

項次	菌株代碼	菌種代碼	菌株名稱
1	FP00001	FT006	Acinetobacter haemolyticus
2	FP00002	FT006	Acinetobacter baumannii
3	FP00003	FT006	Acinetobacter lwoffii
4	FP00004	FT006	Acinetobacter spp.
5	FP00005	FT004	Aeromonas caviae
6	FP00006	FT004	Aeromonas hydrophila
7	FP00007	FT004	Aeromonas sobria
8	FP00008	FT006	Elizabethkingia menigseptica
9	FP00009	FT004	Citrobacter amalonaticus
10	FP00011	FT004	Citrobacter freundii
11	FP00012	FT006	Delftia acidovorans
12	FP00013	FT004	Pantoea agglomerans
13	FP00014	FT004	Enterobacter aerogenes
14	FP00015	FT004	Enterobacter cloacae
15	FP00016	FT004	Enterobacter sakazakii
16	FP00017	FT004	Escherichia coli
17	FP00018	FT003	Haemophilus influenzae
18	FP00019	FT004	Klebsiella oxytoca
19	FP00020	FT004	Klebsiella ozaenae
20	FP00021	FT004	Klebsiella pneumoniae
21	FP00022	FT004	Morganella morganii
22	FP00023	FT002	Neisseria spp.
23	FP00024	FT004	Proteus mirabilis
24	FP00025	FT004	Proteus penneri
25	FP00026	FT004	Proteus vulgaris
26	FP00027	FT004	Providencia rettgeri
27	FP00028	FT004	Providencia stuartii
28	FP00029	FT006	Pseudomonas aeruginosa
29	FP00030	FT006	Pseudomonas putida
30	FP00031	FT006	Pseudomonas spp.
31	FP00032	FT004	Salmonella Group B
32	FP00033	FT004	Salmonella Group D
33	FP00034	FT004	Salmonella spp.
34	FP00035	FT004	Serratia marcescens
35	FP00036	FT006	Shewanella putrefaciens
36	FP00037	FT006	Sphingomonas paucimobilis
37	FP00038	FT006	Stenotrophomonas maltophilia
38	FP00039	FT006	Acinetobacter junii
39	FP00040	FT005	Bordetella bronchiseptica (CDC IVa)
40	FP00041	FT004	Citrobacter diversus
41	FP00042	FT004	Citrobacter spp.
42	FP00043	FT003	Eikenella corrodens
43	FP00044	FT004	Flavimonas oryzae



44	FP00045	FT006	Flavobacterium spp.
45	FP00046	FT003	Haemophilus influenzae type b
46	FP00047	FT006	Methylobacterium mesophilicum
47	FP00048	FT002	Neisseria meningitidis
48	FP00049	FT004	Proteus spp.
49	FP00052	FT007	Bacillus spp.
50	FP00053	FT008	beta-Hemolytic streptococcus not Gr.A,B,C,D,F,G
51	FP00054	FT007	Corynebacterium spp.
52	FP00056	FT008	Enterococcus faecalis
53	FP00057	FT008	Enterococcus faecium
54	FP00059	FT008	Micrococcus spp.
55	FP00060	FT011	Nocardia spp.
56	FP00061	FT008	Staphylococcus aureus
57	FP00062	FT008	Staphylococcus epidermidis
58	FP00063	FT008	Staphylococcus saccharolyticus
59	FP00064	FT008	Staphylococcus saprophyticus
60	FP00065	FT008	Streptococcus pneumoniae
61	FP00066	FT008	Viridans streptococci
62	FP00067	FT008	Aerococcus spp.
63	FP00068	FT008	Enterococcus spp.
64	FP00069	FT008	Streptococcus haemolyticus
65	FP00070	FT001	Bacteroides capillosus
66	FP00071	FT001	Bacteroides caccae
67	FP00072	FT001	Bacteroides eggerthii
68	FP00073	FT001	Bacteroides Fragilis group
69	FP00074	FT001	Bacteroides fragilis
70	FP00075	FT001	Bacteroides ovatus
71	FP00076	FT001	Bacteroides thetaiotaomicron
72	FP00077	FT001	Bacteroides merdae
73	FP00078	FT001	Bacteroides gracilis
74	FP00079	FT001	Bacteroides uniformis
75	FP00080	FT001	Bacteroides vulgatus
76	FP00081	FT001	Bifidobacterium spp.
77	FP00082	FT001	Bifidobacterium wadsworthia
78	FP00083	FT012	Cladosporium spp.
79	FP00084	FT001	Clostridium bifermentans
80	FP00085	FT001	Clostridium difficile
81	FP00086	FT001	Clostridium sordellii
82	FP00087	FT001	Clostridium septicum
83	FP00088	FT001	Clostridium perfringens
84	FP00089	FT001	Clostridium spp.
85	FP00090	FT001	Eubacterium spp.
86	FP00091	FT001	Eubacterium lentum
87	FP00092	FT001	Fusobacterium necrophorum



88	FP00093	FT001	Fusobacterium nucleatum
89	FP00094	FT001	Fusobacterium spp.
90	FP00095	FT001	Fusobacterium varium
91	FP00096	FT001	Fusobacterium necrogenes
92	FP00097	FT001	Fusobacterium mortiferum
93	FP00098	FT008	Gemella morbillorum (Streptococcus morbillonim)
94	FP00099	FT001	Lactobacillus spp. (anaerobe)
95	FP00100	FT001	Mobiluncus curtisii
96	FP00101	FT001	Peptostreptococcus anaerobius
97	FP00102	FT001	Peptostreptococcus asaccharolyticus
98	FP00103	FT001	Peptostreptococcus magnus
99	FP00104	FT001	Peptostreptococcus micros
100	FP00105	FT001	Peptostreptococcus prevotii
101	FP00106	FT001	Peptostreptococcus spp.
102	FP00107	FT001	Prevotella bivia
103	FP00108	FT001	Prevotella buccae
104	FP00109	FT001	Prevotella corporis
105	FP00110	FT001	Prevotella intermedia
106	FP00111	FT001	Prevotella melaninogenica
107	FP00112	FT001	Prevotella oris
108	FP001121	FT013	Rotavirus
109	FP00113	FT001	Prevotella oralis
110	FP00114	FT001	Prevotella spp.
111	FP00115	FT001	Propionibacterium acnes
112	FP001157	FT006	Myroides spp.
113	FP00116	FT001	Propionibacterium granulosum
114	FP00117	FT001	Porphyromonas gingivalis
115	FP00119	FT001	Porphyromonas asaccharolytica
116	FP00120	FT008	Streptococcus intermedius
117	FP00121	FT001	Veillonella parvula
118	FP00122	FT008	Viridans dispar
119	FP00123	FT001	Prevotella denticola
120	FP00124	FT001	Prevotella disiens
121	FP00125	FT001	Prevotella loescheii
122	FP00127	FT001	Propionibacterium spp.
123	FP00128	FT001	Veillonella spp.
124	FP00129	FT011	Candida glabrata
125	FP00130	FT011	Candida guilliermondii
126	FP00131	FT011	Candida intermedia
127	FP00132	FT011	Candida sake
128	FP00133	FT011	Candida holmii
129	FP00134	FT011	Candida krusei
130	FP00135	FT011	Candida lipolytica
131	FP00136	FT011	Candida lusitaniae



132	FP00137	FT011	Candida parapsilosis
133	FP00138	FT011	Candida colliculosa
134	FP00139	FT011	Cryptococcus laurentii
135	FP00140	FT011	Zygosaccharomyces spp.
136	FP00141	FT011	Candida albicans
137	FP00142	FT011	Candida spp.
138	FP00143	FT011	Candida tropicalis
139	FP00144	FT012	Aspergillus spp.
140	FP00145	FT012	Fusarium spp.
141	FP00146	FT012	Unidentified fungus
142	FP00147	FT012	Penicillium spp.
143	FP00148	FT012	Scopulariopsis spp.
144	FP00149	FT007	Streptomyces spp.
145	FP00150	FT011	Torulopsis glabrata
146	FP00151	FT008	Abiotrophia spp.
147	FP00152	FT004	Aeromonas salmonicida
148	FP00153	FT004	Aeromonas spp.
149	FP00154	FT006	Alcaligenes xylosoxidans
150	FP00155	FT012	Aspergillus fumigatus
151	FP00156	FT012	Aspergillus terreus
152	FP00157	FT012	Aspergillus flavus
153	FP00158	FT012	Aspergillus niger
154	FP00159	FT007	Bacillus cereus
155	FP00160	FT007	Bacillus circulans
156	FP00161	FT007	Bacillus coagulans
157	FP00162	FT007	Bacillus polymyxa
158	FP00163	FT007	Bacillus pumilis
159	FP00164	FT007	Bacillus sphaericus
160	FP00165	FT001	Bacteroides splanchnicus
161	FP00166	FT001	Bacteroides distasonis
162	FP00167	FT001	Bacteroides vulgatus
163	FP00168	FT001	Bacteroides stercoris
164	FP00169	FT001	Bacteroides ureolyticus
165	FP00170	FT005	Branhamella (Moraxella) catarrhalis
166	FP00171	FT006	Burkholderia cepacia
167	FP00172	FT001	Campylobacter spp.
168	FP00173	FT003	Capnocytophaga spp.
169	FP00174	FT006	Chryseobacterium indologenes
170	FP00175	FT004	Citrobacter koseri
171	FP00176	FT001	Clostridium hastiforme
172	FP00177	FT001	Clostridium innocuum
173	FP00178	FT001	Clostridium paraputrificum
174	FP00179	FT001	Clostridium ramosum
175	FP00180	FT006	Comamonas testosteroni



176	FP00181	FT007	Corynebacterium group JK
177	FP00182	FT011	Cryptococcus luteolus
178	FP00183	FT011	Cryptococcus neoformans
179	FP00184	FT012	Dematiaceous mold
180	FP00185	FT001	Eubacterium lentum
181	FP00186	FT008	Enterococcus avium
182	FP00187	FT008	Enterococcus casseliflavus
183	FP00188	FT004	Enterobacter spp.
184	FP00189	FT004	Enterobacter gergoviae
185	FP00190	FT001	Eubacterium aerofaciens
186	FP00191	FT001	Eubacterium limosum
187	FP00192	FT008	Group A streptococcus (S. pyogenes)
188	FP00193	FT008	Group B streptococcus (S. agalactiae)
189	FP00194	FT008	Group C streptococcus
190	FP00195	FT008	Streptococcus group D non-enterococcus
191	FP00196	FT008	Group F streptococcus (S. anffinosus)
192	FP00197	FT008	Group G streptococcus
193	FP00198	FT003	Haemophilus parainfluenzae
194	FP00199	FT004	Helicobacter pylori (Campylobacter pylori)
195	FP00200	FT004	Klebsiella spp.
196	FP00201	FT004	Klebsiella terrigena
197	FP00202	FT012	Monilinia sitophila
198	FP00203	FT006	Moraxella osloensis
199	FP00204	FT012	Mucor spp.
200	FP00205	FT002	Neisseria gonorrhoeae
201	FP00206	FT011	Nocardia asteroides
202	FP00207	FT006	Unidentified nonfermentative Gram negative bacillus
203	FP00208	FT006	Ochrobactrum anthropi
204	FP00209	FT012	Paecilomyces spp.
205	FP00210	FT008	Pediococcus spp.
206	FP00211	FT012	Penicillium marneffeii
207	FP00212	FT001	Peptostreptococcus productus
208	FP00213	FT001	Peptostreptococcus tetradius
209	FP00214	FT004	Providencia spp.
210	FP00215	FT006	Pseudomonas fluorescens
211	FP00216	FT006	Pseudomonas mendocina
212	FP00217	FT006	Pseudomonas stutzeri
213	FP00218	FT006	Pseudomonas fluorescens/putida/mendocina
214	FP00219	FT006	Ralstonia pickettii
215	FP00220	FT008	r-Hemolytic streptococcus
216	FP00221	FT011	Rhodotorula glutinis
217	FP00222	FT011	Rhodotorula nibra
218	FP00223	FT011	Rhodotorula spp.
219	FP00224	FT011	Saccharomyces cerevisiae



220	FP00225	FT004	Salmonella Group C1
221	FP00226	FT004	Salmonella Group E1
222	FP00227	FT004	Salmonella typhi
223	FP00228	FT012	Scedosporium apiospermum
224	FP00229	FT004	Serratia liquefaciens
225	FP00230	FT004	Serratia rubidaea
226	FP00231	FT004	Serratia spp.
227	FP00232	FT008	Staphylococcus hominis
228	FP00233	FT008	Staphylococcus not aureus
229	FP00234	FT008	Stomatococcus spp.
230	FP00235	FT008	Streptococcus constellatus
231	FP00236	FT011	Torulopsis Candida
232	FP00237	FT012	Trichophyton rubrum
233	FP00238	FT011	Trichosporon beigeli
234	FP00239	FT011	Trichosporon mucoides
235	FP00240	FT999	Unidentified anaerobic G(+) non-sporeforming bacillus
236	FP00241	FT004	Unidentified Gram negative bacillus
237	FP00242	FT007	Unidentified Gram positive bacillus
238	FP00244	FT999	Unidentified nonfermentative Gram negative bacillus
239	FP00245	FT004	Vibrio cholerae non-O1
240	FP00246	FT004	Vibrio vulnificus
241	FP00247	FT011	Yeast-like
242	FP00248	FT012	Absidia spp.
243	FP00249	FT006	Achromobacter (Vd)
244	FP00250	FT006	Achromobacter spp.
245	FP00251	FT006	Achromobacter xylosoxidans
246	FP00252	FT006	Acinetobacter alcaligenes
247	FP00253	FT006	Acinetobacter anitratus
248	FP00254	FT006	Acinetobacter calcoaceticus
249	FP00255	FT006	Acinetobacter johnsonii
250	FP00256	FT003	Actinobacillus actinomycetemcomitans
251	FP00257	FT003	Actinobacillus capsulatus
252	FP00258	FT003	Actinobacillus equuli
253	FP00259	FT003	Actinobacillus hominis
254	FP00260	FT003	Actinobacillus lignieresii
255	FP00261	FT003	Actinobacillus spp.
256	FP00262	FT003	Actinobacillus suis
257	FP00263	FT001	Actinomyces bovis
258	FP00264	FT001	Actinomyces israelii
259	FP00265	FT001	Actinomyces meyeri
260	FP00266	FT001	Actinomyces naeslundii
261	FP00267	FT001	Actinomyces odontolyticus
262	FP00268	FT001	Actinomyces pyogenes
263	FP00269	FT001	Actinomyces spp.



264	FP00270	FT001	Actinomyces viscosus
265	FP00271	FT004	Aeromonas salmonicida subspp. achromogenes
266	FP00272	FT004	Aeromonas salmonicida subspp. masoucida
267	FP00273	FT004	Aeromonas salmonicida subspp. salmonicida
268	FP00274	FT008	Aerococcus viridans
269	FP00275	FT004	Aeromonas veronii
270	FP00276	FT006	Agrobacterium radiobacter
271	FP00277	FT006	Agrobacterium spp
272	FP00278	FT006	Agrobacterium tumefaciens
273	FP00279	FT006	Alcaligenes xylosoxidans subsp. denitrificans
274	FP00280	FT006	Alcaligenes faecalis
275	FP00281	FT006	Alcaligenes odorans
276	FP00282	FT006	Alcaligenes spp.
277	FP00283	FT012	Alternaria spp.
278	FP00284	FT001	Anaerobiospirillum succiniciproducens
279	FP00285	FT001	Anaerobiospirillum spp.
280	FP00286	FT001	Anaerovibrio lipolyticus
281	FP00287	FT001	Anaerobic bacteria
282	FP00288	FT001	Anaerobic GNB
283	FP00289	FT001	Anaerobic GNC
284	FP00290	FT001	Anaerobic GPB
285	FP00291	FT001	Anaerobic GPC
286	FP00292	FT001	Arachnia propionica
287	FP00293	FT007	Arcanobacterium haemolyticum
288	FP00294	FT007	Arthrobacter globiformis
289	FP00295	FT007	Arthrobacter spp.
290	FP00297	FT007	Bacillus alvei
291	FP00298	FT007	Bacillus amyloliquefaciens
292	FP00299	FT007	Bacillus anthracis
293	FP00300	FT007	Bacillus badius
294	FP00301	FT007	Bacillus brevis
295	FP00303	FT007	Bacillus firmus
296	FP00304	FT007	Bacillus laterosporus
297	FP00305	FT007	Bacillus lentus
298	FP00306	FT007	Bacillus licheniformis
299	FP00307	FT007	Bacillus macerans
300	FP00308	FT007	Bacillus megaterium
301	FP00309	FT007	Bacillus mycoides
302	FP00310	FT007	Bacillus pantothenicus
303	FP00311	FT007	Bacillus stearothermophilus
304	FP00312	FT007	Bacillus subtilis
305	FP00313	FT007	Bacillus thuringiensis
306	FP00314	FT001	Bacteroides caecae
307	FP00315	FT001	Bacteroides capillosus



308	FP00316	FT001	<del>Bacteroides distasonis</del>
309	FP00317	FT001	Bacteroides group CDC F2
310	FP00318	FT001	Bacteroides group fragilis
311	FP00319	FT001	<del>Bacteroides splanchnicus</del>
312	FP00320	FT001	<del>Bacteroides thetaiotaomicron</del>
313	FP00321	FT001	<del>Bacteroides uniformis</del>
314	FP00322	FT001	Bacteroides ureolyticus group
315	FP00323	FT001	<del>Bacteroides vulgatus</del>
316	FP00324	FT001	Bacteroides levii
317	FP00325	FT001	<del>Bacteroides ovatus</del>
318	FP00326	FT001	Bacteroides spp.
319	FP00327	FT001	Bifidobacterium adolescentis
320	FP00328	FT001	Bifidobacterium bifidum
321	FP00329	FT001	Bifidobacterium breve
322	FP00330	FT001	Bifidobacterium dentium
323	FP00331	FT001	Bifidobacterium infantis
324	FP00332	FT012	Blastomyces dermatitidis
325	FP00333	FT012	Blastomyces spp.
326	FP00334	FT005	Bordetella avium
327	FP00335	FT005	Bordetella parapertussis
328	FP00336	FT005	Bordetella pertussis
329	FP00337	FT005	Bordetella spp.
330	FP00338	FT009	Borrelia spp.
331	FP00339	FT005	Branhamella spp.
332	FP00340	FT007	Brevibacterium linens
333	FP00341	FT007	Brevibacterium spp.
334	FP00342	FT008	Brochothrix thermosphacta
335	FP00343	FT005	Brucella abortus
336	FP00344	FT005	Brucella canis
337	FP00345	FT005	Brucella melitensis
338	FP00346	FT005	Brucella ovis
339	FP00347	FT005	Brucella spp.
340	FP00348	FT005	Brucella suis
341	FP00349	FT004	Budvicia aquatica
342	FP00350	FT004	Buttiauxella agrestis
343	FP00351	FT001	Campylobacter sputorum subsp. bubulus
344	FP00352	FT001	Campylobacter fetus subsp. venerealis
345	FP00353	FT001	Campylobacter cinaedi
346	FP00354	FT001	Campylobacter coli
347	FP00355	FT001	Campylobacter concisus
348	FP00356	FT001	Campylobacter cryaerophilus
349	FP00357	FT001	Campylobacter fecalis
350	FP00358	FT001	Campylobacter fennelliae
351	FP00359	FT001	Campylobacter fetus



352	FP00360	FT001	Campylobacter fetus subspp. fetus
353	FP00361	FT001	Campylobacter hyointestinalis
354	FP00362	FT001	Campylobacter jejuni subspp. doylei
355	FP00363	FT001	Campylobacter jejuni subspp. jejuni
356	FP00364	FT001	Campylobacter lari
357	FP00365	FT001	Campylobacter lari spp UPTC
358	FP00366	FT001	Campylobacter mucosalis
359	FP00367	FT001	Campylobacter sputorum
360	FP00368	FT001	Campylobacter upsaliensis
361	FP00369	FT001	Campylobacterium spp.
362	FP00370	FT011	Candida boidinii
363	FP00371	FT011	Candida catenulata
364	FP00372	FT011	Candida ciferrii
365	FP00373	FT011	Candida globobata
366	FP00374	FT011	Candida curvata
367	FP00375	FT011	Candida dattila
368	FP00376	FT011	Candida famata
369	FP00377	FT011	Candida globosa
370	FP00378	FT011	Candida humicola
371	FP00379	FT011	Candida inconspicua
372	FP00380	FT011	Candida kefyr
373	FP00381	FT011	Candida lambica
374	FP00382	FT011	Candida magnoliae
375	FP00383	FT011	Candida maris
376	FP00384	FT011	Candida melibiosica
377	FP00385	FT011	Candida membranaefaciens
378	FP00386	FT011	Candida norvegensis
379	FP00387	FT011	Candida norvegica
380	FP00388	FT011	Candida paratropicalis
381	FP00389	FT011	Candida pelliculosa
382	FP00390	FT011	Candida pseudotropicalis(C.kdf
383	FP00391	FT011	Candida pulcherrima
384	FP00392	FT011	Candida rugosa
385	FP00393	FT011	Candida silvicola
386	FP00394	FT011	Candida sphaerica
387	FP00395	FT011	Candida stellatoidea
388	FP00397	FT011	Candida utilis
389	FP00398	FT011	Candida valida
390	FP00399	FT011	Candida zeylanoides
391	FP00400	FT003	Capnocytophaga gingivalis
392	FP00401	FT003	Capnocytophaga ochracea
393	FP00402	FT003	Capnocytophaga sputigena
394	FP00403	FT003	Cardiobacterium hominis
395	FP00404	FT001	Carnobacterium divergens



396	FP00405	FT001	Carnobacterium piscicola
397	FP00406	FT006	CDC group M-5
398	FP00407	FT006	CDC group M-6
399	FP00408	FT004	Cedecea davisae
400	FP00409	FT004	Cedecea lapagei
401	FP00410	FT004	Cedecea neteri
402	FP00411	FT004	Cedecea spp
403	FP00412	FT012	Cephalosporium spp.
404	FP00413	FT014	Chlamydia psittaci
405	FP00414	FT014	Chlamydia spp.
406	FP00415	FT014	Chlamydia trachomatis
407	FP00416	FT004	Chromobacterium violaceum
408	FP00417	FT006	Chryseomonas luteola
409	FP00418	FT012	Chrysosporium spp.
410	FP00419	FT012	Circinella spp.
411	FP00421	FT012	Cladosporium carrionii
412	FP00422	FT001	Clostridium acetobutylicum
413	FP00423	FT001	Clostridium barati
414	FP00425	FT001	Clostridium botulinum
415	FP00426	FT001	Clostridium butyricum
416	FP00427	FT001	Clostridium cadaveris
417	FP00428	FT001	Clostridium chauvoei
418	FP00429	FT001	Clostridium clostridioforme
419	FP00430	FT001	Clostridium fallax
420	FP00431	FT001	Clostridium glycolicum
421	FP00432	FT001	Clostridium histolyticum
422	FP00433	FT001	Clostridium limosum
423	FP00434	FT001	Clostridium malenominatum
424	FP00435	FT001	Clostridium novyi
425	FP00437	FT001	Clostridium perenne
426	FP00438	FT001	Clostridium putrificum
427	FP00439	FT001	Clostridium sphenoides
428	FP00440	FT001	Clostridium sporogenes
429	FP00441	FT001	Clostridium subterminale
430	FP00442	FT001	Clostridium symbiosum
431	FP00443	FT001	Clostridium tertium
432	FP00444	FT001	Clostridium tetani
433	FP00445	FT001	Clostridium tyrobutyricum
434	FP00446	FT012	Coccidioides spp.
435	FP00447	FT012	Coccidioides brasiliensis
436	FP00448	FT006	Comamonas spp
437	FP00449	FT007	Corynebacterium diphtheriae biotype belfanti
438	FP00450	FT007	Corynebacterium diphtheriae biotype gravis
439	FP00451	FT007	Corynebacterium diphtheriae biotype mitis



440	FP00452	FT007	Corynebacterium pseudodiphtheriticum
441	FP00453	FT007	Corynebacterium pseudotuberculosis
442	FP00454	FT007	Corynebacterium spp.
443	FP00455	FT007	Corynebacterium aquaticum
444	FP00456	FT007	Corynebacterium bovis
445	FP00457	FT007	Corynebacterium cystitidis
446	FP00458	FT007	Corynebacterium diphtheriae
447	FP00459	FT007	Corynebacterium flavesens
448	FP00460	FT007	CDC coryneform group A
449	FP00461	FT007	CDC coryneform group ANF
450	FP00462	FT007	Corynebacterium group B
451	FP00463	FT007	Corynebacterium group C
452	FP00464	FT007	Corynebacterium group D
453	FP00465	FT007	Corynebacterium group E
454	FP00466	FT007	Corynebacterium group F
455	FP00467	FT007	Corynebacterium group G
456	FP00468	FT007	Corynebacterium group I
457	FP00469	FT007	Corynebacterium jeikeium
458	FP00470	FT007	Corynebacterium kutscheri
459	FP00471	FT007	Corynebacterium minutissimum
460	FP00472	FT007	Corynebacterium pilosum
461	FP00473	FT007	Corynebacterium renale
462	FP00474	FT007	Corynebacterium xerosis
463	FP00475	FT007	Corynebacterium striatum
464	FP00476	FT007	Corynebacterium ulcerans
465	FP00477	FT011	Cryptococcus albidus
466	FP00478	FT011	Cryptococcus gastricus
467	FP00479	FT011	Cryptococcus spp.
468	FP00480	FT011	Cryptococcus terreus
469	FP00481	FT011	Cryptococcus uniguttulatus
470	FP00482	FT012	Cunninghamella spp.
471	FP00483	FT012	Curvularia spp.
472	FP00484	FT012	Cusarium spp.
473	FP00485	FT013	Cytomegalovirus
474	FP00487	FT003	CDC group DF-2
475	FP00488	FT004	Edwardsiella hoshinae
476	FP00489	FT004	Edwardsiella ictaluri
477	FP00490	FT004	Edwardsiella spp
478	FP00491	FT004	Edwardsiella tarda
479	FP00492	FT004	CDC group EF-4
480	FP00493	FT008	Enterococcus durans
481	FP00496	FT008	Enterococcus gallinarum
482	FP00497	FT008	Enterococcus malodoratus
483	FP00498	FT008	Enterococcus mundtii



484	FP00499	FT008	Enterococcus sp.
485	FP00500	FT004	Enterobacter spp.
486	FP00501	FT004	Enterobacter amnigenus
487	FP00502	FT004	Enterobacter asburiae
488	FP00503	FT004	Enterobacter intermedium
489	FP00504	FT004	Enterobacter taylorae
490	FP00505	FT008	Enterococcus casseliflavus
491	FP00506	FT008	Enterococcus hirae
492	FP00507	FT012	Epicoccum spp.
493	FP00508	FT012	Epidermophyton floccosum
494	FP00509	FT012	Epidermophyton spp.
495	FP00510	FT004	Erwinia carotovora subspp. betavsculorum
496	FP00511	FT004	Erwinia amylovora
497	FP00512	FT004	Erwinia ananatis
498	FP00513	FT004	Erwinia carotovora
499	FP00514	FT004	Erwinia carotovora subspp. atroseptica
500	FP00515	FT004	Erwinia carotovora subspp. carotovora
501	FP00516	FT004	Erwinia chrysanthemi
502	FP00517	FT004	Erwinia cypripedii
503	FP00518	FT004	Erwinia mallotivora
504	FP00519	FT004	Erwinia nigrifluens
505	FP00520	FT004	Erwinia quercina
506	FP00521	FT004	Erwinia rhapontici
507	FP00522	FT004	Erwinia rubrifaciens
508	FP00523	FT004	Erwinia salicis
509	FP00524	FT004	Erwinia spp.
510	FP00525	FT004	Erwinia stewartii
511	FP00526	FT004	Erwinia uredovora
512	FP00527	FT007	Erysipelothrix spp.
513	FP00528	FT004	Escherichia blattae
514	FP00529	FT004	Escherichia fergusonii
515	FP00531	FT004	Escherichia hermannii
516	FP00532	FT004	Escherichia spp.
517	FP00533	FT004	Escherichia vulneris
518	FP00534	FT001	Eubacterium alactolyticum
519	FP00535	FT004	Ewingella americana
520	FP00536	FT006	Flavobacterium breve
521	FP00537	FT006	Flavobacterium group IIe
522	FP00538	FT006	Flavobacterium group IIh
523	FP00539	FT006	Flavobacterium group Ili
524	FP00540	FT006	Flavobacterium indologenes
525	FP00541	FT006	Flavobacterium meningosepticum
526	FP00542	FT006	Flavobacterium multivorum
527	FP00543	FT006	Flavobacterium odoratum



528	FP00544	FT006	Flavobacterium oryzihabitans
529	FP00545	FT012	Fonsecaea compacta
530	FP00546	FT012	Fonsecaea pedrosi
531	FP00547	FT012	Fonsecaea spp.
532	FP00548	FT003	Francisella spp.
533	FP00549	FT003	Francisella tularensis
534	FP00550	FT001	Fusobacterium gonidiaformans
535	FP00552	FT007	Gardnerella vaginalis
536	FP00553	FT008	Gemella haemolysans
537	FP00554	FT008	Gemella morbillorum
538	FP00555	FT012	Geotrichum candidum
539	FP00556	FT012	Geotrichum capitatum
540	FP00557	FT012	Geotrichum fermentans
541	FP00558	FT012	Geotrichum penicillatum
542	FP00559	FT012	Geotrichum spp.
543	FP00560	FT012	Gliocladium spp.
544	FP00561	FT004	GNB (Unidentified GNB)
545	FP00562	FT005	Unidentified Gram negative coccus (GNC)
546	FP00565	FT008	Unidentified Gram positive coccus (GPC)
547	FP00566	FT006	CDC Group EF-4b
548	FP00567	FT006	CDC Group EO-2
549	FP00568	FT006	CDC Group IV-C2
550	FP00569	FT006	CDC Group IV-E
551	FP00570	FT003	Haemophilus aegyptius
552	FP00571	FT003	Haemophilus aphrophilus
553	FP00572	FT003	Haemophilus avium
554	FP00573	FT003	Haemophilus ducreyi
555	FP00574	FT003	Haemophilus haemoglobinophilus
556	FP00575	FT003	Haemophilus haemolyticus
557	FP00576	FT003	Haemophilus paragallinarum
558	FP00577	FT003	Haemophilus parahaemolyticus
559	FP00578	FT003	Haemophilus paraphrohaemolyticus
560	FP00579	FT003	Haemophilus paraphrophilus
561	FP00580	FT003	Haemophilus pleuropneumoniae
562	FP00581	FT003	Haemophilus segnis
563	FP00582	FT003	Haemophilus spp
564	FP00583	FT004	Hafnia alvei
565	FP00584	FT004	Hafnia spp.
566	FP00585	FT011	Hansenula anomala
567	FP00586	FT011	Hansenula saturnus
568	FP00587	FT004	CDC group HB-5
569	FP00588	FT012	Helminthosporium spp.
570	FP00589	FT013	Herpes Simplex virus
571	FP00590	FT012	Heterosporium spp.



572	FP00591	FT012	Histoplasma capsulatum
573	FP00592	FT012	Histoplasma spp.
574	FP00593	FT004	Kingella denitrificans
575	FP00594	FT004	Kingella kingae
576	FP00595	FT004	Kingella spp.
577	FP00596	FT004	Raoultella planticola
578	FP00597	FT004	Klebsiella rhinoscleromatics
579	FP00598	FT011	Kloeckera apiculata
580	FP00599	FT011	Kloeckera apis
581	FP00600	FT011	Kloeckera japonica
582	FP00601	FT011	Kloeckera spp.
583	FP00602	FT004	Kluyvera ascorbata
584	FP00603	FT004	Kluyvera cryocrescens
585	FP00604	FT004	Kluyvera spp.
586	FP00605	FT006	Koserella trabulsii
587	FP00606	FT001	Lactobacillus casei subsp. alactosus
588	FP00607	FT001	Lactobacillus casei subsp. casei
589	FP00608	FT001	Lactobacillus paracasei
590	FP00609	FT001	Lactobacillus delbrueckii subsp. bulgaricus
591	FP00610	FT001	Lactobacillus helveticus subsp. jugurti
592	FP00611	FT001	Lactobacillus lactis subsp. hordniae
593	FP00612	FT001	Lactobacillus acidophilus
594	FP00613	FT001	Lactobacillus brevis
595	FP00614	FT001	Lactobacillus buchneri
596	FP00615	FT001	Lactobacillus casei
597	FP00616	FT001	Lactobacillus cateniformis
598	FP00617	FT001	Lactobacillus cellobiosus
599	FP00618	FT001	Lactobacillus collinoides
600	FP00619	FT001	Lactobacillus crispatus
601	FP00620	FT001	Lactobacillus curvatus
602	FP00621	FT001	Lactobacillus delbrueckii spp
603	FP00622	FT001	Lactobacillus fermentum
604	FP00623	FT001	Lactobacillus fructivorans
605	FP00624	FT001	Lactobacillus helveticus spp
606	FP00625	FT001	Lactobacillus jensenii
607	FP00626	FT001	Lactobacillus leichmannii
608	FP00627	FT001	Lactobacillus lindneri
609	FP00628	FT001	Lactobacillus minutus
610	FP00629	FT001	Lactobacillus pentosus
611	FP00630	FT001	Lactobacillus plantarum
612	FP00631	FT001	Lactobacillus rhamnosus
613	FP00632	FT001	Lactobacillus salivarius
614	FP00634	FT001	Lactobacillus viridescens
615	FP00635	FT008	Lactococcus garvieae



616	FP00636	FT008	<i>Lactococcus lactis</i> subsp. <i>cremoris</i>
617	FP00637	FT008	<i>Lactococcus lactis</i> subsp. <i>lactis</i>
618	FP00638	FT008	<i>Lactococcus plantarum</i>
619	FP00639	FT008	<i>Lactococcus raffinolactis</i>
620	FP00640	FT004	<i>Leclercia adecarboxylata</i>
621	FP00641	FT004	<i>Legionella anisa</i>
622	FP00642	FT004	<i>Legionella bozemanni</i>
623	FP00643	FT004	<i>Legionella dumoffii</i>
624	FP00644	FT004	<i>Legionella feeleei</i>
625	FP00645	FT004	<i>Legionella jordanis</i>
626	FP00646	FT004	<i>Legionella longbeachae</i>
627	FP00647	FT004	<i>Legionella micdadei</i>
628	FP00648	FT004	<i>Legionella oakridgensis</i>
629	FP00649	FT004	<i>Legionella pneumophila</i>
630	FP00650	FT004	<i>Legionella sainthelensi</i>
631	FP00651	FT004	<i>Legionella</i> spp.
632	FP00652	FT004	<i>Legionella wadsworthii</i>
633	FP00653	FT009	<i>Leptospira biflexa</i>
634	FP00654	FT009	<i>Leptospira interrogans</i>
635	FP00655	FT009	<i>Leptospira</i> spp.
636	FP00656	FT009	<i>Leptotrichia buccalis</i>
637	FP00657	FT009	<i>Leptotrichia</i> spp.
638	FP00658	FT008	<i>Leuconostoc carnosum</i>
639	FP00659	FT008	<i>Leuconostoc citreum</i>
640	FP00660	FT008	<i>Leuconostoc dextranicum</i>
641	FP00661	FT008	<i>Leuconostoc gelidum</i>
642	FP00662	FT008	<i>Leuconostoc lactis</i>
643	FP00663	FT008	<i>Leuconostoc mesenteroides</i>
644	FP00664	FT008	<i>Leuconostoc paramesenteroides</i>
645	FP00665	FT008	<i>Leuconostoc</i> spp.
646	FP00666	FT007	<i>Listeria grayi</i>
647	FP00667	FT007	<i>Listeria innocua</i>
648	FP00668	FT007	<i>Listeria ivanovii</i>
649	FP00669	FT007	<i>Listeria monocytogenes</i>
650	FP00670	FT007	<i>Listeria seeligeri</i>
651	FP00671	FT007	<i>Listeria</i> spp.
652	FP00672	FT007	<i>Listeria welshimeri</i>
653	FP00673	FT008	<i>Kocuria kristinae</i>
654	FP00674	FT008	<i>Micrococcus luteus</i>
655	FP00675	FT008	<i>Micrococcus lylae</i>
656	FP00676	FT008	<i>Micrococcus nishinomiyaensis</i>
657	FP00677	FT008	<i>Kocuria roseus</i>
658	FP00678	FT008	<i>Micrococcus sedentarius</i>
659	FP00679	FT008	<i>Kocuria varians</i>



660	FP00680	FT012	<i>Microsporum audouinii</i>
661	FP00681	FT012	<i>Microsporum canis</i>
662	FP00682	FT012	<i>Microsporum gypseum</i>
663	FP00683	FT012	<i>Microsporum</i> spp.
664	FP00684	FT001	<i>Mobiluncus mulieris</i>
665	FP00685	FT001	<i>Mobiluncus</i> spp.
666	FP00686	FT004	<i>Moellerella</i> spp.
667	FP00687	FT004	<i>Moellerella wisconsensis</i>
668	FP00688	FT006	<i>Moraxella atlantae</i>
669	FP00689	FT006	<i>Moraxella lacunata</i>
670	FP00690	FT006	<i>Moraxella nonliquefaciens</i>
671	FP00691	FT006	<i>Moraxella phenylpyruvica</i>
672	FP00692	FT006	<i>Moraxella</i> spp.
673	FP00693	FT004	<i>Morganella</i> spp.
674	FP00694	FT002	<i>Neisseria cinerea</i>
675	FP00695	FT002	<i>Neisseria elongata</i>
676	FP00696	FT002	<i>Neisseria flava</i>
677	FP00697	FT002	<i>Neisseria flavescens</i>
678	FP00698	FT002	<i>Neisseria lactamica</i>
679	FP00699	FT002	<i>Neisseria mucosa</i>
680	FP00700	FT002	<i>Neisseria perflava</i>
681	FP00701	FT002	<i>Neisseria polysaccharea</i>
682	FP00702	FT002	<i>Neisseria sicca</i>
683	FP00703	FT002	<i>Neisseria subflava</i>
684	FP00704	FT012	<i>Nigrospora</i> spp.
685	FP00705	FT006	<i>Oerskovia</i> spp.
686	FP00706	FT006	<i>Oligella</i> spp.
687	FP00707	FT006	<i>Oligella ureolytica</i>
688	FP00708	FT006	<i>Oligella urethralis</i>
689	FP00710	FT004	<i>Enterobacter agglomerans</i>
690	FP00711	FT004	<i>Pantoea dispersa</i>
691	FP00712	FT003	<i>Pasteurella aerogenes</i>
692	FP00713	FT003	<i>Pasteurella gallinarum</i>
693	FP00714	FT003	<i>Pasteurella haemolytica</i>
694	FP00715	FT003	<i>Pasteurella multocida</i>
695	FP00716	FT003	<i>Pasteurella pneumotropica</i>
696	FP00717	FT003	<i>Pasteurella</i> spp. new species 1
697	FP00718	FT003	<i>Pasteurella</i> spp.
698	FP00719	FT003	<i>Pasteurella ureae</i>
699	FP00720	FT008	<i>Pediococcus acidilactici</i>
700	FP00721	FT008	<i>Pediococcus damnosus</i>
701	FP00722	FT008	<i>Pediococcus halophilus</i>
702	FP00723	FT008	<i>Pediococcus pentosaceus</i>
703	FP00724	FT001	<i>Peptococcus niger</i>



704	FP00725	FT001	Peptococcus spp.
705	FP00726	FT001	Peptostreptococcus asaccharolyticus
706	FP00727	FT001	Peptostreptococcus indolicus
707	FP00728	FT012	Petriellidium spp.(Allescheria
708	FP00729	FT012	Phialophora jeanselmei
709	FP00730	FT012	Phialophora spp.
710	FP00731	FT012	Phialophora verrucosum
711	FP00732	FT011	Pichia carsonii
712	FP00733	FT011	Pichia etchellsii
713	FP00734	FT011	Pichia farinosa
714	FP00735	FT011	Pichia ohmeri
715	FP00736	FT011	Pichia spartinae
716	FP00737	FT011	Pichia spp.
717	FP00738	FT004	Plesiomonas shigelloides
718	FP00739	FT001	Porphyromonas endodontalis
719	FP00740	FT001	Prevotella buccalis
720	FP00741	FT001	Prevotella ruminicola
721	FP00742	FT001	Propionibacterium avidum
722	FP00743	FT001	Propionibacterium propionicus
723	FP00744	FT004	Providencia alcalifaciens
724	FP00745	FT004	Providencia heimbachae
725	FP00746	FT004	Providencia rustigianii
726	FP00747	FT006	Pseudomonas alcaligenes
727	FP00748	FT006	Pseudomonas aureofaciens
728	FP00749	FT006	Pseudomonas cepacia
729	FP00750	FT006	Pseudomonas chlororaphis
730	FP00751	FT006	Pseudomonas diminuta
731	FP00752	FT006	Pseudomonas group Ve-2
732	FP00753	FT006	Pseudomonas mallei
733	FP00754	FT006	Pseudomonas mesophilica
734	FP00755	FT006	Pseudomonas picketti biovar 1
735	FP00756	FT006	Pseudomonas picketti biovar 2
736	FP00757	FT006	Pseudomonas pseudoalcaligenes
737	FP00758	FT006	Pseudomonas pseudomallei
738	FP00759	FT006	Pseudomonas putrefaciens biova
739	FP00762	FT006	Pseudomonas vesicularis
740	FP00763	FT006	Pseudomonas-like
741	FP00764	FT012	Aureobasidium spp.
742	FP00765	FT004	Rahnella aquatilis
743	FP00766	FT011	Rhodotorula equi
744	FP00767	FT011	Rhodotorula minuta
745	FP00768	FT011	Rhodotorula pilimanae
746	FP00769	FT011	Rhodotorula rubra
747	FP00770	FT011	Saccharomyces kluyveri



748	FP00771	FT011	<i>Saccharomyces</i> spp.
749	FP00772	FT011	<i>Saccharomyces uvarum</i>
750	FP00773	FT004	<i>Salmonella arizonae</i>
751	FP00774	FT004	<i>Salmonella choleraesuis</i>
752	FP00775	FT004	<i>Salmonella enteritidis</i>
753	FP00776	FT004	<i>Salmonella gallinarum</i>
754	FP00777	FT004	<i>Salmonella paratyphi</i> A
755	FP00778	FT004	<i>Salmonella pullorum</i>
756	FP00779	FT004	<i>Salmonella typhisuis</i>
757	FP00780	FT004	<i>Salmonella typhmurium</i>
758	FP00781	FT004	<i>Salmonella</i> , other serotypes
759	FP00782	FT012	<i>Sepedonium</i> spp.
760	FP00783	FT004	<i>Serratia ficaria</i>
761	FP00784	FT004	<i>Serratia fonticola</i>
762	FP00785	FT004	<i>Serratia grimesii</i>
763	FP00786	FT004	<i>Serratia odorifera</i>
764	FP00787	FT004	<i>Serratia plymuthica</i>
765	FP00788	FT004	<i>Serratia proteamaculans</i>
766	FP00789	FT004	<i>Serratia proteamaculans</i> subsp. <i>quinovora</i>
767	FP00790	FT004	<i>Serratia rubidaea</i>
768	FP00791	FT004	<i>Shigella</i> spp.
769	FP00792	FT004	<i>Shigella boydii</i> (group C)
770	FP00793	FT004	<i>Shigella dysenteriae</i> (group A)
771	FP00794	FT004	<i>Shigella flexneri</i> (group B)
772	FP00795	FT004	<i>Shigella sonnei</i> (group D)
773	FP00797	FT006	<i>Sphingobacterium spiritivorum</i>
774	FP00798	FT011	<i>Sporobolomyces roseus</i>
775	FP00799	FT011	<i>Sporobolomyces salmonicolor</i>
776	FP00800	FT011	<i>Sporobolomyces</i> spp
777	FP00801	FT012	<i>Sporothrix schenckii</i>
778	FP00802	FT012	<i>Sporothrix</i> spp
779	FP00803	FT008	<i>Staphylococcus arlettae</i>
780	FP00805	FT008	<i>Staphylococcus auricularis</i>
781	FP00806	FT008	<i>Staphylococcus capitis</i>
782	FP00807	FT008	<i>Staphylococcus caprae</i>
783	FP00808	FT008	<i>Staphylococcus carnosus</i>
784	FP00809	FT008	<i>Staphylococcus chromogenes</i>
785	FP00810	FT008	<i>Staphylococcus cohnii</i>
786	FP00811	FT008	<i>Staphylococcus equorum</i>
787	FP00812	FT008	<i>Staphylococcus gallinarum</i>
788	FP00813	FT008	<i>Staphylococcus haemolyticus</i>
789	FP00814	FT008	<i>Staphylococcus hyicus</i>
790	FP00815	FT008	<i>Staphylococcus intermedius</i>
791	FP00816	FT008	<i>Staphylococcus kloosii</i>



792	FP00817	FT008	<i>Staphylococcus lentus</i>
793	FP00818	FT008	<i>Staphylococcus lugdunensis</i>
794	FP00819	FT008	<i>Staphylococcus schleiferi</i>
795	FP00820	FT008	<i>Staphylococcus sciuri</i>
796	FP00821	FT008	<i>Staphylococcus simulans</i>
797	FP00822	FT008	<i>Staphylococcus</i> spp.
798	FP00823	FT008	<i>Staphylococcus warneri</i>
799	FP00824	FT008	<i>Staphylococcus xylosus</i>
800	FP00825	FT012	<i>Stemphylium</i> spp.
801	FP00826	FT008	<i>Stomatococcus mucilaginosus</i>
802	FP00827	FT008	<i>Streptococcus acidominimus</i>
803	FP00828	FT008	<i>Streptococcus adjacens</i>
804	FP00829	FT008	<i>Streptococcus agalactiae</i>
805	FP00830	FT008	<i>Streptococcus alactolyticus</i>
806	FP00831	FT008	<i>Streptococcus anginosus</i>
807	FP00832	FT008	<i>Streptococcus bovis</i> biotype II
808	FP00833	FT008	<i>Streptococcus canis</i>
809	FP00834	FT008	<i>Streptococcus cecorum</i>
810	FP00835	FT008	<i>Streptococcus downei</i>
811	FP00836	FT008	<i>Streptococcus dysgalactiae</i>
812	FP00837	FT008	<i>Streptococcus equi</i> subspp. <i>equi</i>
813	FP00838	FT008	<i>Streptococcus equi</i> subspp. <i>zooepidemicus</i>
814	FP00839	FT008	<i>Streptococcus equisimilis</i>
815	FP00840	FT008	<i>Streptococcus gordonii</i>
816	FP00841	FT008	<i>Streptococcus</i> group E
817	FP00842	FT008	<i>Streptococcus</i> group G
818	FP00843	FT008	<i>Streptococcus</i> group L
819	FP00844	FT008	<i>Streptococcus</i> group P
820	FP00845	FT008	<i>Streptococcus</i> group U
821	FP00846	FT008	<i>Streptococcus milleri</i>
822	FP00847	FT008	<i>Streptococcus mitis</i>
823	FP00848	FT008	<i>Streptococcus mutans</i>
824	FP00849	FT008	<i>Streptococcus non-hemolyticus</i>
825	FP00850	FT008	<i>Streptococcus oralis</i>
826	FP00851	FT008	<i>Streptococcus procinus</i>
827	FP00852	FT008	<i>Streptococcus pyogenes</i>
828	FP00853	FT008	<i>Streptococcus saccharolyticus</i>
829	FP00854	FT008	<i>Streptococcus salivarius</i>
830	FP00855	FT008	<i>Globicatella sanguinis</i>
831	FP00856	FT008	<i>Streptococcus sobrinus</i>
832	FP00857	FT008	<i>Streptococcus</i> spp.
833	FP00858	FT008	<i>Streptococcus suis</i>
834	FP00859	FT008	<i>Streptococcus uberis</i>
835	FP00860	FT008	<i>Streptococcus vestibularis</i>



836	FP00861	FT008	<i>Streptococcus viridans</i>
837	FP00862	FT007	<i>Streptomyces somaliensis</i>
838	FP00863	FT002	<i>Suttonella indologenes</i>
839	FP00864	FT012	<i>Syncephalastrum</i> spp
840	FP00865	FT011	<i>Torulopsis pintolopesii</i>
841	FP00866	FT004	<i>Tatumella ptyseos</i>
842	FP00867	FT011	<i>Torulopsis</i> spp.
843	FP00868	FT012	<i>Trichoderma</i> spp.
844	FP00869	FT012	<i>Trichophyton mentagrophytes</i>
845	FP00870	FT012	<i>Trichophyton schoenleinii</i>
846	FP00871	FT012	<i>Trichophyton</i> spp.
847	FP00872	FT012	<i>Trichophyton tonsurans</i>
848	FP00873	FT012	<i>Trichophyton verrucosum</i>
849	FP00874	FT012	<i>Trichophyton violaceum</i>
850	FP00875	FT011	<i>Trichosporon pullulans</i>
851	FP00876	FT011	<i>Trichosporon</i> spp.
852	FP00877	FT999	<i>Ureaplasma</i> spp.
853	FP00878	FT999	<i>Ureaplasma urealyticum</i>
854	FP00879	FT001	<i>Veillonella alcalescens</i>
855	FP00880	FT004	<i>Vibrio alginolyticus</i>
856	FP00881	FT004	<i>Vibrio cholerae</i>
857	FP00882	FT004	<i>Vibrio damsela</i>
858	FP00883	FT004	<i>Vibrio fluvialis</i>
859	FP00884	FT004	<i>Vibrio furnissii</i>
860	FP00885	FT004	<i>Vibrio harveyi</i>
861	FP00886	FT004	<i>Vibrio hollisae</i>
862	FP00887	FT004	<i>Vibrio metschnikovii</i>
863	FP00888	FT004	<i>Vibrio mimicus</i>
864	FP00889	FT004	<i>Vibrio parahaemolyticus</i>
865	FP00890	FT004	<i>Vibrio</i> spp.
866	FP00891	FT006	<i>Weeksella</i> spp.
867	FP00892	FT006	<i>Weeksella virosa</i>
868	FP00893	FT006	<i>Weeksella zoohelcum</i>
869	FP00894	FT006	<i>Xanthomonas campestris</i>
870	FP00895	FT006	<i>Xanthomonas maltophilia</i>
871	FP00896	FT004	<i>Yersinia aldovae</i>
872	FP00897	FT004	<i>Yersinia enterocolitica</i>
873	FP00898	FT004	<i>Yersinia frederiksenii</i>
874	FP00899	FT004	<i>Yersinia intermedia</i>
875	FP00900	FT004	<i>Yersinia kristensenii</i>
876	FP00901	FT004	<i>Yersinia pestis</i>
877	FP00902	FT004	<i>Yersinia pseudotuberculosis</i>
878	FP00903	FT004	<i>Yersinia ruckeri</i>
879	FP00904	FT004	<i>Yersinia</i> spp.



880	FP00906	FT008	Streptococcus group D
881	FP00907	FT008	Oxacillin Resistant Staphylococcus aureus (ORSA)
882	FP00908	FT007	Corynebacterium striatum
883	FP00909	FT006	Sphingobacterium spp.
884	FP00910	FT008	Coagulase negative staphylococci
885	FP00911	FT007	Diphtheroids
886	FP00912	FT007	Corynebacterium macginleyi
887	FP00913	FT008	Alliococcus otitidis
888	FP00914	FT006	Pseudomonas luteola
889	FP00915	FT006	CDC GROUP IVC-2
890	FP00916	FT006	CDC GROUP EO-2
891	FP00917	FT006	CDC GROUP VB-3
892	FP00918	FT006	Empedobacter brevis
893	FP00919	FT004	pantoea spp.
894	FP00921	FT007	Corynebacterium renale group
895	FP00922	FT006	GNF (Unidentified GNF)
896	FP00923	FT006	Sphingomonas parapaucimobilis
897	FP00924	FT001	Porphyromonas spp.
898	FP00925	FT011	Yeast-non Candida albicans
899	FP00926	FT004	Salmonella Group A
900	FP00927	FT004	Salmonella Group E
901	FP00928	FT004	Salmonella Group C2
902	FP00929	FT004	Enterobacter hormaechei
903	FP00930	FT004	Enterobacter dissolvens
904	FP00931	FT004	Aeromonas jandaei
905	FP00932	FT004	Aeromonas enteropelogenes/ (trout)
906	FP00933	FT004	Acidaminococcus fermentans
907	FP00934	FT006	Brevundimonas diminuta
908	FP00935	FT006	Burkholderia gladioli
909	FP00936	FT006	Pseudomonas gladioli
910	FP00937	FT006	Burkholderia mallei
911	FP00938	FT006	Burkholderia pickettii
912	FP00939	FT006	Burkholderia spp.
913	FP00940	FT006	Brevundimonas vesicularis
914	FP00941	FT004	Chryseobacterium gleum
915	FP00942	FT004	Citrobacter braakii
916	FP00943	FT004	Citrobacter farmeri
917	FP00944	FT004	Citrobacter werkmanii
918	FP00945	FT004	Citrobacter youngae
919	FP00946	FT006	Flavobacterium spiritivorum
920	FP00947	FT008	Helcococcus kunzii
921	FP00948	FT004	Klebsiella ornithinolytica
922	FP00949	FT004	Raoultella ornithinolytica
923	FP00950	FT008	Lactococcus spp.



924	FP00951	FT006	Moraxella urethralis
925	FP00952	FT008	Streptococcus morbillorum
926	FP00953	FT008	Streptococcus group F
927	FP00954	FT006	Sphingobacterium multivorum
928	FP00955	FT001	Tissierella praeacuta
929	FP00956	FT001	Bacteroides praeacuta
930	FP00957	FT004	Vibrio cincinnatiensis
931	FP00958	FT001	Veillonella dispar
932	FP00959	FT004	Yokenella regensburgei
933	FP00960	FT001	Prevotella intermedia
934	FP00961	FT001	Prevotella oulori
935	FP00962	FT004	Photobacterium damsela
936	FP00963	FT007	GPB (Unidentified Gram positive bacillus)
937	FP00964	FT013	SARS virus
938	FP00965	FT006	Myroides odoratus
939	FP00966	FT004	Leminorella spp.
940	FP00967	FT008	Streptococcus fecalis
941	FP00968	FT008	B-Form streptococcus species
942	FP00969	FT007	Coryneform
943	FP00970	FT011	Kodamaea Ohmeri
944	FP00971	FT011	Candida dubliniensis
945	FP00972	FT011	Cryptococcus humicola
946	FP00973	FT007	Mycobacterium tuberculosis
947	FP00974	FT007	Mycobacterium chelonae
948	FP00975	FT007	Mycobacterium fortuitum
949	FP00976	FT001	Campylobacter jejuni
950	FP00977	FT016	Campylobacter coli
951	FP00978	FT008	Non- enterococcus
952	FP00979	FT008	Micrococcus
953	FP00980	FT003	Haemophilus influenzae non-type b
954	FP00981	FT008	$\alpha$ -hemolytic Streptococcus
955	FP00982	FT004	Yersinia bercovieri
956	FP00983	FT004	Yersinia mollaretii
957	FP00984	FT004	Vibrio carchariae
958	FP00985	FT004	Aeromonas schubertii
959	FP00986	FT004	Aeromonas trota
960	FP00987	FT003	Pasteurella bettyae
961	FP00988	FT003	Pasteurella canis
962	FP00989	FT003	Pasteurella dagmatis
963	FP00990	FT004	Pasteurella stomatis
964	FP00991	FT003	Pasteurella trehalosi
965	FP00992	FT003	Pasteurella volantium
966	FP00993	FT003	Capnocytophaga canimorsus
967	FP00994	FT003	Capnocytophaga cynodegmi



968	FP00995	FT004	Kingella oralis
969	FP00996	FT006	Moraxella canis
970	FP00997	FT006	Moraxella lincolnii
971	FP00998	FT006	Achromobacter piechaudii
972	FP00999	FT006	Sphingobacterium(Flavobacterium) spiritivorum
973	FP01000	FT006	Sphingobacterium thalpophilum
974	FP01001	FT006	Sphingobacterium mizutaii
975	FP01002	FT006	Flavobacterium yabuuchiae
976	FP01003	FT006	Comamonas terrigena
977	FP01004	FT004	Citrobacter sedlakii
978	FP01005	FT005	Bordetella holmesii
979	FP01006	FT004	Kingella oralis
980	FP01007	FT006	Acinetobacter radioresistens
981	FP01008	FT008	Staphylococcus muscae
982	FP01009	FT008	Staphylococcus piscifermentans
983	FP01010	FT008	Staphylococcus felis
984	FP01011	FT008	Staphylococcus delphini
985	FP01012	FT008	Staphylococcus caseolyticus
986	FP01013	FT008	Staphylococcus vitulinus
987	FP01014	FT008	Streptococcus mucilaginosus
988	FP01015	FT007	Corynebacterium mycetoides
989	FP01016	FT007	Corynebacterium genitalium
990	FP01017	FT007	Corynebacterium pseudogenitalium
991	FP01018	FT007	Erysipelothrix rhusiopathiae
992	FP01019	FT002	Neisseria parelongata
993	FP01020	FT001	Actinomyces hordeovulneris
994	FP01021	FT001	Actinomyces howellii
995	FP01022	FT001	Bacteroides coagulans
996	FP01023	FT001	Bacteroides forsythus
997	FP01024	FT001	Bacteroides macacae
998	FP01025	FT001	Bacteroides microfusis
999	FP01026	FT001	Bacteroides putredinis
1000	FP01027	FT001	Bacteroides salivarius
1001	FP01028	FT001	Bifidobacterium catenulatum
1002	FP01029	FT001	Bifidobacterium longum
1003	FP01030	FT004	Enterobacter cancerogenus
1004	FP01031	FT001	Clostridium absonum
1005	FP01032	FT001	Clostridium beijerinckii
1006	FP01033	FT001	Clostridium botulinum A
1007	FP01034	FT001	Clostridium botulinum B
1008	FP01035	FT001	Clostridium botulinum E
1009	FP01036	FT001	Clostridium botulinum G
1010	FP01037	FT001	Clostridium butyricum
1011	FP01038	FT001	Clostridium camis



1012	FP01039	FT001	<i>Clostridium celatum</i>
1013	FP01040	FT001	<i>Clostridium chauvoei</i>
1014	FP01041	FT001	<i>Clostridium haemolyticum</i>
1015	FP01042	FT004	<i>Enterobacter nimipressuralis</i>
1016	FP01043	FT001	<i>Clostridium lituseburense</i>
1017	FP01044	FT001	<i>Clostridium novyi</i> A
1018	FP01045	FT001	<i>Clostridium novyi</i> B
1019	FP01046	FT001	<i>Clostridium spiroforme</i>
1020	FP01047	FT007	<i>Corynebacterium matruchotii</i>
1021	FP01048	FT001	<i>Fusobacterium naviforme</i>
1022	FP01049	FT001	<i>Fusobacterium periodonticum</i>
1023	FP01050	FT001	<i>Fusobacterium russii</i>
1024	FP01051	FT001	<i>Fusobacterium simiae</i>
1025	FP01052	FT001	<i>Fusobacterium sulci</i>
1026	FP01053	FT001	<i>Mobiluncus curtisii</i> subsp. <i>holmesii</i>
1027	FP01054	FT001	<i>Prevotella veroralis</i>
1028	FP01055	FT001	<i>Prevotella zoogloiformans</i>
1029	FP01056	FT001	<i>Propionibacterium acidipropionici</i>
1030	FP01057	FT001	<i>Propionibacterium freudenreichii</i>
1031	FP01058	FT001	<i>Propionibacterium jensenii</i>
1032	FP01059	FT001	<i>Propionibacterium lymphophilum</i>
1033	FP01060	FT001	<i>Propionibacterium thoenii</i>
1034	FP01061	FT001	<i>Porphyromonas lerii</i>
1035	FP01062	FT011	<i>Candida iusitaniae</i>
1036	FP01063	FT012	<i>Microsporum cookei</i>
1037	FP01064	FT012	<i>Microsporum distortum</i>
1038	FP01065	FT012	<i>Microsporum nanum</i>
1039	FP01066	FT012	<i>Microsporum vanbreuseghemii</i>
1040	FP01067	FT012	<i>Trichophyton ajelloi</i>
1041	FP01068	FT012	<i>Trichophyton metagrophytes</i>
1042	FP01069	FT012	<i>Trichophyton rubrum</i>
1043	FP01070	FT012	<i>Trichophyton schoenleinii</i>
1044	FP01071	FT012	<i>Trichophyton soudanense</i>
1045	FP01072	FT012	<i>Trichophyton terrestre</i>
1046	FP01073	FT012	<i>Cladosporium</i> type sporulation
1047	FP01074	FT012	<i>Phialophora</i> type sporulation
1048	FP01075	FT012	<i>Coccidioides immitis</i>
1049	FP01076	FT012	<i>Aspergillus glaucus</i>
1050	FP01077	FT012	<i>Aspergillus versicolor</i>
1051	FP01078	FT012	<i>Aspergillus nidulans</i>
1052	FP01079	FT012	<i>Chrysosporium keratinophilum</i>
1053	FP01080	FT012	<i>Chrysosporium pannorum</i>
1054	FP01081	FT012	<i>Chrysosporium pruinsum</i>
1055	FP01082	FT012	<i>Chrysosporium tropicum</i>



1056	FP01083	FT012	Chrysosporium parvum
1057	FP01084	FT012	Epicoccum nigrum
1058	FP01085	FT012	Fusarium solani
1059	FP01086	FT012	Helminthosporium solani
1060	FP01087	FT012	Nigrospora oryzae
1061	FP01088	FT012	Scopulariopsis brevicaulis
1062	FP01089	FT012	Syncephalastrum racemosum
1063	FP01090	FT004	Proteus myxofaciens
1064	FP01091	FT004	Salmonella,group C2,C3
1065	FP01092	FT004	Salmonella group D1,not typhi
1066	FP01093	FT004	Serratia entomophila
1067	FP01094	FT004	Yersinia rohdei
1068	FP01095	FT016	Arcobacter nitrofigilis
1069	FP01096	FT001	Eubacterium brachy
1070	FP01097	FT001	Eubacterium budavi
1071	FP01098	FT001	Eubacterium combesii
1072	FP01099	FT001	Eubacterium cylindroides
1073	FP01100	FT001	Eubacterium dolichum
1074	FP01101	FT001	Eubacterium formicigenerans
1075	FP01102	FT001	Eubacterium nodatum
1076	FP01103	FT001	Eubacterium saburreum
1077	FP01104	FT001	Eubacterium timidum
1078	FP01105	FT001	<del>Eubacterium spp.</del>
1079	FP01106	FT012	Stemphylium sarcinaeforme
1080	FP01107	FT012	Cunninghamella bertholletiae
1081	FP01108	FT007	Mycobacterium, non-tuberculosis (NTM)
1082	FP01109	FT008	Enterococcus raffinosus
1083	FP01110	FT007	Mycobacterium abscessus
1084	FP01111	FT007	Mycobacterium africanum
1085	FP01112	FT007	Mycobacterium avium complex (MAC)
1086	FP01113	FT007	Mycobacterium bovis
1087	FP01114	FT007	Mycobacterium canetti
1088	FP01115	FT007	Mycobacterium kansasii
1089	FP01116	FT007	Mycobacterium marinum
1090	FP01117	FT007	Mycobacterium microti
1091	FP01118	FT012	Unidentified fungus
1092	FP01119	FT013	Human respiratory syncytial virus (RSV)
1093	FP01120	FT013	Norovirus
1094	FP01121	FT013	Rotavirus
1095	FP01122	FT012	Mycelia sterila
1096	FP01123	FT013	Adenovirus
1097	FP01124	FT013	Parainfluenza virus
1098	FP01125	FT007	Corynebacterium propinquum
1099	FP01126	FT004	Escherichia coli (ESBL)



1100	FP01127	FT004	Klebsiella oxytoca (ESBL)
1101	FP01128	FT004	Klebsiella pneumoniae (ESBL)
1102	FP01129	FT004	Proteus mirabilis (ESBL)
1103	FP01130	FT006	Acinetobacter calcoaceticus-Acinetobacter baumannii complex
1104	FP01131	FT006	Roseomonas spp.
1105	FP01132	FT007	Kurthia spp.
1106	FP01133	FT008	Globicatella spp.
1107	FP01134	FT014	Chlamydophila pneumoniae
1108	FP01135	FT013	Influenza virus
1109	FP01136	FT006	Ralstonia spp.
1110	FP01137	FT007	Mycobacterium spp.
1111	FP01138	FT006	Acinetobacter ursingii
1112	FP01139	FT008	Rhodococcus spp.
1113	FP01140	FT008	Enterococcus dispar
1114	FP01141	FT006	Chryseobacterium indoltheticum
1115	FP01142	FT013	Coxsackievirus
1116	FP01143	FT013	Echovirus
1117	FP01144	FT013	Enterovirus
1118	FP01145	FT013	Poliovirus
1119	FP01146	FT006	Shewanella algae
1120	FP01147	FT008	Leuconostoc mesenteroides ssp. cremoris
1121	FP01148	FT012	Beauveria spp.
1122	FP01149	FT006	Myroides spp.
1123	FP01150	FT006	Ochrobactrum spp.
1124	FP01151	FT012	Conidiobolus spp.
1125	FP01152	FT004	Wolinella spp.
1126	FP01153	FT011	Malassezia spp.
1127	FP01154	FT006	Shewanella spp.
1128	FP01155	FT006	Chryseobacterium spp.
1129	FP01156	FT007	Actinomadura spp.
1130	FP01157	FT006	Sphingomonas spp.
1131	FP01158	FT007	Leifsonia aquatica
1132	FP01159	FT008	Leuconostoc pseudomesenteroides
1133	FP01160	FT012	Trichosporon inkin
1134	FP01161	FT007	Corynebacterium amycolatum
1135	FP01162	FT007	Corynebacterium urealyticum
1136	FP01163	FT012	Malassezia furfur
1137	FP01164	FT008	Granulicatella adiacens
1138	FP01165	FT011	Stephanoascus ciferrii
1139	FP01166	FT008	Streptococcus sanguinis
1140	FP01167	FT006	Acinetobacter nosocomialis
1141	FP01168	FT006	Acinetobacter pittii
1142	FP01169	FT004	Roseomonas gilardii
1143	FP01170	FT012	Phialemonium spp.



1144	FP01171	FT004	Brevundimonas species
1145	FP01172	FT007	Dermabacter hominis
1146	FP01173	FT006	Chryseobacterium meningosepticum
1147	FP01174	FT006	Pseudomonas oryzihabitans
1148	FP01175	FT004	Streptococcus parasanguinis
1149	FP01176	FT008	Streptococcus infantarius
1150	FP01177	FT006	Cupriavidus pauculus
1151	FP01178	FT001	Eggerthella lenta
1152	FP01179	FT001	Weissella confuse
1153	FP01180	FT011	Trichosporon asahii
1154	FP01181	FT006	Bergeyella zoohelcum
1155	FP01182	FT008	Parvimonas micra
1156	FP01183	FT007	Corynebacterium glucuronolyticum
1157	FP01184	FT007	Arthrobacter woluwensis
1158	FP01185	FT006	Ralstonia mannitolilytica
1159	FP01186	FT001	Paenibacillus durus
1160	FP01187	FT012	Mold
1161	FP01188	FT001	Bilophila spp.
1162	FP01189	FT007	Microbacterium paraoxydans
1163	FP01190	FT007	Mycobacterium tuberculosis complex
1164	FP01191	FT001	Peptoniphilus harei
1165	FP01192	FT004	Arcobater butzleri
1166	FP01193	FT001	Elizabethkingia miricola
1167	FP01194	FT008	Streptococcus gallolyticus
1168	FP01195	FT006	Pseudomonas mosselii
1169	FP01196	FT006	Burkholderia multivorans
1170	FP01197	FT001	Pseudoflavonifractor capillosus
1171	FP01198	FT008	Rothia mucilaginosa
1172	FP01199	FT004	Proteus hauseri
1173	FP01200	FT013	Pan-Enterovirus
1174	FP01201	FT008	Kocuria species
1175	FP01202	FT008	Rhodococcus erythropolis
1176	FP01203	FT001	Paenibacillus pabuli
1177	FP01204	FT006	Acinetobacter baylyi
1178	FP01205	FT011	Rhizomucor spp.
1179	FP01206	FT004	Roseomonas mucosa
1180	FP01207	FT004	Salmonella Group C
1181	FP01208	FT007	Corynebacterium tuberculostearicum
1182	FP01209	FT011	Candida metapsilosis
1183	FP01210	FT011	Rhodotorula mucilaginosa
1184	FP01211	FT007	Paenibacillus spp.
1185	FP01212	FT006	Pseudomonas putida group
1186	FP01213	FT007	Facklamia hominis
1187	FP01214	FT006	Burkholderia vietnamiensis



1188	FP01215	FT011	Candida haemulonii
1189	FP01216	FT004	Enterobacter cloacae complex
1190	FP01217	FT007	Turicella otitidis
1191	FP01218	FT001	Peptoniphilus species
1192	FP01219	FT006	Acinetobacter guillouiae
1193	FP01220	FT006	Burkholderia cenocepacia
1194	FP01221	FT006	Achromobacter denitrificans
1195	FP01222	FT008	Lactococcus lactis
1196	FP01223	FT001	Actinomyces radingae
1197	FP01224	FT001	Paenibacillus provencensis
1198	FP01225	FT007	Microbacterium spp.
1199	FP01226	FT007	Arcanobacterium pyogenes
1200	FP01227	FT006	Pseudomonas monteilii
1201	FP01228	FT001	Peptoniphilus asaccharolyticus
1202	FP01229	FT007	Corynebacterium resistens
1203	FP01230	FT006	Pseudomonas gessardii
1204	FP01231	FT004	Pseudoxanthomonas mexicana
1205	FP01232	FT001	Prevotella nigrescens
1206	FP01233	FT001	Prevotella timonensis
1207	FP01234	FT012	Candida palmiophila
1208	FP01235	FT011	Debaryomyces species
1209	FP01236	FT006	Acinetobacter parvus
1210	FP01237	FT001	Anaerococcus tetradius
1211	FP01238	FT007	Microbacterium saperdae
1212	FP01239	FT007	Tsukamurella spp.
1213	FP01240	FT012	Candida orthopsilosis
1214	FP01241	FT006	Acinetobacter gyllenbergii
1215	FP01242	FT006	Paracoccus yeei
1216	FP01243	FT002	Gemella spp.
1217	FP01244	FT008	Streptococcus sanguinis
1218	FP01245	FT999	Mycoplasma hominis
1219	FP01246	FT010	Acanthamoeba
1220	FP01247	FT007	Peptostreptococcus intermedius
1221	FP01248	FT012	Chaetomium sp
1222	FP01249	FT012	Geotrichum sp
1223	FP01250	FT012	Exophiala sp
1224	FP01251	FT999	Mycoplasma sp
1225	FP01252	FT004	Acidaminococcus sp
1226	FP01253	FT008	Actinobaculum schaalii
1227	FP01254	FT016	Actinobaculum urinale
1228	FP01255	FT001	Alloscardovia omnicolens
1229	FP01256	FT001	Alistipes sp
1230	FP01257	FT001	Anaerococcus hydrogenalis
1231	FP01258	FT001	Anaerococcus murdochii



1232	FP01259	FT001	Anaerococcus octavius
1233	FP01260	FT001	Anaerococcus sp
1234	FP01261	FT001	Anaerococcus vaginalis
1235	FP01262	FT001	Anaerotruncus colihominis
1236	FP01263	FT003	Atopobium minutum
1237	FP01264	FT003	Atopobium parvulum
1238	FP01265	FT003	Atopobium sp
1239	FP01266	FT001	Bifidobacterium animalis
1240	FP01267	FT001	Bifidobacterium boum
1241	FP01268	FT001	Bulleidia extructa
1242	FP01269	FT004	Butyricimonas virosa
1243	FP01270	FT001	Campylobacter curvus
1244	FP01271	FT001	Campylobacter rectus
1245	FP01272	FT001	Campylobacter showae
1246	FP01273	FT006	Cellulosimicrobium cellulans
1247	FP01274	FT001	Collinsella aerofaciens
1248	FP01275	FT001	Coprobacillus cateniformis
1249	FP01276	FT001	Dialister micraerophilus
1250	FP01277	FT001	Dialister pneumosintes
1251	FP01278	FT001	Eggerthia cateniformis
1252	FP01279	FT001	Eubacterium callanderi
1253	FP01280	FT001	Flavonifractor plautii
1254	FP01281	FT001	Lachnoanaerobaculum orale
1255	FP01282	FT001	Lachnoanaerobaculum saburreum
1256	FP01283	FT001	Lachnoanaerobaculum umeaense
1257	FP01284	FT009	Leptotrichia trevisanii
1258	FP01285	FT001	Megamonas sp
1259	FP01286	FT001	Megasphaera micronuciformis
1260	FP01287	FT001	Olsenella uli
1261	FP01288	FT001	Parabacteroides goldsteinii
1262	FP01289	FT001	Parabacteroides johnsonii
1263	FP01290	FT001	Peptoniphilus gorbachii
1264	FP01291	FT001	Ruminococcus gnavus
1265	FP01292	FT001	Slackia exigua
1266	FP01293	FT001	Solobacterium moorei
1267	FP01294	FT012	Debaryomyces hansenii
1268	FP01295	FT012	Lodderomyces elongisporus
1269	FP01296	FT012	Magnusiomyces capitatus
1270	FP01297	FT012	Debaryomyces etchellsii
1271	FP01298	FT006	Kerstersia gyiorum
1272	FP01299	FT008	Staphylococcus agalactiae
1273	FP01300	FT011	Clavispora lusitaniae
1274	FP01301	FT008	Streptococcus gallolyticus subspp. Pasteurian
1275	FP01302	FT008	Gemella sanguinis



1276	ZP00001	FT999	No Growth
1277	ZP00002	FT999	No Culture
1278	ZP00003	FT999	Normal Flora
1279	ZP00004	FT999	Mixed flora
1280	ZP00005	FT999	No Significant



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[illegible]



[illegible]



[illegible]



[illegible]



	已刪除
	已刪除
氣味類香菌	
糞鏈球菌	
Beta-溶血性鏈球菌	已刪除
Candida dubliniensis	
土生隱球菌	
M.tuberculosis	
M.chelonae	
M.fortuitum	
Campylobacter jejuni	
Campylobacter coli	已刪除
Non-enterococcus	已刪除
Micrococcus	已刪除
Haemophilus influenzae	
Yersinia bercovieri	
Yersinia mollaretii	
Vibrio carchariae	
Aeromonas schubertii	
Aeromonas trota	
Pasteurella bettyae	
Pasteurella canis	
Pasteurella dagmatis	
Pasteurella stomatis	
Pasteurella trehalos	
Pasteurella volantis	
Capnocytophaga canim	
Capnocytophaga cynod	



[illegible]



[illegible]



[illegible]



[illegible]



[illegible]



[illegible]



106/12/11新增	
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107/01/24新增	
107/01/24新增	
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107/01/24新增	



Mixed flora	
No Significant	