protein (*25, 26*), certain N-specific mouse IgGs exhibited a signalto-noise ratio favourable for icELISA (Fig. 1A). A comparison of Vero E6 and human Caco-2 cells revealed that the icELISA is applicable to different cells (Fig. 1B).

Since the icELISA signal directly correlated with viral replication and viral antigen expression, we tested its ability to determine antiviral effects. Human Caco-2 cells were treated with graded concentrations of human interferon (IFN) α 2 or IFN β and infected 3 h later with SARS-CoV-2. At 3 d p.i., viral antigen amounts were quantified by icELISA. In accordance with a recent clinical phase 2 trial (27), IFN β exhibited strong and dose-dependent antiviral activity against SARS-CoV-2 in human cells (Fig. 1C), indicating that the icELISA is applicable for future experiments addressing the efficacy of potential antiviral drugs in human cells.

Despite different start MOIs, similar icELISA signals were observed at 72 h p.i. in Vero E6 (Fig. 1B), indicating multiple rounds of virus amplification and extraordinary fast replication kinetics of SARS-CoV-2 in Vero E6 cells, consistent with previous studies (28). To test if shorter infection periods might result in virus dose-dependent signals, we analysed infected Vero E6 cells after 6, 15, and 22 h. SARS-CoV-2 was readily detectable in Vero E6 cells by icELISA already at 6 h p.i. (Fig. 1D-F).

Taken together, these data indicate that the icELISA allowed rapid identification and quantification
of SARS-CoV-2 replication in Vero E6 and human cells as well as its inhibition by antiviral
compounds.

161 Neutralization tests based on icELISA allow the quantification of SARS-CoV-2 NAbs as early as 6
162 hours post infection

163 Since the icELISA allowed simple and automated quantification of SARS-CoV-2-dependent 164 antigen expression, we tested if an icELISA-based neutralization test (icNT) can be established. We infected Vero E6 cells for 6, 15, and 24 h with graded SARS-CoV-2 infection doses (500, 165 5000 or 50,000 PFU/well) in the absence or presence of 2 convalescent sera in 3 different 166 concentrations (1/20, 1/40, and 1/80 dilution). Using the high infectious dose, viral antigens 167 became dose-dependently detectable by icELISA as early as 6 h p.i. (Fig. 2A). Both immune sera 168 strongly and dose-dependently neutralized the virus-induced signal (Fig. 2B). At 15 h p.i., the 169 intermediate infectious virus dose also became detectable (Fig. 2C) and both human sera 170 efficiently neutralized the infection (Fig. 2D). At 24 h p.i., all infection conditions resulted in a 171 strong icELISA signal (Fig. 2E), indicating that infectious doses as low as 500 PFU/well are 172 detectable by icELISA after 24 h of infection. Even the strong icELISA signal at 24 h p.i. was 173 dose-dependently neutralized by both sera (Fig. 2F). Please note that the neutralizing capacity of 174 given sera dilutions was less pronounced at higher virus doses as compared to lower virus doses, 175 176 as indicated by residual icELISA signals.

Taken together, the icELISA resulted in a time- and virus dose-dependent signal constituting a surrogate for SARS-CoV-2 infection and replication. The fact that the infection and the resulting icELISA signal were neutralized by NAbs present in immune sera indicated that the fast and automated icELISA format is applicable for icNTs.

181 *The icNT results correlate with the standard SARS-CoV-2 neutralization test conducted by staining*

182 *of virus foci and microscopic counting*

Although most SARS-CoV-2 NTs have not been formally validated and certified, classic plaque 183 reduction neutralization tests (PRNT) are currently considered to represent the gold standard for 184 the detection of SARS-CoV-2-specific NAbs. Various commercially available IgM, IgA, and IgG 185 ELISAs have been compared to PRNTs (e.g., (28)). To validate the novel icNT, 20 sera - 10 186 positive for SARS-CoV-2-specific IgG as determined using the Euroimmun ELISA (ELISA ratio: 187 1.26 to 11.39) and 10 ELISA-negative sera (ELISA ratio < 0.9) - were compared side-by-side by 188 189 icNT and standard PRNT using 200 PFU/well (Fig. 3A). One set was processed by icNT, while 190 for the other set virus foci were stained using an antibody-based AEC staining method and manually counted by microscopy. The icNT correctly recognized all 10 positive and all 10 negative 191 192 sera (Fig. 3B), indicating optimal sensitivity and specificity. In total, both methods gave almost congruent results in terms of the highest dilution resulting in 50% neutralization (Pearson's 193 correlation coefficient 0.965), highlighting excellent test performances of the icNT compared to 194 the PRNT. 195

196 The icNT discriminates SARS-CoV-2-neutralizing sera from non-immune sera and provides
197 superior resolution when increased virus doses are used

Standard SARS-CoV-2 NTs base on microscopic counting of plaques or antibody-stained virus 198 foci. To enable plaque/foci recognition and individual counting by trained personnel, a countable 199 number of PFU must be applied in PRNTs. Depending on the PRNT protocol, 100 (19) to 400 200 PFU (20) are used to infect each well. Based on previous experiences with virus neutralization 201 experiments (29, 30), we suspected that lower infectious doses might be more susceptible to NAbs 202 than higher virus doses, e.g., through altered ratios of NAbs and antigenic regions determining 203 neutralization, such as the receptor binding domain (RBD) of the S protein of SARS-CoV-2. To 204 address if the icNT can provide superior resolution in terms of discriminating intermediate from 205

206 strong NAb responses in convalescent sera, we selected prototypical sera and conducted comparative icNTs applying either 200 PFU/well or 40-fold more virus (8000 PFU/well). While 207 all positive sera clearly neutralized the low infectious dose of 200 PFU/well (Fig. 4A-C), the 208 neutralizing capacity of the same sera considerably differed at the more restrictive high virus dose 209 (Fig. 4D-F). Despite their neutralizing capacity at low dose infections, some of the convalescent 210 sera (e.g., serum #19) almost did not show appreciable neutralization in high MOI infection and 211 212 did not reach the typical benchmark of 50% neutralization at a 1/8 serum dilution. Thus, we 213 concluded that the icNT principle allows higher infectious doses which appear to discriminate intermediate from superior SARS-CoV-2 neutralizing antibody titres. 214

215

216 **Discussion**

We established a novel icELISA-based test principle for detection and quantification of SARSCoV-2. Given the excellent signal-to-noise ratio between infected and uninfected cells, the test
was applicable to quantify the efficacy of antiviral compounds, here shown for IFNβ, and SARSCoV-2-specific NAbs present in immune sera.

Compared to icELISA and icNT, standard virus titrations and PRNTs are far more laborious, time consuming, and expensive - not to speak from subjective and expectation biases upon usage for research. The entire icNT can be processed in less than 2 days including the infection period. Given that the protocol includes an early fixation step using 4% (w/v) paraformaldehyde which inactivates SARS-CoV-2, all subsequent steps can be processed outside the biosafety level 3 laboratory. The actual data acquisition is conducted within seconds, using standard ELISA plate readers present in most routine diagnostics departments. The icELISA and icNT provide increased

data quality and precision by generating continuous data sets. Since the detection antibodies can 228 be applied in icELISA and icNTs in relatively high dilutions (1/5000 and 1/2000 of primary and 229 secondary antibody, respectively), the assay is relatively cheap (in our case, around 0.10 € per well 230 for both antibodies and the TMB peroxidase ELISA substrate). The specificity of the icELISA and 231 icNT is provided by the defined SARS-CoV-2 added to the cell cultures on purpose. The primary 232 and secondary detection antibodies just serve to visualize and quantify viral antigens. Thus, SARS-233 234 CoV-2-specific antibodies can be applied for icELISA detection notwithstanding potential cross-235 recognition of other CoVs such as hCoV-HKU1 or hCoV-OC43 - simply because these viruses are not present in the culture. Obviously, such antibodies recognizing conserved residues cannot 236 237 be used for classic antigen-recognizing ELISAs due to their inability to discriminate coronaviruses. More virus (>1 PFU/cell) can be applied to icNTs. Such high-PFU icNTs scrutinize the virus-238 neutralizing capacity of sera more strictly, enabling a higher resolution compared to PRNT assays 239 240 which all rely on low virus numbers. It is tempting to speculate that sera exhibiting superior neutralization in high PFU icNT might be more beneficial in CP therapies. 241

NTs based on pseudo-typed viruses using heterologous expression of the SARS-CoV-2-encoded S by viruses may have certain advantages. However, genetically modified viruses are inapplicable by law in various routine diagnostics departments and usually unavailable in less developed countries. Without the use of genuine infectious viruses, assays relying on pseudo-typed viruses do not fully interrogate the full spectrum of antiviral effects for example if other viral proteins influence the system, e.g., by complement activation (*31*).

Taken together, we propose the icELISA and icNT for the quantification of SARS-CoV-2 replication and its inhibition by NAbs and antiviral compounds. By changing the detection antibody, the test principle is transferable to all other viruses.

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260 Author's contributions

- 261 L.S., V.T.K.L-T., M.E., and D.M. performed research. O.E.A., A.K., A.H. provided essential
- reagents. L.S., V.T.K.L-T., and M.T. analysed data. L.S., V.T.K.L-T., U.D. and M.T. interpreted
- data. V.T.K.L-T and M.T. supervised the project. L.S., V.T.K.L-T., and M.T. wrote the
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Figures and figure legends

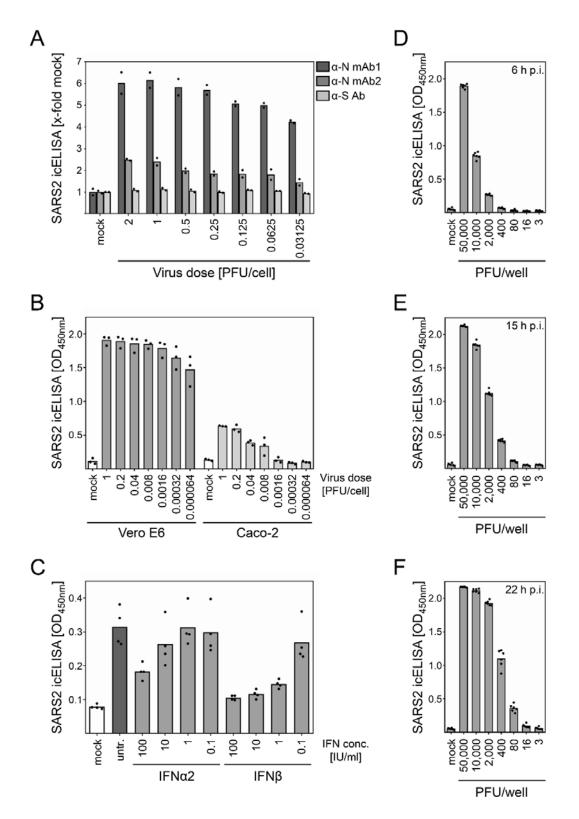


Figure 1: The icELISA test allows quantification of SARS-CoV-2 replication and its inhibition by antiviral compounds

(A) Caco-2 cells were infected with indicated doses of SARS-CoV-2. At 3 d p.i., cells were fixed and detected by icELISA using S- and N-specific primary antibodies. For all further icELISAs, α -N mAb1 was used. (B) Vero E6 and Caco-2 cells were infected with indicated doses of SARS-CoV-2. At 3 d p.i., cells were analysed by icELISA. (C) Caco-2 cells were treated with indicated concentrations of IFN α 2 or IFN β . At 3 h post treatment, cells were infected with SARS-CoV-2 (MOI 0.1). Viral replication was evaluated at 3 d p.i. by icELISA. (D - F) Vero E6 cells were infected with indicated doses of SARS-CoV-2. At 6, 15, and 22 h p.i. (D, E, and F, respectively), cells were analysed by icELISA. Bars depict the mean values. Dots show the values of the individual measurements.

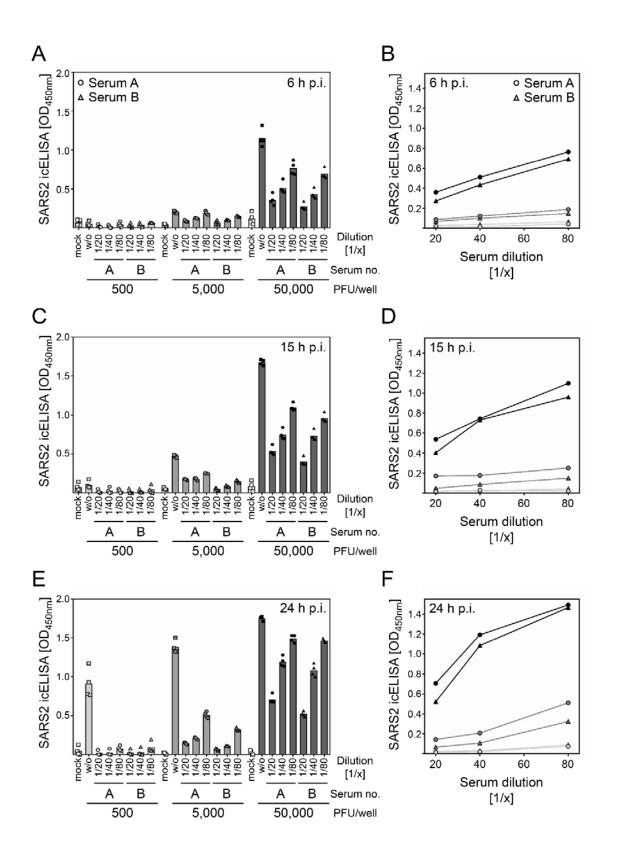
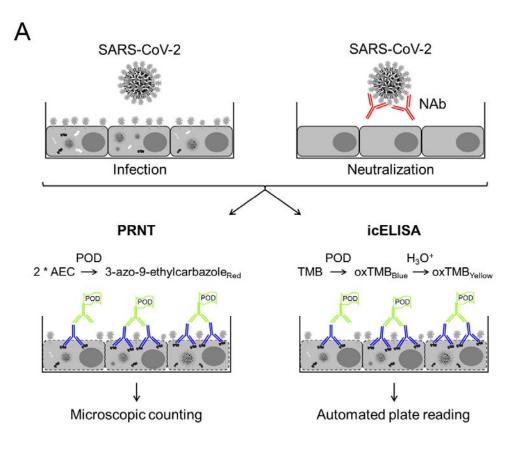


Figure 2: The icNT allows the quantification of SARS-CoV-2 NAbs as early as 6 h p.i.

(A - F) 500, 5000, and 50,000 PFU of SARS-CoV-2 were incubated with indicated dilutions of 2 convalescent sera for 1 h before Vero E6 cells were infected. At 6 h p.i. (A, B), 15 h p.i. (C, D), and 24 h p.i. (E, F), cells were analysed by icELISA to evaluate the neutralizing capacity of the sera. (A, C, E) Bars depict the mean values. Dots show the values of the individual measurements. (B, D, F) All mean values of the different serum dilutions and virus doses were depicted in one diagram to compare the influence of the input virus amount on the course of the curve. Light grey, 500 PFU. Grey, 5000 PFU. Dark grey, 50,000 PFU.



В

Serum	50% neutralisation titre [dilution 1/x]		
[no]	PRNT	icNT	
1	256	>256	
2	0	0	
3	>256	>256	
4	0	0	
5	128	128	
6	0	0	
7	>256	>256	
8	0	0	
9	>256	>256	
10	0	0	
11	>256	>256	
12	0	0	
13	128	128	
14	0	0	
15	>256	>256	
16	0	0	
17	>256	>256	
18	0	0	
19	128	256	
20	0	0	
	Pearson's <i>r</i> : 0.965		

Figure 3: The icNT results correlate with the standard SARS-CoV-2 neutralization test

(A) Schematic representation of AEC stain-based classic NT and icELISA-based icNT. NAb, neutralizing antibody. PRNT, plaque reduction neutralization test. POD, peroxidase. AEC, 3-amino-9-ethylcarbazole. TMB, tetramethylbenzidine. (B) 200 PFU of SARS-CoV-2 were incubated with different dilutions (1/8, 1/16, 1/32, 1/64, 1/128, 1/256) of 20 serum samples (10 negatively and 10 positively tested for SARS-CoV-2 IgG by Euroimmun ELISA) for 1 h before Vero E6 were infected. At 20 and 48 h p.i., neutralizing capacity was evaluated by icELISA and AEC staining with subsequent microscopic counting, respectively. The highest dilution capable to neutralize 50% of input virus was determined and results of PRNT and icNT were compared.

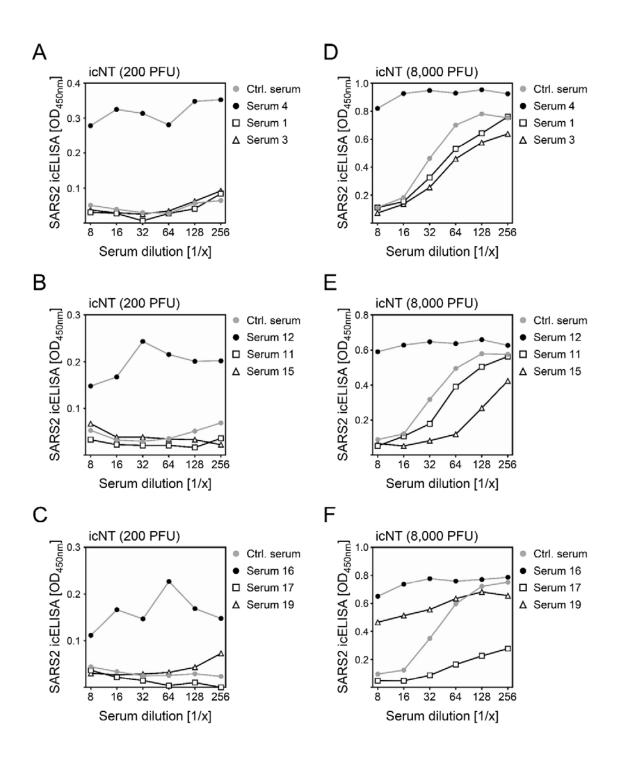


Figure 4: The icNT provides superior resolution upon usage of increased virus doses

(A - F) 200 or 8,000 PFU of SARS-CoV-2 were incubated with indicated dilutions of serum samples for 1 h before Vero E6 cells were infected. At 20 and 16 h p.i. (for 200 and 8,000 PFU, respectively), neutralizing capacity was evaluated by icELISA. Samples measured on the same plate are depicted in one diagram. The control serum was used as reference for all plates.

icNT icELISA Assay Protocol Trilling Lab

SARS-CoV-2 icNT (steps 1 to 13) and icELISA (steps 6 to 13)

Materials:

- Vero E6 cells
- DMEM + Glucose/Glutamine/Pyruvate supplemented with 10% FCS, Pen/Strep
- Virus stock SARS-CoV-2 (titer: at least 2x 10^5 PFU/ml)
- Primary and secondary antibody (α-N mAb, α-mouse IgG POD-coupled)
- Paraformaldehyde (PFA)
- 1x PBS (PBS), 2x PBS
- Triton-X-100
- Tween-20
- Fetal Calf Serum (FCS)
- Tetramethylbenzidine (TMB)
- 0.5 M HCI
- Distilled or deoinized water

Equipment:

- 96-well microplate
- 37°C CO₂ incubator
- Multichannel pipette
- Microplate reader

Notes:

- The wells of the microplate should not be allowed to dry at any point during the assay procedure.
- It is recommended to use a plate shaker during the incubation steps of the icELISA.

REAGENT	REAGENT PREPATION INSTRUCTION	WHEN TO PREPARE?
Wash Buffer 🥒	500µl Tween-20 + 1 liter PBS	At start of experiment
8% (w/v)	1.) Dissolve a 2x PBS tablet in 500 ml	At start of experiment
Paraformaldehyde	distilled or deionized water	(store in aliquots
Solution	2.) Add 80 gr PFA	at -20°C)
	 Use 4 M NaOH to dissolve the PFA in PBS 	
	Use HCl for pH adjustment (pH 7,4)	
	 Fill up with distilled or deionized water (1 liter) 	
Permeabilization	For one 96-well microplate:	Prepare immediately
Buffer	Dilute 250µl Triton-X-100 in 24.75 ml PBS.	before use
	Mix well by vortexing.	
Blocking Solution	3% (v/v) FCS in PBS	Prepare immediately
		before use
Incubation Buffer	1% (v/v) FCS in PBS	Prepare immediately
		before use

1

icNT icELISA Assay Protocol Trilling Lab

- 1.) Seed Vero E6 cells into 96-well microplates the day before.
 - → Perform the test at least in duplicate, better in quadruplicate.
 - → Include on every plate virus control and NT with control sera as references.
 - a. Cells should be 80 100% confluent at the day of infection.
 - b. Plates can be prepared outside of the BSL3 lab.
- 2.) Dilute the control serum and the sera to be tested 1/4, 1/8, 1/16, 1/32, 1/64, 1/128, 1/256.
 - → Final serum dilutions are 1/8, 1/16, 1/32, 1/64, 1/128, 1/256, 1/512 after adding the virus solution.
 - → Add further dilutions if needed.
 - a. Calculate the required total volume of the different dilutions (50µl per well) including at least 20% excess volume.
 - b. Prepare double the amount of the required volume of 1/4 serum dilution in medium (to do 2-fold serial dilutions).
 - c. Prepare stepwise the other dilutions.
 - d. Discard the excess amount of the last dilution.
- 3.) Prepare the virus solution (5.5 ml per 96-well microplate).
 - a. Calculate the required amount of total virus (8,000 10,000 PFU per well in 50µl).
 - b. Prepare the virus solution by diluting the required amount of virus stock in medium.
- 4.) Neutralisation: Incubation of virus with serum.
 - Add virus solution to the prepared serum dilutions (50µl serum dilution + 50µl virus solution per well).
 - b. Incubate for 60 90 min at 37 °C.
- 5.) Infection.
 - a. Aspirate the medium of the Vero E6 cells in the 96-well microplate.
 - b. Prevent cells from drying out during aspiration step.
 - c. Pipette the virus-serum solution into the wells (100µl per well).
 - d. Include the virus control and the control sera on every plate.
- 6.) Fix cells to microplate.
 - a. At 16 24 h p.i., add an equal volume of 8% paraformaldehyde solution to the wells
 - containing culture media (= 4% paraformaldehyde solution)
 - b. Incubate for at least 2 h (or overnight) at room temperature.

→ For experiments using non-BSL3 organisms, 15 min of fixation is sufficient.

7.) Replace the lids with new ones. Wipe the plates thoroughly with 4% Dismozon. Plates can now be discharged from the BSL3 laboratory.

2

icNT icELISA Assay Protocol Trilling Lab

- 8.) Gently aspirate the fixing solution from the microplate. Wash the microplate 3 times with 300µl PBS per well. Add 200µl PBS to the wells. The microplate (sealed with parafilm) can now be stored at 4°C for several days.
- 9.) Permeabilize cells.
 - a. Prepare permeabilization buffer (for one microplate: dilute 250µl Triton-X-100 in 24.75 ml PBS).
 - b. Aspirate PBS and add 200µl of freshly prepared permeabilization buffer to each well.
 - c. Incubate for 30 min.
- 10.) Blocking.
 - Aspirate permeabilization buffer and add 100µl of blocking solution (3% FCS in PBS) to each well.
 - b. Incubate for 1-2 h.
- 11.) Incubation with primary antibody.
 - Prepare primary antibody (α-SARS-CoV-2-N, 1:5000) by diluting stock antibody in the required volume of incubation buffer (1% FCS in PBS).
 - b. Aspirate blocking solution and add 50µl of antibody solution to each well.
 - c. Incubate for 2 h at room temperature or overnight at 4°C (sealed with parafilm).
- 12.) Incubation with secondary antibody.
 - a. Prepare antibody solution by diluting stock antibody in the required volume of incubation buffer (1% FCS in PBS).
 - Aspirate primary antibody solution. Wash the microplate 3 times with 250µl wash buffer per well.
 - c. Aspirate the wash buffer and add 50µl of secondary antibody solution to each well.
 - d. Incubate for 1-2 h at room temperature.
 - e. Aspirate secondary antibody solution. Wash 4 times with 250µl wash buffer per well.
- 13.) Signal measurement.
 - a. Prepare the required amount of TMB and 0.5 M HCl.
 - b. Aspirate the last wash. Make sure that you have completely removed the liquid.
 - c. Add 100µl TMB (blue color development).
 - d. Stop the reaction with 100µl of 0.5 M HCl (by the time the uninfected wells [mock] are starting to turn blue as well).
 - e. Record data at 450 nm absorbance, 620 nm reference using a microplate reader.
 - 3