

Extra INSTAND EQA Scheme (340) - April 2020

Virus Genome Detection SARS-CoV-2

Final Evaluation of Results

Submitted by 463 out of 487 Laboratories
from 36 Countries

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³IQVD GmbH - Institut für Qualitätssicherung in der Virusdiagnostik

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Disclosure

Heinz Zeichhardt is share-holder
of

GBD Gesellschaft für Biotechnologische Diagnostik mbH, Berlin
and

IQVD GmbH - Institut für Qualitätssicherung in der Virusdiagnostik, Berlin

EQA Network

Scientific umbrella

German Association for Prevention of Virus Diseases (DVV e.V.)

Society of Virology (GfV e.V.)

German Society for Hygiene and Microbiology (DGHM e.V.)

Partners

Berlin

Teams of

GBD Gesellschaft für Biotechnologische Diagnostik mbH, Berlin

and

IQVD GmbH - Institut für Qualitätssicherung in der Virusdiagnostik, Berlin

and

38 INSTAND Expert Laboratories

incl.

Robert Koch-Institut

Paul-Ehrlich-Institut

National Reference and Consiliary Labs

and

and

Düsseldorf

INSTAND-Team

EMPIR
AntiMicroResist

EMRP
European Metrology Research Programme
Programme of EURAMET
The EMRP is jointly funded by the EMRP participating countries within EURAMET and the European Union



EURAMET
European Association of National Metrology Institutes



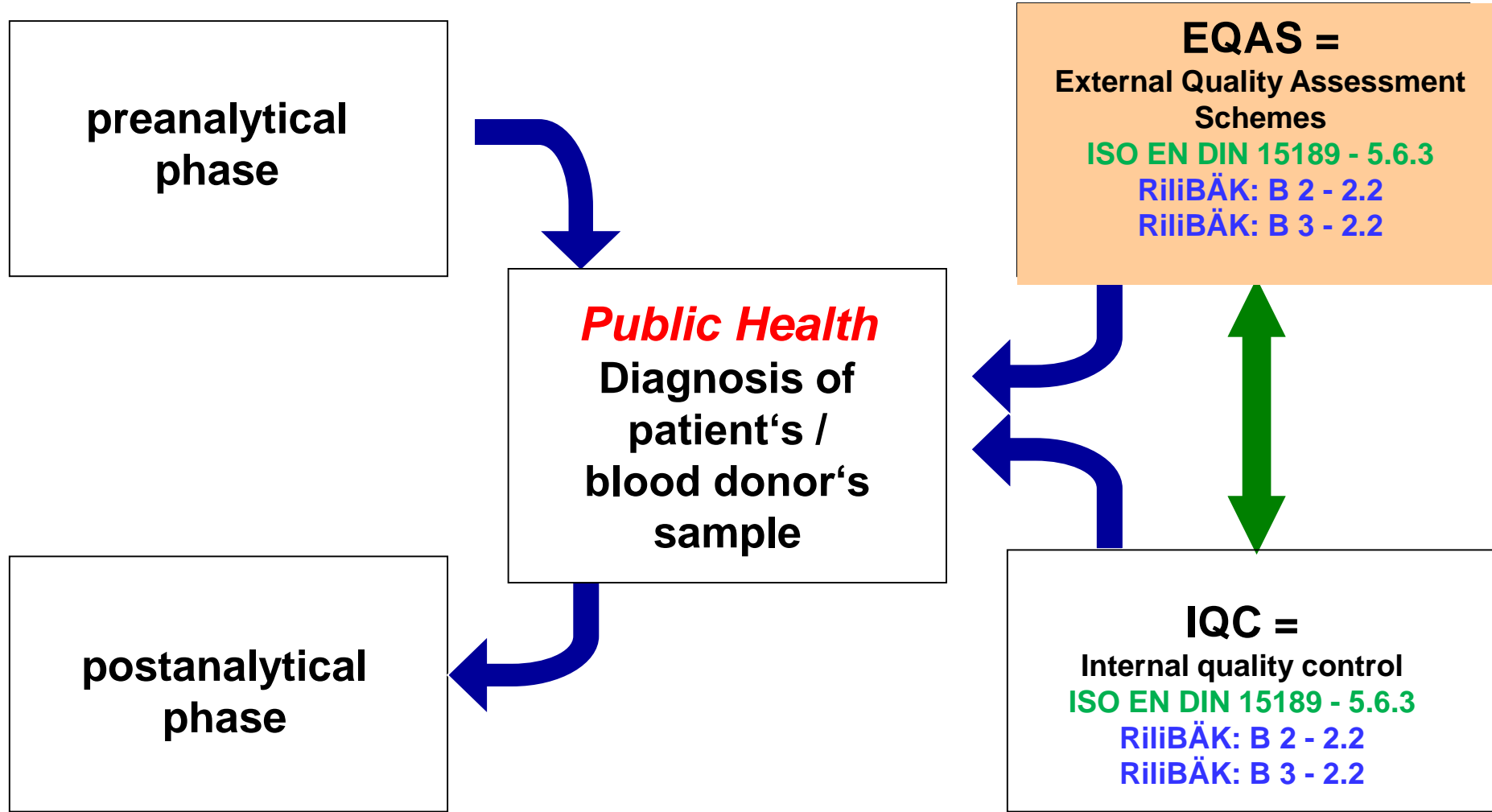
JRC – Joint Research Centre (BE)

➔ LGC – National Measurement Laboratory (UK)

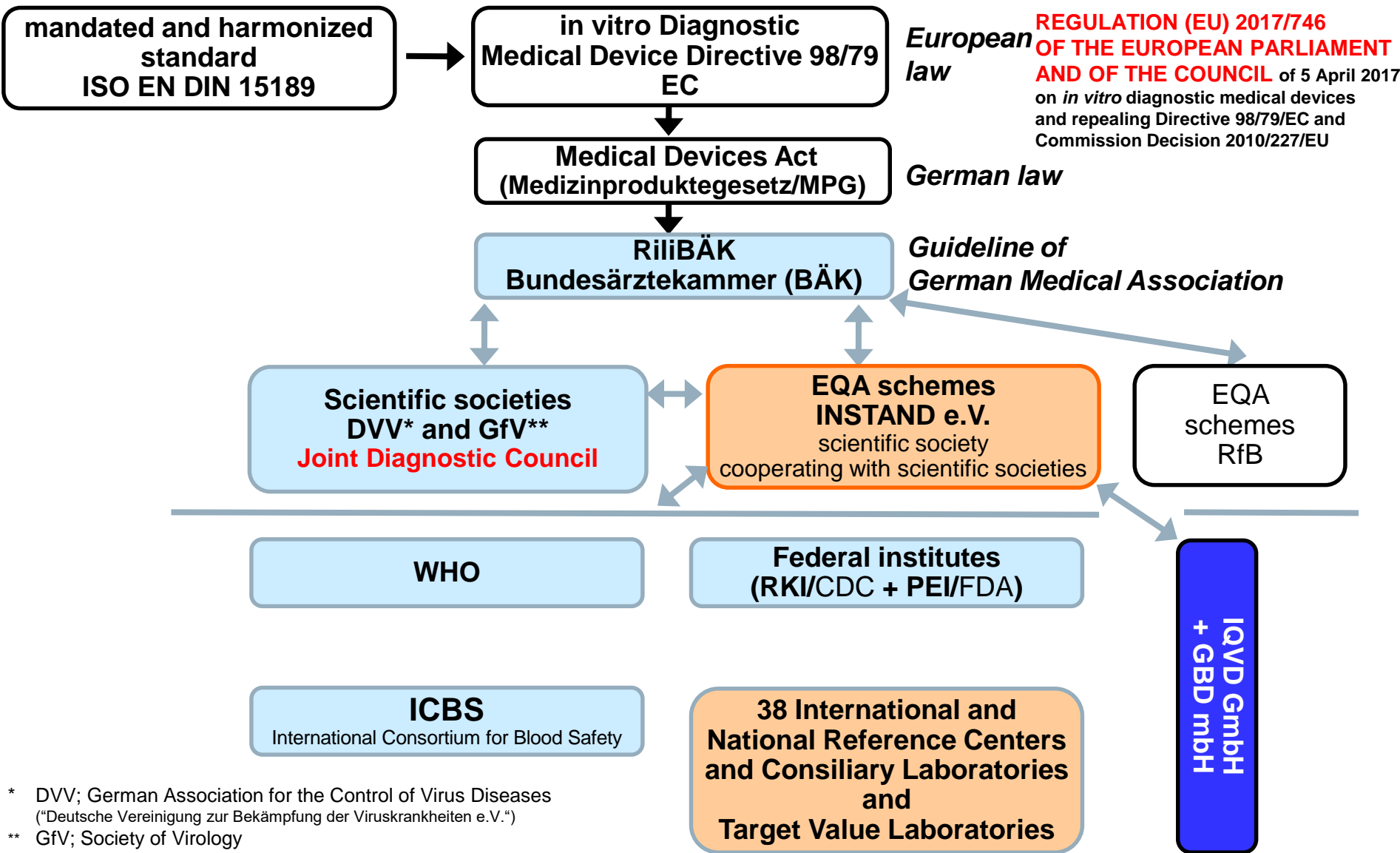
NIB – National Institute of Biology (SI)

PTB - Physikalisch-Technische Bundesanstalt (DE)

The Role of Quality Control for Public Health and Quality Improvement of Diagnostics and Blood Safety



INSTAND e.V. - Quality Assurance in (Virus) Diagnostics



* DVV; German Association for the Control of Virus Diseases ("Deutsche Vereinigung zur Bekämpfung der Viruskrankheiten e.V.")
 ** GfV; Society of Virology ("Gesellschaft für Virologie e.V.")

INSTAND EQA Schemes (78 Schemes) in Virus Immunology and Genome Detection 2020

Serology and Antigen Detection		Virus Genome Detection and Typing	
HIV-1/2	Herpes simplex Viruses	HIV-1 (RNA) HIV-2 (RNA)	Measles Rubella Mumps Viruses
HIV-1 p24 Ag	Varicella Zoster Virus	HIV-1 Resistance + Tropism	Adenoviruses
HTLV-1/2	Epstein Barr Virus	Hepatitis A Virus	Norovirus Rotavirus
Hepatitis A Virus	Resp. Sync. Virus Ag	Hepatitis B Virus + Genotyping + Resistance	CoV incl. MERS CoV SARS-CoV-2 NEW
Hepatitis B Virus, Prg. I	Influenza A and B Ag + A/H1N1 pdm 2009 + A/H5N1+A/H7N9	Hepatitis C Virus + Genotyping + Resistance	Enteroviruses + Enterovirus (WHO/RKI) Parechovirus NEW
Hepatitis B Virus, Prg. II	Rubella Virus Measles Virus Mumps Virus	Hepatitis D Virus Hepatitis E Virus	Human Rhinoviruses Resp. Syncytial Virus Hum. Metapneumovirus Parainfluenza viruses
Hepatitis C Virus	TBE Virus	Torque Teno Virus NEW	Influenza A and B incl avian
Hepatitis D Virus	Hantavirus	Cytomegalovirus + Resistance	BK Virus JC Virus
Hepatitis E Virus	Dengue Virus Chikungunya Virus Zika Virus	Epstein-Barr Virus Herpes simplex Viruses HHV-6 NEW HHV-8 NEW	Dengue Virus West Nile Virus Chikungunya Virus Zika Virus
Parvovirus B19	Rabies Virus	Varicella Zoster Virus	Hum. Papilloma Viruses
Cytomegalovirus	Borna Virus NEW	Parvovirus B19	Rabies Virus
SARS-CoV-2 NEW	BSE (PrPsc) (2002-2007)	Multiplex: Respiratory Viruses (2 programs)	Multiplex: NEW Viral Meningitis/Encephalitis
		Multiplex: Gastrointestinal Viruses	Borna Viruses NEW

Extra INSTAND EQA Scheme (340) - April 2020

Virus Genome Detection SARS-CoV-2

Cooperation partner

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Dr. Andreas Kummrow

Dr. Anabell Plauth

Dr. Samreen Falak

Final Evaluation

Based on the Results of 463 out of 487 Laboratories

Sample no.	Sample properties	Sample evaluated
340059	SARS-CoV-2 / 1 : 1 000 diluted ^{\$}	no
340060	HCoV OC43 / 1 : 2 500 diluted / specificity control	no
340061	SARS-CoV-2 / 1 : 1 000 000 diluted ^{\$}	yes
340062	CoV negative / specificity control	yes
340063	SARS-CoV-2 / 1 : 10 000 diluted ^{\$}	yes
340064	SARS-CoV-2 / 1 : 100 000 diluted ^{\$}	no
340065	HCoV 229E / 1 : 2 500 diluted specificity control	yes

^{\$} The SARS-CoV-2 positive samples 340059, 340061, 340063 and 340064 represent different dilution steps from a dilution series of a lysate from cells, infected with SARS-CoV-2 (inactivated).

Final Evaluation

Based on the Results of 463 out of 487 Laboratories

Results of sensitivity panel

- qualitative results
ct/cp/cq/CN values
- quantitative results in copies/ml
qPCR and dPCR
- what can we learn from quantification of CMV and HIV-1?

Results of specificity panel

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340064	SARS-CoV-2 / 1 : 100 000 diluted ^{\$}	no
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Final Evaluation

Based on the Results of 463 out of 487 Laboratories

Sample no.	Sample properties	Expected qualitative result for SARS-CoV-2	Gene region	Correct results per reported results differentiated by gene region	<i>Correct results per reported results differentiated by gene region (reduced by the no. of incorrect result assignments (mix-ups) for samples 340064 and 340065)</i>	Reported Ct/Cp/Cq/CN- results differentiated by gene region median (min – max)
Sample 340059 ^{s,*}	SARS-CoV-2 1 : 1 000 diluted sample not evaluated ^s	positive	E	373/373 (100%)	n.a.	22.6 (16.8-34.0)
			N	165/167 (98.8%)		23.6 (17.9-34.9)
			ORF1a	45/46 (97.8%)		22.2 (20.8-28.7)
			ORF1ab	48/48 (100%)		21.8 (10.9-29.1)
			RdRP	185/185 (100%)		23.8 (10.0-34.5)
			S	100/100 (100%)		21.8 (17.5-27.8)
			n.s. ^s	64/64 (100%)		22.6 (9.4-33.0)
total			980/983 (99.7%) ^s		22.8	
Sample 340063 [*]	SARS-CoV-2 1 : 10 000 diluted	positive	E	368/371 (99.2%)	n.a.	25.7 (19.8-35.3)
			N	165/167 (98.8%)		26.9 (20.0-39.6)
			ORF1a	45/46 (97.8%)		25.5 (24.0-31.5)
			ORF1ab	45/47 (95.7%)		24.7 (14.5-33.4)
			RdRP	184/184 (100%)		27.2 (13.4-38.3)
			S	100/100 (100%)		25.0 (18.0-30.0)
			n.s. ^s	64/64 (100%)		25.7 (12.8-35.9)
total			971/983 (98.8%)		26.0	
Sample 340064 ^{s,*}	SARS-CoV-2 1 : 100 000 diluted sample not evaluated ^s	positive	E	356/373 (95.4%)	356/360 (98.9%)	29.2 (17.7-36.0)
			N	146/166 (88.0%)	145/147 (98.6%)	30.2 (20.0-41.5)
			ORF1a	44/46 (95.7%)	44/46 (95.7%)	28.8 (27.3-34.0)
			ORF1ab	41/48 (85.4%)	41/41 (100%)	28.8 (18.4-37.0)
			RdRP	168/185 (90,8%)	168/173 (97.1%)	30.5 (16.0-41.4)
			S	97/100 (97.0%)	97/97 (100%)	28.6 (20.0-34.2)
			n.s. ^s	64/64 (100%)	64/64 (100%)	29.0 (16.5-40.0)
total			916/983 (93.2%) ^s		29.5	
Sample 340061 [*]	SARS-CoV-2 1 : 1 000 000 diluted	positive	E	364/373 (97.6%)	n.a.	32.1 (15.0-40.0)
			N	153/167 (91.6%)		33.3 (20.0-40.7)
			ORF1a	44/46 (95.7%)		31.7 (30.6-36.0)
			ORF1ab	42/48 (87.5%)		31.5 (20.0-39.3)
			RdRP	158/185 (85.4%)		33.5 (19.5-42.8)
			S	97/100 (97.0%)		31.5 (20.0-39.0)
			n.s. ^s	56/64 (87.5%)		31.9 (19.3-37.5)
total			914/983 (93.0%)		32.4	

Final Evaluation

Based on the Results of 463 out of 487 Laboratories

Sample no.	Sample properties	Expected qualitative result for SARS-CoV-2	Gene region	Correct results per reported results differentiated by gene region	Correct results per reported results differentiated by gene region <i>(reduced by the no. of incorrect result assignments (mix-ups) for samples 340064 and 340065)</i>	Reported Ct/Cp/Cq/CN- results differentiated by gene region median (min – max)
Sample 340059 ^{s,*}	SARS-CoV-2 1 : 1 000 diluted sample not evaluated ^s	positive	E	373/373 (100%)	n.a.	22.6 (16.8-34.0)
			N	165/167 (98.8%)		23.6 (17.9-34.9)
			ORF1a	45/46 (97.8%)		22.2 (20.8-28.7)
			ORF1ab	48/48 (100%)		21.8 (10.9-29.1)
			RdRP	185/185 (100%)		23.8 (10.0-34.5)
			S	100/100 (100%)		21.8 (17.5-27.8)
		n.s. ^s	64/64 (100%)	22.6 (9.4-33.0)		
total	980/983 (99.7%) ^s	22.8				
Sample 340063 [*]	SARS-CoV-2 1 : 10 000 diluted	positive	E	368/371 (99.2%)	n.a.	25.7 (19.8-35.3)
			N	165/167 (98.8%)		26.9 (20.0-39.6)
			ORF1a	45/46 (97.8%)		25.5 (24.0-31.5)
			ORF1ab	45/47 (95.7%)		24.7 (14.5-33.4)
			RdRP	184/184 (100%)		27.2 (13.4-38.3)
			S	100/100 (100%)		25.0 (18.0-30.0)
		n.s. ^s	64/64 (100%)	25.7 (12.8-35.9)		
total	971/983 (98.8%)	26.0				
Sample 340064 ^{s,*}	SARS-CoV-2 1 : 100 000 diluted sample not evaluated ^s	positive	E	356/373 (95.4%)	356/360 (98.9%)	29.2 (17.7-36.0)
			N	146/166 (88.0%)	145/147 (98.6%)	30.2 (20.0-41.5)
			ORF1a	44/46 (95.7%)	44/46 (95.7%)	28.8 (27.3-34.0)
			ORF1ab	41/48 (85.4%)	41/41 (100%)	28.8 (18.4-37.0)
			RdRP	168/185 (90,8%)	168/173 (97.1%)	30.5 (16.0-41.4)
			S	97/100 (97.0%)	97/97 (100%)	28.6 (20.0-34.2)
		n.s. ^s	64/64 (100%)	64/64 (100%)	29.0 (16.5-40.0)	
total	916/983 (93.2%) ^s	915/929 (98.5%) ^s	29.5			
Sample 340061 [*]	SARS-CoV-2 1 : 1 000 000 diluted	positive	E	364/373 (97.6%)	n.a.	32.1 (15.0-40.0)
			N	153/167 (91.6%)		33.3 (20.0-40.7)
			ORF1a	44/46 (95.7%)		31.7 (30.6-36.0)
			ORF1ab	42/48 (87.5%)		31.5 (20.0-39.3)
			RdRP	158/185 (85.4%)		33.5 (19.5-42.8)
			S	97/100 (97.0%)		31.5 (20.0-39.0)
		n.s. ^s	56/64 (87.5%)	31.9 (19.3-37.5)		
total	914/983 (93.0%)	32.4				

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Sample 340061 (Result (qual)) - SARS-CoV-2 positive (inactivated), 1 : 1 000 000 diluted

Reagent	Testkit	Total (N=983)	below detection limit/negative (N=58)	intermediate (N=11)	positive (N=914)	Quota (93%)	Ct/Cp/ Ct/CN Median	Ct/Cp/ Cq/CN Min	Ct/Cp/ Cq/CN Max
E-Gene									
ALTONA DIAGNOSTIC	RealStar SARS-CoV-2 RT-PCR Kit 1.0	48	0	0	48	100,00%	31.7	28.7	36.0
ANATOLIA GENEWORKS	Bosphore Novel Coronavirus Detection Kit v2	4	1	0	3	75,00%	29.4	28.5	30.2
OTHER MANUFACTURERS		10	0	0	10	100,00%	33.0	32.4	36.5
BOSCH HEALTHCARE SOLUTIONS	Vivalytic VRI Multiplex Test	2	0	0	2	100,00%			
CEPHEID	Xpert Xpress SARS-CoV-2	11	0	0	11	100,00%	31.1	30.5	35.1
IN HOUSE (dPCR)		1	0	0	1	97,87%			
IN HOUSE		47	1	0	46	100,00%	32.9	25.7	37.5
ELITech	GeneFinder COVID-19 Plus	8	2	0	6	75,00%	31.2	30.4	35.5
EURO IMMUN	EURORealTime SARS-CoV-2	1	0	0	1	100,00%	33.7	33.7	33.7
EUROFINS TECHNOLOGIES	VirSeek SARS-CoV-2 Screen	1	0	0	1	100,00%	35.5	35.5	35.5
GENETIC SIGNATURES	EasyScreen SARS-CoV-2 Detection Kit	1	0	0	1	100,00%	32.2	32.2	32.2
GERBION	virellaSARS-CoV-2 seqc	5	0	0	5	100,00%	33.0	31.8	33.0
IMMUNDIAGNOSTIK	MutaPLEX Coronavirus	5	1	0	4	80,00%	33.5	31.4	37.9
LIFERIVER	Novel Coronavirus (2019-nCoV) Real Time Multiplex RT-PCR Kit	3	0	0	3	100,00%	30.0	27.0	30.5
LUMINEX	NxTAG CoV	1	0	0	1	100,00%			
MIKROGEN	ampliCube Coronavirus Panel	1	0	0	1	100,00%	32.0	32.0	32.0
MIKROGEN	ampliCube Coronavirus SARS-CoV-2 RUO	3	0	0	3	100,00%	31.4	30.8	35.3
Priv. Inst. f. Immunol. u. Mol.genetik	AmpliGnost CoV-2 E-Gen	4	0	0	4	100,00%	29.9	28.0	32.2
QIAGEN	QIAstat-Dx Respiratory 2019-nCoV Panel	3	0	0	3	100,00%	33.6	28.1	38.6
R-BIOPHARM	RIDA GENE SARS-CoV-2 RUO	55	1	0	54	98,18%	33.3	15.0	37.1
ROCHE DIAGNOSTICS	COBAS SARS-CoV-2 Test	39	0	0	39	100,00%	32.4	30.5	34.9
SEEGENE	Allplex 2019 n-CoV Assay	51	0	1	50	98,04%	31.1	19.2	36.3
TIB MOLBIOL	LightMix Modular SARS and Wuhan CoV E-gene	68	2	0	66	97,06%	32.7	29.6	40.0
TIB MOLBIOL	LightMix Sarbeco E-gene	1	0	0	1	100,00%	32.0	32.0	32.0
		373	8	1	364	97,59%	32.1	15.0	40.0

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Sample 340060 [§]	HCoV OC43 1 : 2 500 diluted specificity control sample not evaluated [§]	negative	E	368/373 (98.7%)	n.a.	-
			N	162/166 (97.6%)		-
			ORF1a	45/46 (97.8%)		-
			ORF1ab	45/48 (93.8%)		-
			RdRP	178/181 (98.3%)		-
			S	99/100 (99.0%)		-
			n.s. [§]	64/64 (100%)		-
total			961/983 (97.8%) [§]		-	
Sample 340065	HCoV 229E 1 : 2 500 diluted specificity control	negative	E	355/373 (95.2%)	354/360 (98.3%)	-
			N	146/166 (88.0%)	144/147 (98.0%)	-
			ORF1a	44/46 (95.7%)	44/46 (95.7%)	-
			ORF1ab	41/48 (85.4%)	40/41 (97.6%)	-
			RdRP	165/182 (90.7%)	165/170 (97.1%)	-
			S	93/100 (93.0%)	93/97 (95.9%)	-
			n.s. [§]	64/64 (100%)	64/64 (100%)	-
total			908/983 (92.4%)	904/929 (97.3%)	-	
Sample 340062	CoV negative specificity control	negative	E	371/373 (99.5%)	n.a.	-
			N	164/167 (98.2%)		-
			ORF1a	46/46 (100%)		-
			ORF1ab	47/48 (97.9%)		-
			RdRP	178/182 (97.8%)		-
			S	99/100 (99.0%)		-
			n.s. [§]	64/64 (100%)		-
total			969/983 (98.6%)		-	
Success rate for all 4 evaluated samples [§]					428/461 (92.8%) [§]	
Success rate for all 4 evaluated samples (corrected) [§]					428/439 (97.5%) [§]	

Thank you very much!

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