Center for Genomic Epidemiology

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ResFinder-4.1 Server - Results

Input Files: scaffolds.fasta

Warning:

One or more resistance genes have unknown phenotypes. The species specific summary table does not take this into account.

Antimicrobial	Class	WGS-predicted phenotype	Genetic background
vancomycin	glycopeptide	No resistance	
mupirocin	pseudomonic acid	No resistance	
obramycin	aminoglycoside	No resistance	
rirginiamycin m	streptogramin a	No resistance	
sepamicin	aminoglycoside	No resistance	
virginiamycin s	streptogramin b	No resistance	
outirosin	aminoglycoside	No resistance	
ampicillin	beta-lactam	Resistant	blaTEM-1B (blaTEM-1B_AY458016), blaCTX-M-15 (blaCTX-M-15_AY044436)
astromicin	aminoglycoside	No resistance	
ividomycin	aminoglycoside	No resistance	
sulfamethoxazole	folate pathway antagonist	Resistant	sul1 (sul1_AY115475), sul1 (sul1_U12338), sul1 (sul1_DQ914960), sul1 (sul1_AY522923 sul2 (sul2 HQ840942)
emocillin	beta-lactam	No resistance	(48.2_1.48.66.49)
rimethoprim	folate pathway antagonist	Resistant	dfrA7 (dfrA7_AB161450)
oleandomycin	macrolide	No resistance	
unknown macrolide	macrolide	Resistant	mdf(A) (mdf(A)_Y08743)
florfenicol	phenicol	No resistance	- () () () () () () () () () (
luoroquinolone	under_development		
quinupristin	streptogramin b	No resistance	
ceftriaxone	beta-lactam	Resistant	blaCTX-M-15 (blaCTX-M-15_AY044436)
cephalothin			blaTEM-1B (blaTEM-1B AY458016)
·	beta-lactam	Resistant	DIATEM-IB (DIATEM-IB_AT456016)
incomycin	lincosamide	No resistance	
spectinomycin	aminocyclitol	No resistance	
quinupristin+dalfopristin	streptogramin a	No resistance	
paromomycin	aminoglycoside	No resistance	
clindamycin	lincosamide	No resistance	
amoxicillin+clavulanic acid	beta-lactam	No resistance	
gentamicin	aminoglycoside	No resistance	
piperacillin+tazobactam	beta-lactam	No resistance	
tiamulin	pleuromutilin	No resistance	
ribostamycin	aminoglycoside	No resistance	
erythromycin	macrolide	No resistance	
kanamycin	aminoglycoside	No resistance	
teicoplanin	glycopeptide	No resistance	
amikacin	aminoglycoside	No resistance	
igecycline	tetracycline	No resistance	
icarcillin+clavulanic acid	beta-lactam	No resistance	
cephalotin	beta-lactam	No resistance	
cefoxitin	beta-lactam	No resistance	
hygromycin	aminoglycoside	No resistance	
penicillin	beta-lactam	No resistance	
neomycin	aminoglycoside	No resistance	
dalfopristin	streptogramin a	No resistance	
piperacillin	beta-lactam	Resistant	blaTEM-1B (blaTEM-1B_AY458016), blaCTX-M-15 (blaCTX-M-15_AY044436)
elithromycin	macrolide	No resistance	, , , , , , , , , , , , , , , , , ,
amoxicillin	beta-lactam	Resistant	blaTEM-1B (blaTEM-1B_AY458016), blaCTX-M-15 (blaCTX-M-15_AY044436)
meropenem	beta-lactam	No resistance	- (
sisomicin	aminoglycoside	No resistance	
J. J	arminglycoside	140 100101010100	

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Antimicrobial	Class	WGS-predicted phenotype	Genetic background
cefepime	beta-lactam	Resistant	blaCTX-M-15 (blaCTX-M-15_AY044436)
butiromycin	aminoglycoside	No resistance	
doxycycline	tetracycline	Resistant	tet(A) (tet(A)_AJ517790)
unknown fluoroquinolone	fluoroquinolone	Resistant	mdf(A) (mdf(A)_Y08743)
fusidic acid	steroid antibacterial	No resistance	
ciprofloxacin	fluoroquinolone	No resistance	
colistin	polymyxin	No resistance	
imipenem	beta-lactam	No resistance	
arbekacin	aminoglycoside	No resistance	
nalidixic acid	fluoroquinolone	No resistance	
metronidazole	nitroimidazole	No resistance	
cefixime	beta-lactam	No resistance	
unknown phenicol	phenicol	Resistant	mdf(A) (mdf(A)_Y08743)
pristinamycin ia	streptogramin b	No resistance	
tylosin	macrolide	No resistance	
cefotaxime+clavulanic acid	beta-lactam	No resistance	
rifampicin	rifamycin	No resistance	
fosfomycin	fosfomycin	No resistance	
ceftazidime	beta-lactam	Resistant	blaCTX-M-15 (blaCTX-M-15_AY044436)
fortimicin	aminoglycoside	No resistance	
carbomycin	macrolide	No resistance	
ticarcillin	beta-lactam	Resistant	blaTEM-1B (blaTEM-1B_AY458016), blaCTX-M-15 (blaCTX-M-15_AY044436)
azithromycin	macrolide	No resistance	
kasugamycin	aminoglycoside	No resistance	
chloramphenicol	phenicol	No resistance	
ampicillin+clavulanic acid	beta-lactam	No resistance	
cefotaxime	beta-lactam	Resistant	blaCTX-M-15 (blaCTX-M-15_AY044436)
unknown tetracycline	tetracycline	Resistant	mdf(A) (mdf(A)_Y08743)
piperacillin+clavulanic acid	beta-lactam	No resistance	
ceftazidime+avibactam	beta-lactam	No resistance	
apramycin	aminoglycoside	No resistance	
spiramycin	macrolide	No resistance	
unknown rifamycin	rifamycin	Resistant	mdf(A) (mdf(A)_Y08743)
dibekacin	aminoglycoside	No resistance	
ertapenem	beta-lactam	No resistance	
tetracycline	tetracycline	Resistant	tet(A) (tet(A)_AJ517790)
linezolid	oxazolidinone	No resistance	
netilmicin	aminoglycoside	No resistance	
minocycline	tetracycline	No resistance	
aztreonam	beta-lactam	Resistant	blaCTX-M-15 (blaCTX-M-15_AY044436)
unknown beta-lactam	beta-lactam	No resistance	
pristinamycin iia	streptogramin a	No resistance	
streptomycin	aminoglycoside	Resistant	aph(6)-ld (aph(6)-ld_M28829), aph(3")-lb (aph(3")-lb_AF321551)

Fluoroquinolone

Resistance gene	Identity	Alignment Length/Gene Length	Position in reference	Contig or Depth	Position in contig	Phenotype	PMID	Accession no.	
mdf(A)	98.4590429846	1233/1233	11233	NODE_4_length _388939_cov_80 .679974	307306308538	unknown macrolide,unkno wn aminoglycoside,u nknown tetracycline,unkn own fluoroquinolone,u nknown phenicol,unknow n rifamycin	9079913	<u>Y08743</u>	Md E 1 Ch anc bu w

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Aminoglycoside

Resistance gene	Identity	Alignment Length/Gene Length	Position in reference	Contig or Depth	Position in contig	Phenotype	PMID	Accession no.	
mdf(A)	98.4590429846	1233/1233	11233	NODE_4_length _388939_cov_80 _679974	307306308538	unknown macrolide,unkno wn aminoglycoside,u nknown tetracycline,unkn own fluoroquinolone,u nknown phenicol,unknow n rifamycin	9079913	<u>Y08743</u>	Mdf(// ss ss ii Ery Te Ri Ka Chlo and (but i whi res
aph(3")-lb	100.0	804/804	1804	NODE_1_length _1628642_cov_7 8.003180	799877800680	streptomycin	12029529	<u>AF321551</u>	Alter
aph(6)-ld	100.0	837/837	1837	NODE_1_length _1628642_cov_7 8.003180	799041799877	streptomycin	2653965	<u>M28829</u>	Alter

Rifamycin

Resistance gene	Identity	Alignment Length/Gene Length	Position in reference	Contig or Depth	Position in contig	Phenotype	PMID	Accession no.	
mdf(A)	98.4590429846	1233/1233	11233	NODE_4_length _388939_cov_80 .679974	307306308538	unknown macrolide,unkno wn aminoglycoside,u nknown tetracycline,unkn own fluoroquinolone,u nknown phenicol,unknow n rifamycin	9079913	<u>Y08743</u>	Mdf(

Macrolide

Resistance gene	Identity	Alignment Length/Gene Length	Position in reference	Contig or Depth	Position in contig	Phenotype	PMID	Accession no.	
mdf(A)	98.4590429846	1233/1233	11233	NODE_4_length _388939_cov_80 _679974	307306308538	unknown macrolide,unkno wn aminoglycoside,u nknown tetracycline,unkn own fluoroquinolone,u nknown phenicol,unknow n rifamycin	9079913	<u>Y08743</u>	En T I I Chil and bu w

Folate pathway antagonist

Resistance gene	Identity	Alignment Length/Gene Length	Position in reference	Contig or Depth	Position in contig	Phenotype	PMID	Accession no.
sul1	100.0	761/840	1761	NODE_1_length _1628642_cov_7 8.003180	806890807650	sulfamethoxazole	unpublished	<u>U12338</u>
dfrA7	100.0	474/474	1474	NODE_1_length _1628642_cov_7 8.003180	808221808694	trimethoprim	unpublished	AB161450
sul1	100.0	761/828	1761	NODE_1_length _1628642_cov_7 8.003180	806890807650	sulfamethoxazole	16339607	AY522923
sul2	100.0	816/816	1816	NODE_1_length _1628642_cov_7 8.003180	800741801556	sulfamethoxazole	22888274	HQ840942
sul1	100.0	761/840	1761	NODE_1_length _1628642_cov_7 8.003180	806890807650	sulfamethoxazole	18753343	<u>AY115475</u>
sul1	100.0	761/882	1761	NODE_1_length _1628642_cov_7 8.003180	806890807650	sulfamethoxazole	17158944	DQ914960

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Tetracycline

Resistance gene	Identity	Alignment Length/Gene Length	Position in reference	Contig or Depth	Position in contig	Phenotype	PMID	Accession no.	
tet(A)	100.0	1200/1200	11200	NODE_17_lengt h_4054_cov_26. 052763	9562155	doxycycline,tetra cycline	12654659	AJ517790	
mdf(A)	98.4590429846	1233/1233	11233	NODE_4_length _388939_cov_80 .679974	307306308538	unknown macrolide,unkno wn aminoglycoside,u nknown tetracycline,unkn own fluoroquinolone,u nknown phenicol,unknow n rifamycin	9079913	<u>Y08743</u>	Ch and bi

Phenicol

Resistance gene	Identity	Alignment Length/Gene Length	Position in reference	Contig or Depth	Position in contig	Phenotype	PMID	Accession no.	
mdf(A)	98.4590429846	1233/1233	11233	NODE_4_length _388939_cov_80 _679974	307306308538	unknown macrolide,unkno wn aminoglycoside,u nknown tetracycline,unkn own fluoroquinolone,u nknown phenicol,unknow n rifamycin	9079913	<u>Y08743</u>	Er To F K Chlo and but wl

Beta-lactam

Resistance gene	Identity	Alignment Length/Gene Length	Position in reference	Contig or Depth	Position in contig	Phenotype	PMID	Accession no.	
blaCTX-M-15	100.0	876/876	1876	NODE_11_lengt h_88402_cov_75 .628805	6354364418	amoxicillin,ampici llin,aztreonam,ce fepime,cefotaxim e,ceftazidime,ceft riaxone,piperacilli n,ticarcillin	11470367, 26169409	AY044436	
blaTEM-1B	100.0	861/861	1861	NODE_11_lengt h_88402_cov_75 .628805	5986160721	amoxicillin,ampici llin,cephalothin,pi peracillin,ticarcilli n	15388431	AY458016	1

Download acquired AMR gene results:

Results as text Hit in genome sequences Resistance gene sequences Results as tabseperated file

Selected %ID threshold for ResFinder: 90 % Selected minimum length for ResFinder: 60 %

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Technical problems

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