

Fcγ-Receptor-Based Enzyme-Linked Immunosorbent Assays for Sensitive, Specific, and Persistent Detection of Anti-SARS-CoV-2 Nucleocapsid Protein IgG Antibodies in Human Sera

[®] Christina Deschermeier, ^a Christa Ehmen, ^a Ronald von Possel, ^{b,c} Carolin Murawski, ^a Ben Rushton, ^a John Amuasi, ^d Nimako Sarpong, ^e Oumou Maiga-Ascofaré, ^{e,f} Raphael Rakotozandrindrainy, ^g Danny Asogun, ^h Yemisi Ighodalo, ^h Lisa Oestereich, ^{b,f} Sophie Duraffour, ^{b,f} Meike Pahlmann, ^{b,f} Petra Emmerich^{b,c}

^aDiagnostics Development Laboratory, Bernhard Nocht Institute for Tropical Medicine, Hamburg, Germany
^bDepartment for Virology, Bernhard Nocht Institute for Tropical Medicine, Hamburg, Germany
^cDepartment of Tropical Medicine and Infectious Diseases, Center of Internal Medicine II, University of Rostock, Rostock, Germany
^dGlobal Health and Infectious Disease Research Group, Kumasi Centre for Collaborative Research in Tropical Medicine, Kumasi, Ghana
^eInfectious Disease Epidemiology Research Group, Kumasi Centre for Collaborative Research in Tropical Medicine, Kumasi, Ghana
^eGerman Center for Infection Research (DZIF), Hamburg-Lübeck-Borstel-Riems, Hamburg, Germany
^gDepartment for Microbiology and Parasitology, University of Antananarivo, Antananarivo, Madagascar
^hInstitute of Lassa Fever Research and Control, Irrua Specialist Teaching Hospital, Irrua, Nigeria

ABSTRACT Sensitive and specific serological tests are mandatory for epidemiological studies evaluating severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) prevalence as well as coronavirus disease 2019 (COVID-19) morbidity and mortality rates. The accuracy of results is challenged by antibody waning after convalescence and by cross-reactivity induced by previous infections with other pathogens. By employing a patented platform technology based on capturing antigen-antibody complexes with a solid-phase-bound Fcy receptor (FcyR) and truncated nucleocapsid protein as the antigen, two SARS-CoV-2 IgG enzyme-linked immunosorbent assays (ELISAs), featuring different serum and antigen dilutions, were developed. Validation was performed using a serum panel comprising 213 longitudinal samples from 35 COVID-19 patients and a negative-control panel consisting of 790 pre-COVID-19 samples from different regions of the world. While both assays show similar diagnostic sensitivities in the early convalescent phase, ELISA 2 (featuring a higher serum concentration) enables SARS-CoV-2 IgG antibody detection for a significantly longer time postinfection (≥15 months). Correspondingly, analytical sensitivity referenced to indirect immunofluorescence testing (IIFT) is significantly higher for ELISA 2 in samples with a titer of \leq 1:640; for high-titer samples, a prozone effect is observed for ELISA 2. The specificities of both ELISAs were excellent not only for pre-COVID-19 serum samples from Europe, Asia, and South America but also for several challenging African sample panels. The SARS-CoV-2 IgG FcyR ELISAs, methodically combining antigen-antibody binding in solution and isotype-specific detection of immune complexes, are valuable tools for seroprevalence studies requiring the (long-term) detection of anti-SARS-CoV-2 IgG antibodies in populations with a challenging immunological background and/or in which spike-proteinbased vaccine programs have been rolled out.

KEYWORDS immunoassay, infectious disease, immunoglobulins, laboratory methods and tools, viral diseases

Reliable serological assays are mandatory for monitoring the progression of the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) pandemic, particularly in regions of the world where access to medical care and molecular testing capacities are limited. To enable a conclusive estimation of SARS-CoV-2 seroprevalence, a SARS-CoV-2 IgG assay has

Editor Yi-Wei Tang, Cepheid

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Address correspondence to Christina Deschermeier, descherm@bnitm.de. The authors declare a conflict of interest. Research Funding: C. Deschermeier, German Federal Ministry of Education and Research (grant no. 01Kl20210) and European Regional Development Fund (FRDF) (grant no. BWF/H/ 52228/2012/13.10.10-1/3.4,6) to institution; P. Emmerich, German Federal Ministry of Education and Research (grant no. 01Kl20210) to institution; L. Oestereich: Leibniz Association (grant no. J59/20218) to institution. Patents: C. Deschermeier, FP3207375; P. Emmerich, FP2492689, FP3207375, Other financial or nonfinancial interests: C. Deschermeier and B. Rushton are cofounders and shareholders of Panadea Diagnostics GmbH. Role of Sponsor: The funding organizations played no role in the design of study, choice of enrolled patients, review and interpretation of data, preparation of manuscript, or final approval of manuscript. No other Potential Conflicts of Interest have been declared.

Received 21 January 2022 Returned for modification 13 February 2022 Accepted 14 April 2022 Published 16 May 2022

to fulfill several crucial requirements. First, due to postconvalescence antibody waning (1), high assay sensitivity in coronavirus disease 2019 (COVID-19) patient samples collected more than 6 months after infection is needed. Second, although SARS-CoV-2 IgG assay specificity in general is good to excellent in sample panels from Europe, Asia, and the United States (2), several African serum panels were recently shown to strongly challenge assay specificity due to a different immunological background caused by endemic infectious diseases such as Plasmodium falciparum malaria and others (3–6). Therefore, if use in these settings is intended, critical assessment and (where necessary) optimization of assay specificity are required. Third, with the progressive rollout of country-specific vaccine programs, humoral immune responses induced by natural SARS-CoV-2 infection must be differentiated from vaccine responses. If spike-based vaccines like, e.g., Comirnaty (BioNTech/Pfizer) or Vaxzevria (AstraZeneca) are administered, assays employing the SARS-CoV-2 nucleocapsid protein (NCP) (7) as the antigen still allow the unequivocal detection of antibodies induced by natural SARS-CoV-2 infection. Notably, highly sensitive assays will be required to detect the apparently reduced humoral anti-NCP response that has been observed in vaccine breakthrough infections (8). In contrast, vaccines based on inactivated SARS-CoV-2 (e.g., the Sinopharm vaccine) potentially elicit antibodies against all structural virus proteins. In principle, natural infection in individuals immunized with those vaccines could be proven by the detection of antibodies targeting nonstructural proteins like open reading frame 3 (ORF3) or ORF8 (9). Nevertheless, several SARS-CoV-2 lineages lacking the functional expression of these proteins have been described (10, 11).

During the last few years, we have employed a patented platform technology (12, 13) methodically combining antigen-antibody binding in solution with isotype-specific detection of in vitro-formed immune complexes to develop several highly sensitive and specific plate-based enzyme-linked immunosorbent assays (ELISAs) for the detection of both IgM and IgG antibodies directed against viral pathogens such as dengue virus (DENV) (14), Lassa virus (LASV) (15), Crimean-Congo hemorrhagic fever virus (CCHFV) (16), and Zika virus (ZIKV) (17, 18). Briefly, for the detection of IgG, patient sera are coincubated with soluble, labeled recombinant antigen (and, if necessary, unlabeled competitor molecules suppressing cross-reactive antibody binding) on a 96-well plate coated with the recombinantly produced immunoglobulin-like (IgI) domain of Homo sapiens Fcg receptor IIA_{H131} (HsFcyRIIA_{H131}) (mediating the isotype-specific binding of IgG-antigen immune complexes). Thereby, the use of natively folded antigen in the liquid phase guarantees the preservation of conformational epitopes and prevents nonspecific interactions with hydrophobic amino acid stretches that may be exposed in partially unfolded or denatured proteins. If necessary, even high serum concentrations can be applied without increasing the assay background (19). Captured immune complexes are then detected by a colorimetric reaction that can be quantified using a conventional ELISA reader.

Here, we utilize this technology to develop ELISA protocols for the detection of antibodies directed against the SARS-CoV-2 NCP and present detailed validation data on reproducibility, diagnostic and analytical sensitivities, specificity, and interference.

MATERIALS AND METHODS

Generation of prokaryotic expression vectors (pOPIN-J-CoV-N2b/SB/N3). Amplicons encoding the N2b/SB/N3 domain (7) of coronavirus (CoV) NCPs were generated by PCR (see Table S1 in the supplemental material) using either random-primed cDNA transcribed from virus RNA or synthesized cDNA (codon optimized for expression in *Escherichia coli*; GenScript) as a template and inserted by in-fusion cloning (TaKaRa) into pOPIN-J (20) cut with Hindlll/Kpnl. Insert sequences were verified using Sanger sequencing (Seqlab).

Recombinant expression and purification of recombinant antigens. Expression vectors encoding tandem-affinity-tagged, N-terminally truncated CoV NCPs (Fig. 1A) were transformed into *E. coli* BL21 RIG cells. Upon induction with 1 mM isopropyl- β -D-thiogalactopyranoside (IPTG), recombinant fusion proteins were expressed for 3 h at 30°C or 37°C. Harvested bacteria were lysed, and fusion proteins were purified from the soluble fraction of the bacterial lysate (Fig. 1B) by Ni-nitrilotriacetic acid (NTA) chromatography under native conditions followed by on-column cleavage of the tandem His₆–glutathione *S*-transferase (GST) tag (Fig. 1C) as described previously (16).

Biotinylation of recombinant antigens. Purified, recombinant antigens were covalently coupled to biotin using EZ-Link sulfo-NHS (*N*-hydroxysuccinimide)-biotin reagent (Thermo Fisher) at a 30-fold molar excess according to the manufacturer's instructions. Biotinylated proteins were stored in $0.5 \times$ phosphate-buffered saline (PBS)–50% glycerol at -20° C.



D	CoV-2 NCP∆246	CoV-1 NCP∆247	MERS NCP∆238	HKU1 NCP∆256	OC43 NCP∆258	NL63 NCP∆222	229E NCP∆235
SARS-CoV-2 NCP∆246	100.0	88.6	40.8	26.4	25.3	20.5	19.9
SARS-CoV-1 NCP∆247	94.3	100.0	39.8	26.4	25.3	19.1	19.7
MERS NCPA238	53.6	51.9	100.0	27.8	29.9	19.4	20.0
HKU1 NCP∆256	43.5	43.0	44.8	100.0	59.6	18.6	18.6
OC43 NCP∆258	43.3	42.3	46.4	71.5	100.0	20.2	17.6
NL63 NCP∆222	37.5	35.4	35.0	34.6	37.3	100.0	38.2
229E NCP∆235	34.7	33.7	37.2	31.4	30.6	62.4	100.0

FIG 1 Prokaryotic expression and purification of N-terminally truncated SARS-CoV-2 NCP. (A) Schematic representation of the fusion protein His_6 -GST-3C-SARS-CoV-2 NCP Δ 246 (calculated molecular weight, 47.3 kDa). (B) Total lysates preinduction (pre) and postinduction (post) and soluble (supernatant [SN]) and insoluble (pellet [P]) lysate fractions. (C) Eluate (E) and matrix (M) after on-column cleavage. (D) Amino acid sequence comparison of truncated CoV NCPs. Above the diagonal/light gray shading are identity scores (percent), and below the diagonal/dark gray shading are similarity scores (percent). MERS, Middle East respiratory syndrome.

Human sera. This study complies with the Declaration of Helsinki. Written informed consent was obtained from all individuals or, in the case of minors, from parents or legal guardians before enrollment. Data privacy protection was guaranteed by the anonymization of samples.

(i) COVID-19 patient sera. Longitudinal serum samples (range of days after the onset of symptoms [days postonset {dpo}]], 10 to 446; number of serum samples per patient, 1 to 13) from 35 patients (median age, 45 years; age range, 19 to 64 years; 23 females and 12 males) with PCR-confirmed SARS-CoV-2 infection collected in Germany between March 2020 and May 2021 were analyzed. All patients displayed ambulatory mild disease (21). Fifteen patients received a COVID-19 vaccination after more than 6 months of convales-cence (10 BioNTech/Pfizer, 2 Moderna, and 3 AstraZeneca). The collection of samples was approved by the Medical Association, Hamburg, Germany (no. 2020-10162-BO-ff). World Health Organization (WHO) international standard plasma 20/136 (1,000 binding antibody units [BAU]/mL) was obtained from the National Institute for Biological Standards and Control (NIBSC).

(ii) *A priori* SARS-CoV-2 IgG-negative control sera. The study was performed using stored human serum/plasma samples from symptom-free donors collected before the COVID-19 pandemic in Germany (2004 to 2015), Ghana (1999 and 2014 to 2015), Madagascar (2010), Nigeria (2018), Colombia (2014), and Lao People's Democratic Republic (PDR) (2014) (Table S2) (3). The collection of samples was approved by the Ethics Committees of the Kwame Nkrumah University of Science and Technology (Kumasi, Ghana) (CHRPE/AP/427/ 13, 2013), the Comité d'Ethique de la Vice Primature Chargée de la Santé Publique (Antananarivo, Madagascar) (no. 051-CE/MINSAN, 2009), the Irrua Specialist Teaching Hospital (Irrua, Nigeria) (ISTH/HREC/20171019/28, 2017), the Hospital Rosario Pumarejo de Lopez (Valledupar, Colombia) (2013), Lao People's Democratic Republic (no. 030/NECHR), and the Medical Association, Hamburg, Germany (no. PV4608, 2013).

SARS-CoV-2 IgG indirect immunofluorescence testing (IIFT). Immunofluorescence analysis detecting IgG antibodies targeting SARS-CoV-2 full-virus antigens was performed as described previously (22). Briefly, SARS-CoV-2-infected Vero-E6 cells (ATCC CRL-1586) were fixed with acetone-methanol and incubated with serial dilutions of patient sera. After washing, the detection of bound IgG was performed using a fluorescein isothiocyanate (FITC)-labeled anti-IgG antibody (Sifin).

SARS-CoV-2 IgG ELISAs 1 and 2. For the SARS-CoV-2 IgG Fc_yR ELISA, Nunc MaxiSorp ELISA plates were coated with 5 μ g/mL HsFc_yR-IgI (CD32 [Fc_yRIIA_{H131}]) (14) in PBS (pH 7.4). After blocking for 2 h at room

temperature with 1 × PBS (pH 7.4)–0.25% bovine serum albumin (BSA)–0.05% Tween 20, plates were stabilized with a liquid plate sealer (Candor) and stored at 4°C until use. Human sera (ELISA 1, 1:50 in 1 × PBS [pH 7.4]–0.05% ProClin 300–0.01% phenol red; ELISA 2, pure sera) and biotinylated recombinant antigen (ELISA 1, 1:10,000; ELISA 2, 1:25,000) (in conjugate dilution buffer [CDB] [1 × PBS {pH 7.4}–1% BSA–0.5% fetal bovine serum–1% Nonidet P-40–0.1% ProClin 300]) were mixed 1:1 on the *Hs*FcyR-IgI-coated plates and coincubated overnight at 4°C. After washing with 100 mmol/L Tris-HCI (pH 7.4)–150 mmol/L NaCI–0.05% Tween 20–0.005% ProClin 300, horseradish peroxidase (HRP)-labeled streptavidin (0.1 μ g/mL in CDB) was applied to the wells for 1 h at 4°C. After washing again, tetramethylbenzidine (TMB; KPL) was added for 20 min at room temperature, and the reaction was stopped by the addition of 1 N sulfuric acid (Merck). The TMB reaction product was quantified by measuring the absorbances at 450 nm (A450) and 620 nm (A620) on a SPECTROStar Nano ELISA (BMG Labtech) and calculating the difference A450-A620. The Euroimmun anti-SARS-CoV-2 NCP ELISA (IgG) was performed and evaluated according to the manufacturer's instructions.

Common cold CoV IgG ELISAs. IgG antibodies interacting with the N-terminally truncated NCPs of OC43, HKU1, NL63, and 229E were detected in human sera as described previously (3). Briefly, IgG Fc₇R ELISA 1 described above was performed using biotinylated OC43 NCPΔ258, HKU NCPΔ256, NL63 NCPΔ222, and 229E NCPΔ258 (Table S1), respectively, as the antigens.

Interference testing. To evaluate interference by hemolytic, icteric, and lipemic serum conditions as well as high biotin and rheumatoid factor (RF) concentrations, assay mixtures were spiked with hemoglobin (Sigma), bilirubin (Sigma), triglycerides (Sigma), biotin (Sigma), and RF (Lee Biosolutions), respectively. Simulated serum concentrations were 5 mg/mL and 10 mg/mL for hemoglobin, 0.5 mg/mL for bilirubin, and 5 mg/mL for triglycerides. Biotin interference was tested at simulated serum concentrations of between 0.1 ng/mL and 100,000 ng/mL; simulated RF serum concentrations were 50 IU/mL, 100 IU/mL, and 1,000 IU/mL. Assays were performed by employing three COVID-19 patient serum samples covering a range of A_{450} - A_{620} readings ("low," "medium," and "high") and at least three prepandemic negative-control serum samples.

Sequence alignments and protein structure visualization. Multiple-amino-acid sequence alignments and identity/similarity scores were generated using the MUSCLE algorithm in MacVector (version 12.7.5). The three-dimensional (3D) structure of truncated SARS-CoV-2 NCP (comprising the dimerization domain and the C-terminal tail) was predicted using AlphaFold2 (23). Predicted structures were visualized with UCSF Chimera (version 1.13.1) (24) and aligned to the experimentally determined structure of the dimerization domain (25) using the MatchMaker utility.

Data analysis. Receiver operating characteristic (ROC) analyses were performed with MedCalc (version 19.2.1); statistical analyses (calculation of 95% confidence intervals [CIs] according to the modified Wald method and Fisher's exact test) were done using GraphPad Prism QuickCalcs. To avoid statistical bias, a maximum of 1 sample per patient per category was included for calculations of diagnostic and analytical sensitivities. If several samples from an individual patient belonged to the same category, the sample taken on the latest day after the onset of symptoms was included in the analysis.

RESULTS

Production of N-terminally truncated SARS-CoV-2 NCP for use as an ELISA antigen. SARS-CoV-2 NCP Δ 246 was recombinantly expressed in *E. coli* and purified from the soluble fraction of the bacterial lysate (Fig. 1A to C). The amino acid sequence identities of SARS-CoV-2 NCP Δ 246 with the respective NCP fragments of other human-pathogenic betacoronaviruses ranged between 25.3% for OC43 and 88.6% for SARS-CoV-1 (Fig. 1D; see also Fig. S1 in the supplemental material).

Development of Fc₇**R-based SARS-CoV-2 IgG ELISA protocols.** Assay protocols were developed based on the Fc₇**R-based** platform technology patented by the Bernhard Nocht Institute for Tropical Medicine (BNITM) (12). Thereby, human sera are coincubated with a biotinylated antigen in a 96-well plate coated with the recombinantly produced immunoglobulin-like extracellular domain of human Fc₇RIIA (CD32 H131) (14). Subsequently, bound IgG-antigen complexes are detected by the application of HRP-labeled streptavidin and the colorimetric HRP substrate TMB. Two assay versions with different serum and antigen dilutions were validated. In assay version 1 ("ELISA 1"), a final in-well serum dilution of 1:100 (as utilized in the previously developed CCHFV and ZIKV IgG Fc₇R ELISAs [16, 17]) was employed, while in assay version 2 ("ELISA 2"), a high serum concentration (1:2, final in well) was applied. Biotinylated antigen was titrated at final dilutions in well of 1:20,000, 1:50,000, and 1:75,000 (data not shown); based on the obtained signal-to-noise ratios, final in-well antigen dilutions of 1:20,000 and 1:50,000 for ELISAs 1 and 2, respectively, were chosen.

Reproducibility. Intra- and inter-assay variations were assessed using four SARS-CoV-2 lgG-positive serum samples and four *a priori* SARS-CoV-2 lgG-negative serum samples (Table S3). Both ELISAs generated highly reproducible results, with mean intra-assay coefficients of variation (CVs) of <5% and mean inter-assay CVs of <10% for positive signals.

Detection range and linearity. The detection ranges of the two ELISAs were determined using serial dilutions of WHO international standard plasma 20/136 simulating



FIG 2 Detection range. WHO international standard plasma 20/136 (1,000 BAU/mL) was serially diluted with serum dilution buffer (SDB) (1 × PBS [pH 7.4], 0.05% ProClin 300, 0.01% phenol red) to simulate samples with antibody concentrations ranging between 0.1 BAU/mL and 1,000 BAU/mL. Simulated samples were tested with ELISA 1 (open circles) (final in-well dilutions of 1:100 for samples and 1:20,000 for the conjugate) and ELISA 2 (filled circles) (final in-well dilutions of 1:2 for samples and 1:50,000 for the conjugate). Dotted lines indicate A_{450} - A_{620} values of 0.2 and 0.4, respectively. Solid lines indicate the linear regression fit.

samples with antibody concentrations of between 0.1 BAU/mL and 1,000 BAU/mL (Fig. 2). Assuming an assay cutoff of an A_{450} - A_{620} of 0.2 (see below), ELISA 2 showed a significantly lower detection limit (0.25 U/mL) than ELISA 1 (10.4 U/mL) but exhibited a strong prozone effect at high antibody concentrations (Fig. 2). In particular, signal saturation was observed for ELISA 2 for antibody concentrations of between 7.5 BAU/mL and 75 U/mL; higher antibody concentrations induced a continuous signal drop to an A_{450} - A_{620} of 0.725 for 1,000 BAU/mL.

Diagnostic sensitivity. To determine the diagnostic sensitivities of the two SARS-CoV-2 IgG F_{C7}R ELISAs, 213 longitudinal serum samples obtained from 35 German patients with PCR-confirmed SARS-CoV-2 infection and 139 serum samples from healthy German blood donors (HDs), collected before the COVID-19 pandemic, were analyzed (Fig. 3). ROC analyses were performed separately for sample sets collected at different time points after the onset of symptoms (Fig. S2). By applying the lowest possible cutoff value generating 100% specificity in the negative HD panel (A_{450} - A_{620} cutoff = 0.2), diagnostic sensitivities were 90.5% (95% CI, 69.9 to 98.5%) and 100.0% (95% CI, 81.8 to 100.0%) in samples collected 1 to 2 months after the onset of symptoms for ELISAs 1 and 2, respectively (Table 1). While the diagnostic sensitivity of ELISA 1 (sera, 1:100; conjugate, 1:20,000) was found to drop quickly over time in convalescent patients, ELISA 2 (sera, 1:2; conjugate, 1:50,000) enabled the detection of anti-SARS-CoV-2 NCP IgG antibodies for a much longer time period after acute infection (Fig. 3 and Table 1). With the assays being based on NCP antigen, the time course of the humoral response to SARS-CoV-2 infection could still be monitored in individuals having received spike-protein-based vaccines (filled dots in Fig. 3).

Analytical sensitivity. Immunofluorescence analysis of COVID-19 patient samples using SARS-CoV-2-infected Vero cells revealed the highest IgG antibody titers between 1 and 2 months after the onset of symptoms (Fig. 4A and B). Later in the convalescent phase, the median IgG antibody titers dropped considerably but persisted at detectable levels in most patients for at least 12 months; spike-based vaccines strongly boosted SARS-CoV-2 IgG IIFT titers (filled dots in Fig. 4A and B). To determine the analytical sensitivities of the two SARS-CoV-2 IgG Fc₇R ELISAs in relation to IIFT (serving as a highly sensitive gold-standard antibody binding assay), 109 samples from 35 patients with PCR-confirmed SARS-CoV-2 infection taken prior to an eventual vaccination with a spike-based vaccine were stratified according to their respective IgG IIFT titers (\leq 1 sample/patient/category; median/range, 123/13 to 443 dpo). While samples with a high IIFT titer of SARS-CoV-2 IgG antibodies (\geq 1,280) were readily detected as positive by both ELISAs, the analytical sensitivity was significantly higher for ELISA 2 when analyzing samples with an IgG IIFT titer of between 160 and 640 (Fig. 4C and D and Table 2). In samples with a high SARS-CoV-2 IgG IIFT titer (\geq 5,120), a prozone effect was observed for SARS-CoV-2 IgG Fc₇R ELISA 2 (Fig. 4D).

Comparative testing of a subset of 82 IIFT positive samples from 26 COVID-19 patients (dpo range: 10 to 178) with a commercially available, NCP-based indirect ELISA (Euroimmun SARS-CoV-2 NCP IgG) revealed a high concordance with SARS-CoV-2 Fc₂R IgG ELISA 1 (Fig. S3).



FIG 3 Diagnostic sensitivity. Longitudinal serum samples from 35 COVID-19 patients were analyzed using ELISA 1 (A and C) and ELISA 2 (B and D). (A and B) Trajectories (individual patients; 1 to 13 samples/patient; n = 213). (C and D) Stratification according to months after the onset of symptoms (≤ 1 sample/patient/category; n = 146). Dotted lines indicate A_{450} - A_{620} values of 0.2 and 0.4. HD, healthy donors (Germany) (n = 139). Filled dots indicate postvaccination samples (spike-based vaccine).

Specificity. The specificities of the two SARS-CoV-2 IgG $F_{C\gamma}R$ ELISAs in comparison with the Euroimmun SARS-CoV-2 IgG NCP ELISA (Table 3; Fig. S4) were evaluated using *a priori* SARS-CoV-2 IgG-negative serum panels acquired from symptom-free donors before the COVID-19 pandemic in Europe (Germany), Africa (Ghana, Madagascar, and Nigeria), Asia

TABLE 1 Diagnostic sensitivity^a

mpo	dpo	No. of samples	A ₄₅₀ -A ₆₂₀ cutoff	No. of positive samples, % sensitivity (95% Cl) for ELISA 1 (sera, 1:100)	No. of positive samples, % sensitivity (95% Cl) for ELISA 2 (sera, 1:2)	Р
<1	10–28	19	0.200	14, 73.7 (50.9–88.5)	19, 100.0 (80.2–100.0)	0.0463
			0.400	11, 57.9 (36.2–76.9)	17, 89.5 (67.4–98.3)	0.0625
1–2	29–56	21	0.200	19, 90.5 (69.9–98.5)	21, 100.0 (81.8–100.0)	0.4878
			0.400	19, 90.5 (69.9–98.5)	20, 95.2 (75.6–100.0)	1.0000
2–5	57–140	34	0.200	28, 82.3 (66.1–92.0)	32, 94.1 (79.9–99.3)	0.2585
			0.400	26, 76.5 (59.8–87.8)	31, 91.2 (76.3–97.7)	0.1863
5–8	141–224	27	0.200	17, 63.0 (44.2–78.5)	24, 88.9 (71.1–96.8)	0.0537
			0.400	13, 48.1 (30.7–66.0)	23, 85.2 (66.9–94.7)	0.0084
8–12	225–336	24	0.200	8, 33,3 (17,8–53,4)	17, 70.8 (50.6–85.3)	0.0199
			0.400	3, 12.5 (3.5–31.8)	16, 66.7 (46.6–82.2)	0.0003
12–15	337-396	17	0.200	5, 29,4 (13,0-53,4)	14, 82, 3 (58, 2–94, 6)	0.0049
			0.400	2, 11.8 (2.0–35.6)	14, 82.3 (58.2–94.6)	0.0001

^aLongitudinal serum samples obtained from 35 COVID-19 patients were stratified according to months postonset (mpo) (\leq 1 sample/patient/category). Sensitivities as well as 95% confidence intervals (Cls) were calculated for two alternative cutoff values (A_{450} - A_{620} = 0.2 and 0.4). dpo, days postonset. *P* values were determined using Fisher's exact test.



IgG IIFT titer IgG IIFT titer

FIG 4 Analytical sensitivity. (A and B) Longitudinal serum samples from 35 COVID-19 patients were analyzed by IgG IIFT. Filled dots indicate postvaccination samples (spike-based vaccine). (A) Trajectories (individual patients; 1 to 13 samples/patient; n = 213; (B) stratification according to months after the onset of symptoms (≤ 1 sample/ patient/category; n = 146). (C and D) Results of ELISA 1 (C) and ELISA 2 (D) for prevaccination samples stratified according to IIFT titers (≤ 1 sample/patient/category; n = 109). Dotted lines indicate A_{450} - A_{620} values of 0.2 and 0.4.

(Lao PDR), and South America (Colombia) (Table S2) (3). While false-positive rates were low for both the commercially available ELISA and the newly developed tests with the European serum panel, both SARS-CoV-2 IgG FcyR ELISAs showed superior specificity in application to African serum panels (Table 3; Fig. S4).

Although IgG antibodies interacting with the corresponding NCP fragments of the betacoronaviruses OC43 and/or HKU1 (Table S1) were detected in 244 (31%) of 790 tested serum samples (Table 3; Fig. S5) (3), no obvious cross-reactivity of these antibodies with the SARS-CoV-2 antigen has been observed. Indeed, false-positive signals obtained for a small number of Nigerian serum samples with SARS-CoV-2 IgG Fc₂R ELISA 1 could not be suppressed with an excess of unlabeled OC43 NCP, although high levels of antibodies interacting with this protein were present in these samples (data not shown).

Interference. For SARS-CoV-2 IgG FcyR ELISA 1 (sera, 1:100, final in well; antigen, 1:20,000, final in well), no significant influence on the obtained A_{450} - A_{620} values was observed for final serum concentrations of 10 mg/mL hemoglobin, 0.5 mg/mL bilirubin, and 5 mg/mL triglycerides and biotin concentrations of up to 100,000 ng/mL. While a rheumatoid factor serum concentration of 50 IU/mL did not influence the signal height, concentrations of 100 IU/mL and 1,000 IU/mL suppressed positive signals by ca. 30% and 50%, respectively, without increasing the assay background.

For SARS-CoV-2 IgG FcyR ELISA 2 (sera, 1:2, final in well; antigen, 1:50,000, final in well), although a final biotin serum concentration of 100,000 ng/mL induced a significant signal loss for positive sera, a concentration of 3,500 ng/mL did not significantly influence A_{450} - A_{620} values. Hemoglobin serum concentrations of up to 5 mg/mL did not influence assay performance, but signals were impacted at 10 mg/mL. Rheumatoid factor serum concentrations of 50 IU/mL and 1,000 IU/mL reduced the intensities of positive signals by ca. 40%, and 50%, respectively, without increasing the assay background. The influence of bilirubin and

lgG IIFT titer	No. of samples	A ₄₅₀ -A ₆₂₀ cutoff	No. of positive samples, % sensitivity (95% CI), for ELISA 1 (sera, 1:100)	No. of positive samples, % sensitivity (95% CI), for ELISA 2 (sera, 1:2)	Р
≤80	7	0.200	0, 0.0 (0.0–40.4)	4, 57.1 (25.0–84.2)	0.0699
		0.400	0, 0.0 (0.0–40.4)	1, 14.3 (0.5–53.3)	1.0000
160	17	0.200	6, 35.3 (17.2–58.8)	13, 76.5 (52.2–90.9)	0.0366
		0.400	3, 17.6 (5.4–41.8)	13, 76.5 (52.2–90.9)	0.0016
320	22	0.200	14, 63.6 (42.9–80.4)	21, 95.4 (76.5–100.0)	0.0212
		0.400	9, 40.9 (23.2–61.3)	21, 95.4 (76.5–100.0)	0.0002
640	25	0.200	20, 80.0 (60.4–91.6)	24, 96.0 (78.9–100.0)	0.1895
		0.400	17, 68.0 (48.3–82.9)	24, 96.0 (78.9–100.0)	0.0232
1,280	19	0.200	19, 100.0 (80.2–100.0)	19, 100.0 (80.2–100.0)	1.0000
,		0.400	19, 100.0 (80.2–100.0)	19, 100.0 (80.2–100.0)	1.0000
2,560	11	0.200	10, 90.9 (60.1–100.0)	11, 100.0 (70.0–100.0)	1.0000
,		0.400	10, 90.9 (60.1–100.0)	10, 90.9 (60.1–100.0)	1.0000
≥5.120	8	0.200	8, 100,0 (62,8–100,0)	8, 100,0 (62,8–100,0)	1.0000
	-	0.400	8, 100.0 (62.8–100.0)	8, 100.0 (62.8–100.0)	1.0000

TABLE 2 Analytical sensitivity^a

^{*a*}Longitudinal serum samples obtained from 35 COVID-19 patients were stratified according to SARS-CoV-2 IgG IIFT titers (\leq 1 sample/patient/category). Sensitivities as well as 95% confidence intervals (CIs) were calculated for two alternative cutoff values (A_{450} - A_{620} = 0.2 and 0.4). *P* values were determined using Fisher's exact test.

triglycerides could not be studied for ELISA 2 due to the strong impact of the solvent used (chloroform) on antigen integrity.

DISCUSSION

Assay protocol. The SARS-CoV-2 IgG ELISAs developed and validated in this work employ a patented platform technology (12, 14-17) in which immune complexes between a labeled, recombinant antigen and human serum IgG antibodies are formed in solution and subsequently captured by an isotype-specific FcyR immunoglobulin-like domain. Thereby, the use of a soluble, natively folded antigen has a direct positive impact on sensitivity and specificity because conformational epitopes are preserved and nonspecific interactions with hydrophobic amino acid stretches that may be exposed in incorrectly folded proteins cannot occur. With the main intended use of our assays being the monitoring of seroconversion and seroprevalence in archived samples, development was focused on the optimization of sensitivity and specificity rather than a minimized time to result. Nevertheless, preliminary results indicate that upon adjustments of conjugate and serum concentrations, shortening the antigen-sample incubation step from overnight to 2 h might be possible, with only minor losses in sensitivity and no increase in the assay background (data not shown). Importantly, the assays, which are procurable via the European Virus Archive Goes Global (EVAg) webpage (26), can be performed manually, and results are generated using a standard absorbance ELISA reader, facilitating their use also in settings where no highly specialized technical equipment is available.

Choice of antigen. As an antigen, a SARS-CoV-2 NCP fragment comprising the dimerization and C-terminal tail domains of SARS-CoV-2 NCP (see Fig. S1 in the supplemental material) was used. Similar fragments of SARS-CoV-1 NCP have previously been shown to allow the sensitive detection of anti-SARS-CoV-1 IgG antibodies in patient sera (27, 28) while generating fewer false-positive responses than the full-length protein in negative-control sera (28). As for other viral nucleocapsid proteins (16, 28), the cost-efficient production of large quantities of natively folded SARS-CoV-2 full-length NCP and subdomains is possible in *E. coli* (7).

Assay sensitivity. Complete validation was performed for two SARS-CoV-2 IgG Fc γ R ELISA protocols employing different final serum and conjugate dilutions (ELISA 1, sera at 1:100 and conjugate at 1:20,000; ELISA 2, sera at 1:2 and conjugate at 1:50,000). ELISA 1 displayed diagnostic and analytical sensitivities similar to those of a commercially available, manually performed, plate-based indirect ELISA (Euroimmun SARS-CoV-2 NCP IgG ELISA) (29), thus showing a significant waning of signal intensity in COVID-19 patients at approximately 6

Panel	No. of samples	No. (%) of betacoronavirus IgG- positive samples	A ₄₅₀ -A ₆₂₀ cutoff	No. of negative samples, % specificity (95% Cl), for ELISA 1 (sera, 1:100)	No. of negative samples, % specificity (95% CI), for ELISA 2 (sera, 1:2)	No. of negative/bl samples, % specificity (95% Cl), for Euroimmun NCP IgG
Germany	139	51 (36.7)	0.200	139, 100.0 (96.8–100.0)	139, 100.0 (96.8–100.0)	139, 100.0 (96.8–100.0)
			0.400	139, 100.0 (90.8-100.0)	139, 100.0 (90.8-100.0)	
Ghana A	131	33 (25.2)	0.200	128, 97.7 (93.2–99.5)	129, 98.5 (94.3–99.9)	97, 74.0 (65.9–80.8)
			0.400	129, 98.5 (94.3–99.9)	131, 100.0 (96.6–100.0)	
Ghana B	145	34 (23.4)	0.200	142, 97.9 (93.8–99.6)	145, 100.0 (96.9–100.0)	132, 91.0 (85.1–94.8)
			0.400	145, 100.0 (96.9–100.0)	145, 100.0 (96.9–100.0)	
Madagascar	166	30 (18.1)	0.200	165, 99.4 (96.3–100.0)	166, 100.0 (97.3–100.0)	164, 98.8 (95.4–100.0)
5			0.400	166, 100.0 (97.3–100.0)	166, 100.0 (97.3–100.0)	
Nigeria	149	79 (53.0)	0.200	143, 96.0 (91.3–98.3)	149, 100.0 (97.0–100.0)	107, 71.8 (64.1–78.4)
5			0.400	146, 98.0 (94.0–99.6)	149, 100.0 (97.0–100.0)	
Colombia	40	12 (30.0)	0.200	40, 100.0 (89.6–100.0)	40, 100.0 (89.6–100.0)	40, 100.0 (89.6–100.0)
			0.400	40, 100.0 (89.6–100.0)	40, 100.0 (89.6–100.0)	·, · · · · · · · · · · · · · · · ,
Lao PDR	20	5 (25.0)	0.200	20, 100.0 (81.0–100.0)	20, 100.0 (81.0–100.0)	20, 100,0 (81,0–100,0)
200.01		- (2010)	0.400	20, 100.0 (81.0–100.0)	20, 100.0 (81.0–100.0)	20, 10010 (01.0 10010)

TABLE 3 Specificity of SARS-CoV-2 IgG ELISAs^a

^aSerum/plasma samples collected from symptom-free donors before 2019 in Europe, Africa, South America, and Asia were analyzed with SARS-CoV-2 IgG Fc₃R ELISAs 1 and 2 and the Euroimmun anti-SARS-CoV-2 NCP ELISA (IgG). Specificities and 95% confidence intervals (CIs) were calculated for two alternative cutoff values (A_{450} - A_{620} = 0.2 and 0.4). bl, borderline.

months postinfection. In contrast, ELISA 2 allowed the persistent detection of anti-NCP IgG antibodies for more than 12 months postinfection for most patients. Thus, SARS-CoV-2 IgG Fc₂R ELISA 2 displays a sensitivity comparable to that of Elecsys anti-SARS-CoV-2 (Roche), a commercially available NCP-based automated electrochemiluminescence immunoassay system detecting total Ig through double-antigen binding (30, 31).

Assay specificity. As we have shown recently, unacceptably high false-positive rates of up to 61% are obtained with several commercially available indirect SARS-CoV-2 NCP IgG ELISAs in African sample panels collected before the COVID-19 pandemic, although the very same assays displayed excellent specificity in European samples (3). Therefore, the specificity of the newly developed SARS-CoV-2 IgG FcyR ELISAs was challenged with an extensive panel of 790 pre-COVID-19 negative-control samples originating from Europe (n = 139), South America (n = 40), Asia (n = 20), and Africa (n = 591). Both ELISAs were found to be highly specific (>96%), even in the most challenging sample panel, originating from Nigeria. For applications requiring maximum specificity in African samples, we recommend using a slightly elevated cutoff value of an A_{450} - A_{620} of 0.4 (instead of 0.2 for samples from European residents). As for the commercially available assays evaluated in our previous study (3), the presence of IgG antibodies elicited by previous infections with the common cold CoVs OC43, HKU1, NL63, and 229E did not per se compromise SARS-CoV-2 IgG FcyR ELISA specificity. Indeed, in concordance with data from previous studies (1, 32), those antibodies were found in a significant proportion of donor sera of each origin (Table 3; Fig. S5). Three Nigerian samples generated false-positive signals in ELISA 1 but not ELISA 2. Although high levels of anti-OC43 IgG antibodies were detected in all three serum samples, signals in the SARS-CoV-2 IgG FcyR ELISA could not be suppressed by the addition of excess OC43 NCP, ruling out OC43 cross-reactivity as a cause. A possible explanation for these false-positive signals occurring in ELISA 1 but not ELISA 2 could be the presence of high levels of IgG antibodies elicited by another, as-yet-unidentified member of the Coronaviridae family binding with low affinity to SARS-CoV-2 NCP.

Caveats and limitations. When applying the SARS-CoV-2 IgG Fc₇R ELISAs presented in this study in seroepidemiological research projects, several caveats and limitations have to be considered. First of all, due to the use of SARS-CoV-2 NCP as the antigen, humoral immune responses elicited by spike-based vaccines (e.g., BioNTech/Pfizer and AstraZeneca) are not detected by these assays. Correspondingly, detection of natural

SARS-CoV-2 infection is possible in naive individuals and probands vaccinated with spikebased vaccines but not with vaccines based on inactivated full virus (e.g., Sinopharm and Sinovac). The assay version should be chosen according to the respective research questions: while ELISA 1 is most suitable to monitor the development of the humoral immune response during the acute and early convalescent phases of the disease, ELISA 2 allows the detection of SARS-CoV-2-specific antibodies for a much longer time period postinfection and therefore is the assay version of choice for large-scale seroprevalence studies (30). Nevertheless, this assay shows a strong prozone effect in serum samples with very high anti-NCP IgG titers. Therefore, misleadingly low readings could occur in certain situations, such as during early convalescent stages of severe infections (1) or upon reinfections/vaccinations strongly boosting the anti-NCP response. If these scenarios are likely, the parallel performance of ELISAs 1 and 2 is strongly recommended. Furthermore, ELISA 2 is (due to the direct use of pure, undiluted serum samples) more susceptible to interfering substances than ELISA 1. Therefore, serum quality should be monitored and critically assessed, particularly when using this assay version. Further potential limitations, relevant for both assay versions, are the differential affinities of the recombinant capture molecule CD32 H131 for the various lgG subclasses (33) and the potential masking of relevant epitopes by the biotin label. Nevertheless, the dominant IgG subclasses induced by SARS-CoV-2 infection, IgG1 and IgG3 (34, 35), have been shown to strongly interact with CD32 H131 in a previous study (33).

SUPPLEMENTAL MATERIAL

Supplemental material is available online only. SUPPLEMENTAL FILE 1, PDF file, 7.4 MB. SUPPLEMENTAL FILE 2, PDF file, 0.1 MB. SUPPLEMENTAL FILE 3, PDF file, 0.1 MB.

ACKNOWLEDGMENTS

We thank M. Panning for providing CoV RNAs, J. Blessmann for providing serum samples from Lao PDR, and S. Kann for providing serum samples from Colombia. Furthermore, we thank A. Karstedt, R. Ferraz-Jansen, T. Kopp, F. Geisinger, U. Fröhlke, and N. Pekarek for expert technical assistance.

C.D. received funding from the German Federal Ministry of Education and Research (grant no. 01Kl20210) and the European Regional Development Fund (ERDF) (grant no. BWF/H/ 52228/2012/13.10.10-1/3.4,6) to the institution, P.E. received funding from the German Federal Ministry of Education and Research (grant no. 01Kl20210) to the institution, and L.O. received funding from the Leibniz Association (grant no. J59/20218) to the institution. The funding organizations played no role in the design of the study, choice of enrolled patients, review and interpretation of data, preparation of the manuscript, or final approval of the manuscript.

C.D. and P.E. are inventors on patent EP3207375, and P.E. is an inventor on patent EP2492689. C.D. and B.R. are cofounders and shareholders of Panadea Diagnostics GmbH. We declare no other potential conflicts of interest.

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