

dPCR Microbial DNA Detection Assays

Assay/target list

The dPCR Microbial DNA Detection Assays are designed to test for the presence of bacterial, fungal, parasitic, viral, antibiotic resistance or virulence factor genes using digital PCR. The assays can be run on the QIAcuity® in about two hours with minimal hands-on time. The assays target over 800 bacterial, fungal, parasitic, viral, antibiotic resistance or virulence factor genes. The assay product consists of a tube containing a primer pair and a hydrolysis probe and is configurable for the dye (fluorophore). Selectable dyes are FAM, HEX, ROX, TAMRA, Cy5 and ATTO700 to enable a mix and match of up to 12 targets/assay in a dPCR multiplex reaction. However, each assay

supplies enough material to perform 200 x 40 µL dPCR reactions in Nanoplates 26k or 666 x 12 µL dPCR reactions in Nanoplates 8.5k. A reaction is run by combining the assays with the QIAcuity Probe PCR Kit (for DNA targets), the QIAcuity High Multiplex Probe PCR Kit (for DNA targets) or the QIAcuity OneStep Advanced Probe Kit (for RNA or RNA+DNA targets).

More information on the dPCR Microbial DNA Detection Assays can be found at geneglobe.qiagen.com/applications/digital-pcr.

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Table 1. Bacterial identification targets/assays

Target species (target region, NCBI taxonomy)	Application field	GeneGlobe® catalog number	Wet-lab tested
<i>Abiotrophia defectiva</i> (46125)	Human pathogens	DMA00001	
<i>Acetobacter</i> spp. (16S, 434)	Food production	DMA00753	YES
<i>Achromobacter xylosoxidans</i> (85698)	Human pathogens	DMA00002	
<i>Acidaminococcus fermentans</i> (951)	Gastrointestinal pathogens	DMA00003	YES
<i>Acinetobacter baumannii</i> (557601)	Human pathogens	DMA00004	YES
<i>Acinetobacter calcoaceticus</i> (471)	Human pathogens, Multiple drug resistance (MDR)	DMA00005	YES
<i>Acinetobacter haemolyticus</i> (29430)	Human pathogens, MDR	DMA00006	YES
<i>Actinobacillus hominis</i> (719)	Human microbiome	DMA00007	
<i>Actinobaculum schaalii</i> (59505)	Human pathogens	DMA00741	YES
<i>Actinomyces gerencseriae</i> (52769)	Human pathogens	DMA00009	
<i>Actinomyces graevenitzi</i> (55565)	Infectious diseases	DMA00010	
<i>Actinomyces israelii</i> (1659)	Infectious diseases	DMA00011	YES
<i>Actinomyces lingnae</i> (114702)	Infectious diseases	DMA00012	
<i>Actinomyces naeslundii</i> (1655)	Infectious diseases	DMA00013	YES
<i>Actinomyces urogenitalis</i> (103621)	Infectious diseases	DMA00016	
<i>Actinomyces viscosus</i> (1656)	Infectious diseases	DMA00017	YES
<i>Aerococcus christensenii</i> (87541)	Human microbiome	DMA00018	
<i>Aerococcus urinae</i> (1376)	Infectious diseases	DMA00475	
<i>Aerococcus viridans</i> (1377)	Infectious diseases	DMA00019	
<i>Aeromonas hydrophila</i> (644)	Gastrointestinal infections	DMA00020	
<i>Aeromonas veronii</i> (654)	Gastrointestinal pathogens	DMA00021	
<i>Aggregatibacter actinomycetemcomitans</i> (714)	Infectious diseases	DMA00022	
<i>Aggregatibacter segnis</i> (739)	Infectious diseases	DMA00023	
<i>Akkermansia muciniphila</i> (239935)	Human microbiome	DMA00024	YES
<i>Alcaligenes faecalis</i> (511)	Sepsis, MDR	DMA00025	
<i>Aliarcobacter butzleri</i> (28197)	Foodborne pathogens, MDR	DMA00033	
<i>Alicyclobacillus acidoterrestris</i> (1450)	Food production	DMA00765	YES
<i>Alicyclobacillus</i> spp (29330)	Food production	DMA00766	YES
<i>Alistipes putredinis</i> (28117)	Human microbiome	DMA00026	
<i>Allorevotella tannerae</i> (76122)	Human microbiome	DMA00273	
<i>Anaerobutyricum hallii</i> (39488)	Human microbiome	DMA00141	YES
<i>Anaerococcus hydrogenalis</i> (33029)	Human microbiome	DMA00027	
<i>Anaerococcus lactolyticus</i> (33032)	Human microbiome	DMA00028	
<i>Anaerococcus prevotii</i> (33034)	Human microbiome	DMA00029	
<i>Anaeroglobus geminatus</i> (156456)	Human microbiome	DMA00030	
<i>Anaerostipes caccae</i> (105841)	Human microbiome	DMA00031	

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<i>Anaerotruncus colihominis</i> (169435)	Human microbiome	DMA00032	
<i>Arachnia propionica</i> (1750)	Human microbiome	DMA00276	
<i>Arcobacter skirrowii</i> (28200)	Gastrointestinal infections	DMA00034	
<i>Bacillus anthracis</i> (1392)	Human pathogens	DMA00040	YES
<i>Bacillus atrophaeus</i> (1452)	Human pathogens	DMA00739	YES
<i>Bacillus cereus</i> (1396)	Probiotics, Food production, Infectious diseases	DMA00041	YES
<i>Bacillus licheniformis</i> (1402)	Food production, Infectious diseases	DMA00042	YES
<i>Bacillus subtilis</i> (1423)	Probiotics, Gastrointestinal infections	DMA00043	YES
<i>Bacterial vaginosis-associated bacterium 2</i>	Omit, Human pathogens	DMA00727	YES
<i>Bacterioides</i> sp. (4_3_47FAA, 457394)	Human microbiome	DMA00055	
<i>Bacteroides caccae</i> (47678)	Human microbiome	DMA00044	YES
<i>Bacteroides eggerthii</i> (28111)	Human microbiome	DMA00048	
<i>Bacteroides fragilis</i> (817)	Human microbiome	DMA00049	YES
<i>Bacteroides intestinalis</i> (329854)	Human microbiome	DMA00050	YES
<i>Bacteroides stercoris</i> (46506)	Human pathogens	DMA00056	
<i>Bacteroides thetaiotaomicron</i> (818)	Human microbiome	DMA00057	YES
<i>Bifidobacterium adolescentis</i> (1680)	Human microbiome, Probiotics	DMA00060	YES
<i>Bifidobacterium bifidum</i> (1681)	Human microbiome, Probiotics	DMA00061	YES
<i>Bifidobacterium breve</i> (1685)	Human microbiome, Probiotics	DMA00062	YES
<i>Bifidobacterium dentium</i> (1689)	Human microbiome	DMA00063	YES
<i>Bifidobacterium longum</i> (216816)	Human microbiome, Probiotics	DMA00064	YES
<i>Bifidobacterium pseudocatenulatum</i> (28026)	Human microbiome	DMA00065	YES
<i>Bifidobacterium scardovii</i> (158787)	Human microbiome	DMA00066	YES
<i>Blautia coccooides</i> (1532)	Human microbiome	DMA00404	YES
<i>Blautia hydrogenotrophica</i> (53443)	Human microbiome	DMA00067	
<i>Blautia obeum</i> (40520)	Human microbiome	DMA00289	
<i>Bordetella bronchiseptica</i> (518), <i>Bordetella holmesii</i> (35814), <i>Bordetella parapertussis</i> (519), <i>Bordetella pertussis</i> (520)	Human pathogens	DMA00068	
<i>Bordetella parapertussis</i> (519)	Human pathogens	DMA00402	YES
<i>Bordetella pertussis</i> (520)	Human pathogens	DMA00401	YES
<i>Borrelia burgdorferi</i> (139)	Human pathogens	DMA00792	YES
<i>Brevibacillus agri</i> (51101)	Human pathogens	DMA00069	
<i>Brevibacillus brevis</i> (1393)	Infectious diseases, Environmental Microbes	DMA00070	
<i>Brevibacterium casei</i> (33889)	Food production, Infectious diseases	DMA00071	
<i>Brevundimonas diminuta</i> (293)	Infectious diseases	DMA00072	
<i>Brevundimonas vesicularis</i> (41276)	Infectious diseases	DMA00073	



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<i>Brucella anthropi</i> (529)	Environmental microbes, Infectious diseases	DMA00240	
<i>Burkholderia cepacia</i> (292)	Human pathogens	DMA00074	
<i>Burkholderia gladioli</i> (28095)	Food production, Infectious diseases	DMA00075	
<i>Burkholderia mallei</i> (13373)	Infectious diseases	DMA00076	
<i>Butyrivibrio pullicaecorum</i> (501571)	Human microbiome	DMA00077	
<i>Butyrivibrio crossotus</i> (45851)	Human microbiome	DMA00078	
<i>Butyrivibrio fibrisolvens</i> (831)	Human microbiome	DMA00079	
<i>Campylobacter</i> (194)	Infectious diseases, Food production, Human pathogens	DMA00355	YES
<i>Campylobacter coli</i> (195)	Gastrointestinal infections	DMA00080	YES
<i>Campylobacter concisus</i> (199)	Human microbiome	DMA00081	YES
<i>Campylobacter fetus</i> (196)	Food production, Infectious diseases	DMA00082	YES
<i>Campylobacter gracilis</i> (824)	Gastrointestinal infections	DMA00083	YES
<i>Campylobacter jejuni</i> (197)	Gastrointestinal infections	DMA00084	YES
<i>Campylobacter jejuni</i> (197; 195; 201)	Food production	DMA00850	YES
<i>Campylobacter rectus</i> (203)	Infectious diseases	DMA00085	YES
<i>Campylobacter showae</i> (204)	Human microbiome	DMA00086	YES
<i>Campylobacter sputorum</i> (206)	Food production, Human microbiome	DMA00087	YES
<i>Campylobacter upsaliensis</i> (28080)	Gastrointestinal infections	DMA00088	YES
<i>Campylobacter ureolyticus</i> (827)	Gastrointestinal infections	DMA00058	
<i>Capnocytophaga gingivalis</i> (1017)	Human microbiome	DMA00094	
<i>Capnocytophaga granulosa</i> (45242)	Human microbiome	DMA00095	
<i>Capnocytophaga ochracea</i> (1018)	Human microbiome	DMA00096	
<i>Capnocytophaga sputigena</i> (1019)	Human microbiome	DMA00097	
<i>Cardiobacterium hominis</i> (2718)	Human microbiome, Infectious diseases	DMA00098	
<i>Catellibacterium marimammalium</i> (300419)	Human microbiome	DMA00099	
<i>Catenibacterium mitsuokai</i> (100886)	Human microbiome	DMA00100	
<i>Catonella morbi</i> (43997)	Human microbiome	DMA00101	
<i>Chlamydia pneumoniae</i> (83558)	Infectious diseases	DMA00103	
<i>Chlamydia psittaci</i> (83554)	Infectious diseases	DMA00104	
<i>Chlamydia trachomatis</i> (813)	Infectious diseases	DMA00102	
<i>Citrobacter freundii</i> (546)	Human microbiome, Wastewater, Food production	DMA00105	
<i>Citrobacter youngae</i> (133448)	Infectious diseases	DMA00106	
<i>Clostridioides difficile</i> (1496)	MDR, Sepsis, Infectious diseases, Wastewater	DMA00107	YES
<i>Clostridium botulinum</i> (1491)	Food production	DMA00849	YES
<i>Clostridium perfringens</i> (1502)	Human pathogens	DMA00109	YES
<i>Clostridium septicum</i> (1504)	Human pathogens	DMA00110	

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<i>Clostridium</i> sp. (M62/1, 411486)	Human microbiome	DMA00112	YES
<i>Clostridium</i> sp. (SS2/1, 411484)	Human microbiome	DMA00113	YES
<i>Clostridium tetani</i> (1513)	Human pathogens	DMA00114	YES
<i>Collinsella aerofaciens</i> (74426)	Human microbiome	DMA00115	
<i>Coprococcus comes</i> (410072)	Human microbiome	DMA00116	
<i>Coprococcus eutactus</i> (33043)	Human microbiome	DMA00117	
<i>Corynebacterium aurimucosum</i> (169292)	Human microbiome; Infectious diseases	DMA00118	YES
<i>Corynebacterium diphtheriae</i> (1717)	Human pathogens	DMA00119	YES
<i>Corynebacterium diphtheriae</i> (rpoB, 1717)	Infectious diseases	DMA00717	YES
<i>Corynebacterium durum</i> (61592)	Human microbiome	DMA00120	YES
<i>Corynebacterium matruchotii</i> (43768)	Human microbiome	DMA00121	YES
<i>Corynebacterium pseudodiphtheriticum</i> (37637)	Human microbiome, Infectious diseases	DMA00122	YES
<i>Corynebacterium riegelii</i> (156976)	Human pathogens	DMA00742	YES
<i>Corynebacterium ulcerans</i> rpoB (65058)	Human pathogens	DMA00794	YES
<i>Coxiella burnetii</i> (777)	Human pathogens, Infectious diseases	DMA00123	
<i>Cronobacter</i> (413496)	Food production, Infectious diseases	DMA00694	YES
<i>Cronobacter sakazakii</i> (28141)	Food production	DMA00847	YES
<i>Cutibacterium acnes</i> (1747)	Human microbiome	DMA00275	
<i>Cylindrospermum raciborskii</i> (77022)	Wastewater	DMA00813	YES
<i>Desulfovibrio desulfuricans</i> (876)	Infectious diseases	DMA00124	
<i>Desulfovibrio piger</i> (901)	Human microbiome	DMA00125	
<i>Desulfovibrio vulgaris</i> (881)	Human microbiome	DMA00126	
<i>Dialister invisus</i> (218538)	Human microbiome	DMA00127	
<i>Dialister pneumosintes</i> (39950)	Infectious diseases	DMA00128	
<i>Dolichospermum ucrainicum</i> (748770)	Wastewater	DMA00764	YES
<i>Dorea formicigenerans</i> (39486)	Human microbiome	DMA00129	
<i>Eggerthella sinensis</i> (242230)	Human pathogens	DMA00130	
<i>Ehrlichia canis</i> (944)	Animal pathogens	DMA00131	
<i>Eikenella corrodens</i> (539)	Human microbiome	DMA00132	
<i>Elizabethkingia meningoseptica</i> (238)	Human pathogens	DMA00133	
<i>Enterobacter cloacae</i> 1 (16S, 550)	Human pathogens, Wastewater, MDR	DMA00134	YES
<i>Enterobacter cloacae</i> 2 (ompC, 550)	Human pathogens, Wastewater, MDR	DMA00449	YES
<i>Enterococcus casseliflavus</i> (37734)	Infectious diseases	DMA00135	YES
<i>Enterococcus faecalis</i> 1 (16S, 1351)	Probiotics, Wastewater, Gastrointestinal infections	DMA00136	YES
<i>Enterococcus faecalis</i> 2 (23S, 1351)	Probiotics, Wastewater, Infectious diseases	DMA00446	YES
<i>Enterococcus italicus</i> (246144)	Food production	DMA00138	YES



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<i>Eperythrozoon wenyoni</i> (65123)	Animal disease	DMA00806	
<i>Erysiphe necator</i> (cytB, 52586)	Food production	DMA00777	YES
<i>Erysipelothrix rhusiopathiae</i> (1648)	Food production	DMA00139	
<i>Escherichia coli</i> (16S, 562)	Human microbiome, Wastewater	DMA00140	YES
<i>Escherichia coli</i> (stx1, 562)	Human microbiome, Wastewater	DMA00696	YES
<i>Escherichia coli</i> (stx2, 562)	Human microbiome, Wastewater	DMA00697	YES
<i>Escherichia coli</i> (eae, 562)	Human microbiome, Wastewater	DMA00698	YES
<i>Escherichia coli</i> (rfbE, 562)	Human microbiome, Wastewater	DMA00699	YES
<i>Escherichia coli</i> O157:H7 (83334)	Food production	DMA00851	YES
<i>Eubacterium infirmum</i> (56774)	Human microbiome	DMA00142	YES
<i>Eubacterium rectale</i> (39491)	Human microbiome	DMA00143	YES
<i>Eubacterium siraeum</i> (39492)	Human microbiome	DMA00145	
<i>Eubacterium ventriosum</i> (39496)	Human microbiome	DMA00146	YES
<i>Exiguobacterium aurantiacum</i> (33987)	Infectious diseases	DMA00147	
<i>Faecalibacterium prausnitzii</i> (853)	Human microbiome	DMA00148	YES
<i>Fannyhessea vaginae</i> (82135)	Human microbiome	DMA00039	YES
<i>Filifactor alocis</i> (143361)	Human microbiome, Human pathogens	DMA00149	
<i>Finegoldia magna</i> (1260)	Human microbiome	DMA00150	YES
<i>Francisella tularensis</i> (263)	Infectious diseases	DMA00151	
<i>Fructilactobacillus lindneri</i> (53444)	Probiotics, Fermentation and degradation	DMA00385	
<i>Fusobacterium mortiferum</i> (850)	Human microbiome	DMA00152	YES
<i>Fusobacterium necrophorum</i> (859)	Human microbiome, Infectious diseases	DMA00153	YES
<i>Fusobacterium nucleatum</i> (851)	Human microbiome	DMA00154	YES
<i>Fusobacterium periodonticum</i> (860)	Human microbiome, Infectious diseases	DMA00155	YES
<i>Fusobacterium varium</i> (856)	Human microbiome	DMA00156	YES
<i>Gardnerella vaginalis</i> 409-05 (Sialidase A)	Human microbiome, Infectious diseases	DMA00408	YES
<i>Gardnerella vaginalis</i> 409-05 (16S, 553190)	Human microbiome, Infectious diseases	DMA00157	YES
<i>Gardnerella vaginalis</i> (Sialidase A, 553190)	Human microbiome, Infectious diseases	DMA00408	
<i>Gemella bergeri</i> (84136)	Infectious diseases	DMA00158	
<i>Gemella haemolysans</i> (1379)	Human microbiome	DMA00159	
<i>Gemella morbillorum</i> (29391)	Infectious diseases	DMA00160	
<i>Geobacillus stearothermophilus</i> (1422)	Food production	DMA00161	
<i>Gleimia europaea</i> (66228)	Human pathogens	DMA00008	
<i>Gluconobacter</i> spp./ <i>Acetobacter</i> spp. (441)	Food production	DMA00768	YES
<i>Granulicatella adiacens</i> (46124)	Infectious diseases	DMA00162	
<i>Granulicatella elegans</i> (137732)	Human microbiome, Infectious diseases	DMA00163	
<i>Haemophilus ducreyi</i> (730)	Human pathogens, Infectious diseases	DMA00164	YES

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<i>Haemophilus influenzae</i> (727)	Human microbiome, Infectious diseases, Sepsis	DMA00165	YES
<i>Haemophilus parainfluenzae</i> (729)	Human microbiome, Infectious diseases, Sepsis	DMA00166	YES
<i>Hafnia alvei</i> (569)	Human microbiome, Food production	DMA00167	
<i>Helicobacter cinaedi</i> (213)	Infectious diseases	DMA00168	YES
<i>Helicobacter fennelliae</i> (215)	Infectious diseases	DMA00169	YES
<i>Helicobacter pylori</i> (210)	Infectious diseases	DMA00170	YES
<i>Inquilinus limosus</i> (171674), <i>Inquilinus ginsengisoli</i> (363840)	Human pathogens	DMA00171	
<i>Kingella denitrificans</i> (502)	Human pathogens, Infectious diseases	DMA00172	
<i>Kingella kingae</i> (504)	Infectious diseases	DMA00173	
<i>Klebsiella aerogenes</i> (548)	Infectious diseases	DMA00448	YES
<i>Klebsiella granulomatis</i> (39824)	Human pathogens	DMA00174	YES
<i>Klebsiella oxytoca</i> (571)	Human pathogens	DMA00740	YES
<i>Klebsiella pneumoniae</i> (573)	Human pathogens, Infectious diseases	DMA00367	YES
<i>Lachnoanaerobaculum saburreum</i> (467210)	Human microbiome	DMA00144	
<i>Lachnobacterium bovis</i> (140626)	Food production	DMA00176	
<i>Lactocaseibacillus casei</i> (1582)	Probiotics, Human microbiome, Food production	DMA00178	YES
<i>Lactocaseibacillus rhamnosus</i> (47715)	Probiotics, Human microbiome, Food production	DMA00186	
<i>Lactiplantibacillus plantarum</i> (1590)	Probiotics, Human microbiome	DMA00185	YES
<i>Lactobacillus acidophilus</i> (1579)	Probiotics, Human microbiome, Food production	DMA00177	YES
<i>Lactobacillus brevis</i> (1580)	Probiotics	DMA00381	
<i>Lactobacillus crispatus</i> (47770)	Probiotics, Human microbiome, Food production	DMA00179	YES
<i>Lactobacillus delbrueckii</i> (1584)	Probiotics, Human microbiome, Food production	DMA00180	
<i>Lactobacillus gasseri</i> (1596)	Probiotics, Human microbiome, Food production	DMA00182	
<i>Lactobacillus iners</i> (147802)	Probiotics, Human microbiome, Food production	DMA00183	YES
<i>Lactobacillus jensenii</i> (109790)	Human microbiome	DMA00184	
<i>Lactobacillus Pan (A/AB)</i> (16S, 1578)	Food production	DMA00759	YES
<i>Lactobacillus Pan (B/AB)</i> (16S, 1578)	Food production	DMA00760	YES
<i>Lactococcus garvieae</i> (1363)	Food production	DMA00189	
<i>Lactococcus lactis</i> (1358)	Probiotics, Human microbiome, Food production	DMA00190	
<i>Lancefieldella parvula</i> (1382)	Human microbiome	DMA00037	YES
<i>Lancefieldella rimae</i> (1383)	Human microbiome	DMA00038	YES
<i>Latilactobacillus curvatus</i> (28038)	Probiotics	DMA00384	
<i>Lautropia mirabilis</i> (47671)	Human microbiome, Infectious diseases	DMA00191	



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<i>Legionella</i> (445)	Environmental microbes, Infectious diseases	DMA00705	YES
<i>Legionella longbeachae</i> (450)	Environmental microbes, Infectious diseases	DMA00363	YES
<i>Legionella pneumophila</i> (16S, 446)	Infectious diseases	DMA00192	YES
<i>Legionella pneumophila</i> (2 mip region, 446)	Environmental microbes, Infectious diseases	DMA00706	YES
<i>Legionella pneumophila</i> serotyp 1 (66976)	Wastewater	DMA00778	YES
<i>Leifsonia aquatica</i> serotype (144185)	Infectious diseases	DMA00193	
<i>Lentilactobacillus buchneri</i> (1581)	Probiotics, Food production, Biocontrol	DMA00382	
<i>Leptospira interrogans</i> (173)	Human pathogens, Infectious diseases	DMA00194	YES
<i>Leptotrichia buccalis</i> (40542)	Human microbiome	DMA00196	
<i>Leptotrichia wadei</i> (157687)	Human microbiome	DMA00198	
<i>Ligilactobacillus salivarius</i> (1624)	Human microbiome	DMA00187	
<i>Limosilactobacillus fermentum</i> (1613)	Probiotics, Human microbiome, Food production	DMA00181	
<i>Limosilactobacillus vaginalis</i> (1633)	Human microbiome	DMA00188	YES
<i>Listeria</i> (1637)	Infectious diseases, Food production, Human pathogens	DMA00695	YES
<i>Listeria grayi</i> (1641)	Probiotics, Food production, Environmental microbes	DMA00707	
<i>Listeria monocytogenes</i> (1639)	Infectious diseases, Food production, Human pathogens	DMA00199	YES
<i>Loigolactobacillus coryniformis</i> (1610)	Probiotics	DMA00383	
<i>Lysinibacillus fusiformis</i> (28031)	Infectious diseases, Sepsis	DMA00200	
<i>Mageeibacillus indolicus</i> (884684)	Human microbiome	DMA00409	
<i>Massilia timonae</i> (47229)	Infectious diseases, Human pathogens	DMA00201	
<i>Megasphaera cerevisiae</i> (39029)	Food production	DMA00386	
<i>Megasphaera genomsp. type_1</i> (699192)	Human microbiome, Infectious diseases, Human pathogens	DMA00405	
<i>Megasphaera genomsp. type_2</i> (699193)	Human microbiome, Infectious diseases, Human pathogens	DMA00406	
<i>Megasphaera micronuciformis</i> (187326)	Infectious diseases, Human pathogens	DMA00202	
<i>Megasphaera</i> phylotype 2	Infectious diseases, Human pathogens		YES
<i>Megasphaera</i> sp. (DJF_B143, 537288)	Food production	DMA00203	
<i>Methanobrevibacter smithii</i> (2173)	Human microbiome	DMA00491	
<i>Methylobacterium fujisawaense</i> (107400)	Infectious diseases	DMA00204	
<i>Methylobacterium mesophilicum</i> (39956)	Infectious diseases	DMA00205	
<i>Methylobacterium zatmanii</i> (29429)	Human microbiome, Infectious diseases	DMA00206	
<i>Microbacterium binotii</i> (462710)	Human microbiome, Infectious diseases	DMA00207	
<i>Micrococcus luteus</i> (1270)	Human microbiome, Environmental microbe	DMA00208	
<i>Mitsuokella multacida</i> (52226)	Human microbiome	DMA00209	
<i>Mobiluncus</i> (2050)	Human microbiome, Infectious diseases	DMA00413	

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<i>Mobiluncus curtisii</i> (2051)	Human microbiome	DMA00210	YES
<i>Mobiluncus mulieris</i> (2052)	Human microbiome, Infectious diseases	DMA00211	YES
<i>Mobiluncus</i> spp. (47293)	Infectious diseases, Human pathogens	DMA00723	YES
<i>Mogibacterium timidum</i> (35519)	Human microbiome	DMA00212	
<i>Moraxella catarrhalis</i> (480)	Human pathogens	DMA00213	YES
<i>Moraxella lacunata</i> (477)	Human pathogens	DMA00214	YES
<i>Morganella morganii</i> (582)	Human microbiome, Infectious diseases	DMA00215	
<i>Mycobacterium avium</i> (1764)	Human pathogens, Infectious diseases, MDR	DMA00217	
<i>Mycobacterium intracellulare</i> (1767)	Human pathogens	DMA00219	
<i>Mycobacterium kansasii</i> (1786)	Human pathogens	DMA00220	
<i>Mycobacterium marinum</i> (1781)	Animal disease	DMA00779	YES
<i>Mycobacterium tuberculosis</i> (1773)	Human pathogens	DMA00221	YES
<i>Mycobacterium tuberculosis variant africanum</i> (33894)	Human pathogens	DMA00216	
<i>Mycobacteroides abscessus</i> (36809)	Infectious diseases, MDR	DMA00414	
<i>Mycobacteroides chelonae</i> (1774)	Wastewater, Infectious diseases	DMA00218	
<i>Mycolicibacterium fortuitum</i> (1766)	Infectious diseases	DMA00415	
<i>Mycoplasma genitalium</i> (2097)	Human pathogen, Infectious diseases	DMA00222	YES
<i>Mycoplasma hominis</i> (16S, 2098)	Human pathogen, Infectious diseases	DMA00223	YES
<i>Mycoplasma hominis</i> (YidC, 2098)	Human pathogen, Infectious diseases	DMA00410	YES
<i>Mycoplasma orale</i> (2121)	Human microbiome	DMA00224	
<i>Mycoplasma pneumoniae</i> (2104)	Infectious diseases	DMA00225	YES
<i>Neisseria bacilliformis</i> (267212)	Human pathogens, Infectious diseases	DMA00226	
<i>Neisseria elongata</i> (495)	Infectious diseases	DMA00227	
<i>Neisseria flavescens</i> (484)	Human microbiome, Infectious diseases	DMA00228	
<i>Neisseria gonorrhoeae</i> (485)	Infectious diseases	DMA00229	YES
<i>Neisseria lactamica</i> (486)	Human microbiome	DMA00230	
<i>Neisseria meningitidis</i> (487)	Human pathogens	DMA00231	YES
<i>Neisseria mucosa</i> (488)	Human microbiome, Infectious diseases	DMA00232	YES
<i>Neisseria sicca</i> (490)	Human microbiome, Infectious diseases	DMA00233	
<i>Neisseria subflava</i> (28449)	Human microbiome	DMA00234	
<i>Neorickettsia risticii</i> (950)	Animal pathogens	DMA00235	
<i>Nocardia asteroides</i> (1824)	Human pathogens, Infectious diseases	DMA00236	
<i>Nocardia farcinica</i> (37329)	Infectious diseases	DMA00237	
<i>Nocardioides</i> sp. (NS/27, 223904)	Environmental microbes	DMA00238	
<i>Novosphingobium</i> sp. (K39, 233904)	Environmental microbes	DMA00239	
<i>Oenococcus</i> spp. (46254)	Food production	DMA00767	YES



Target species (target region, NCBI taxonomy)	Application field	GeneGlobe catalog number	Wet-lab tested
<i>Oribacterium sinus</i> (237576)	Infectious diseases	DMA00241	
<i>Oxalobacter formigenes</i> (847)	Human microbiome, probiotics	DMA00490	
<i>Paenibacillus larvae</i> (1464)	Animal pathogens	DMA00242	
<i>Paenibacillus macerans</i> (44252)	Environmental microbes	DMA00243	
<i>Paenibacillus thiaminolyticus</i> (49283)	Infectious diseases	DMA00244	
<i>Paenoclostridium sordellii</i> (1505)	Infectious diseases	DMA00111	
<i>Pan Bacteria 1</i> (2)	Bacterial load	DMA00347	YES
<i>Pan Bacteria 3</i> (2)	Bacterial load	DMA00348	YES
<i>Pantoea ananatis</i> (553)	Infectious diseases	DMA00245	
<i>Pantoea stewartii</i> (66269)	Food production	DMA00801	YES
<i>Pasteurella multocida</i> (747)	Human pathogens	DMA00251	
<i>Pectinatus cerevisiiphilus</i> (86956)	Food production	DMA00387	
<i>Pectinatus frisingensis</i> (865)	Food production	DMA00388	
<i>Pediococcus acidilactici</i> (1254)	Probiotics, Human microbiome	DMA00252	
<i>Pediococcus damnosus</i> (51663)	Food production	DMA00389	
<i>Pediococcus pentosaceus</i> (1255)	Food production, Fermentation and degradation	DMA00253	
<i>Peptoniphilus asaccharolyticus</i> (1258)	Human microbiome, Infectious diseases	DMA00254	
<i>Peptostreptococcus anaerobius</i> (1261)	Human microbiome, Infectious diseases	DMA00255	
<i>Peptostreptococcus</i> sp. (1257)	Human microbiome, Infectious diseases, MDR	DMA00412	
<i>Peptostreptococcus stomatis</i> (341694)	Human microbiome, Infectious diseases	DMA00256	
<i>Phocaeicola coprocola</i> (310298)	Human microbiome	DMA00045	
<i>Phocaeicola coprophilus</i> (387090)	Human microbiome	DMA00046	
<i>Phocaeicola dorei</i> (357276)	Human microbiome	DMA00047	
<i>Phocaeicola vulgatus</i> (821)	Gastrointestinal infections	DMA00059	YES
<i>Plesiomonas shigelloides</i> (703)	Human pathogens, Infectious diseases	DMA00257	YES
<i>Porphyromonas asaccharolytica</i> (28123)	Infectious diseases	DMA00259	
<i>Porphyromonas endodontalis</i> (28124)	Infectious diseases	DMA00260	
<i>Porphyromonas gingivalis</i> (837)	Infectious diseases	DMA00261	
<i>Prevotella</i> (838)	Human microbiome, Infectious diseases	DMA00411	
<i>Prevotella bivia</i> (28125)	Human pathogens, Infectious diseases	DMA00262	YES
<i>Prevotella buccalis</i> (28127)	Human microbiome, Infectious diseases	DMA00263	
<i>Prevotella copri</i> (165179)	Human microbiome, Infectious diseases	DMA00264	
<i>Prevotella denticola</i> (28129)	Human microbiome, Fermentation and degradation	DMA00265	
<i>Prevotella disiens</i> (28130)	Human microbiome, Infectious diseases	DMA00266	YES
<i>Prevotella intermedia</i> (28131)	Human microbiome, Infectious diseases	DMA00267	
<i>Prevotella loescheii</i> (840)	Human microbiome, Infectious diseases	DMA00268	

Target species (target region, NCBI taxonomy)	Application field	GeneGlobe catalog number	Wet-lab tested
<i>Prevotella melaninogenica</i> (28132)	Human microbiome, Human pathogens	DMA00269	
<i>Prevotella nigrescens</i> (28133)	Infectious diseases	DMA00270	
<i>Prevotella oralis</i> (28134)	Human microbiome, Infectious diseases	DMA00271	YES
<i>Prevotella oris</i> (28135)	Human microbiome, Infectious diseases	DMA00272	
<i>Prevotella veroralis</i> (28137)	Human microbiome	DMA00274	
<i>Proteus mirabilis</i> (ureR, 584)	Human pathogens	DMA00745	YES
<i>Proteus mirabilis</i> (intergenic region, 584)	Infectious diseases	DMA00474	
<i>Proteus vulgaris</i> (585)	Human pathogens	DMA00743	YES
<i>Providencia stuartii</i> (588)	Human pathogens	DMA00744	YES
<i>Pseudanabaena</i> spp. (MIB, 1152)	Wastewater	DMA00763	YES
<i>Pseudoleptotrichia goodfellowii</i> (157692)	Human microbiome	DMA00197	
<i>Pseudomonas aeruginosa</i> (287)	Infectious diseases, MDR	DMA00278	YES
<i>Pseudomonas putida</i> (303)	Biocontrol, Fermentation and degradation	DMA00279	YES
<i>Pseudomonas straminea</i> (47882)	Environmental microbes	DMA00280	YES
<i>Pseudoramibacter alactolyticus</i> (113287)	Infectious diseases	DMA00281	
<i>Rahnella aquatilis</i> (34038)	Infectious diseases, Sepsis	DMA00282	
<i>Ralstonia pickettii</i> (329)	Environmental microbes	DMA00283	
<i>Rhodococcus hoagii</i> (43767)	Human pathogens, Infectious diseases	DMA00284	
<i>Rickettsia rickettsii</i>	Human pathogens	DMA00802	YES
<i>Rothia dentocariosa</i> (2047)	Human microbiome	DMA00285	
<i>Rothia kristinae</i> (37923)	Human microbiome	DMA00175	
<i>Rothia mucilaginosa</i> (43675)	Human microbiome, Infectious diseases	DMA00286	
<i>Ruminococcus bromii</i> (40518)	Human microbiome	DMA00287	
<i>Ruminococcus gnavus</i> (33038)	Human microbiome	DMA00288	
<i>Ruminococcus torques</i> (33039)	Human microbiome	DMA00290	
<i>Saccharomyces cerevisiae</i> (COG6, 4932)	Food production	DMA00758	YES
<i>Saccharomyces</i> Pan (A/AB) (4930)	Food production	DMA00761	YES
<i>Saccharomyces</i> Pan (B/AB) (4930)	Food production	DMA00762	YES
<i>Salmonella</i> (590)	Human pathogens, Wastewater, Food production	DMA00704	YES
<i>Salmonella enterica</i> (28901)	Human pathogens	DMA00291	YES
<i>Salmonella enterica</i> serovar <i>Enteritidis</i> (149539)	Food production	DMA00848	YES
<i>Salmonella enterica</i> serovar <i>Typhimurium</i> (90371)	Food production	DMA00846	YES
<i>Salmonella enterica</i> subsp. <i>enterica</i> (59201)	Human pathogens, Infectious diseases	DMA00360	YES
<i>Scardovia wiggisiae</i> (230143)	Infectious diseases	DMA00399	
<i>Schaalia odontolytica</i> (1660)	Infectious diseases	DMA00014	YES
<i>Schaalia radingae</i> (131110)	Infectious diseases	DMA00015	



Target species (target region, NCBI taxonomy)	Application field	GeneGlobe catalog number	Wet-lab tested
<i>Selenomonas infelix</i> (135082)	Human pathogens, Human microbiome	DMA00292	
<i>Selenomonas noxia</i> (135083)	Human microbiome, Infectious diseases	DMA00293	
<i>Selenomonas sputigena</i> (69823)	Human microbiome, Infectious diseases	DMA00294	
<i>Serratia marcescens</i> (615)	Human pathogens	DMA00749	YES
<i>Shigella dysenteriae</i> (622)	Human pathogens	DMA00295	YES
<i>Shigella sonnei</i> (624)	Gastrointestinal infections	DMA00361	YES
<i>Shuttleworthia satellites</i> (177972)	Human microbiome	DMA00296	
<i>Sneathia sanguinegens</i> (40543)	Human microbiome	DMA00297	
<i>Sneathia vaginalis</i> (187101)	Human microbiome	DMA00195	
<i>Solobacterium moorei</i> (102148)	Infectious diseases, Sepsis	DMA00298	
<i>Sphingomonas bisphenolicum</i> (296544)	Fermentation and degradation	DMA00300	
<i>Sphingomonas paucimobilis</i> (13689)	Human pathogens, Omit	DMA00299	
<i>Sporobacter termitidis</i> (44749)	Environmental microbes, Fermentation and degradation	DMA00301	
<i>Staphylococcus aureus</i> (1280)	Human pathogens, Omit	DMA00302	YES
<i>Staphylococcus caprae</i> (29380)	Human microbiome, Omit	DMA00303	YES
<i>Staphylococcus epidermidis</i> (1282)	Sepsis, MDR	DMA00304	YES
<i>Staphylococcus lugdunensis</i> (28035)	Human microbiome, Infectious diseases	DMA00416	
<i>Staphylococcus saprophyticus</i> (29385)	Infectious diseases	DMA00305	
<i>Stenotrophomonas maltophilia</i> (40324)	Infectious diseases, MDR	DMA00306	
<i>Streptobacillus moniliformis</i> (34105)	Infectious diseases	DMA00307	
<i>Streptococcus agalactiae</i> (1311)	Infectious diseases	DMA00308	
<i>Streptococcus anginosus</i> (1328)	Human microbiome, Infectious diseases	DMA00309	
<i>Streptococcus australis</i> (113107)	Human microbiome	DMA00310	
<i>Streptococcus constellatus</i> (76860)	Human microbiome	DMA00311	
<i>Streptococcus downei</i> (1317)	Human microbiome, Infectious diseases	DMA00312	
<i>Streptococcus gordonii</i> (1302)	Human microbiome	DMA00313	
<i>Streptococcus infantis</i> (68892)	Human pathogens, Human microbiome	DMA00314	
<i>Streptococcus mitis</i> (28037)	Human microbiome, Omit	DMA00315	YES
<i>Streptococcus mutans</i> (1309)	Human microbiome	DMA00316	
<i>Streptococcus oralis</i> (1303)	Human microbiome	DMA00317	YES
<i>Streptococcus pneumoniae</i> (1313)	Human pathogens, Infectious diseases	DMA00318	YES
<i>Streptococcus pyogenes</i> (1314)	Human pathogens, Infectious diseases	DMA00319	YES
<i>Streptococcus salivarius</i> (1304)	Human microbiome, Probiotics	DMA00320	YES
<i>Streptococcus sanguinis</i> (1305)	Human microbiome, Infectious diseases	DMA00321	
<i>Streptococcus sinensis</i> (176090)	Infectious diseases	DMA00322	
<i>Streptococcus sobrinus</i> (1310)	Infectious diseases	DMA00400	
<i>Streptococcus suis</i> (1307)	Food production	DMA00323	

Target species (target region, NCBI taxonomy)	Application field	GeneGlobe catalog number	Wet-lab tested
<i>Streptomyces bikiniensis</i> (1896)	Environmental microbes, Biocontrol	DMA00324	
<i>Streptomyces griseus</i> (1911)	Environmental microbes, Biocontrol	DMA00325	
<i>Streptomyces somaliensis</i> (78355)	Human pathogens	DMA00326	
<i>Subdoligranulum variabile</i> (214851)	Human microbiome	DMA00327	
<i>Tannerella forsythia</i> (28112)	Infectious diseases	DMA00328	
<i>Treponema denticola</i> (158)	Human pathogens, Omit	DMA00329	
<i>Tannerella forsythia</i> (28112)	Infectious diseases	DMA00328	
<i>Thermoactinomyces</i> (2023)	Environmental microbes, Infectious diseases, Food production	DMA00370	
<i>Treponema denticola</i> (158)	Human pathogens, Infectious diseases	DMA00329	
<i>Treponema pallidum</i> (tpp47, 160)	Human pathogens, Omit	DMA00729	YES
<i>Treponema pallidum</i> (16S, 160)	Human pathogens	DMA00330	
<i>Treponema socranskii</i> (53419)	Infectious diseases	DMA00331	
<i>Trichomonas vaginalis</i> (5722)	Human pathogens, Infectious diseases	DMA00332	YES
<i>Tropheryma whippelii</i> (2039)	Human pathogens	DMA00854	YES
<i>Turicibacter sanguinis</i> (154288)	Human pathogens	DMA00333	
<i>Tyzzera nexilis</i> (29361)	Human microbiome	DMA00108	
<i>Ureaplasma parvum</i> (134821)	Human microbiome	DMA00334	
<i>Ureaplasma urealyticum</i> (2130)	Human pathogens	DMA00335	
<i>Varibaculum cambriense</i> (184870)	Infectious diseases	DMA00336	
<i>Veillonella dispar</i> (39778)	Human microbiome, Omit	DMA00337	
<i>Veillonella parvula</i> (29466)	Human microbiome, Omit	DMA00338	
<i>Vibrio alginolyticus</i> (663)	Human pathogens, Omit	DMA00339	
<i>Vibrio cholerae</i> (666)	Human pathogens, Omit	DMA00340	YES
<i>Vibrio parahaemolyticus</i> (670)	Environmental microbes, Infectious diseases	DMA00341	YES
<i>Vibrio vulnificus</i> (672)	Environmental microbes	DMA00342	YES
<i>Weissella confusa</i> (1583)	Environmental microbes, Biocontrol, Fermentation and degradation	DMA00343	
<i>Xylella fastidiosa</i> (2371)	Plant pathogens	DMA00719	YES
<i>Yersinia enterocolitica</i> (630)	Human pathogens	DMA00344	YES
<i>Yersinia pestis</i> (632)	Human pathogens	DMA00345	YES

Table 2. Fungal identification targets/assays

Target species (target region, NCBI taxonomy)	GeneGlobe catalog number	Wet-lab tested
<i>Alternaria alternata</i> (5599)	DMA00481	
<i>Aspergillus</i> (5052), <i>Candida</i> (1535326)	DMA00346	YES



Target species (target region, NCBI taxonomy)	GeneGlobe catalog number	Wet-lab tested
<i>Aspergillus</i> (5052), <i>Penicillium</i> (5073)	DMA00369	YES
<i>Aspergillus flavus</i> (1) (ITS1-5.8S)	DMA00035	
<i>Aspergillus flavus</i> (2) (ITS1-5.8S)	DMA00398	YES
<i>Aspergillus fumigatus</i> (746128)	DMA00036	YES
<i>Aspergillus niger</i> (5061)	DMA00365	YES
<i>Aspergillus terreus</i> 1 (ITS, 33178)	DMA00489	YES
<i>Aspergillus terreus</i> 2 (cadA, 33178)	DMA00477	YES
<i>Aspergillus versicolor</i> (46472)	DMA00482	YES
<i>Brettanomyces bruxellensis</i> (5007)	DMA00718	YES
<i>Candida albicans</i> (5476)	DMA00089	YES
<i>Candida auris</i> (498019)	DMA00793	YES
<i>Candida dubliniensis</i> (42374)	DMA00407	YES
<i>Candida glabrata</i> / <i>Nakaseomyces glabratus</i> (28S, 5478)	DMA00090	YES
<i>Candida parapsilosis</i> (5480)	DMA00092	YES
<i>Candida tropicalis</i> (5482)	DMA00093	YES
<i>Cryptococcus gattii</i> (515176)	DMA00781	YES
<i>Curvularia lunata</i> (5503)	DMA00483	
<i>Enterocytozoon hepatopenaei</i> (646526)	DMA00451	
<i>Epidermophyton floccosum</i> (34391)	DMA00478	
Fungi (4751)	DMA00391	YES
Fungi (4751)	DMA00393	YES
Fungi (4751)	DMA00394	YES
<i>Fusarium acuminatum</i> (5515)	DMA00824	YES
<i>Fusarium avenaceum</i> (40199)	DMA00822	YES
<i>Fusarium culmorum</i> (5932)	DMA00826	YES
<i>Fusarium graminearum</i> (5518)	DMA00820	YES
<i>Fusarium redolens</i> (48865)	DMA00825	YES
<i>Fusarium solani</i> (169388)	DMA00479	
<i>Fusarium</i> spp (29916)	DMA00834	YES
<i>Fusarium sporotrichioides</i> (5514)	DMA00823	YES
<i>Lachancea thermotolerans</i> (381046)	DMA00799	YES
<i>Metschnikowia pulcherrima</i> (27326)	DMA00800	YES
<i>Microsporum audouinii</i> (34393)	DMA00460	
<i>Microsporum canis</i> (63405)	DMA00461	
<i>Mucor hiemalis</i> (64493)	DMA00368	YES
<i>Nakaseomyces glabratus</i> (28S, 37769)	DMA00780	YES
<i>Neoscytalidium dimidiatum</i> (108428)	DMA00480	

Target species (target region, NCBI taxonomy)	GeneGlobe catalog number	Wet-lab tested
<i>Pichia kudriavzevii</i> (4909)	DMA00091	YES
<i>Pneumocystis jirovecii</i> (42068)	DMA00258	
<i>Sarocladium strictum</i> (5046)	DMA00476	YES
<i>Scopulariopsis brevicaulis</i> (561152)	DMA00484	
<i>Starmerella bacillaris</i> (1247836)	DMA00821	YES
<i>Starmerella stellata</i> (45594)	DMA00798	YES
<i>Stenocarpella macrospora</i> (371126)	DMA00819	YES
<i>Stenocarpella maydis</i> (238245)	DMA00803	YES
<i>Torulaspora delbrueckii</i> (4950)	DMA00805	YES
<i>Trichophyton benhamiae</i> (63400)	DMA00804	YES
<i>Trichophyton indotinae</i> (2739387)	DMA00812	YES
<i>Trichophyton interdigitale</i> (101480)	DMA00463	
<i>Trichophyton mentagrophytes</i> (523103)	DMA00464	
<i>Trichophyton rubrum</i> (5551)	DMA00450	
<i>Trichophyton tonsurans</i> (34387)	DMA00462	
<i>Trichophyton verrucosum</i> (63417)	DMA00465	

Table 3. Parasite identification targets/assays

Target species (target region, NCBI taxonomy)	Taxonomy	GeneGlobe catalog number	Wet-lab tested
<i>Ancylostoma duodenale</i> (51022)	Metazoa	DMA00466	
<i>Anisakis pegreffii</i> (mt cox2, 303229)	Nematoda	DMA00828	YES
<i>Anisakis simplex</i> (mt cox2, 6269)	Nematoda	DMA00829	YES
<i>Ascaris lumbricoides</i> (6252)	Metazoa	DMA00467	
<i>Blastocystis hominis</i> (12968)	Stramenopiles	DMA00354	
<i>Cryptosporidium hominis</i> (237895), <i>Cryptosporidium parvum</i> (5807), <i>Cryptosporidium cuniculus</i> (564424)	Alveolata	DMA00356	YES
<i>Cyclospora cayetanensis</i> (ITS, 88456)	Apicomplexa	DMA00774	YES
<i>Dientamoeba fragilis</i> (43352)	Excavata	DMA00357	
<i>Endolimax nana</i> (110788)	Amoebozoa	DMA00468	
<i>Entamoeba histolytica</i> (5759)	Amoebozoa	DMA00358	YES
<i>Entamoeba muris</i> (110766)	Amoebozoa	DMA00469	YES
<i>Enterobius vermicularis</i> (51028)	Metazoa	DMA00366	
<i>Giardia intestinalis</i> (5741)	Metamonada	DMA00359	YES
<i>Hysterothylacium aduncum</i> (mt cox 2, 118886)	Nematoda	DMA00833	YES
<i>Ichthyophthirius multifiliis</i> (18S, 5932)	Ciliophora	DMA00827	YES



Target species (target region, NCBI taxonomy)	Taxonomy	GeneGlobe catalog number	Wet-lab tested
<i>Leishmania</i> spp._Pan (5658)	Euglenozoa	DMA00771	YES
<i>Leishmania brasiliensis</i> (5660)	Euglenozoa	DMA00772	YES
<i>Naegleria fowleri</i> ITS (5763)	Percolozoa	DMA00776	YES
<i>Necator americanus</i> (51031)	Metazoa	DMA00470	
<i>Pentatrichomonas hominis</i> (5728)	Metamonada	DMA00471	
<i>Plasmodium falciparum</i> (5833)	Apicomplexa	DMA00735	YES
<i>Plasmodium knowlesi</i> (5850)	Apicomplexa	DMA00738	YES
<i>Plasmodium malariae</i> (5858)	Apicomplexa	DMA00773	YES
<i>Plasmodium ovale</i> (36330)	Apicomplexa	DMA00736	YES
<i>Plasmodium vivax</i> (5855)	Apicomplexa	DMA00737	YES
<i>Pseudoterranova bulbosa</i> (mt cox2, 172674)	Nematoda	DMA00831	YES
<i>Pseudoterranova decipiens</i> (mt cox2, 6271)	Nematoda	DMA00830	YES
<i>Pseudoterranova krabbei</i>	Nematoda	DMA00832	YES
<i>Plasmodium vivax</i> (5855)	Apicomplexa	DMA00737	YES
<i>Rodentolepis nana</i> (102285)	Metazoa	DMA00362	
<i>Trichuris trichiura</i> (36087)	Metazoa	DMA00364	
<i>Trypanosoma brucei</i> (5691)	Euglenozoa	DMA00752	YES
<i>Trypanosoma cruzi</i> (5693)	Euglenozoa	DMA00791	YES

Table 4. Viral identification targets/assays

Target species (target region, NCBI taxonomy)	GeneGlobe catalog number	Wet-lab tested
Adenovirus (10535)	DMA00371	YES
African swine fever virus (10497)	DMA00716	YES
African swine fever virus (10497)	DMA00716	YES
African swine fever virus (10497)	DMA00716	YES
Baculovirus penaei (384357)	DMA00424	
Bocaparvovirus (1507401)	DMA00456	
Bourbon Virus (1618189)	DMA00789	YES
Cauliflower mosaic virus (10641)	DMA00814	YES
Chikungunya virus (37124)	DMA00748	YES
Cucumber mosaic virus (12305)	DMA00865	YES
Decapod penstylidensovirus 1 (1513224)	DMA00420	
Dengue virus (pan) (12637)	DMA00788	YES
Dengue virus Serotype 1	DMA00731	YES
Dengue virus Serotype 2	DMA00732	YES

**Target species (target region,
NCBI taxonomy)**
GeneGlobe catalog number
Wet-lab tested

Dengue virus Serotype 3	DMA00733	YES
Dengue virus Serotype 4	DMA00734	YES
Eastern equine encephalitis virus (gp3, 11021)	DMA00754	YES
Ebola virus (1570291)	DMA00815	YES
Enterovirus (12059)	DMA00487	YES
Gill associated virus genotype 2 (557593)	DMA00418	
Heartland Virus (1216928)	DMA00790	YES
Hepatitis A (208726)	DMA00807	YES
Hepatitis B (10407)	DMA00808	YES
Hepatitis C (3052230)	DMA00809	YES
Hepatitis D (12721)	DMA00810	YES
Hepatitis E virus	DMA00750	YES
Hepatopancreatic parvovirus (139034)	DMA00419	
Herpes simplex virus type 1 (glycoprotein G, 10298)	DMA00724	YES
Herpes simplex virus type 2 (glycoprotein D, 10310)	DMA00725	YES
Human alphaherpesvirus (glycoprotein H, 10335)	DMA00397	
Human astrovirus 1 (12456)	DMA00485	
Human betaherpesvirus 5 (10359)	DMA00835	YES
Human betaherpesvirus 6A (32603)	DMA00837	YES
Human betaherpesvirus 6B (32604)	DMA00838	YES
Human betaherpesvirus 7 (10372)	DMA00839	YES
Human gammaherpesvirus 8 (37296)	DMA00836	YES
Human coronavirus (229E, 11137)	DMA00430	
Human coronavirus (HKU1, 290028)	DMA00452	
Human coronavirus (NL63, 277944)	DMA00453	
Human coronavirus (OC43, 31631)	DMA00431	
Human coronavirus SARS-CoV-2/ Severe acute respiratory syndrome coronavirus 2 (N1, 2697049)	DMA00710	
Human coronavirus SARS-CoV-2/ Severe acute respiratory syndrome coronavirus 2 (N2, 2697049)	DMA00711	
Human herpesvirus 1 (glycoprotein I, 10298)	DMA00395	
Human herpesvirus 1 and 2 (10298 and 10310)	DMA00726	YES
Human herpesvirus 2 (glycoprotein G, 10310)	DMA00396	
Human immunodeficiency virus (12721)	DMA00811	YES
Human metapneumovirus (162145)	DMA00372	YES
Human orthorubulavirus 2 (2560525)	DMA00376	YES
Human orthorubulavirus 4 (2560526)	DMA00454	YES
Human parainfluenza 1 virus (12730)	DMA00375	YES



Target species (target region, NCBI taxonomy)	GeneGlobe catalog number	Wet-lab tested
Human parainfluenza 3 (11216)	DMA00377	YES
Human parvovirus B19 (10798)	DMA00862	YES
Human respiratory syncytial virus 1 (matrix protein, 12814)	DMA00379	YES
Human respiratory syncytial virus 2 (matrix protein M, 12814)	DMA00715	
Human rhinovirus sp. (169066)	DMA00378	
Human rotavirus A (10941)	DMA00488	YES
Human T-cell leukemia virus type I (11908)	DMA00867	YES
Human T-cell leukemia virus 2 (11909)	DMA00868	YES
Human T-cell leukemia virus type I (2) (11908)	DMA00869	YES
Human T-cell leukemia virus 2 (2) (11909)	DMA00870	YES
Influenza A (1) (matrix protein M1, 11320)	DMA00712	
Influenza A (3) (matrix protein M1, 11320)	DMA00713	
Influenza A Pan (11320)	DMA00855	YES
Influenza A virus (11320)	DMA00373	YES
Influenza A virus (1/2) (11320)	DMA00858	YES
Influenza A virus (2/2) (11320)	DMA00859	YES
Influenza B (11520)	DMA00714	
Influenza B Pan (11520)	DMA00856	YES
Influenza B virus (11520)	DMA00374	YES
Japanese encephalitis virus (NS1, 11072)	DMA00757	YES
Laem Singh virus (691504)	DMA00422	
Mayaro virus (59301)	DMA00457	
Measles morbillivirus	DMA00786	YES
Middle East respiratory syndrome-related coronavirus (1335626)	DMA00844	YES
Monkeypox virus (10244)	DMA00747	YES
Monkeypox virus CB, Clade Ia	DMA00770	YES
Monkeypox virus CB, Clade Ia+Ib	DMA00852	YES
Monkeypox virus CB, Clade Ib	DMA00853	YES
Monkeypox virus, Clade II (10244)	DMA00746	YES
Mourilyan virus (325212)	DMA00423	
Mumps orthorubulavirus	DMA00787	YES
Norovirus GI (122928)	DMA00472	YES
Norovirus GII (122929)	DMA00473	YES
Orthomareburgvirus marburgense Pan (1/2) (3052505)	DMA00874	YES
Orthomareburgvirus marburgense Pan (2/2) (3052505)	DMA00875	YES
Orthospovirus tomatomaculae (3052585)	DMA00863	YES
Oropouche virus (118655)	DMA00459	

Target species (target region, NCBI taxonomy)	GeneGlobe catalog number	Wet-lab tested
Penaeid shrimp infectious myonecrosis virus (282786)	DMA00421	
Pepper mild mottle virus PMMoV (12239)	DMA00722	YES
Potato leafroll virus (12045)	DMA00871	YES
Potato virus S (12169)	DMA00872	YES
Potato virus X (12183)	DMA00873	YES
Potato virus Y (12216)	DMA00866	YES
Rabies lyssavirus (11292)	DMA00816	YES
Rift Valley fever virus (11588)	DMA00817	YES
Saint Louis encephalitis virus (11080)	DMA00818	YES
Sapovirus (1906670)	DMA00486	
Sapovirus GI (515176)	DMA00782	YES
Sapovirus GII (515177)	DMA00783	YES
Sapovirus GIV (515179)	DMA00784	YES
Sapovirus GVI (1471435)	DMA00785	YES
Severe acute respiratory syndrome-related coronavirus (694009)	DMA00845	YES
Spawner-isolated mortality virus (207679)	DMA00425	
Taura syndrome virus (142102)	DMA00426	
Tobacco Mosaic virus (12242)	DMA00860	YES
Tomato yellow leaf curl virus	DMA00864	YES
Varicella-zoster virus/Human alphaherpesvirus 3 (ORF38, 10335)	DMA00728	YES
Venezuelan equine encephalomyelitis virus (11036)	DMA00756	YES
West Nile virus (11082)	DMA00455	
Western equine encephalitis virus gp2 (11039)	DMA00755	YES
White spot syndrome virus (342409)	DMA00427	
Yellow fever virus (11089)	DMA00458	
Yellow head virus 1 Replicase polyprotein 1ab (96029)	DMA00428	
Yellow head virus 2 Replicase polyprotein 1ab (96029)	DMA00429	
Zika Virus (64320)	DMA00841	YES

Table 5. Antibiotic resistance gene detection targets/assays

Antibiotic resistance gene	Target info	GeneGlobe catalog number	Wet-lab tested
<i>aac(6)-Ib (1)</i>	Bacterial fluoroquinolone-acetylating aminoglycoside 6'-N-acetyltransferase AAC(6')-Ib-cr5 gene	DMA00492	
<i>aac(6)-Ib (2)</i>	Bacterial aminoglycoside N-acetyltransferase AAC(6')-Ib4 gene	DMA00579	YES
<i>aacC1</i>	Bacterial AAC(3)-I family aminoglycoside 3-N-acetyltransferase gene	DMA00493	



Antibiotic resistance gene	Target info	GeneGlobe catalog number	Wet-lab tested
<i>aacC2</i>	Bacterial aminoglycoside N-acetyltransferase AAC(3)-IId gene	DMA00494	
<i>aacC4</i>	Bacterial aminoglycoside N-acetyltransferase AAC(3)-Iva gene	DMA00495	
<i>aadA1</i>	Bacterial ANT(3'')-Ia family aminoglycoside nucleotidyltransferase AadA1 gene	DMA00496	
<i>aadB</i>	Bacterial aminoglycoside nucleotidyltransferase ANT(2'')-Ia gene	DMA00497	
ACC-1 group	Bacterial cephalosporin-hydrolyzing class C beta-lactamase ACC-1 gene	DMA00531	YES
ACC-3 group	Bacterial AMPC cephalosporinase precursor protein ACC-3 gene	DMA00532	
ACT 5/7 group	Bacterial cephalosporin-hydrolyzing class C beta-lactamase ACT-5 gene	DMA00533	
ACT-1 group	Bacterial cephalosporin-hydrolyzing class C beta-lactamase ACT-1 gene	DMA00534	
<i>ant-Ia</i>	Bacterial aminoglycoside nucleotidyltransferase ANT(2'')-Ia gene	DMA00580	
<i>aph(3)</i>	Bacterial APH(3')-VI family aminoglycoside O-phosphotransferase gene	DMA00581	
<i>aphA6</i>	Bacterial aminoglycoside O-phosphotransferase APH(3')-Via gene	DMA00498	
<i>BES-1</i>	Bacterial class A extended spectrum beta-lactamase BES-1 gene	DMA00500	YES
<i>BIC-1</i>	Bacterial carbapenem-hydrolyzing class A beta-lactamase BIC-1 gene	DMA00501	
<i>BIC-1</i>	Bacterial carbapenem-hydrolyzing class A beta-lactamase BIC-1 gene	DMA00501	
<i>bla CTX-M (CTX-M-2, -M-8 and -M-25)</i>	Bacterial CTX-M family class A beta-lactamase CTX-M gene	DMA00843	YES
<i>blaMOX-1</i>	Bacterial CMY-1/MOX family class C beta-lactamase MOX-1 gene	DMA00541	
<i>blaVIM-1</i> group	Bacterial subclass B1 metallo-beta-lactamase VIM1 gene	DMA00528	YES
<i>blaVIM-13</i>	Bacterial subclass B1 metallo-beta-lactamase VIM-13 gene	DMA00529	
<i>blaVIM-7</i>	Bacterial subclass B1 metallo-beta-lactamase VIM-7 gene	DMA00530	
<i>ccrA</i>	Bacterial subclass B1 metallo-beta-lactamase CfiA gene	DMA00522	
<i>CFE-1</i>	Bacterial CMY-2 family cephalosporin-hydrolyzing class C beta-lactamase CFE-1 gene	DMA00535	YES
<i>CMY-10</i> group	Bacterial CMY-1/MOX family class C beta-lactamase MOX-14 gene	DMA00536	
<i>CTX-M-1</i> group	Bacterial class A extended-spectrum beta-lactamase CTX-M-15 gene	DMA00502	YES
<i>CTX-M-8</i> group	Bacterial extended-spectrum beta-lactamase CTX-M-8 gene	DMA00503	
<i>CTX-M-9</i> group	Bacterial class A extended-spectrum beta-lactamase CTX-M-9 gene	DMA00504	
<i>dfra17</i>	Bacterial trimethoprim-resistant dihydrofolate reductase DfrA17 gene	DMA00582	
<i>dha</i>	Bacterial class C beta-lactamase DHA-1 gene	DMA00537	YES
<i>E. coli gyrA (S83L)</i>	Bacterial DNA topoisomerase (ATP-hydrolyzing) subunit A (gyrA) gene	DMA00591	

Antibiotic resistance gene	Target info	GeneGlobe catalog number	Wet-lab tested
<i>E. coli gyrA</i> (S83W)	"Bacterial DNA topoisomerase (ATP-hydrolyzing) subunit A (gyrA) gen"	DMA00592	
<i>ereB</i>	Bacterial EreB family erythromycin esterase gene	DMA00556	YES
<i>ermA</i>	Bacterial 23S rRNA (adenine(2058)-N(6))-methyltransferase Erm(A) gene	DMA00567	YES
<i>ermB</i>	Bacterial adenine(2058)-N(6))-dimethyltransferase Erm(B) gene	DMA00568	YES
<i>ermC</i>	Bacterial erythromycin resistance protein ErmC gene	DMA00569	
<i>fosA</i> (<i>Klebsiella</i> sp.)	Bacterial FosA family fosfomycin resistance glutathione transferase gene	DMA00583	YES
<i>fosA</i> (<i>Pseudomonas</i> sp.)	Bacterial FosA family fosfomycin resistance glutathione transferase gene	DMA00584	YES
<i>fox</i>	Bacterial cephalosporin-hydrolyzing class C beta-lactamase FOX gene	DMA00538	YES
<i>GES-1</i>	Bacterial class A extended-spectrum beta-lactamase GES-1 gene	DMA00505	YES
<i>IMI & NMC-A</i>	Bacterial carbapenem-hydrolysing class A beta-lactamase NmcA gene	DMA00506	
<i>IMP-1</i> group	Bacterial class B beta-lactamase IMP-1 gene	DMA00523	
<i>IMP-12</i> group	Bacterial class B1 metallo-beta-lactamase IMP-12 gene	DMA00524	
<i>IMP-2</i> group	Bacterial subclass B1 metallo-beta-lactamase IMP-2	DMA00525	
<i>IMP-5</i> group	Bacterial subclass B1 metallo-beta-lactamase IMP-5	DMA00507	
<i>KPC-1</i>	Bacterial carbapenem-hydrolyzing class A beta-lactamase KPC-2 gene	DMA00539	
<i>LAT-1</i>	Bacterial CMY-2 family cephalosporin-hydrolyzing class C beta-lactamase LAT-1 gene	DMA00526	
<i>mcr1</i>	Colistin resistance MCR-1, polymyxin resistance	DMA00796	YES
<i>mecA</i>	Methicillin resistance, MRSA	DMA00795	YES
<i>mecC</i>	Bacterial PBP2a family beta-lactam-resistant peptidoglycan transpeptidase gene MecA	DMA00499	YES
<i>mefA</i>	Bacterial macrolide efflux MFS transporter (mefA) gene	DMA00570	
<i>MIR-1</i>	Bacterial cephalosporin-hydrolyzing class C beta-lactamase MIR-1 gene	DMA00540	
<i>msrA</i>	Bacterial ABC-F type ribosomal protection protein Msr(A) gene	DMA00571	
<i>oprJ</i>	Bacterial multidrug efflux transporter outer membrane subunit OprJ gene	DMA00572	YES
<i>oprM</i>	Bacterial multidrug efflux RND transporter outer membrane channel subunit OprM gene	DMA00573	YES
<i>OqxA</i>	Bacterial multidrug efflux RND transporter periplasmic adaptor subunit OqxA gene	DMA00585	YES
<i>OqxB</i>	Bacterial multidrug efflux RND transporter permease subunit OqxB gene	DMA00586	
<i>OXA-10</i> group	Bacterial oxacillin-hydrolyzing class D beta-lactamase OXA-10 gene	DMA00542	YES
<i>OXA-18</i>	Bacterial class D extended-spectrum beta-lactamase OXA-18 gene	DMA00543	
<i>OXA-2</i> group	Bacterial OXA-2 family class D beta-lactamase OXA-2 gene	DMA00544	
<i>OXA-23</i> group	Bacterial OXA-23 family carbapenem-hydrolyzing class D beta-lactamase OXA-967 gene	DMA00545	YES



Antibiotic resistance gene	Target info	GeneGlobe catalog number	Wet-lab tested
OXA-24 group	Bacterial carbapenem-hydrolyzing class D beta-lactamase OXA-24 gene	DMA00546	
OXA-45	Bacterial oxacillin-hydrolyzing class D extended-spectrum beta-lactamase OXA-45 gene	DMA00547	
OXA-48 group	Bacterial OXA-48 family class D beta-lactamase OXA-1055 gene	DMA00548	YES
OXA-50 group	Bacterial OXA-50 family oxacillin-hydrolyzing class D beta-lactamase OXA-50 gene	DMA00549	
OXA-51 group	Bacterial OXA-51 family carbapenem-hydrolyzing class D beta-lactamase OXA-51 gene	DMA00550	
OXA-54	Bacterial OXA-48 family carbapenem-hydrolyzing class D beta-lactamase OXA-54 gene	DMA00551	
OXA-55	Bacterial carbapenem-hydrolyzing class D beta-lactamase OXA-55 gene	DMA00552	
OXA-58 group	Bacterial carbapenem-hydrolyzing class D beta-lactamase OXA-58 gene	DMA00553	YES
OXA-60	Bacterial carbapenem-hydrolyzing class D beta-lactamase OXA-60 gene	DMA00554	
OXA-62	Bacterial carbapenem-hydrolyzing class D beta-lactamase OXA-62 gene	DMA00555	
Per-1 group	Bacterial class A extended-spectrum beta-lactamase PER-1	DMA00508	YES
Per-2 group	Bacterial class A extended-spectrum beta-lactamase PER-14 gene	DMA00509	
QepA	Bacterial fluoroquinolone efflux MFS transporter QepA1 gene	DMA00557	YES
QnrA	Bacterial quinolone resistance pentapeptide repeat protein QnrA1 gene	DMA00558	YES
QnrB	Plasmid-mediated quinolone resistance-conferring gene	DMA00797	YES
QnrB-1 group	Bacterial quinolone resistance pentapeptide repeat protein QnrB1 gene	DMA00559	YES
QnrB-31 group	Bacterial quinolone resistance pentapeptide repeat protein QnrB31 gene	DMA00560	
QnrB-4 group	Bacterial quinolone resistance pentapeptide repeat protein QnrB4 gene	DMA00561	
QnrB-5 group	Bacterial quinolone resistance pentapeptide repeat protein QnrB5 gene	DMA00562	
QnrB-8 group	Bacterial quinolone resistance pentapeptide repeat protein QnrB8 gene	DMA00563	
QnrC	Bacterial QnrC family quinolone resistance pentapeptide repeat protein gene	DMA00564	
QnrD	Bacterial quinolone resistance pentapeptide repeat protein QnrD1 gene	DMA00565	
QnrS	Bacterial quinolone resistance pentapeptide repeat protein QnrS1 gene	DMA00566	YES
SFC-1	Bacterial carbapenem-hydrolyzing class A beta-lactamase SFC-1 gene	DMA00510	
SFO-1	Bacterial class A extended-spectrum beta-lactamase SFO-1 gene	DMA00511	
SHV (156D)	Bacterial SHV family beta-lactamase gene	DMA00513	
SHV (156G)	Bacterial SHV family beta-lactamase gene	DMA00514	
SHV (238G240E)	Bacterial class A broad-spectrum beta-lactamase SHV-11 gene	DMA00515	

Antibiotic resistance gene	Target info	GeneGlobe catalog number	Wet-lab tested
SHV (238G240K)	Bacterial class A broad-spectrum beta-lactamase SHV-11 gene	DMA00516	
SHV (238S240E)	Bacterial class A broad-spectrum beta-lactamase SHV-11 gene	DMA00517	
SHV (238S240K)	Bacterial class A broad-spectrum beta-lactamase SHV-11 gene	DMA00518	
SHV-1	Bacterial class A broad-spectrum beta-lactamase SHV-1 gene	DMA00512	
SME-1	Bacterial SME family beta-lactamase gene	DMA00519	
<i>sul1</i>	Bacterial quaternary ammonium compound efflux SMR transporter QacE delta 1 gene	DMA00587	YES
<i>sul2</i>	Bacterial sulfonamide-resistant dihydropteroate synthase Sul2 gene	DMA00588	
TEM	Bacterial class A broad-spectrum beta-lactamase TEM-1 gene	DMA00589	YES
<i>tetA</i>	Bacterial tetracycline efflux MFS transporter Tet(A) gene	DMA00574	YES
<i>tetA</i> (C)	Tetracyclin efflux pump	DMA00769	YES
<i>tetB</i>	Bacterial tetracycline efflux MFS transporter Tet(B) gene	DMA00575	YES
<i>tetK</i>	<i>tetK</i> efflux pump, tetracycline resistance	DMA00751	YES
<i>tetM</i>	Bacterial tetracycline resistance ribosomal protection protein Tet(M) gene	DMA00590	
<i>tetS</i>	Bacterial tetracycline resistance ribosomal protection protein Tet(S) gene	DMA00593	
<i>tetW</i>	Bacterial tetracycline resistance ribosomal protection protein Tet(W) gene	DMA00842	YES
TLA-1	Bacterial class A extended-spectrum beta-lactamase TLA-1 gene	DMA00520	
<i>vanA</i>	Bacterial D-alanine--(R)-lactate ligase VanA gene	DMA00594	
<i>vanA</i> (VRE)	Bacterial D-alanine--(R)-lactate ligase VanA gene	DMA00578	
<i>vanB</i>	Bacterial D-alanine--(R)-lactate ligase VanB gene	DMA00576	YES
<i>vanC</i>	Bacterial D-alanine--D-serine ligase VanC1 gene	DMA00577	
VEB	Bacterial class A beta-lactamase VEB gene	DMA00521	

Table 6. Virulence factor gene detection targets/assays

Virulence gene	Target info	GeneGlobe catalog number	Wet-lab tested
<i>aatA</i>	Bacterial outer membrane protein (<i>aatA</i>) gene	DMA00595	
<i>ace</i> (<i>E. faecalis</i>)	Bacterial collagen-binding MSCRAMM adhesin Ace gene	DMA00596	YES
<i>ace</i> (<i>V. cholerae</i>)	Bacterial accessory cholera enterotoxin gene	DMA00597	YES
<i>ail</i>	Bacterial attachment invasion locus protein gene	DMA00598	
<i>aslA</i>	Bacterial arylsulfatase <i>aslA</i> gene	DMA00599	
<i>badA</i>	Bacterial trimeric autotransporter adhesin BapA gene	DMA00600	
<i>bepB</i>	Bacterial T4SS effector adenyltransferase BepC gene	DMA00601	
<i>bsh</i>	Bacterial linear amide C-N hydrolase gene	DMA00602	



Virulence gene	Target info	GeneGlobe catalog number	Wet-lab tested
<i>chuS</i>	Bacterial hematin-forming heme oxygenase ChuS gene	DMA00603	
<i>cloSI</i>	Bacterial alpha-clostripain gene	DMA00604	
<i>clpB</i>	Bacterial type VI secretion system ATPase TssH gene	DMA00605	
<i>clpC</i>	Bacterial ATP-dependent Clp protease ATP-binding subunit gene	DMA00606	
<i>cpsL</i>	Bacterial capsular polysaccharide repeat unit transporter cpsL gene	DMA00607	
<i>CT456</i>	Bacterial translocated actin-recruiting phosphoprotein gene	DMA00608	
<i>ctxA</i>	Bacterial cholera enterotoxin catalytic subunit CtxA gene	DMA00609	
<i>eae</i>	Bacterial intimin type gamma protein gene	DMA00610	
<i>efaA</i>	Bacterial metal ABC transporter substrate-binding protein gene	DMA00611	YES
<i>Enterotoxigenic E. coli, LT</i>	<i>E. coli</i> heat-labile enterotoxin LT subunit B gene	DMA00612	
<i>Enterotoxigenic E. coli, ST</i>	<i>E. coli</i> heat-stable toxin I (estA1) gene	DMA00613	
<i>fimH</i>	Bacterial type 1 fimbriae D-mannose specific adhesin FimH gene	DMA00614	YES
<i>flaB</i>	Bacterial flagellin B gene	DMA00615	
<i>fleN</i>	Bacterial flagellar synthesis regulator FleN gene	DMA00616	
<i>flgG</i>	Bacterial flagellar hook protein FlgE gene	DMA00617	
<i>flgK</i>	Bacterial flagellar hook-associated protein FlgK gene	DMA00618	
<i>flhA</i>	Bacterial flagellar biosynthesis protein FlhA gene	DMA00619	
<i>flaB</i>	Bacterial flagellin B gene	DMA00615	
<i>fleN</i>	Bacterial flagellar synthesis regulator FleN gene	DMA00616	
<i>flgG</i>	Bacterial flagellar hook protein FlgE gene	DMA00617	
<i>flgK</i>	Bacterial flagellar hook-associated protein FlgK gene	DMA00618	
<i>flhA</i>	Bacterial flagellar biosynthesis protein FlhA gene	DMA00619	
<i>fliA</i>	Bacterial flagellar biosynthesis sigma factor fliA gene	DMA00620	
<i>fliC2</i>	Bacterial flagellin fliC2 gene	DMA00621	
<i>fliF</i>	Bacterial flagellar M-ring protein FliF gene	DMA00622	
<i>fliG</i>	Bacterial flagellar motor switch protein FliG gene	DMA00623	
<i>fliN</i>	Bacterial flagellar motor switch protein FliN gene	DMA00624	
<i>babA (H. pylori)</i>	Bacterial adhesin-binding fucosylated histo-blood group antigen babA gene	DMA00625	
<i>cagA (H. pylori)</i>	Bacterial cytotoxin-associated protein A (cagA) gene	DMA00626	
<i>dupA (H. pylori)</i>	Bacterial duodenal ulcer-promoting protein dupA gene	DMA00627	
<i>iceA (H. pylori)</i>	Bacterial ulcer-associated restriction endonuclease protein iceA gene	DMA00628	
<i>vacA (H. pylori)</i>	Bacterial autotransporter vacuolating cytotoxin VacA gene	DMA00629	YES
<i>virB (H. pylori)</i>	Bacterial P-type DNA transfer ATPase VirB11 gene	DMA00630	
<i>virD (H. pylori)</i>	Bacterial sodium-calcium antiporter virD gene	DMA00631	
<i>hap</i>	Bacterial autotransporter outer membrane beta-barrel domain-containing protein gene	DMA00632	

Virulence gene	Target info	GeneGlobe catalog number	Wet-lab tested
<i>hla</i>	Bacterial alpha-hemolysin <i>hla</i> gene	DMA00633	
<i>hlyB</i>	Bacterial sphingomyelinase <i>hlyB</i> gene	DMA00634	
<i>hlyB</i>	Bacterial gamma-hemolysin component B (<i>hlyB</i>) gene	DMA00635	YES
<i>hlyC</i>	Bacterial bi-component gamma-hemolysin <i>HlyCB</i> subunit C gene	DMA00636	
<i>htpB</i>	Bacterial molecular chaperone GroEL (<i>htpB</i>) gene	DMA00637	
<i>iap</i>	Bacterial invasion associated endopeptidase (<i>iap</i>) gene	DMA00638	
<i>icmK</i>	Bacterial intracellular multiplication protein (<i>IcmK</i>) gene	DMA00639	
<i>inlA</i>	Bacterial internalin A gene	DMA00640	
<i>inv</i>	Bacterial invasin <i>Inv</i> gene	DMA00641	
<i>invA</i>	Bacterial type III secretion system export apparatus protein <i>InvA</i> gene	DMA00642	YES
<i>ipaH</i>	<i>Bacterial invasion plasmid antigen ipaH gene</i>	DMA00643	
<i>iucA</i>	Bacterial aerobactin siderophore biosynthesis protein <i>iucA</i> gene	DMA00644	
<i>iucB</i>	Bacterial aerobactin siderophore biosynthesis protein <i>iucB</i> gene	DMA00645	
<i>iucC</i>	Bacterial aerobactin synthase <i>iucC</i> gene	DMA00646	
<i>iucD</i>	Bacterial L-lysine N6-monooxygenase <i>iucD</i> gene	DMA00647	
<i>iutA</i>	Bacterial ferric aerobactin receptor <i>iutA</i> gene	DMA00648	
<i>lasI</i>	Bacterial acyl-homoserine-lactone synthase <i>LasI</i> gene	DMA00649	
<i>lbpA</i>	Bacterial lactoferrin/transferrin family TonB-dependent receptor gene	DMA00650	
<i>lepA</i>	Bacterial interaptin <i>lepA</i> gene	DMA00651	
<i>lukF</i>	Bacterial Pantone-Valentine bi-component leukocidin subunit F gene	DMA00652	
<i>lytA</i>	Bacterial autolysin (N-acetylmuramoyl-L-alanine amidase) (Mureinhydrolase, Mucopolysaccharide aminohydrolase, Cell wall hydrolase) gene	DMA00653	
<i>mbtA</i>	Bacterial bifunctional salicyl-AMP ligase/salicyl-S-ArCP synthetase <i>mbtA</i> gene	DMA00654	
<i>mbtD</i>	Bacterial mycobactin polyketide synthase <i>MbtD</i> gene	DMA00655	
<i>mbtE</i>	Bacterial non-ribosomal peptide synthetase <i>mbtE</i> gene	DMA00656	
<i>neuC</i>	Bacterial UDP-N-acetylglucosamine 2-epimerase <i>neuC</i> gene	DMA00657	
<i>oipA</i>	Bacterial outer inflammatory protein <i>OipA</i> gene	DMA00658	
<i>phoP</i>	Bacterial two-component system response regulator <i>PhoP</i> gene	DMA00659	YES
<i>pilG</i>	Bacterial pilus assembly protein <i>PilG</i> gene	DMA00660	
<i>pirA/pirB</i>	Bacterial toxin <i>pirA/pirB</i> gene	DMA00661	
<i>pkn5</i>	Chlamydia serine/threonine protein kinase gene	DMA00662	
<i>plcA</i>	Bacterial 1-phosphatidylinositol phosphodiesterase gene	DMA00663	
<i>ply</i>	Bacterial pneumolysin (<i>ply</i>) gene	DMA00664	YES
<i>psaA</i>	Bacterial adhesin <i>PsaA</i> gene	DMA00665	

Virulence gene	Target info	GeneGlobe catalog number	Wet-lab tested
<i>psaB</i>	Bacterial chaperone protein PsaB gene	DMA00666	
<i>psaC</i>	Bacterial fimbrial biogenesis usher protein PsaC gene	DMA00667	
<i>ptxA</i>	Bacterial pertussis toxin ADP-ribosyltransferase subunit PtxA gene	DMA00668	YES
<i>rpoS</i>	Bacterial serine/threonine protein kinase gene	DMA00669	
<i>slo</i>	Bacterial cholesterol-dependent cytolysin streptolysin O <i>slo</i> gene	DMA00670	
<i>spa</i>	Bacterial staphylococcal protein A <i>spa</i> gene	DMA00671	
<i>spaA</i>	Bacterial isopeptide-forming domain-containing fimbrial protein gene	DMA00672	
<i>spaB</i>	Bacterial carboxypeptidase regulatory-like domain-containing protein gene	DMA00673	
<i>spaO</i>	Bacterial SPI-1 type III secretion system protein SpaO gene	DMA00674	
<i>spaP</i>	Bacterial SPI-1 type III secretion system export apparatus protein SpaP gene	DMA00675	
<i>speB</i>	Bacterial cysteine proteinase exotoxin SpeB gene	DMA00676	
<i>stx1B</i>	Bacterial shiga toxin Stx1a subunit B gene	DMA00677	YES
<i>stx2A</i>	Bacterial shiga toxin Stx2c subunit A gene	DMA00678	YES
<i>stxA</i>	Bacterial shiga toxin Stx1 subunit A gene	DMA00679	YES
<i>stxB</i>	Bacterial shiga toxin Stx1a subunit B gene	DMA00680	YES
<i>tbpA</i>	Bacterial lactoferrin/transferrin family TonB-dependent receptor gene	DMA00681	
<i>tcdA</i>	Bacterial toxin A (<i>tcdA</i>) gene	DMA00682	
<i>tcdB</i>	Bacterial glycosylating toxin TcdB gene	DMA00683	
<i>ureA</i>	Bacterial urease subunit alpha <i>ureA</i> gene	DMA00684	
<i>ureI</i>	Bacterial acid-activated urea channel <i>ureI</i> gene	DMA00685	
<i>vasH</i>	Bacterial sigma-54 dependent transcriptional regulator VasH gene	DMA00686	
<i>waaC</i>	Bacterial lipopolysaccharide heptosyltransferase I <i>waaC</i> gene	DMA00687	
<i>wbkA</i>	Bacterial glycosyltransferase family 4 protein gene	DMA00688	YES
<i>wzt</i>	Bacterial ABC transporter ATP-binding protein gene	DMA00689	
<i>ybtT</i>	Bacterial yersiniabactin biosynthesis thioesterase YbtT gene	DMA00690	
YE2522	Bacterial flagellin A gene	DMA00691	
<i>ystA</i>	Bacterial enterotoxin <i>ystA</i> gene	DMA00692	
<i>zot</i>	Bacterial zonula occludens toxin ZOT gene	DMA00693	

Table 7. Control assays

Control assays	Target info	GeneGlobe catalog number	Wet-lab tested
Escherichia phage MS2 (12022)	Wastewater	DMA00775	YES
HF183 marker	human-specific HF183 Bacteroides/ Phocaeicola spp. 16S rRNA genetic marker	DMA00721	YES
RNaseP (g)	human ribonuclease P/MRP Subunit P30 gene RPP30	DMA00720	YES

Ordering Information

Product	Contents	Cat. no.
dPCR Microbial DNA Detection Assays	One tube Microbial DNA dPCR Assay, 200 rxn for 40 µL reaction in Nanoplate 26k, 666 rxn for 12 µL reaction for Nanoplate 8.5k	250207*
	10 QIAcuity Nanoplates 26k 24-well, 11 Nanoplate Seals	250001
QIAcuity Probe PCR Kit (1 mL, 5 mL, 25 mL)	1 mL, 5 mL, or 25 mL Master Mix for the QIAcuity dPCR instrument; water	250101 250102 250103
QIAcuity OneStep Advanced Probe Kit (1 mL, 5 mL)	1 mL or 5 mL OneStep Master Mix for the QIAcuity dPCR instrument, RT Mix, Internal Control RNA, Enhancer GC, water	250131 250132
QIAcuity High Multiplex Probe PCR Kit (1 mL, 5 mL)	1 mL or 5 mL 4x concentrated QIAcuity High Multiplex Probe PCR Master Mix, 2 x 1.9 mL RNase-free Water, QN Internal Control DNA	250133 250134
QIAcuity UCP Probe PCR Kit (1 mL, 5 mL)	1 mL or 5 mL ultra clean production Master Mix for the QIAcuity dPCR instrument; water	250121 250122

* For the respective Assay ID, see the tables above; full assay cat. no. will be available after dye configuration.

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