

Supplementary Material

Potential risks of antibiotic resistant bacteria and genes in bioremediation of petroleum hydrocarbon contaminated soils

Colin J. Cunningham^a, Maria S. Kuyukina*^{b, c}, Irena B. Ivshina^{b, c}, Alexandr I. Konev^b, Tatyana A. Peshkur^d, Charles W. Knapp^d

^a Sustainability Fife Ltd, Glenrothes, UK

^b Microbiology and Immunology Department, Perm State University, 15 Bukirev Street, 614990 Perm, Russia

^c Institute of Ecology and Genetics of Microorganisms, Ural Branch of the Russian Academy of Sciences, 13 Golev Street, 614081 Perm, Russia

^d Department of Civil and Environmental Engineering, University of Strathclyde, Glasgow, UK

*Corresponding author.

Professor Maria S. Kuyukina

Microbiology and Immunology Department

Perm State University, 15 Bukirev Street, 614990 Perm, Russia

Tel.: (+7)3422808114, Fax: (+7)3422809211

E-mail address: kuyukina@iegm.ru

Table S1. Abundance of ARGs in most important hydrocarbon-degrading *Actinobacteria* genera (totally 13 genera, 851 species)

ARGs	Target antibiotics	Resistance mechanism/gene family	<i>Actinobacteria</i> genera – number of ARG-carrying species	Total number of ARG-carrying species
<i>marR</i>	Multidrug	Antibiotic target alteration, antibiotic efflux/resistance-nodulation-cell division (RND) antibiotic efflux pump	<i>Actinomyces</i> – 43; <i>Arthrobacter</i> – 21; <i>Corynebacterium</i> – 100; <i>Dietzia</i> – 8; <i>Gordonia</i> – 29; <i>Micrococcus</i> – 4; <i>Micromonospora</i> – 52; <i>Mycobacterium</i> – 76; <i>Nocardioides</i> – 21; <i>Nocardia</i> – 78; <i>Pseudonocardia</i> – 9; <i>Rhodococcus</i> – 30; <i>Streptomyces</i> – 314	785
<i>acrR (tetR)</i>	Multidrug	Antibiotic target alteration, antibiotic efflux/resistance-nodulation-cell division (RND) antibiotic efflux pump	<i>Actinomyces</i> – 41; <i>Arthrobacter</i> – 17; <i>Corynebacterium</i> – 93; <i>Dietzia</i> – 12; <i>Gordonia</i> – 30; <i>Micrococcus</i> – 3; <i>Micromonospora</i> – 52; <i>Mycobacterium</i> – 76; <i>Nocardioides</i> – 21; <i>Pseudonocardia</i> – 9; <i>Rhodococcus</i> – 27; <i>Streptomyces</i> – 317	698
<i>rarD</i>	Chloramphenicol	Antibiotic efflux/predicted chloramphenicol resistance permease	<i>Actinomyces</i> – 31; <i>Arthrobacter</i> – 20; <i>Corynebacterium</i> – 74; <i>Dietzia</i> – 8; <i>Gordonia</i> – 11; <i>Micrococcus</i> – 3; <i>Micromonospora</i> – 51; <i>Mycobacterium</i> – 9; <i>Nocardioides</i> – 18; <i>Nocardia</i> – 66; <i>Pseudonocardia</i> – 8; <i>Rhodococcus</i> – 26; <i>Streptomyces</i> – 262	587
<i>aph</i>	Aminoglycosides	Antibiotic inactivation/aminoglycoside O-phosphotransferase (APH)	<i>Actinomyces</i> – 26; <i>Arthrobacter</i> – 13; <i>Corynebacterium</i> – 55; <i>Dietzia</i> – 5; <i>Gordonia</i> – 7; <i>Micrococcus</i> – 2; <i>Micromonospora</i> – 48; <i>Mycobacterium</i> – 48; <i>Nocardia</i> – 70; <i>Pseudonocardia</i> – 9; <i>Rhodococcus</i> – 13; <i>Streptomyces</i> – 277	573
<i>sulI</i>	Sulfonamides and sulfones	Antibiotic target replacement/sulfonamide resistant dihydropteroate synthase	<i>Actinomyces</i> – 4; <i>Arthrobacter</i> – 17; <i>Corynebacterium</i> – 54; <i>Dietzia</i> – 7; <i>Gordonia</i> – 26; <i>Micrococcus</i> – 3; <i>Micromonospora</i> – 41; <i>Mycobacterium</i> – 63; <i>Nocardioides</i> – 17; <i>Nocardia</i> – 62; <i>Pseudonocardia</i> – 8; <i>Rhodococcus</i> – 18; <i>Streptomyces</i> – 253	573
<i>msrA</i>	MLSB	Antibiotic target protection/ABC-F ATP-binding cassette ribosomal protection protein	<i>Actinomyces</i> – 39; <i>Arthrobacter</i> – 12; <i>Corynebacterium</i> – 98; <i>Dietzia</i> – 8; <i>Gordonia</i> – 28; <i>Micrococcus</i> – 2; <i>Micromonospora</i> – 44; <i>Mycobacterium</i> – 15; <i>Nocardioides</i> – 21; <i>Nocardia</i> – 31; <i>Pseudonocardia</i> – 5; <i>Rhodococcus</i> – 23;	541

			<i>Streptomyces</i> – 200	
<i>carB</i>	MLSB	Antibiotic target alteration/23S ribosomal RNA methyltransferase	<i>Actinomyces</i> – 31; <i>Arthrobacter</i> – 19; <i>Corynebacterium</i> – 91; <i>Dietzia</i> – 7; <i>Gordonia</i> – 27; <i>Micrococcus</i> – 4; <i>Micromonospora</i> – 49; <i>Mycobacterium</i> – 73; <i>Nocardioides</i> – 19; <i>Nocardia</i> – 48; <i>Pseudonocardia</i> – 8; <i>Rhodococcus</i> – 17; <i>Streptomyces</i> – 120	513
<i>folP</i>	Sulfonamides	Antibiotic target alteration/antibiotic resistant dihydropteroate synthase	<i>Actinomyces</i> – 12; <i>Arthrobacter</i> – 16; <i>Corynebacterium</i> – 78; <i>Dietzia</i> – 4; <i>Gordonia</i> – 26; <i>Micrococcus</i> – 3; <i>Micromonospora</i> – 17; <i>Mycobacterium</i> – 39; <i>Nocardioides</i> – 8; <i>Nocardia</i> – 43; <i>Pseudonocardia</i> – 6; <i>Rhodococcus</i> – 16; <i>Streptomyces</i> – 190	458
<i>pbp</i>	β -Lactam antibiotics	Antibiotic target alteration/penicillin-binding protein	<i>Actinomyces</i> – 2; <i>Arthrobacter</i> – 3; <i>Corynebacterium</i> – 67; <i>Gordonia</i> – 24; <i>Nocardioides</i> – 20; <i>Pseudonocardia</i> – 8; <i>Streptomyces</i> – 314	438
<i>folA</i>	Trimethoprim	Antibiotic target alteration/antibiotic resistant dihydrofolate reductase	<i>Actinomyces</i> – 11; <i>Arthrobacter</i> – 12; <i>Corynebacterium</i> – 46; <i>Gordonia</i> – 24; <i>Micrococcus</i> – 1; <i>Micromonospora</i> – 35; <i>Mycobacterium</i> – 24; <i>Nocardioides</i> – 12; <i>Nocardia</i> – 52; <i>Pseudonocardia</i> – 4; <i>Rhodococcus</i> – 16; <i>Streptomyces</i> – 172	409
<i>emrB/qacA</i>	Multidrug	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Actinomyces</i> – 16; <i>Arthrobacter</i> – 6; <i>Corynebacterium</i> – 33; <i>Gordonia</i> – 24; <i>Micrococcus</i> – 2; <i>Micromonospora</i> – 10; <i>Mycobacterium</i> – 19; <i>Nocardioides</i> – 10; <i>Nocardia</i> – 30; <i>Pseudonocardia</i> – 2; <i>Rhodococcus</i> – 11; <i>Streptomyces</i> – 159	322
<i>mdtH</i>	Multidrug	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Actinomyces</i> – 1; <i>Corynebacterium</i> – 14; <i>Gordonia</i> – 10; <i>Micrococcus</i> – 1; <i>Micromonospora</i> – 7; <i>Mycobacterium</i> – 5; <i>Nocardioides</i> – 1; <i>Nocardia</i> – 36; <i>Pseudonocardia</i> – 4; <i>Rhodococcus</i> – 15; <i>Streptomyces</i> – 179	273
<i>acrB</i>	Multidrug	Antibiotic efflux/resistance-nodulation-cell division (RND) antibiotic efflux pump	<i>Arthrobacter</i> – 13; <i>Corynebacterium</i> – 5; <i>Micrococcus</i> – 3; <i>Micromonospora</i> – 44; <i>Mycobacterium</i> – 6; <i>Nocardioides</i> – 2; <i>Nocardia</i> – 1; <i>Rhodococcus</i> – 1; <i>Streptomyces</i> – 173	248
<i>mefA</i>	MLSB	Antibiotic efflux/ major facilitator	<i>Micrococcus</i> – 1; <i>Nocardia</i> – 2; <i>Streptomyces</i> – 236	239

		superfamily (MFS) antibiotic efflux pump		
<i>mecA</i>	Penams (penicillins)	Antibiotic target replacement/methicillin resistant penicillin-binding protein	<i>Micrococcus</i> – 1; <i>Streptomyces</i> – 236	237
<i>ampC</i>	Cephalosporins, penams (penicillins)	Antibiotic inactivation/beta-lactamase	<i>Actinomyces</i> – 2; <i>Arthrobacter</i> – 2; <i>Corynebacterium</i> – 14; <i>Dietzia</i> – 5; <i>Gordonia</i> – 10; <i>Micromonospora</i> – 9; <i>Mycobacterium</i> – 55; <i>Nocardioides</i> – 8; <i>Nocardia</i> – 56; <i>Pseudonocardia</i> – 3; <i>Rhodococcus</i> – 4; <i>Streptomyces</i> – 67	235
<i>vgb</i>	Streptogramins	Antibiotic inactivation/virginiamycin B lyase	<i>Arthrobacter</i> – 1; <i>Micromonospora</i> – 30; <i>Nocardioides</i> – 7; <i>Nocardia</i> – 6; <i>Pseudonocardia</i> – 2; <i>Streptomyces</i> – 130	176
<i>vanB</i>	Glycopeptide antibiotics	Antibiotic target alteration/vancomycin resistant ligase	<i>Actinomyces</i> – 2; <i>Corynebacterium</i> – 3; <i>Gordonia</i> – 1; <i>Micromonospora</i> – 51; <i>Mycobacterium</i> – 25; <i>Nocardioides</i> – 2; <i>Nocardia</i> – 2; <i>Pseudonocardia</i> – 5; <i>Streptomyces</i> – 27	118
<i>acrA</i>	Multidrug	Antibiotic efflux/resistance-nodulation-cell division (RND) antibiotic efflux pump	<i>Actinomyces</i> – 19; <i>Arthrobacter</i> – 4; <i>Corynebacterium</i> – 7; <i>Micrococcus</i> – 1; <i>Micromonospora</i> – 22; <i>Mycobacterium</i> – 2; <i>Nocardioides</i> – 2; <i>Pseudonocardia</i> – 1; <i>Rhodococcus</i> – 1; <i>Streptomyces</i> – 55	114
<i>pbp5</i>	β -Lactam antibiotics	Antibiotic target alteration/penicillin-binding protein	<i>Rhodococcus</i> – 1; <i>Streptomyces</i> – 94	95
<i>tetM</i>	Tetracyclines	Antibiotic target protection/ribosomal protection protein	<i>Actinomyces</i> – 1; <i>Corynebacterium</i> – 3; <i>Micromonospora</i> – 2; <i>Mycobacterium</i> – 1; <i>Nocardia</i> – 1; <i>Streptomyces</i> – 67	75
<i>floR</i>	Phenicol	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Mycobacterium</i> – 48; <i>Streptomyces</i> – 1	49
<i>tnpA</i>	Multidrug	ISNCY family-transposase	<i>Corynebacterium</i> – 12; <i>Gordonia</i> – 1; <i>Micrococcus</i> – 1; <i>Micromonospora</i> – 1; <i>Nocardia</i> – 4; <i>Rhodococcus</i> – 9; <i>Streptomyces</i> – 21	49
<i>bacA</i>	Peptide antibiotics	Antibiotic target alteration/undecaprenyl	<i>Actinomyces</i> – 15; <i>Dietzia</i> – 1; <i>Gordonia</i> – 2; <i>Micromonospora</i> – 6; <i>Mycobacterium</i> – 10; <i>Nocardia</i> – 1;	46

		pyrophosphate phosphatase	<i>Pseudonocardia</i> – 1; <i>Rhodococcus</i> – 1; <i>Streptomyces</i> – 9	
<i>aac</i>	Aminoglycosides	Antibiotic inactivation/ aminoglycoside acetyltransferase	<i>Mycobacterium</i> – 19; <i>Pseudonocardia</i> – 2; <i>Streptomyces</i> – 20	41
<i>mdtL</i>	Multidrug	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Corynebacterium</i> – 5; <i>Dietzia</i> – 1; <i>Gordonia</i> – 2; <i>Mycobacterium</i> – 2; <i>Nocardia</i> – 3; <i>Pseudonocardia</i> – 1; <i>Rhodococcus</i> – 4; <i>Streptomyces</i> – 11	29
<i>fosB</i>	Fosfomycin	Antibiotic inactivation/thiol transferase	<i>Gordonia</i> – 1; <i>Mycobacterium</i> – 6; <i>Nocardia</i> – 1; <i>Pseudonocardia</i> – 1; <i>Rhodococcus</i> – 1; <i>Streptomyces</i> – 18	28
<i>penA</i>	β -Lactam antibiotics	Antibiotic target alteration/ penicillin-binding protein	<i>Actinomyces</i> – 2; <i>Corynebacterium</i> – 3; <i>Micrococcus</i> – 1; <i>Mycobacterium</i> – 3; <i>Nocardia</i> – 4; <i>Rhodococcus</i> – 3; <i>Streptomyces</i> – 7	23
<i>tetA</i>	Tetracyclines	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Corynebacterium</i> – 7; <i>Gordonia</i> – 2; <i>Micrococcus</i> – 1; <i>Mycobacterium</i> – 3; <i>Nocardioides</i> – 1; <i>Nocardia</i> – 2; <i>Pseudonocardia</i> – 1; <i>Rhodococcus</i> – 2; <i>Streptomyces</i> – 3	22
<i>tetC</i>	Tetracyclines	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Corynebacterium</i> – 3; <i>Gordonia</i> – 1; <i>Mycobacterium</i> – 7; <i>Nocardioides</i> – 1; <i>Nocardia</i> – 2; <i>Pseudonocardia</i> – 1; <i>Rhodococcus</i> – 2; <i>Streptomyces</i> – 2	19
<i>mdtG</i>	Fosfomycin	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Arthrobacter</i> – 1; <i>Corynebacterium</i> – 1; <i>Mycobacterium</i> – 5; <i>Nocardia</i> – 1; <i>Rhodococcus</i> – 2; <i>Streptomyces</i> – 7	17
<i>pmrA</i>	Fluoroquinolones	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Rhodococcus</i> – 2; <i>Streptomyces</i> – 15	17
<i>tetD</i>	Tetracyclines	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Arthrobacter</i> – 1; <i>Corynebacterium</i> – 1; <i>Dietzia</i> – 1; <i>Micrococcus</i> – 1; <i>Mycobacterium</i> – 1; <i>Nocardia</i> – 1; <i>Pseudonocardia</i> – 1; <i>Rhodococcus</i> – 2; <i>Streptomyces</i> – 8	17
<i>lmrA</i>	MLSB	Antibiotic target alteration, antibiotic efflux/ATP- binding cassette (ABC) antibiotic efflux pump	<i>Actinomyces</i> – 1; <i>Corynebacterium</i> – 3; <i>Dietzia</i> – 1; <i>Nocardia</i> – 1; <i>Rhodococcus</i> – 1; <i>Streptomyces</i> – 9	16
<i>mepA</i>	Multidrug	Antibiotic efflux/multidrug and toxic compound extrusion (MATE) transporter	<i>Corynebacterium</i> – 11; <i>Mycobacterium</i> – 1; <i>Nocardia</i> – 1; <i>Streptomyces</i> – 3	16

<i>strA</i>	Aminoglycosides	Antibiotic inactivation/ aminoglycoside phosphotransferase	<i>Corynebacterium</i> – 4; <i>Streptomyces</i> – 12	16
<i>cphA</i>	Carbapenems	Antibiotic inactivation/beta- lactamase	<i>Corynebacterium</i> – 1; <i>Mycobacterium</i> – 6; <i>Nocardioides</i> – 2; <i>Nocardia</i> – 1; <i>Pseudonocardia</i> – 1; <i>Streptomyces</i> – 4	15
<i>vanA</i>	Glycopeptide antibiotics	Antibiotic target alteration/ vancomycin resistant ligase	<i>Corynebacterium</i> – 3; <i>Gordonia</i> – 4; <i>Rhodococcus</i> – 1; <i>Streptomyces</i> – 7	15
<i>vatD</i>	Streptogramins	Antibiotic inactivation/ streptogramin vat acetyltransferase	<i>Nocardia</i> – 4; <i>Rhodococcus</i> – 2; <i>Streptomyces</i> – 7	13
<i>tolC</i>	Multidrug	Antibiotic efflux/many multidrug efflux pumps	<i>Arthrobacter</i> – 1; <i>Micromonospora</i> – 2; <i>Mycobacterium</i> – 2; <i>Streptomyces</i> – 7	12
<i>cmx</i>	Phenicols	Antibiotic efflux/major facilitator superfamily (MFS) antibiotic efflux pump	<i>Corynebacterium</i> – 9; <i>Streptomyces</i> – 1	10
<i>mdtA</i>	Multidrug	Antibiotic efflux/resistance- nodulation-cell division (RND) antibiotic efflux pump	<i>Actinomyces</i> – 1; <i>Arthrobacter</i> – 1; <i>Mycobacterium</i> – 2; <i>Streptomyces</i> – 6	10
<i>oprD</i>	Multidrug	Reduced permeability to antibiotic/outer membrane porin	<i>Arthrobacter</i> – 1; <i>Mycobacterium</i> – 3; <i>Streptomyces</i> – 6	10
<i>cmr</i>	Phenicols	Antibiotic efflux/major facilitator superfamily (MFS) antibiotic efflux pump	<i>Corynebacterium</i> – 3; <i>Mycobacterium</i> – 4; <i>Nocardioides</i> – 1; <i>Streptomyces</i> – 1	9
<i>pbp2a</i>	Penams (penicillins)	Antibiotic target replacement/methicillin resistant penicillin-binding protein	<i>Corynebacterium</i> – 9	9
<i>fabK</i>	Triclosan	Antibiotic target alteration/3- oxoacyl-acyl carrier protein reductase	<i>Mycobacterium</i> – 2; <i>Streptomyces</i> – 6	8
<i>strB</i>	Aminoglycosides	Antibiotic inactivation/ aminoglycoside phosphotransferase	<i>Corynebacterium</i> – 4; <i>Nocardia</i> – 1; <i>Streptomyces</i> – 3	8
<i>tetB</i>	Tetracyclines	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Corynebacterium</i> – 8	8
<i>tetO</i>	Tetracyclines	Antibiotic target protection/	<i>Nocardia</i> – 1; <i>Streptomyces</i> – 7	8

		ribosomal protection protein		
<i>msrC</i>	MLSB	Antibiotic target protection/ABC-F ATP-binding cassette ribosomal protection protein	<i>Micrococcus</i> – 1; <i>Mycobacterium</i> – 1; <i>Nocardia</i> – 3; <i>Rhodococcus</i> – 2	7
<i>catA1</i>	Chloramphenicol	Antibiotic inactivation/ chloramphenicol acetyltransferase	<i>Corynebacterium</i> – 1; <i>Rhodococcus</i> – 5	6
<i>ermA</i>	MLSB	Antibiotic target alteration/23S ribosomal RNA methyltransferase	<i>Corynebacterium</i> – 4; <i>Nocardia</i> – 1	5
<i>ermB</i>	MLSB	Antibiotic target alteration/23S ribosomal RNA methyltransferase	<i>Mycobacterium</i> – 3; <i>Nocardia</i> – 1	4
<i>aacC</i>	Aminoglycosides	Antibiotic inactivation/ aminoglycoside acetyltransferase	<i>Streptomyces</i> – 3	3
<i>cfiA</i>	Carbapenems	Antibiotic inactivation/beta-lactamase	<i>Corynebacterium</i> – 1; <i>Mycobacterium</i> – 1; <i>Nocardia</i> – 1	3
<i>spcN</i>	Aminoglycosides	Antibiotic inactivation/ aminoglycoside phosphotransferase	<i>Streptomyces</i> – 3	3
<i>aphA1</i>	Aminoglycosides	Antibiotic inactivation/ aminoglycoside phosphotransferase	<i>Corynebacterium</i> – 2	2
<i>aac(6')-II</i>	Aminoglycosides	Antibiotic inactivation/ aminoglycoside acetyltransferase	<i>Streptomyces</i> – 2	2
<i>aadA1</i>	Aminoglycosides	Antibiotic inactivation/ aminoglycoside nucleotidyltransferase	<i>Corynebacterium</i> – 2	2
<i>bla1</i>	Penams (penicillins)	Antibiotic inactivation/beta-lactamase	<i>Mycobacterium</i> – 2	2
<i>emrD</i>	Multidrug	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Mycobacterium</i> – 1; <i>Rhodococcus</i> – 1	2
<i>qacB</i>	Fluoroquinolones	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Corynebacterium</i> – 1; <i>Streptomyces</i> – 1	2
<i>tetW</i>	Tetracyclines	Antibiotic target protection/ ribosomal protection protein	<i>Actinomyces</i> – 1; <i>Corynebacterium</i> – 1	2

<i>vanYB</i>	Glycopeptide antibiotics	Antibiotic target alteration/vancomycin resistant ligase	<i>Nocardia</i> – 1; <i>Rhodococcus</i> – 1	2
<i>aacC4</i>	Aminoglycosides	Antibiotic inactivation/aminoglycoside acetyltransferase	<i>Streptomyces</i> – 1	1
<i>aadA2</i>	Aminoglycosides	Antibiotic inactivation/aminoglycoside nucleotidyltransferase	<i>Corynebacterium</i> – 1	1
<i>aadA9</i>	Aminoglycosides	Antibiotic inactivation/aminoglycoside nucleotidyltransferase	<i>Corynebacterium</i> – 1	1
<i>ceoA</i>	Multidrug	Antibiotic efflux/resistance-nodulation-cell division (RND) antibiotic efflux pump	<i>Mycobacterium</i> – 1	1
<i>cmrA</i>	Phenicols	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Rhodococcus</i> – 1	1
<i>dfrA1</i>	Diaminopyrimidine antibiotic	Antibiotic target replacement/trimethoprim resistant dihydrofolate reductase	<i>Corynebacterium</i> – 1	1
<i>ermC</i>	MLSB	Antibiotic target alteration/23S ribosomal RNA methyltransferase	<i>Nocardia</i> – 1	1
<i>fosB2</i>	Fosfomycin	Antibiotic inactivation/thiol transferase	<i>Rhodococcus</i> – 1	1
<i>mdtE</i>	Multidrug	Antibiotic efflux/resistance-nodulation-cell division (RND) antibiotic efflux pump	<i>Mycobacterium</i> – 1	1
<i>nisB</i>	Nisin	Antibiotic efflux/nisin dehydratase	<i>Streptomyces</i> – 1	1
<i>picA</i>	Macrolides	Antibiotic efflux/unknown	<i>Streptomyces</i> – 1	1
<i>pikR1</i>	MLSB	Antibiotic target alteration/23S ribosomal RNA methyltransferase	<i>Streptomyces</i> – 1	1

<i>pikR2</i>	MLSB	Antibiotic target alteration/23S ribosomal RNA methyltransferase	<i>Streptomyces</i> – 1	1
<i>qacH</i>	Fluoroquinolones	Antibiotic efflux/small multidrug resistance (SMR) antibiotic efflux pump	<i>Corynebacterium</i> – 1	1
<i>sulA</i>	Sulfonamides	Antibiotic target replacement/ sulfonamide resistant dihydropteroate synthase	<i>Mycobacterium</i> – 1	1
<i>sul2</i>	Sulfonamides	Antibiotic target replacement/ sulfonamide resistant dihydropteroate synthase	<i>Nocardia</i> – 1	1
<i>tetL</i>	Tetracyclines	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Nocardia</i> – 1	1
<i>yceL</i>	Fosfomicin	Antibiotic efflux/ major facilitator superfamily (MFS) antibiotic efflux pump	<i>Streptomyces</i> – 1	1

MLSB - Macrolide-Lincosamide-Streptogramin B.