

AIN16239.1 CP008943.1:2128421-2134498 <i>Yersinia pseudotuberculosis</i>	Terminase-SS ? ? Prim_Zn_Ribbon Toprim-DNAG DUF927 CoLE7*
ARJ44344.1 CP019706.1:3892353-3895726 <i>Pantoea alhagi</i>	? ? IAT_beta DUF2474 CoLE7*
AVJ18069.1 CP023268.1:2803821-2809366 <i>Serratia sp. MYb239</i>	Trp_halogenase Rieske KshA_C CoLE7* SUKH_5 DUF2511
EGV2247329.1 AASWUB020000123.1:12-929 <i>Escherichia coli</i>	CoLE7*
EHM1384157.1 ABAHQB010000007.1:106091-111545 <i>Salmonella enterica</i>	? ? CoLE7* RecA ?
EJA1006716.1 ABFNUN010000356.1:1-492 <i>Escherichia coli</i>	CoLE7*
HBI6868227.1 DAEUWM010000006.1:161966-166518 <i>Enterobacter cancerogenus</i>	Virulence_RhuM ? CoLE7* ? Acetylase
HCB3587581.1 DAHACD010000003.1:111260-115636 <i>Citrobacter koseri</i>	GFO_IDH_MocA_C GFO_IDH_MocA Acetylase T6SS_HCP CoLE7* ? CoLE7*
MBB1420127.1 JACJEW010000145.1:1-1646 <i>Pseudoalteromonas sp. SG44-1</i>	CoLE7* SUKH-SMI1 Imm36
MBB7476818.1 AATMOP020000290.1:1-743 <i>Escherichia coli</i>	CoLE7*
MBH1929607.1 JADULK010000003.1:344160-348769 <i>Serratia rubidaea</i>	MatE MatE_DUF805 CoLE7* ? LysR_substrate
MBK1426233.1 JAEKFW010000001.1:4290224-4293376 <i>Yersinia pseudotuberculosis</i>	? Acetylase CoLE7* ? DOC
MBK4703974.1 JAEPIT010000141.1:1-2215 <i>Enterobacter cloacae</i>	CoLE7* HAD Lysozyme
NIY87957.1 JAAT0Q010000075.1:13305-18777 <i>Vibrio campbellii</i>	WYL HSP70 CoLE7* Glyco_transf_8 Xre_MbcA_ParS_C
PAO06760.1 NPNR010000001.1:891873-896259 <i>Enterobacter cloacae</i>	? ? CoLE7* ? MdoG TAT_signal
PLR41191.1 PJZF010000002.1:362050-365153 <i>Chimaeribacter californicus</i>	? ? CoLE7* ? IMS IMS_HHH IMS_C DUF4113
SQJ18331.1 LS483492.1:2118021-2123577 <i>Serratia rubidaea</i>	DUF2511 SUKH_5 CoLE7* Rieske KshA_C Trp_halogenase
TAT65213.1 RQJF010000074.1:1-2851 <i>Enterobacter cloacae</i>	CoLE7* ? MFS_2
UJD81973.1 CP042353.1:4317521-4322265 <i>Serratia rubidaea</i>	Phage_portal ? CoLE7* His-Kinase Receiver
VFS25351.1 CAADIW010000020.1:12633-15827 <i>Enterobacter cancerogenus</i>	Peptidase_M90 ? CoLE7* ? ?
WP_172513334.1 NZ_BLPJ020000002.1:167816-175250 <i>Enterobacter roggenkampii</i>	Malic_M malic ADH_zinc_N ADH_N ? CoLE7* ABC_tran ABC_tran_Xtn ABC_tran
WP_072102241.1 NZ_CWJN01000153.1:64-1502 <i>Yersinia pseudotuberculosis</i>	? ? CoLE7*
WP_188723241.1 NZ_BMKJ010000001.1:608753-613592 <i>Franconibacter daqui</i>	OppC_N BPD_transp_1 ? TRAM_2 RADICAL-SAM UPF0004 CoLE7* ?
WP_072071386.1 NZ_CGBP010000094.1:73-1016 <i>Yersinia similis</i>	? CoLE7*
WP_193030490.1 NZ_AP022508.1:3085983-3091387 <i>Enterobacter bugandensis</i>	Phos_pyr_kin ? CoLE7* HK Collagen YadA_anchor
WP_034896424.1 NZ_JRUQ010000055.1:45670-50268 <i>Erwinia typographi</i>	? CoLE7* ? Bax1-I DUF1615
WP_232298346.1 NZ_JRUQ010000056.1:43505-46529 <i>Erwinia typographi</i>	Shikimate_dh_N ? ? ? CoLE7*
WP_139281492.1 NZ_AP024862.1:547682-551075 <i>Vibrio aerogenes</i>	? ? PAAR_motif CoLE7* ? ?
WP_208290946.1 NZ_JAGFDW010000164.1:1-1163 <i>Enterobacter sichuanensis</i>	CoLE7*