Antimicrobial resistance - 1: The plasmid er	ncodes ARGs that confe	r resistance to the drug	class {label}.			
AM1000 aminocoumarin antibiotic			AM1002 carbapenem		AM1003 cephalosporin	
M1004 cephamycin AM1005 diaminopyrimidine antib			1		ng agents and antise	
AM1007 fluoroquinolone antibiotic	AM1008 fusidane antibiotic		AM1009 glycopept		AM1010 lincosamide antibiotic	
AM1011 macrolide antibiotic			AM1013 mupirocin		AM1014 nitrofurar	n antibiotic
AM1015 nucleoside antibiotic	1015 nucleoside antibiotic AM1016 oxazolidinone antibiotic		AM1017 penam	AM1018 penem	AM1019 peptide a	ntibiotic
AM1020 phenicol antibiotic AM1021 phosphonic acid antibiotic		acid antibiotic	AM1022 pleuromut		AM1023 rifamycir	
AM1024 streptogramin antibiotic AM1025 sulfonamide			AM1026 sulfone an		AM1027 tetracycli	
Antimicrobial resistance - 2: The plasmid er						
AM2000 ErmB AM2001 IMP-1	AM2002 IMP-4	AM2003 MecA	AM2004 NDM-1	AM2005 OXA-10		
AM2006 OXA-181	AM2007 OXA-23		AM2008 OXA-24	11112000 0111110	AM2009 OXA-48	
AM2010 OXA-58	AM2011 QacE	AM2012 QnrA1	AM2013 QnrB19		AM2014 QnrS1	AM2015 SHV-1
AM2016 TEM-1 AM2017 Tet(M)	THILDIT QUEE	AM2018 VIM-1	AM2019 VIM-2	AM2020 VIM-4	AM2021 VanA	AM2022 VanB
Antimicrobial resistance - 3: The plasmid er	acodes the antibiotic resi			71112020 VIIVI 4	711VIZOZI Vali71	71112022 Valid
AM3000 ABC Transporter	AM3001 Acetyltrans	ferase		coside Modifying En	zvme	
AM3003 Antibiotic Inactivation	AM3004 Chloramphe		AM3005 Class A B		AM3006 Class B F	Reta_Lactamase
AM3007 Class C Beta-Lactamase	AM3008 Class D Bet		AM3009 Gene Mod		AM3010 Gylcoper	
AM3011 MFS Transporter	AM3012 Nucleotidyl		AM3013 Other Effl		AM3010 Gylcoper	
AM3015 Quinolone Resistance	AM3016 RND Antib				AM3014 Thosphot	
AM3019 Tetracycline MFS Efflux		Ribosomal Protection	AM3017 Target Pro	AM3021 rRNA Mo		me macuvation
				AMI3021 IKNA MI	euryttransferase	
Antimicrobial resistance - 4: The plasmid ca AM4000 amikacin	AM4001 aminoglyco				AM4002	
	0,	side	AM4002 apramycir		AM4003 arsenate	
AM4004 arsenic	AM4005 arsenite		AM4006 azithromy		AM4007 beta-lacta	
AM4008 bleomycin	AM4009 cadmium	1251044	AM4010 chloramph		AM4011 clindamy	cın
AM4012 colistin	AM4013 copper	AM4014 erythromy		AM4015 florfenico	AM4019 hygromy	
AM4016 fosfomycin		AM4017 gentamicin		AM4018 glycopeptide		
AM4020 kanamycin	AM4021 lincosamide		AM4022 linezolid		AM4023 macrolide	
AM4024 mercury	AM4025 mupirocin		AM4026 nickel	AM4027 organome		
AM4028 oxazolidinone	AM4029 phenicol		AM4030 pleuromutilin		AM4031 quaternary ammonium	
AM4032 quinolone	AM4033 rifamycin		AM4034 silver	AM4035 spiramyc	in	
AM4036 streptogramin	AM4037 streptogram	in b	AM4038 streptomy	cin	AM4039 streptothi	ricin
AM4040 sulfonamide	AM4041 telithromyc	in	AM4042 tellurium		AM4043 tetracycline	
AM4044 tiamulin	AM4045 tigecycline		AM4046 tobramyci	n	AM4047 trimethop	orim
AM4048 tylosin	AM4049 vancomycir		AM4050 virginiam	ycin		
Bacterial host range - 1: The plasmid is host	ed by bacteria in the phy	/lum {label}.				
HO1000 Actinomycetota	HO1001 Bacillota	HO1002 Bacteroido	ta	HO1003 Campylol	oacterota	
HO1004 Chlamydiota	HO1005 Cyanobacter		HO1006 Deinococc		HO1007 Fusobacte	riota
HO1008 Methanobacteriota	HO1009 Mycoplasma		HO1010 Pseudomo	nadota	HO1011 Spirochae	
HO1012 Thermodesulfobacteriota	HO1013 Verrucomica					
Bacterial host range - 2: The plasmid is host						
HO2000 Actinomycetes	HO2001 Alphaprotec		HO2002 Bacilli	HO2003 Bacteroid	ia	
HO2004 Betaproteobacteria	HO2005 Clostridia		HO2006 Cyanophy		HO2007 Cytophag	ia
HO2008 Deinococci			HO2010 Erysipelot		HO2011 Flavobact	
HO2012 Fusobacteriia		1 1		HO2014 Halobacteria		
Bacterial host range - 3: The plasmid is host		eopaciena	I HO2014 Halobacte	ria	HO2015 Spirochae	tia
	ted by bacteria in the ord		HO2014 Halobacter	ria	HO2015 Spirochae	tia
		ler {label}.		ia	1	
HO3000 Aeromonadales	HO3001 Alteromona	er {label}. dales	HO3002 Bacillales		HO3003 Bacteroid	ales
HO3000 Aeromonadales HO3004 Bifidobacteriales	HO3001 Alteromona HO3005 Burkholderi	er { label }. dales ales	HO3002 Bacillales HO3006 Campylob	acterales	HO3003 Bacteroid HO3007 Cytophag	ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte	er {label}. dales ales rales	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot	acterales richales	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri	ales ales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria	ler {label}. dales ales rales ales	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacter	acterales richales riales	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomio	ales ales ales crobiales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira	er {label}. dales ales rales dles	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte HO3018 Lactobacil	acterales richales riales lales	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell	ales ales ales crobiales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera	er {label}. dales ales rales des des	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte HO3018 Lactobacil HO3022 Micrococc	acterales richales riales lales ales	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella	ales ales ales crobiales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales	er {label}. dales ales rales ales les	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocales	acterales richales riales lales ales	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell	ales ales ales crobiales ales ales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales HO3028 Peptostreptococcales	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales HO3029 Pseudomona	er {label}. dales ales rales ales les les	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocales HO3030 Rhodobaci	acterales richales riales lales ales erales	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell HO3031 Rhodospi	ales ales ales crobiales ales ales ales ales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales HO3028 Peptostreptococcales HO3032 Sphingomonadales	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales HO3029 Pseudomona HO3033 Spirochaeta	er {label}. dales ales rales ales les les les	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocales	acterales richales riales lales ales erales	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell	ales ales ales crobiales ales ales ales ales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales HO3028 Peptostreptococcales HO3032 Sphingomonadales Bacterial host range - 4: The plasmid is host	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales HO3029 Pseudomona HO3033 Spirochaetal	er {label}. dales ales rales ales les les les adales	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte: HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocales HO3030 Rhodobact HO3034 Thiotricha	acterales richales riales lales ales serales	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell HO3031 Rhodospi HO3035 Vibrional	ales ales ales crobiales ales ales ales ales ales ales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales HO3028 Peptostreptococcales HO3032 Sphingomonadales Bacterial host range - 4: The plasmid is host HO4000 Acetobacteraceae	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales HO3029 Pseudomona HO3033 Spirochaetal ted by bacteria in the fan	ler {label}. dales ales rales ales les les les adales les adales adales es adales es adales	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte: HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocale: HO3030 Rhodobact HO3034 Thiotricha	acterales richales riales lales ales serales les ceae	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell HO3031 Rhodospi HO3035 Vibrional	ales ales ales crobiales ales ales ales ales ales ales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales HO3028 Peptostreptococcales HO3032 Sphingomonadales Bacterial host range - 4: The plasmid is host HO4000 Acetobacteraceae HO4004 Bacteroidaceae	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales HO3029 Pseudomona HO3033 Spirochaetal ted by bacteria in the fan HO4001 Aeromonada HO4005 Bifidobacter	er {label}. dales ales rales ales les les les adales adales es adales es adales es adales	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte: HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocale: HO3030 Rhodobact HO3034 Thiotricha HO4002 Alcaligena HO4006 Borreliace	acterales richales riales lales ales serales les ceae	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell HO3031 Rhodospi HO3035 Vibrional	ales ales ales crobiales ales ales ales ales ales ales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales HO3028 Peptostreptococcales HO3032 Sphingomonadales Bacterial host range - 4: The plasmid is host HO4000 Acetobacteraceae HO4004 Bacteroidaceae HO4008 Burkholderiaceae	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales HO3029 Pseudomona HO3033 Spirochaetal ted by bacteria in the fan HO4001 Aeromonada HO4005 Bifidobacter HO4009 Campylobac	er {label}. dales dales ales rales ales les les les dales es dales	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte: HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocale: HO3030 Rhodobact HO3034 Thiotricha HO4002 Alcaligena HO4006 Borreliace HO4010 Clostridiae	acterales richales riales lales ales serales les ceae ae	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell HO3031 Rhodospi HO3035 Vibrional	ales ales ales crobiales ales ales ales ales ales ales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales HO3028 Peptostreptococcales HO3032 Sphingomonadales Bacterial host range - 4: The plasmid is host HO4000 Acetobacteraceae HO4004 Bacteroidaceae HO4008 Burkholderiaceae HO4012 Corynebacteriaceae	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales HO3029 Pseudomona HO3033 Spirochaetal ed by bacteria in the fan HO4001 Aeromonada HO4005 Bifidobacter HO4009 Campylobac HO4013 Enterobacte	er {label}. dales dales ales rales ales les les les dales es dales	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte: HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocale: HO3030 Rhodobact HO3034 Thiotricha HO4002 Alcaligena HO4006 Borreliace HO4010 Clostridiac HO4014 Enterococc	acterales richales riales lales ales serales les ceae ae ceae ae ceae	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell HO3031 Rhodospi HO3035 Vibrional HO4003 Bacillacea HO4007 Brucellace HO4011 Comamon HO4015 Erwiniacea	ales ales ales crobiales ales ales ales ales ales ales ales
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HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales HO3028 Peptostreptococcales HO3032 Sphingomonadales Bacterial host range - 4: The plasmid is host HO4000 Acetobacteraceae HO4004 Bacteroidaceae HO4008 Burkholderiaceae HO4012 Corynebacteriaceae HO4016 Fusobacteriaceae HO4020 Lactobacillaceae	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales HO3029 Pseudomona HO3033 Spirochaetal ted by bacteria in the fan HO4001 Aeromonada HO4005 Bifidobacter HO4009 Campylobac HO4013 Enterobacte HO4017 Haloferacac HO4021 Legionellace	er {label}. dales dales ales rales ales les les les dales es dales dales es dales dales es dales da	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte: HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocale: HO3030 Rhodobacil HO3034 Thiotricha HO4002 Alcaligena HO4006 Borreliace HO4010 Clostridiac HO4014 Enterococc HO4018 Helicobacil HO4022 Leptospira	acterales richales riales lales ales serales les ceae ae eeae eraceae eraceae ceae	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell HO3031 Rhodospi HO3035 Vibrional HO4003 Bacillacea HO4007 Brucellac HO4011 Comamon HO4015 Erwiniacea HO4019 Lachnosp	ales ales ales crobiales ales ales ales ales ales ales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales HO3028 Peptostreptococcales HO3032 Sphingomonadales Bacterial host range - 4: The plasmid is host HO4000 Acetobacteraceae HO4004 Bacteroidaceae HO4008 Burkholderiaceae HO4012 Corynebacteriaceae HO4016 Fusobacteriaceae HO4020 Lactobacillaceae HO4024 Lysobacteraceae	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales HO3029 Pseudomona HO3033 Spirochaetal ed by bacteria in the fan HO4001 Aeromonada HO4005 Bifidobacter HO4009 Campylobac HO4013 Enterobacte HO4017 Haloferacac HO4021 Legionellace HO4025 Methylobac	er {label}. dales dales ales rales ales les les les dales es dales dales es dales dales es dales da	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte: HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocale: HO3030 Rhodobacil HO3034 Thiotricha HO4002 Alcaligena HO4006 Borreliace HO4010 Clostridiac HO4014 Enterococc HO4018 Helicobacil HO4022 Leptospira HO4026 Microbact	acterales richales riales lales ales serales les ceae ae ceae eraceae ceae eriaceae eriaceae	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell HO3031 Rhodospi HO3035 Vibrional HO4003 Bacillacea HO4007 Brucellac HO4011 Comamon HO4015 Erwiniacea HO4019 Lachnosp HO4023 Listeriacea HO4027 Micrococe	ales ales ales ales crobiales ales ales ales ales ales ales ales
HO3000 Aeromonadales HO3004 Bifidobacteriales HO3008 Deinococcales HO3012 Flavobacteriales HO3016 Kitasatosporales HO3020 Leptospirales HO3024 Mycobacteriales HO3028 Peptostreptococcales HO3032 Sphingomonadales Bacterial host range - 4: The plasmid is host HO4000 Acetobacteraceae HO4004 Bacteroidaceae HO4008 Burkholderiaceae HO4012 Corynebacteriaceae HO4016 Fusobacteriaceae HO4020 Lactobacillaceae	HO3001 Alteromona HO3005 Burkholderi HO3009 Enterobacte HO3013 Fusobacteria HO3017 Lachnospira HO3021 Lysobactera HO3025 Neisseriales HO3029 Pseudomona HO3033 Spirochaetal ted by bacteria in the fan HO4001 Aeromonada HO4005 Bifidobacter HO4009 Campylobac HO4013 Enterobacte HO4017 Haloferacac HO4021 Legionellace	ler {label}. dales ales rales ales les les les adales es adales es adales es adales es acae eriaceae eriaceae eriaceae eriaceae eriaceae eriaceae eriaceae eriaceae	HO3002 Bacillales HO3006 Campylob HO3010 Erysipelot HO3014 Halobacte: HO3018 Lactobacil HO3022 Micrococc HO3026 Nostocale: HO3030 Rhodobacil HO3034 Thiotricha HO4002 Alcaligena HO4006 Borreliace HO4010 Clostridiac HO4014 Enterococc HO4018 Helicobacil HO4022 Leptospira	acterales richales riales lales ales serales les ceae ae ceae eraceae ceae eriaceae eriaceae	HO3003 Bacteroid HO3007 Cytophag HO3011 Eubacteri HO3015 Hyphomi HO3019 Legionell HO3023 Moraxella HO3027 Pasteurell HO3031 Rhodospi HO3035 Vibrional HO4003 Bacillacea HO4007 Brucellac HO4011 Comamon HO4015 Erwiniacea HO4019 Lachnosp	ales ales ales ales crobiales ales ales ales ales ales ales ales

HO4036 Paenibacillaceae	HO4037 Paracoccace	296	HO4038 Pasteurella	rceae	HO4039 Pectobac	teriaceae
HO4040 Peptostreptococcaceae	HO4041 Phyllobacteriaceae		HO4042 Piscirickettsiaceae		HO4043 Prevotellaceae	
HO4044 Pseudomonadaceae	HO4045 Rhizobiaceae		HO4046 Roseobacteraceae		HO4047 Sphingomonadaceae	
HO4048 Staphylococcaceae	HO4049 Streptococcaceae		HO4050 Streptomycetaceae		HO4051 Vibriona	
HO4052 Yersiniaceae	Tro to 15 Sureptococc			HO4030 Streptomycetaceae		cuc
Bacterial host range - 5: The plasmid is hos	ted by bacteria in the ge	nus {lahel}				
HO5000 Acetobacter	HO5001 Acinetobaci		HO5002 Aeromona	6	HO5003 Agrobact	arium
HO5004 Bacillus	HO5001 Acmetobacter HO5005 Bacteroides		HO5006 Bordetella		HO5007 Agrobact	Cituiii
HO5004 Bachius HO5008 Borreliella	HO5009 Bradyrhizol		HO5000 Bordetena HO5010 Brucella		HO5007 Borrella HO5011 Burkhold	ani a
		Diulii		1		
HO5012 Campylobacter HO5016 Corynebacterium	HO5013 Citrobacter HO5017 Cronobacte		HO5014 Clostridioi HO5018 Ensifer	HO5019 Enteroba	HO5015 Clostridiu	ım
HO5020 Enterococcus	HO5021 Erwinia	HO5022 Escherichia		HO5023 Faecaliba		.,
HO5024 Fusobacterium	HO5025 Helicobacte		HO5026 Klebsiella	,	HO5027 Komagat	
HO5028 Lacticaseibacillus	HO5029 Lactiplantib	pacillus	HO5030 Lactobacil	****	HO5031 Lactococ	
HO5032 Legionella	HO5033 Leptospira		HO5034 Leuconoste		HO5035 Ligilacto	
HO5036 Listeria	HO5037 Mesorhizob		HO5038 Methyloba		HO5039 Microcoo	
HO5040 Moraxella	HO5041 Morganella		HO5042 Mycobacte	erium	HO5043 Mycobac	teroides
HO5044 Mycolicibacterium	HO5045 Neisseria		HO5046 Nocardia		HO5047 Nostoc	
HO5048 Novosphingobium	HO5049 Pantoea	HO5050 Paraburkho		HO5051 Paracocci		
HO5052 Phaeobacter	HO5053 Phocaeicola	<u> </u>	HO5054 Photobacte	erium	HO5055 Pisciricke	ettsia
HO5056 Prescottella	HO5057 Prevotella		HO5058 Priestia		HO5059 Proteus	
HO5060 Providencia	HO5061 Pseudomon	as	HO5062 Ralstonia		HO5063 Rhizobiu	m
HO5064 Rhodococcus	HO5065 Salmonella		HO5066 Segatella		HO5067 Serratia	
HO5068 Shigella	HO5069 Sinorhizobi		HO5070 Sphingobii		HO5071 Sphingor	
HO5072 Staphylococcus	HO5073 Stenotropho	omonas	HO5074 Streptococ		HO5075 Streptom	yces
HO5076 Tritonibacter	HO5077 Vibrio	HO5078 Xanthomor	nas	HO5079 Yersinia	•	
Ecological host - 1: The plasmid can be foun	d in ecosystems associa	ted with hosts of the pl	hylum {label }.	•		•
EH1000 Arthropoda EH1001 Bacillario	phyta	EH1002 Chordata	EH1003 Streptophy	ta		
Ecological host - 2: The plasmid can be foun	id in ecosystems associa	ted with hosts of the cl	ass {label}.			
EH2000 Actinopteri	EH2001 Arachnida		EH2002 Aves		EH2003 Insecta	
EH2004 Magnoliopsida	EH2005 Malacostrac	a	EH2006 Mammalia			
Ecological host - 3: The plasmid can be foun	d in ecosystems associa	ted with hosts of the or				
EH3000 Anseriformes	EH3001 Artiodactyla		EH3002 Carnivora		EH3003 Decapoda	l
EH3004 Fabales EH3005 Galliform		EH3006 Hymenopte	era	EH3007 Ixodida	EH3008 Poales	
EH3009 Primates	EH3010 Rodentia	, , , , , , , , , , , , , , , , , , , ,	EH3011 Rosales	EH3012 Salmonife		
EH3013 Solanales						1
Ecological host - 4: The plasmid can be foun	d in ecosystems associa	ted with hosts of the fa	mily {label}.			
EH4000 Bovidae EH4001 Canidae	EH4002 Fabaceae	EH4003 Hominidae		EH4004 Penaeidae	a,	
EH4005 Phasianidae	EH4006 Poaceae	EH4007 Salmonidae		EH4008 Solanacea		EH4009 Suidae
Ecological host - 5: The plasmid can be foun				ZII.000 Bolundeet		
EH5000 Bos EH5001 Canis			enus {lahel}			
Elisoot Edins	I E.H. YOU / CTAILING	EH5003 Homo				
Ecological host - 6: The plasmid can be foun	EH5002 Gallus	EH5003 Homo	EH5004 Sus			
Ecological host - 6: The plasmid can be foun	nd in ecosystems associa		EH5004 Sus			
EH6000 Homo sapiens EH6001 Bacteria	d in ecosystems associa EH6002 Eukaryota	ted with hosts of the {I	EH5004 Sus abel}.			
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for	in ecosystems associa EH6002 Eukaryota und in pathogens associa	ted with hosts of the {I attention with the {label} has	EH5004 Sus abel}.	EC1003 Food	FC1004 Human	
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds	ted with hosts of the {I ted with hosts of the {I ted with the {Iabel} hated with the {Iabel} hated EC1002 Environment	EH5004 Sus abel}.	EC1003 Food	EC1004 Human	
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other	ted with hosts of the {I ated with the {label} ha EC1002 Environment EC1007 Rodent	EH5004 Sus abel}. abitat.	EC1003 Food	EC1004 Human	
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for Ecosystem range - 2: The Ecos	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the	abel}. abitat. nt PLSDB scheme.		1	
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture	abel}. abitat. nt PLSDB scheme.	EC2003 estuary	EC2004 food	
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine	abel Sus abel Sus abel Sus abel Sus abel Sus abel Sus abitat. PLSDB scheme. EC2008 saline		1	
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for Ecosystem range - 3: The Plasmid can be for Ecosystem range - 3: The plasmid can be for Ecosystem range - 3: The plasmid can be for Ecosystem range - 3: The plasmid can be for Ecosystem range - 3: The plasmid can be for Ecosystem range - 3: The plasmid can be for Ecosystem range - 3: The plasmid can be for Ecosystem range - 3: The plasmid can b	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine stem, according to the	abel Sus abel Sus abel Sus abel Sus abel Sus abel Sus abitat. PLSDB scheme. EC2008 saline	EC2003 estuary EC2009 sea	EC2004 food EC2010 spring	
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine	EH5004 Sus abel}. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme.	EC2003 estuary EC2009 sea EC3003 aquatic	EC2004 food EC2010 spring EC3004 arthropod	a: insects
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy c EC3006 engineered	ted with hosts of the {I atted with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine stem, according to the EC3002 annelida	EH5004 Sus abel }. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environment	EC2003 estuary EC2009 sea EC3003 aquatic	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair	a: insects
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system EC3009 freshwater	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy c EC3006 engineered EC3010 gut	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine stem, according to the EC3002 annelida EC3011 host-associa	EH5004 Sus abel }. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmentated	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents	a: insects
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system EC3009 freshwater EC3013 large intestine	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy c EC3006 engineered	ted with hosts of the {I atted with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine stem, according to the EC3002 annelida EC3011 host-associa EC3015 mammals	EH5004 Sus abel }. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environment	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine	a: insects
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system EC3009 freshwater EC3013 large intestine EC3019 non-marine saline and alkaline	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy c EC3006 engineered EC3010 gut EC3014 larva	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine stem, according to the EC3002 annelida EC3011 host-associa EC3015 mammals EC3020 oral cavity	EH5004 Sus abel }. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmentated	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human EC3021 palsa	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine EC3022 peat	a: insects
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system EC3009 freshwater EC3013 large intestine EC3019 non-marine saline and alkaline EC3023 peat moss	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy c EC3006 engineered EC3010 gut	ted with hosts of the {I atted with the {Iabel} ha EC1002 Environment EC1007 Rodent Stem, according to the EC2002 cell culture EC2007 marine Stem, according to the EC3002 annelida EC3001 host-associa EC3015 mammals EC3020 oral cavity EC3025 porifera	EH5004 Sus abel }. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmentated EC3016 mammals:	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human EC3021 palsa EC3026 reproduct	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine EC3022 peat ive system	a: insects
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system EC3009 freshwater EC3013 large intestine EC3019 non-marine saline and alkaline	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy c EC3006 engineered EC3010 gut EC3014 larva	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine stem, according to the EC3002 annelida EC3011 host-associa EC3015 mammals EC3020 oral cavity	EH5004 Sus abel }. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmentated EC3016 mammals:	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human EC3021 palsa EC3026 reproduct communities (contig	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine EC3022 peat ive system mixture)	a: insects EC3018 modelec
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system EC3009 freshwater EC3013 large intestine EC3019 non-marine saline and alkaline EC3023 peat moss	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy c EC3006 engineered EC3010 gut EC3014 larva	ted with hosts of the {I atted with the {Iabel} ha EC1002 Environment EC1007 Rodent Stem, according to the EC2002 cell culture EC2007 marine Stem, according to the EC3002 annelida EC3001 host-associa EC3015 mammals EC3020 oral cavity EC3025 porifera	EH5004 Sus abel }. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmentated EC3016 mammals:	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human EC3021 palsa EC3026 reproduct	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine EC3022 peat ive system mixture)	a: insects
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system EC3009 freshwater EC3013 large intestine EC3019 non-marine saline and alkaline EC3023 peat moss EC3027 rock-dwelling (endoliths)	d in ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy c EC3006 engineered EC3010 gut EC3014 larva	ted with hosts of the {I atted with the {Iabel} ha EC1002 Environment EC1007 Rodent Stem, according to the EC2002 cell culture EC2007 marine Stem, according to the EC3002 annelida EC3001 host-associa EC3015 mammals EC3020 oral cavity EC3025 porifera	EH5004 Sus abel }. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmentated EC3016 mammals:	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human EC3021 palsa EC3026 reproduct communities (contig	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine EC3022 peat ive system mixture)	a: insects EC3018 modelec
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system EC3009 freshwater EC3013 large intestine EC3019 non-marine saline and alkaline EC3023 peat moss EC3027 rock-dwelling (endoliths) EC3030 simulated communities (microbial masses)	din ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy EC3006 engineered EC3010 gut EC3014 larva EC3024 plants	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine stem, according to the EC3002 annelida EC3011 host-associa EC3015 mammals EC3020 oral cavity EC3025 porifera EC3028 roots	EH5004 Sus abel }. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmented EC3016 mammals: EC3029 simulated of EC3031 soil	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human EC3021 palsa EC3026 reproduct communities (contig	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine EC3022 peat ive system mixture)	a: insects EC3018 modeled EC3033 sponge
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system EC3009 freshwater EC3013 large intestine EC3019 non-marine saline and alkaline EC3027 rock-dwelling (endoliths) EC3030 simulated communities (microbial material ecas)	din ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy EC3006 engineered EC3010 gut EC3014 larva EC3024 plants aixture) sourced from {label}.	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine stem, according to the EC3002 annelida EC3011 host-associa EC3015 mammals EC3020 oral cavity EC3025 porifera EC3028 roots EC3035 stomach	EH5004 Sus abel }. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmented EC3016 mammals: EC3029 simulated of EC3031 soil	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human EC3021 palsa EC3026 reproduct communities (contig	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine EC3022 peat ive system mixture)	a: insects EC3018 modeled EC3033 sponge EC3038 vagina
EH6000 Homo sapiens EH6001 Bacteria Ecosystem range - 1: The plasmid can be for EC1000 Arthropod EC1005 Mammal Ecosystem range - 2: The plasmid can be for EC2000 anthropogenic EC2005 host associated Ecosystem range - 3: The plasmid can be for EC3000 algae EC3001 anaerobic EC3005 digestive system EC3009 freshwater EC3013 large intestine EC3019 non-marine saline and alkaline EC3023 peat moss EC3027 rock-dwelling (endoliths) EC30304 simulated communities (microbial material property - 1: The plasmid is seen CH1000 Isolate CH1001 Metagene	din ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy EC3006 engineered EC3010 gut EC3014 larva EC3024 plants aixture) sourced from {label}.	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine stem, according to the EC3002 annelida EC3011 host-associa EC3015 mammals EC3020 oral cavity EC3025 porifera EC3028 roots EC3035 stomach	EH5004 Sus abel}. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmentated EC3016 mammals: EC3029 simulated c EC3031 soil EC3036 terrestrial	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human EC3021 palsa EC3026 reproduct communities (contig	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine EC3022 peat ive system mixture) el EC3037 tundra	a: insects EC3018 modeled EC3033 sponge EC3038 vagina
EH6000 Homo sapiens	din ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy EC3006 engineered EC3010 gut EC3014 larva EC3024 plants mixture) sourced from {label}.	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine EC3002 annelida EC3011 host-associa EC3015 mammals EC3020 oral cavity EC3025 porifera EC3028 roots EC3035 stomach	EH5004 Sus abel}. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmented EC3016 mammals: EC3029 simulated of EC3031 soil EC3036 terrestrial	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human EC3021 palsa EC3026 reproduct communities (contig EC3032 solar pane	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine EC3022 peat ive system mixture) el EC3037 tundra	a: insects EC3018 modeled EC3033 sponge EC3038 vagina
EH6000 Homo sapiens	din ecosystems associa EH6002 Eukaryota und in pathogens associa EC1001 Birds EC1006 Other und in the {label} ecosy EC2001 aquatic EC2006 location und in the {label} ecosy EC3006 engineered EC3010 gut EC3014 larva EC3024 plants aixture) courced from {label}. come opology, gram-staining in the electric pathogeness associated as	ted with hosts of the {I ated with the {Iabel} ha EC1002 Environment EC1007 Rodent stem, according to the EC2002 cell culture EC2007 marine EC3002 annelida EC3011 host-associa EC3015 mammals EC3020 oral cavity EC3025 porifera EC3028 roots EC3035 stomach	EH5004 Sus abel}. abitat. nt PLSDB scheme. EC2008 saline IMG/M scheme. EC3007 environmented EC3016 mammals: EC3029 simulated of EC3031 soil EC3036 terrestrial me-assembled genome sts, and phage-plasmic	EC2003 estuary EC2009 sea EC3003 aquatic ntal EC3012 hydrother human EC3021 palsa EC3026 reproduct communities (contig EC3032 solar pane	EC2004 food EC2010 spring EC3004 arthropod EC3008 floodplair mal vents EC3017 marine EC3022 peat ive system mixture) el EC3037 tundra	a: insects EC3018 modeled EC3033 sponge EC3038 vagina

CH2002 The plasmid co	ntains direct terminal	repeat structures		CH2003 Fragmente	d plasmid		
	The plasmid contains direct terminal repeat structures. The plasmid is hosted by Gram-negative bacteria.				id is hosted by Gram	n-positive bacteria	
CH2004 The plasmid is nosted by Gram-negative bacteria. CH2006 The plasmid contains inverted terminal repeat structures.			CH2007 Phage-plas		CH2008 circular p	lasmid	
CH2009 linear plasmid				CH2007 Fliage-plas	siiiu	CH2008 circular plasmid	
Heavy metal resistance	- 1. The placmid and	odes genes that confer	recistance to flahall				
ME1000 Aluminium (Al				ME1002 A	1-1	ME1002 D:	(D:)
	,			ME1002 Arsenic (A		ME1003 Bismuth (Bi)	
ME1004 Cadmium (Cd)			(Cr)	ME1006 Cobalt (Co	0)	ME1007 Copper (Cu)	
	E1008 Gallium (Ga) ME1009 Gold (Au)			ME1010 Iron (Fe)		ME1011 Lead (Pb)	
ME1012 Magnesium (M	012 Magnesium (Mg) ME1013 Manganese		· /	ME1014 Mercury (C/	ME1015 Molybdenum (Mo)	
ME1016 Nickel (Ni)			Se) ME1018 Silver (Ag)		ME1019 Tellurium (Te)		
ME1020 Tungsten (W)		ME1021 Vanadium ((V)	ME1022 Zinc (Zn)			
Heavy metal resistance	- 2: The plasmid enc	odes the metal resistance	ce gene {label}.	•			
ME2000 G2alt	ME2001 acn	ME2002 acr3	ME2003 actP	ME2004 actR	ME2005 aioA (ao	xB)	ME2006 aioE
ME2007 aioR (aoxR)	1	ME2008 arsA	ME2009 arsB	ME2010 arsC	ME2011 arsD	ME2012 arsH	ME2013 arsM
ME2014 arsP	ME2015 arsR	ME2016 arsT	ME2017 baeR	ME2018 baeS	ME2019 bhsA (co		ME2020 cadC
ME2021 cadD	ME2022 cadX	ME2023 chrA	ME2024 chrA1	ME2025 chrB	ME2026 chrB1	ME2027 chrF	ME2028 chrR
ME2021 cadD ME2029 cmtR	ME2030 comR (yo		ME2024 CITAT			ME2027 CHIT	ME2034 copC
		~	1		ME2032 copA		
ME2035 copD	ME2036 copF	ME2037 copG	ME2038 copM	ME2039 copR	ME2040 copS	ME2041 copY (ter	(Y)
ME2042 copZ	ME2043 corA	ME2044 corC	ME2045 corR	ME2046 corR (coal		ME2047 corS	
ME2048 corT (coaT)		ME2049 crdR	ME2050 csoR	ME2051 ctpV	ME2052 cueA	ME2053 cueP	
ME2054 cusA (ybdE)		ME2055 cusB	ME2056 cusC (ylcB	<u> </u>	ME2057 cusF (cus		
ME2058 cusR (ylcA)		ME2059 cusS	ME2060 cutA	ME2061 czcA	ME2062 czcB	ME2063 czcC	ME2064 czcD
ME2065 czcR	ME2066 czcS	ME2067 dmeF	ME2068 dmeR	ME2069 dpsA	ME2070 dsbA	ME2071 dsbC	ME2072 fbpB
ME2073 fbpC	ME2074 fecD	ME2075 fecE	ME2076 fetA (ybbL	.)	ME2077 fetB (ybb	oM)	
ME2078 fieF (yiip)		ME2079 fptA	ME2080 fpvA	ME2081 gesA	ME2082 gesB	ME2083 gesC	ME2084 glpF
ME2085 golS	ME2086 golT	ME2087 hmrR	ME2088 hoxN	ME2089 irlR	ME2090 irlS	ME2091 klaB (kil	
ME2092 klaC (telB)	1/122000 g011	ME2093 kmtR	ME2094 mco	ME2095 mdtA	ME2096 merA	ME2097 merB	ME2098 merC
ME2099 merD	ME2100 merE	ME2101 merF	ME2102 merG	ME2103 merP	ME2104 merR	ME2105 merR1	ME2106 merR2
ME2107 merT	ME2108 mgtA	ME2109 mntH (yfeF		ME2110 modA	ME2111 modB	ME2112 modC	ME2113 mreA
ME2114 ncrA	ME2115 ncrB	ME2116 ncrC	ME2117 ncrY	ME2118 nikA	ME2119 nikB	ME2120 nikC	ME2121 nikD
ME2122 nikE	ME2123 nirD	ME2124 nixA	ME2125 nrsD (nreB	·	ME2126 nrsS	ME2127 pbrA	
ME2128 pbrB (pbrC)		ME2129 pbrR	ME2130 pbrT	ME2131 pcoA	ME2132 pcoB	ME2133 pcoC	ME2134 pcoD
ME2135 pcoE	ME2136 pcoR	ME2137 pcoS	ME2138 pfr	ME2139 pgpA (ltpg	gpA)	ME2140 pitA	ME2141 pmrA
ME2142 pmrB	ME2143 pmrC	ME2144 pmrG	ME2145 pstA	ME2146 pstB	ME2147 pstC	ME2148 pstS	
ME2149 rcnA (yohM)		ME2150 rcnB (yohN	<u>i)</u>	ME2151 rcnR (yoh	L)	ME2152 recG	ME2153 ricR
ME2154 robA	ME2155 ruvB	ME2156 silA	ME2157 silB	ME2158 silC	ME2159 silE	ME2160 silF	ME2161 silP
ME2162 silR	ME2163 silS	ME2164 sitA	ME2165 sitB	ME2166 sitC	ME2167 sitD	ME2168 smtB (zia	
ME2169 soxS	ME2170 tcrA	ME2171 tcrB	ME2172 terY	ME2173 tcrZ	ME2174 tehA	ME2175 tehB	ME2176 terA
ME2177 terB	ME2178 terC	ME2179 terD	ME2180 terE	ME2181 terW	ME2182 terZ	ME2183 troB	ME2184 tunR
ME2185 tupC	ME2178 tere	ME2177 terb ME2187 ybtP	ME2188 ybtQ	ME2189 yfeA	ME2192 tel2 ME2190 yfeB	ME2191 yfeD	ME2192 yfmP
ME2193 ygiW	ME2194 yhcN	ME2195 yieF	ME2196 yjaA	ME2197 yodD	ME2198 yqjH	ME2199 zevB	ME2200 ziaR
ME2201 zinT (yodA)		ME2202 zitB (ybgR)		ME2203 zntR (yhd	M)	ME2204 znuC (ye	bM)
ME2205 zraR (hydH)		ME2206 zraS (hydG		ME2207 zur (yjbK)			
Incompatibility group -							
IN1000 IncA/C2	IN1001 IncB/O/K/	Z					IN1006 IncHI1A
IN1007 IncHI1B	IN1008 IncHI2	IN1009 IncHI2A	IN1010 IncI	IN1011 IncI1	IN1012 IncI2	IN1013 IncL/M	IN1014 IncN
IN1015 IncN2	IN1016 IncP1	IN1017 IncP6	IN1018 IncQ1	IN1019 IncR	IN1020 IncU	IN1021 IncX1	IN1022 IncX2
IN1023 IncX3	IN1024 IncX4	IN1025 IncX5	IN1026 IncY		1		
Incompatibility group -				ding to the Plasmid M	ILST scheme		
IN2000 IncA/C	IN2001 IncF	IN2002 IncHI1	IN2003 IncHI2	IN2004 IncI1	IN2005 IncN		
Incompatibility group -							
						INI200C II. PIC	IN12007 I PII
IN3000 Inc11	IN3001 Inc13	IN3002 Inc18	IN3003 IncC	IN3004 IncFIA	IN3005 IncFIB	IN3006 IncFIC	IN3007 IncFII
IN3008 IncHI1A	IN3009 IncHI1B	IN3010 IncHI2A	IN3011 IncI-gamma		IN3012 IncI1	IN3013 IncI1/B/O	
IN3014 IncI2	IN3015 IncK2/Z	IN3016 IncL/M	IN3017 IncN	IN3018 IncP	IN3019 IncQ1	IN3020 IncR	IN3021 IncU
IN3022 IncX1	IN3023 IncX3	IN3024 IncX4	IN3025 IncY				
Mobility - 1: Three plas	mid mobility categori	es.					
MO1000 conjugative pla	smid	MO1001 mobilizable	e plasmid	MO1002 non-mobi	lizable plasmid		
Mobility - 2: The plasm	id is transferable and	belongs to mobilization	type {label}, according	g to the CONJScan so	heme.	•	
MO2000 MOBB	MO2001 MOBC	MO2002 MOBF	MO2003 MOBH	MO2004 MOBP1		MO2005 MOBP2	
MO2006 MOBP3		MO2007 MOBQ	MO2008 MOBT	MO2009 MOBV		1 3.102000 MODI 2	
	id is transferable and				rheme		
Mobility - 3. The place	io is nansitiable allu	ociongs to mounization				1402006 MODO	MO3007 MOBT
Mobility - 3: The plasm		MO2002 MODE	1 M(O2002 M/ODII				
MO3000 MOBB	MO3001 MOBC	MO3002 MOBF	MO3003 MOBH	MO3004 MOBM	MO3005 MOBP	MO3006 MOBQ	MO3007 MOBT
	MO3001 MOBC MO3009 MOB_Ui	nknown		I	I		

MO4000 MPF_B	MO4001 MPF_C		MO4002 MPF_F		MO4003 MPF FA	
MO4004 MPF_FATA	MO4005 MPF_G		MO4006 MPF_I		MO4007 MPF T	
Mobility - 5: The plasmid encodes a complet	nd belongs to the matin		{label}, according to	_	me.	
MO5000 MPF F MO5001 MPF I	MO5002 MPF_T	MO5003 MPF Unk		(label), according to	the WOD typer sene	inc.
Mobility - 6: The plasmid encodes the {label			nown			
MO6000 T4SS ATPase F_traU	MO6001 T4SS ATPa		MO6002 T4SS ATF	Pace virb/		
MO6003 Type IV coupling protein t4cp1	WOOOOT 1433 ATT		oupling protein t4cp2	asc viio+		
MO6005 Type IV coupling protein teach		WIO0004 Type IV co	oupling protein t-tep2			
Risk index - 1: {label} risk based on the num	han of incontion common	and hambanad by the mla	amid			
RI1000 Minimal RI1001 Low	RI1002 Moderate	RI1003 High	Siliia.			
Risk index - 2: {label} risk based on the distr			i.			
RI2000 Minimal RI2001 Low	RI2002 Moderate	RI2003 High	1 '1			
Risk index - 3: {label} risk based on the num			olasmid.			
RI3000 Minimal RI3001 Low	RI3002 Moderate	RI3003 High				
Risk index - 4: {label} risk based on the num			T			
RI4000 Minimal RI4001 Low	RI4002 Moderate	RI4003 High				
Risk index - 5: {label} risk based on the num			d by the plasmid.			
RI5000 Minimal RI5001 Low	RI5002 Moderate	RI5003 High				
Risk index -6: {label} risk based on the taxon						
RI6000 Minimal RI6001 Low	RI6002 Moderate	RI6003 High				
Risk index -7: Combined {label} risk level.						
RI7000 Minimal RI7001 Low	RI7002 Moderate	RI7003 High				
Virulence factor - 1: The plasmid carries vir						
VF1000 Adherence		al activity/Competitive			VF1002 Biofilm	
VF1003 Effector delivery system	VF1004 Exoenzyme	- *	VF1005 Exotoxin	VF1006 Immune n	nodulation	VF1007 Invasion
VF1008 Motility VF1009 Nutritions	al/Metabolic factor		VF1010 Other	VF1011 Regulation	n	
VF1012 Stress survival						
Virulence factor - 2: The plasmid carries the	virulence factor {label}	}.				
VF2000 <alpha>-Hemolysin</alpha>	VF2001 AAFs	VF2002 AAI/SCI-II	T6SS	VF2003 ACE T6S	S	VF2004 AS
VF2005 AatA, AIDA-I type	VF2006 Adhesive fir		VF2007 Aerobactin		VF2008 Afa/Dr fai	
VF2009 Afimbrial adhesin AFA-I	1 VI 2000 Fidnesive in	VF2010 Agf	VF2011 AggR	VF2012 Agglutinii		
VF2013 Allantion utilization	VF2014 Anthrax tox		VF2015 Antigen 43		Песерия	VF2016 ApeE
VF2017 AtxA VF2018 Aureolysi		VF2019 Autolysin	V1 2013 Antigen 43	VF2020 BFP	VF2021 Bcf	VI ZOTO APCE
VF2017 AXA VF2018 Auteoryst VF2022 Bee (biofilm enhancer in enterococci		VF2019 Autolysiii VF2023 BimA	VF2024 Bsa T3SS	V12020 B11	VI 2021 BCI	
VF2025 Bsa T3SS secreted effectors)	VF2026 BslA	VF2024 Bsa 1333 VF2027 C3610	VF2028 CDT	VF2029 CNF-1	
VF2030 CS31A capsule-like antigen		VF2031 Cah, AIDA		VF2032 Capsule	VF2033 Clumping	factor
VF2030 CS31A capsule-like allugeli VF2034 ClyA VF2035 Col10	VF2036 Col5	VF2037 Call, AIDA VF2037 Colibactin	-1 type	VF2032 Capsule VF2038 Colicin B	VF2033 Ciumping	Tactor
	VF2030 Colicin Ia	VF2037 Collbactill	VE2041 Caliain II	VF2038 COIICIII B	VE2042 Colinin V	
VF2039 Colicin E1			VF2041 Colicin Ib		VF2042 Colicin K	
VF2043 Colicin N	VF2044 Colicin S4		VF2045 Colicin U		VF2046 Colicin Y	
VF2047 Contact-dependent inhibition CDI sy			VF2048 Csu fimbria		VF2049 Curli fiber	'S
VF2050 Cytolysin	VF2051 Direct heme			VF2052 Dispersin		
VF2053 Dot/Icm T4SS secreted effectors		VF2054 Dr adhesins		VF2055 ECP	VF2056 EHS	VF2057 ESX-1
VF2058 ESX-3 VF2059 ESX-5	VF2060 ETT2	VF2061 EaeH	VF2062 Ebp pili		VF2063 Efa-1/Lif	
VF2064 EhaA, AIDA-I type	VF2065 EhaB, AIDA		VF2066 Emp	VF2067 Ent	VF2068 Enterobac	tin
VF2069 Enterobactin synthesis and transport		VF2070 Enterotoxin	SenB/TieB	VF2071 EpeA	VF2072 EspC	
VF2073 EspI, SPATE	VF2074 EspP	VF2075 Etp	VF2076 EtpA	VF2077 F1C fimbi	riae	
VF2078 F9 fimbriae	VF2079 Ferrous iron	1	VF2080 Flagella		VF2081 FnBPs	
VF2082 GacS/GacA two-component system		VF2083 Gsp	VF2084 HBL	VF2085 Hbp (hem	oglobin-binding prot	ease)
VF2086 Heat-labile toxin (LT)	VF2087 Heat-stable	toxin (ST)	VF2088 HemO clus	ster	VF2089 Hemolysis	n
VF2090 Hyaluronate lyase	VF2091 Hyaluronic	acid (HA) capsule		VF2092 Ibes	VF2093 IcsA (Vir	G)
VF2094 IcsP (SopA)	VF2095 IgA1 protea	se	VF2096 Insecticida	crystalline toxins	· · · · · · · · · · · · · · · · · · ·	
VF2097 Intercellular adhesion proteins		VF2098 Intimin	VF2099 Iron-regula			1
VF2100 Iron/manganese transport		VF2101 K1 capsule		VF2102 LPS	VF2103 Ler	
VF2104 Locus for diffuse adherence (lda), afi	mbrial adhesin	1	VF2105 Lpf	VF2106 LukED	-	1
VF2107 Lvh (Legionella vir homologs) type l			1 r-	VF2108 MgtBC	VF2109 Mig-14	VF2110 Mig-5
VF2111 MisL VF2112 MsbB2	VF2113 OmpD	VF2114 P fimbriae		VF2115 Paa	VF2116 PagN	12110 11119 0
VF2117 PagR-XO1	VF2118 Pef	VF2114 Printipliac	s flagella	VF2120 Pet	VF2121 Pic	
VF2117 1 agr-XO1 VF2122 PilA-type pili (PGS1, pilin gene clus		, , z , j , i criticious	VF2123 PilB-type p		VF2124 Pix pilus	I
VF2122 I IIA-type piii (I GS1, piiiii gene clus VF2125 Pyoverdine	VF2126 Quorum-ser	neina	VF2123 Tilb-type p	VF2128 Rck	VF2124 Fix pilus VF2129 RcsAB	VF2130 RmpA
VF2131 RpoS VF2132 S fimbriae	•		onella centrisome islan		VF2129 RCSAB VF2134 SCI-I T6S	1
VF2131 RpoS VF2132 S IIIIIO1146 VF2135 SDr VF2136 SE	VF2137 Saf	`			V1:2134 SCI-1 108	u .
VI:Z133 3D1 VFZ130 3E	1 V C Z L 3 / 3 a f	VF2138 Sal	VF2139 Salmocheli			
		VE2141 C-4	VE0140 C	anna amil A DCDEE		
VF2140 Salmochelin siderophore	1	VF2141 Sat	VF2142 Serine prot		WE2140 C B	
VF2140 Salmochelin siderophore VF2143 Sfp fimbriae	VF2144 ShET2	VF2145 ShdA	VF2146 SigA	VF2147 SinH	VF2148 SpvB	VE0154 T000
VF2140 Salmochelin siderophore	1					VF2154 T2SS

VF2155 T3SS	VF2156 T3SS1	VF2157 T3SS1 secreted effectors			VF2158 T3SS2		
VF2159 T3SS2 secreted effectors			VF2160 T6SS	VF2161 T6SS-1	VF2162 T6SS-II	VF2163 T6SS-III	
VF2164 TFS3a	VF2165 TFS3b	VF2166 TTS1	VF2167 TTS2	VF2168 TTSS	VF2169 TTSS (SP	I-1 encode)	
VF2170 TTSS (SPI-2 encode) VF2171 TTSS secrete			ed effectors		VF2172 TTSS-1 secreted effectors		
VF2173 TTSS-2 secreted effectors			VF2174 Tcf	VF2175 Thermonuc	elease nuc VF2176 Tia/Hek VF2177		VF2177 ToxB
VF2178 TraJ	VF2179 Tsh	VF2180 Type 1 fimbriae VF2181 Type 3 fin		oriae VF2182 Type I fimbriae			
VF2183 Type IV pili VF2184 Type IV secr			retion system		VF2185 Type VII s	secretion system	
VF2186 Vacuolating autotransporter gene Vat			VF2187 VirF	VF2188 VirK	VF2189 Ybt	VF2190 Yersiniaba	ctin
VF2191 Yersiniabactin siderophore							