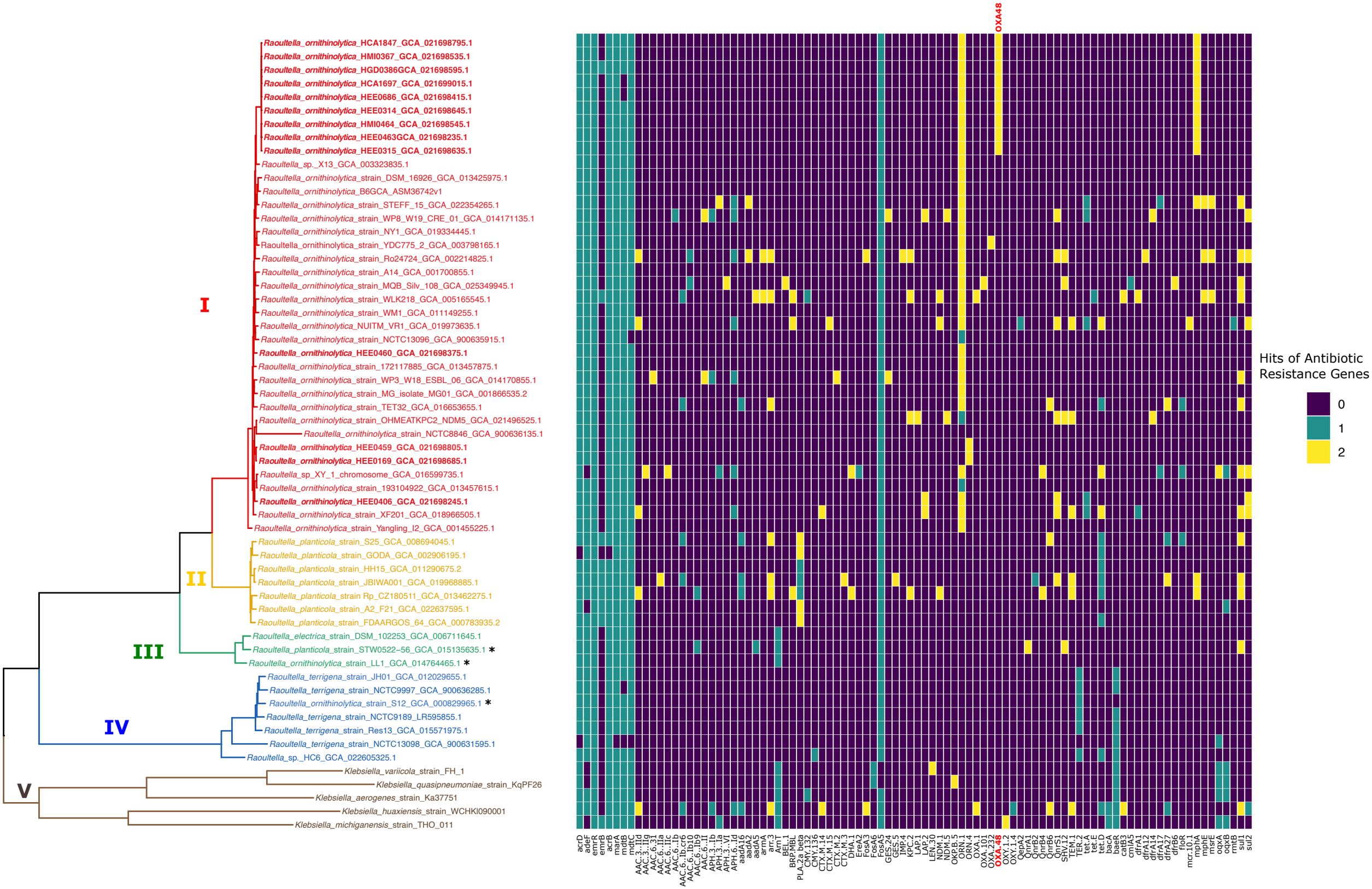
OXA48
Raoultella_ornithinolytica_HCA1847_GCA_021698795.1
Raoultella_ornithinolytica_HMI0367_GCA_021698535.1
Raoultella_ornithinolytica_HGD0386GCA_021698595.1
Raoultella_ornithinolytica_HCA1697_GCA_021699015.1
Raoultella_ornithinolytica_HEE0686_GCA_021698415.1
Raoultella_ornithinolytica_HEE0314_GCA_021698645.1
Raoultella_ornithinolytica_HMI0464_GCA_021698545.1
Raoultella_ornithinolytica_HEE0463GCA_021698235.1
Raoultella_ornithinolytica_HEE0315_GCA_021698635.1
Raoultella_sp._X13_GCA_003323835.1
Raoultella_ornithinolytica_strain_DSM_16926_GCA_013425975.1
Raoultella_ornithinolytica_B6GCA_ASM36742v1
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Raoultella_ornithinolytica_strain_WLK218_GCA_005165545.1
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Raoultella_ornithinolytica_HEE0460_GCA_021698375.1
Raoultella_ornithinolytica_strain_172117885_GCA_013457875.1
Raoultella_ornithinolytica_strain_WP3_W18_ESBL_06_GCA_014170855.1
Raoultella_ornithinolytica_strain_MG_isolate_MG01_GCA_001866535.2
Raoultella_ornithinolytica_strain_TET32_GCA_016653655.1
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Raoultella_ornithinolytica_strain_NCTC8846_GCA_900636135.1
Raoultella_ornithinolytica_HEE0459_GCA_021698805.1
Raoultella_ornithinolytica_HEE0169_GCA_021698685.1
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Raoultella_ornithinolytica_strain_Yangling_I2_GCA_001455225.1
Raoultella_planticola_strain_S25_GCA_008694045.1
Raoultella_planticola_strain_GODA_GCA_002906195.1
Raoultella_planticola_strain_HH15_GCA_011290675.2
Raoultella_planticola_strain_JBIWA001_GCA_019968885.1
Raoultella_planticola_strain Rp_CZ180511_GCA_013462275.1
Raoultella_planticola_strain_A2_F21_GCA_022637595.1
Raoultella_planticola_strain_FDAARGOS_64_GCA_000783935.2
Raoultella_electrica_strain_DSM_102253_GCA_006711645.1
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Raoultella_ornithinolytica_strain_LL1_GCA_014764465.1 *
Raoultella_terrigena_strain_JH01_GCA_012029655.1
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Raoultella_ornithinolytica_strain_S12_GCA_000829965.1 *
Raoultella_terrigena_strain_NCTC9189_LR595855.1
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Raoultella_terrigena_strain_NCTC13098_GCA_900631595.1
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Klebsiella_variicola_strain_FH_1
Klebsiella_quasipneumoniae_strain_KqPF26
Klebsiella_aerogenes_strain_Ka37751
Klebsiella_huaxiensis_strain_WCHKl090001
Klebsiella_michiganensis_strain_THO_011
I
II
III
IV
V
Hits of Antibiotic Resistance Genes
0
1
2
acrD
adeF
emrR
emrB
acrB
mdtA
mdtB
mdtC
AAC.3..IId
AAC.3..IIg
AAC.6.31
AAC.6..IIa
AAC.6..IIc
AAC.6..Ib
AAC.6..Ib.cr6
AAC.6..Ib10
AAC.6..Ib9
AAC.6..II
APH.3..Ib
APH.3..Ia
APH.3..VI
APH.6..Id
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aadA2
aadA5
armA
arr.3
AmT
BER.1
BRP.MBL
PLA.2a.beta
CMY.132
CMY.136
CTX.M.14
CTX.M.15
CTX.M.2
CTX.M.3
DHA.1
EreA2
FosA3
FosA6
FosA5
GES.24
GES.5
IMP.4
KPC.2
LAP.1
LAP.2
LEN.30
NDM.1
NDM.5
OKP.B.5
ORN.1
ORN.4
OXA.1
OXA.211
OXA.232
OXA.48
OXY.1.2
OXY.1.4
QepA2
QnrA1
QnrB2
QnrB4
QnrB6
QnrS1
SHV.12
TEM.1
TER.2
tet.A
tet.E
tet.D
bacA
baeR
catB3
cmlA5
dfrA1
dfrA12
dfrA14
dfrA17
dfrA27
dfrB6
floR
mcr.10.1
mphA
mphE
msrE
oqxA
oqxB
rmtB
sul1
sul2