INSTAND

Listing and evaluation of the results

Survey of 18 November 2020

Adviser:

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530 Bacterial Genome Detection - Chlamydia trachomatis & Neisseria gonorrhoeae

Chlamydia trachomatis +

Result (qual)		Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
	(1)	2025301	negative (62)	negative (62)	SV		+
	(1)	2025302	positive (61)	positive (61)	SV		+
	(1)	2025303	positive (61)	positive (61)	SV		+
	(1)	2025304	positive (61)	positive (61)	SV		+

Neisseria gonorrhoeae +

Result (qual)		Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria	
	(1)	2025301	positive (61)	positive (61)	SV		+	
	(1)	2025302	negative (62)	negative (62)	SV		+	
	(1)	2025303	positive (61)	positive (61)	SV		+	
	(1)	2025304	positive (61)					only educative

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

531 Bacterial Genome Detection - Chlamydia trachomatis

Chlamydia trachomatis +

Result (qual)		Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
	(1)	2025311	positive (61)	positive (61)	SV		+
	(1)	2025312	positive (61)	positive (61)	SV		+
	(1)	2025313	positive (61)	positive (61)	SV		+
	(1)	2025314	negative (62)	negative (62)	SV		+

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

532 Bacterial Genome Detection - Bordetella pertussis

Bordetella pertussis +

Result (qual)		Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria	
	(1)	2025321	positive (61)	positive (61)	SV		+	
	(1)	2025322	negative (62)	negative (62)	SV		+	
	(1)	2025323	negative (62)					only educative
	(1)	2025324	positive (61)	positive (61)	SV		+	

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

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533 Bacterial Genome Detection - Helicobacter pylori

Helicobacter pylori   +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025331	positive (61)	positive (61)	SV		+
(1)	2025332	negative (62)	negative (62)	SV		+
(1)	2025333	negative (62)	negative (62)	SV		+
(1)	2025334	negative (62)				only educative

(1) To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

534 Bacterial Genome Detection - EHEC / STEC

EHEC / STEC   +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025341	negative (62)	negative (62)	SV		+
(1)	2025342	positive (61)	positive (61)	SV		+
(1)	2025343	positive (61)	positive (61)	SV		+
(1)	2025344	negative (62)	negative (62)	SV		+

(1) To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

535 Bacterial Genome Detection - Borrelia burgdorferi

Borrelia burgdorferi   +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025351	positive (61)	positive (61)	SV		+
(1)	2025352	negative (62)	negative (62)	SV		+
(1)	2025353	positive (61)	positive (61)	SV		+
(1)	2025354	negative (62)				only educative

(1) To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

536 Bacterial Genome Detection - Legionella pneumophila

Legionella pneumophila   +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025361	positive (61)	positive (61)	SV		+
(1)	2025362	negative (62)	negative (62)	SV		+
(1)	2025363	positive (61)	positive (61)	SV		+
(1)	2025364	positive (61)	positive (61)	SV		+

(1) To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

537   Bacterial Genome Detection - Salmonella enterica

Salmonella enterica +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025371	positive (61)	positive (61)	SV		+
(1)	2025372	negative (62)	negative (62)	SV		+
(1)	2025373	negative (62)	negative (62)	SV		+
(1)	2025374	positive (61)	positive (61)	SV		+

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

538   Bacterial Genome Detection - Listeria spp.

Listeria spp. +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025381	negative (62)	negative (62)	SV		+
(1)	2025382	positive (61)	positive (61)	SV		+
(1)	2025383	positive (61)	positive (61)	SV		+
(1)	2025384	positive (61)	positive (61)	SV		+

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

539   Bacterial Genome Detection - MRSA or cMRSA

MRSA / cMRSA +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria	
(1)	2025391	positive (61)					only educative
(1)	2025392	negative (62)	negative (62)	SV		+	
(1)	2025393	positive (61)	positive (61)	SV		+	
(1)	2025394	positive (61)	positive (61)	SV		+	

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

540   Bacterial Genome Detection - Chlamydia pneumoniae

Chlamydia pneumoniae +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025401	negative (62)	negative (62)	SV		+
(1)	2025402	positive (61)	positive (61)	SV		+
(1)	2025403	positive (61)	positive (61)	SV		+
(1)	2025404	negative (62)	negative (62)	SV		+

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

Listing and evaluation of the results

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Laboratory Mustermann

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541 Bacterial Genome Detection - Mycoplasma pneumoniae

Mycoplasma pneumoniae +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025411	negative (62)	negative (62)	SV		+
(1)	2025412	positive (61)	positive (61)	SV		+
(1)	2025413	positive (61)				only educative
(1)	2025414	positive (61)	positive (61)	SV		+

(1) To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

542 Bacterial Genome Detection - Coxiella burnetii & Bacillus anthracis

Coxiella burnetii +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025421	positive (61)	positive (61)	SV		+
(1)	2025422	negative (62)	negative (62)	SV		+
(1)	2025423	negative (62)	negative (62)	SV		+
(1)	2025424	positive (61)	positive (61)	SV		+

Bacillus anthracis +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025421	positive (61)	positive (61)	SV		+
(1)	2025422	positive (61)	positive (61)	SV		+
(1)	2025423	negative (62)	negative (62)	SV		+
(1)	2025424	negative (62)	negative (62)	SV		+

(1) To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

543 Bacterial Genome Detection - Francisella tularensis & Brucella spp.

Francisella tularensis +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025431	positive (61)	positive (61)	SV		+
(1)	2025432	negative (62)	negative (62)	SV		+
(1)	2025433	positive (61)	positive (61)	SV		+
(1)	2025434	negative (62)	negative (62)	SV		+

Brucella spp. +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025431	positive (61)	positive (61)	SV		+
(1)	2025432	positive (61)	positive (61)	SV		+
(1)	2025433	negative (62)	negative (62)	SV		+
(1)	2025434	negative (62)	negative (62)	SV		+

(1) To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

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544 Bacterial Genome Detection - Carbapenemase-Genes

Carbapenemases +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025441	positive (61)	positive (61)	SV		+
(1)	2025442	negative (62)	negative (62)	SV		+
(1)	2025443	positive (61)	positive (61)	SV		+
(1)	2025444	positive (61)				only educative

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

545 Bacterial Genome Detection - Clostridium difficile (Toxin Gene)

Clostridium difficile toxin-genes +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025451	negative (62)	negative (62)	SV		+
(1)	2025452	positive (61)	positive (61)	SV		+
(1)	2025453	positive (61)	positive (61)	SV		+
(1)	2025454	negative (62)	negative (62)	SV		+

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

546 Bacterial Genome Detection - VRE (Vancomycin Resistant Enterococci)

Vancomycin restant enterococci (VRE) +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025461	positive (61)	positive (61)	SV		+
(1)	2025462	negative (62)	negative (62)	SV		+
(1)	2025463	negative (62)	negative (62)	SV		+
(1)	2025464	positive (61)	positive (61)	SV		+

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

547 Bacterial Genome Detection - Urogenital Panel

Mycoplasma hominis +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025471	negative (62)	negative (62)	SV		+
(1)	2025472	positive (61)	positive (61)	SV		+
(1)	2025473	negative (62)	negative (62)	SV		+
(1)	2025474	negative (62)	negative (62)	SV		+

Mycoplasma genitalium +

Result (qual)	Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
(1)	2025471	negative (62)	negative (62)	SV		+
(1)	2025472	negative (62)	negative (62)	SV		+
(1)	2025473	negative (62)	negative (62)	SV		+
(1)	2025474	negative (62)	negative (62)	SV		+

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Ureaplasma parvum

Result (qual)		Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
	(1)	2025471	positive (61)	positive (61)	SV		+
	(1)	2025472	negative (62)	negative (62)	SV		+
	(1)	2025473	negative (62)	negative (62)	SV		+
	(1)	2025474	negative (62)	negative (62)	SV		+

Ureaplasma urealyticum

Result (qual)		Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
	(1)	2025471	negative (62)	negative (62)	SV		+
	(1)	2025472	negative (62)	negative (62)	SV		+
	(1)	2025473	negative (62)	negative (62)	SV		+
	(1)	2025474	negative (62)	negative (62)	SV		+

Trichomonas vaginalis

Result (qual)		Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
	(1)	2025471	negative (62)	negative (62)	SV		+
	(1)	2025472	negative (62)	negative (62)	SV		+
	(1)	2025473	positive (61)	positive (61)	SV		+
	(1)	2025474	negative (62)	negative (62)	SV		+

Gardnerella vaginalis

Result (qual)		Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
	(1)	2025471	negative (62)	negative (62)	SV		+
	(1)	2025472	negative (62)	negative (62)	SV		+
	(1)	2025473	negative (62)	negative (62)	SV		+
	(1)	2025474	positive (61)	positive (61)	SV		+

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

560	**Fungal Genome Detection - Pneumocystis jirovecii**

Pneumocystis jirovecii

Result (qual)		Sample	Your specification(s)	Correct specification(s)	TV-Type		Meets criteria
	(1)	2025601	positive (61)	positive (61)	SV		+
	(1)	2025602	negative (62)	negative (62)	SV		+
	(1)	2025603	positive (61)	positive (61)	SV		+
	(1)	2025604	negative (62)	negative (62)	SV		+

(1)   To obtain a certificate, at least 3 of the 4 samples must be correctly determined (samples classified as educative were not considered).

530 Bacterial Genome Detection - Chlamydia trachomatis & Neisseria gonorrhoeae

Chlamydia trachomatis (N = 240, Rate of success: 97,9%)

Sample 2025301

Collective	positive (61)	negative (62)	total
all participant	3	237 ●	240

Sample 2025302

Collective	positive (61)	total
all participant	240 ●	240

Sample 2025303

Collective	positive (61)	negative (62)	questionable (63)	total
all participant	238 ●	1	1	240

Sample 2025304

Collective	positive (61)	negative (62)	total
all participant	239 ●	1	240

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Neisseria gonorrhoeae (N = 238, Rate of success: 97,1%)

Sample 2025301

Collective	positive (61)	negative (62)	questionable (63)	total
all participant	236 ●	1	1	238

Sample 2025302

Collective	positive (61)	negative (62)	total
all participant	3	235 ●	238

Sample 2025303

Collective	positive (61)	negative (62)	questionable (63)	total
all participant	234 ●	3	1	238

Sample 2025304 not evaluated

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

531 Bacterial Genome Detection - Chlamydia trachomatis

Chlamydia trachomatis (N = 52, Rate of success: 100%)

Sample 2025311

Collective	positive (61)	total
all participant	52 ●	52

Sample 2025312

Collective	positive (61)	total
all participant	52 ●	52

Sample 2025313

Collective	positive (61)	total
all participant	52 ●	52

Sample 2025314

Collective	negative (62)	total
all participant	52 ●	52

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

532   Bacterial Genome Detection - Bordetella pertussis

Bordetella pertussis (N = 146, Rate of success: 97,9%)

Sample 2025321

Collective	positive (61)	negative (62)	total
all participant	145 ●	1	146

Sample 2025322

Collective	positive (61)	negative (62)	total
all participant	2	144 ●	146

Sample 2025323 not evaluated

Sample 2025324

Collective	positive (61)	negative (62)	Inhibition (64)	total
all participant	144 ●	1	1	146

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

533   Bacterial Genome Detection - Helicobacter pylori

Helicobacter pylori (N = 45, Rate of success: 88,9%)

Sample 2025331

Collective	positive (61)	negative (62)	total
all participant	44 ●	1	45

Sample 2025332

Collective	positive (61)	negative (62)	total
all participant	1	44 ●	45

Sample 2025333

Collective	positive (61)	negative (62)	total
all participant	5	40 ●	45

Sample 2025334 not evaluated

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

534   Bacterial Genome Detection - EHEC / STEC

EHEC / STEC (N = 127, Rate of success: 91,3%)

Sample 2025341

Collective	positive (61)	negative (62)	total
all participant	7	119 ●	126

Sample 2025342

Collective	positive (61)	negative (62)	total
all participant	126 ●	1	127

Sample 2025343

Collective	positive (61)	negative (62)	total
all participant	125 ●	2	127

Sample 2025344

Collective	positive (61)	negative (62)	total
all participant	2	125 ●	127

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

535 Bacterial Genome Detection - Borrelia burgdorferi

Borrelia burgdorferi (N = 98, Rate of success: 93,9%)

Sample 2025351

Collective	positive (61)	negative (62)	total
all participant	92 ●	6	98

Sample 2025352

Collective	negative (62)	total
all participant	98 ●	98

Sample 2025353

Collective	positive (61)	total
all participant	98 ●	98

Sample 2025354 not evaluated

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

536 Bacterial Genome Detection - Legionella pneumophila

Legionella pneumophila (N = 109, Rate of success: 95,4%)

Sample 2025361

Collective	positive (61)	negative (62)	total
all participant	106 ●	3	109

Sample 2025362

Collective	negative (62)	total
all participant	109 ●	109

Sample 2025363

Collective	positive (61)	negative (62)	total
all participant	107 ●	2	109

Sample 2025364

Collective	positive (61)	negative (62)	total
all participant	107 ●	2	109

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

537 Bacterial Genome Detection - Salmonella enterica

Salmonella enterica (N = 24, Rate of success: 100%)

Sample 2025371

Collective	positive (61)	total
all participant	24 ●	24

Sample 2025372

Collective	negative (62)	total
all participant	24 ●	24

Sample 2025373

Collective	negative (62)	total
all participant	24 ●	24

Sample 2025374

Collective	positive (61)	total
all participant	24 ●	24

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

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538   Bacterial Genome Detection - Listeria spp.

Listeria spp. (N = 42, Rate of success: 97,6%)

Sample 2025381

Collective	positive (61)	negative (62)	total
all participant	1	41 ●	42

Sample 2025382

Collective	positive (61)	total
all participant	42 ●	42

Sample 2025383

Collective	positive (61)	total
all participant	42 ●	42

Sample 2025384

Collective	positive (61)	total
all participant	42 ●	42

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

539   Bacterial Genome Detection - MRSA or cMRSA

MRSA / cMRSA (N = 277, Rate of success: 96,4%)

Sample 2025391 not evaluated

Sample 2025393

Collective	positive (61)	negative (62)	total
all participant	273 ●	3	276

Sample 2025392

Collective	positive (61)	negative (62)	total
all participant	5	272 ●	277

Sample 2025394

Collective	positive (61)	negative (62)	total
all participant	275 ●	2	277

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

540   Bacterial Genome Detection - Chlamydia pneumoniae

Chlamydia pneumoniae (N = 119, Rate of success: 97,5%)

Sample 2025401

Collective	positive (61)	negative (62)	total
all participant	1	118 ●	119

Sample 2025402

Collective	positive (61)	total
all participant	119 ●	119

Sample 2025403

Collective	positive (61)	negative (62)	total
all participant	118 ●	1	119

Sample 2025404

Collective	positive (61)	negative (62)	total
all participant	3	116 ●	119

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Bacterial Genome Detection - Mycoplasma pneumoniae

Mycoplasma pneumoniae (N = 136, Rate of success: 95,6%)

Sample 2025411

Collective	positive (61)	negative (62)	total
all participant	3	133 ●	136

Sample 2025412

Collective	positive (61)	negative (62)	total
all participant	134 ●	2	136

Sample 2025413 not evaluated

Sample 2025414

Collective	positive (61)	negative (62)	total
all participant	135 ●	1	136

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Bacterial Genome Detection - Coxiella burnetii & Bacillus anthracis

Coxiella burnetii (N = 47, Rate of success: 93,6%)

Sample 2025421

Collective	positive (61)	negative (62)	total
all participant	46 ●	1	47

Sample 2025422

Collective	positive (61)	negative (62)	total
all participant	2	45 ●	47

Sample 2025423

Collective	negative (62)	total
all participant	47 ●	47

Sample 2025424

Collective	positive (61)	negative (62)	total
all participant	46 ●	1	47

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Bacillus anthracis (N = 24, Rate of success: 100%)

Sample 2025421

Collective	positive (61)	total
all participant	24 ●	24

Sample 2025422

Collective	positive (61)	total
all participant	24 ●	24

Sample 2025423

Collective	negative (62)	total
all participant	24 ●	24

Sample 2025424

Collective	negative (62)	total
all participant	24 ●	24

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Bacterial Genome Detection - Francisella tularensis & Brucella spp.

Francisella tularensis (N = 27, Rate of success: 92,6%)

Sample 2025431

Collective	positive (61)	negative (62)	total
all participant	25 ●	2	27

Sample 2025432

Collective	negative (62)	total
all participant	27 ●	27

Sample 2025433

Collective	positive (61)	total
all participant	27 ●	27

Sample 2025434

Collective	negative (62)	total
all participant	27 ●	27

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Brucella spp. (N = 25, Rate of success: 100%)

Sample 2025431

Collective	positive (61)	total
all participant	25 ●	25

Sample 2025432

Collective	positive (61)	total
all participant	25 ●	25

Sample 2025433

Collective	negative (62)	total
all participant	25 ●	25

Sample 2025434

Collective	negative (62)	total
all participant	25 ●	25

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

544 Bacterial Genome Detection - Carbapenemase-Genes

Carbapenemases (N = 80, Rate of success: 98,8%)

Sample 2025441

Collective	positive (61)	total
all participant	80 ●	80

Sample 2025442

Collective	positive (61)	negative (62)	total
all participant	1	79 ●	80

Sample 2025443

Collective	positive (61)	total
all participant	80 ●	80

Sample 2025444 not evaluated

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

545 Bacterial Genome Detection - Clostridium difficile (Toxin Gene)

Clostridium difficile toxin-genes (N = 165, Rate of success: 97%)

Sample 2025451

Collective	negative (62)	questionable (63)	total
all participant	164 ●	1	165

Sample 2025452

Collective	positive (61)	negative (62)	total
all participant	162 ●	3	165

Sample 2025453

Collective	positive (61)	negative (62)	total
all participant	164 ●	1	165

Sample 2025454

Collective	positive (61)	negative (62)	total
all participant	2	163 ●	165

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

546 Bacterial Genome Detection - VRE (Vancomycin Resistant Enterococci)

Vancomycin restant enterococci (VRE) (N = 63, Rate of success: 98,4%)

Sample 2025461

Collective	positive (61)	total
all participant	63 ●	63

Sample 2025462

Collective	negative (62)	total
all participant	63 ●	63

Sample 2025463

Collective	positive (61)	negative (62)	total
all participant	1	62 ●	63

Sample 2025464

Collective	positive (61)	total
all participant	63 ●	63

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Bacterial Genome Detection - Urogenital Panel

Mycoplasma hominis (N = 71, Rate of success: 94,4%)

Sample 2025471

Collective	positive (61)	negative (62)	total
all participant	2	69 ●	71

Sample 2025472

Collective	positive (61)	total
all participant	71 ●	71

Sample 2025473

Collective	positive (61)	negative (62)	total
all participant	1	70 ●	71

Sample 2025474

Collective	positive (61)	negative (62)	total
all participant	3	68 ●	71

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Mycoplasma genitalium (N = 75, Rate of success: 98,7%)

Sample 2025471

Collective	positive (61)	negative (62)	total
all participant	1	74 ●	75

Sample 2025472

Collective	negative (62)	total
all participant	75 ●	75

Sample 2025473

Collective	negative (62)	total
all participant	75 ●	75

Sample 2025474

Collective	negative (62)	total
all participant	75 ●	75

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Ureaplasma parvum (N = 53, Rate of success: 100%)

Sample 2025471

Collective	positive (61)	total
all participant	53 ●	53

Sample 2025472

Collective	negative (62)	total
all participant	53 ●	53

Sample 2025473

Collective	negative (62)	total
all participant	53 ●	53

Sample 2025474

Collective	negative (62)	total
all participant	53 ●	53

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Ureaplasma urealyticum (N = 57, Rate of success: 89,5%)

Sample 2025471

Collective	positive (61)	negative (62)	total
all participant	6	51 ●	57

Sample 2025472

Collective	negative (62)	total
all participant	57 ●	57

Sample 2025473

Collective	negative (62)	total
all participant	57 ●	57

Sample 2025474

Collective	negative (62)	total
all participant	57 ●	57

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Trichomonas vaginalis (N = 63, Rate of success: 92,1%)

Sample 2025471

Collective	positive (61)	negative (62)	total
all participant	1	62 ●	63

Sample 2025472

Collective	positive (61)	negative (62)	total
all participant	1	62 ●	63

Sample 2025473

Collective	positive (61)	negative (62)	total
all participant	59 ●	4	63

Sample 2025474

Collective	positive (61)	negative (62)	total
all participant	1	62 ●	63

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

Gardnerella vaginalis (N = 17, Rate of success: 94,1%)

Sample 2025471

Collective	negative (62)	total
all participant	17 ●	17

Sample 2025472

Collective	negative (62)	total
all participant	17 ●	17

Sample 2025473

Collective	negative (62)	total
all participant	17 ●	17

Sample 2025474

Collective	positive (61)	negative (62)	total
all participant	16 ●	1	17

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective

560   Fungal Genome Detection - Pneumocystis jirovecii

Pneumocystis jirovecii (N = 111, Rate of success: 95,5%)

Sample 2025601

Collective	positive (61)	negative (62)	total
all participant	107 ●	4	111

Sample 2025602

Collective	positive (61)	negative (62)	total
all participant	1	110 ●	111

Sample 2025603

Collective	positive (61)	total
all participant	111 ●	111

Sample 2025604

Collective	positive (61)	negative (62)	Inhibition (64)	total
all participant	1	109 ●	1	111

The point corresponds to the correct result, the horizontal bar corresponds to your specification, the vertical bar corresponds to your collective