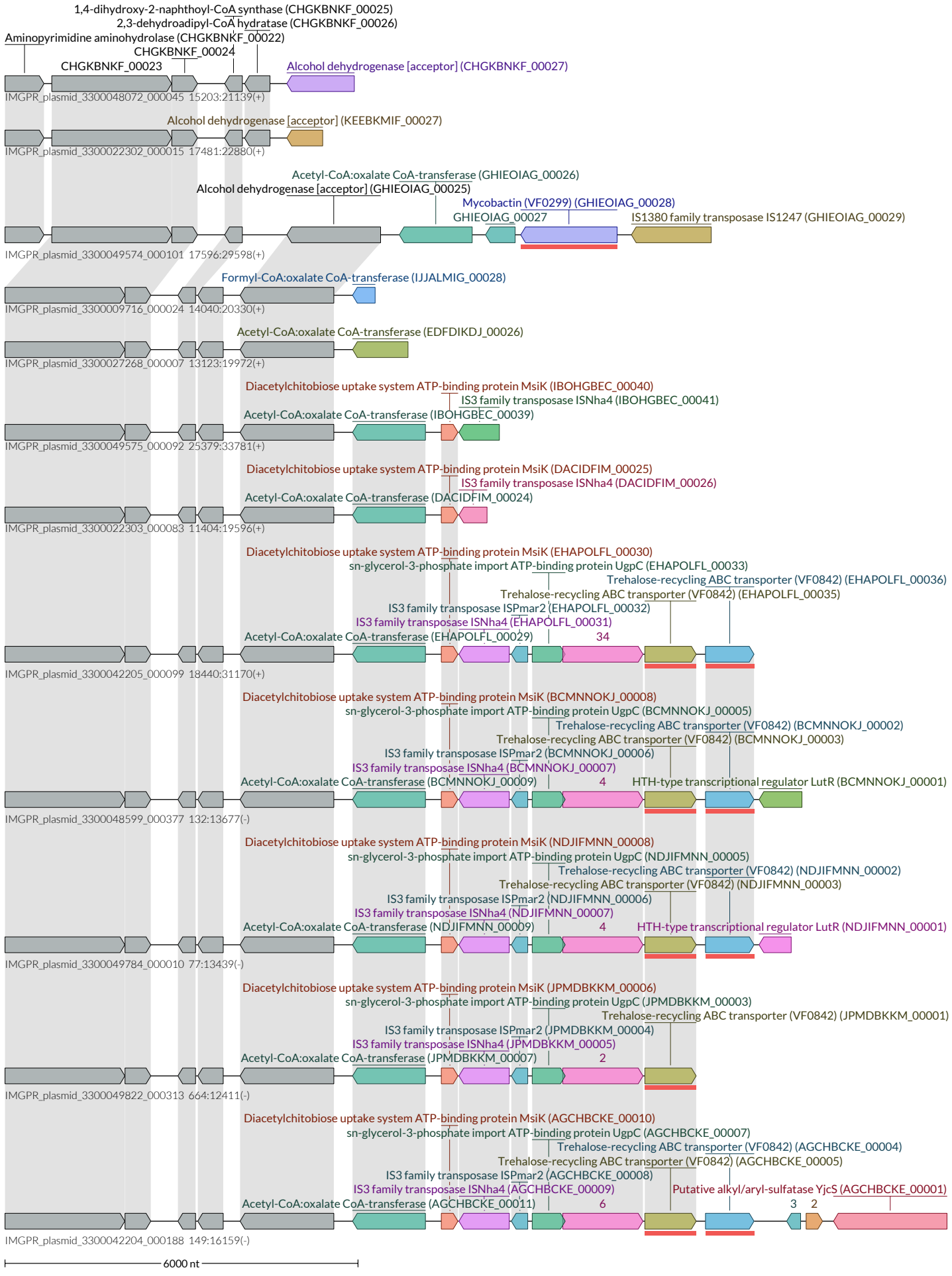
1,4-dihydroxy-2-naphthoyl-CoA synthase (CHGKBNKF_00025)
2,3-dehydroadipyl-CoA hydratase (CHGKBNKF_00026)
Aminopyrimidine aminohydrolase (CHGKBNKF_00022)
CHGKBNKF_00024
CHGKBNKF_00023
Alcohol dehydrogenase [acceptor] (CHGKBNKF_00027)
IMGPR_plasmid_3300048072_000045 15203:21139(+)
Alcohol dehydrogenase [acceptor] (KEEBKMIF_00027)
IMGPR_plasmid_3300022302_000015 17481:22880(+)
Acetyl-CoA:oxalate CoA-transferase (GHIEOIAG_00026)
Alcohol dehydrogenase [acceptor] (GHIEOIAG_00025)
Mycobactin (VF0299) (GHIEOIAG_00028)
GHIEOIAG_00027
IS1380 family transposase IS1247 (GHIEOIAG_00029)
IMGPR_plasmid_3300049574_000101 17596:29598(+)
Formyl-CoA:oxalate CoA-transferase (IJJALMIG_00028)
IMGPR_plasmid_3300009716_000024 14040:20330(+)
Acetyl-CoA:oxalate CoA-transferase (EDFDIKDJ_00026)
IMGPR_plasmid_3300027268_000007 13123:19972(+)
Diacetylchitobiose uptake system ATP-binding protein MsiK (IBOHGBEC_00040)
IS3 family transposase ISNha4 (IBOHGBEC_00041)
Acetyl-CoA:oxalate CoA-transferase (IBOHGBEC_00039)
IMGPR_plasmid_3300049575_000092 25379:33781(+)
Diacetylchitobiose uptake system ATP-binding protein MsiK (DACIDFIM_00025)
IS3 family transposase ISNha4 (DACIDFIM_00026)
Acetyl-CoA:oxalate CoA-transferase (DACIDFIM_00024)
IMGPR_plasmid_3300022303_000083 11404:19596(+)
Diacetylchitobiose uptake system ATP-binding protein MsiK (EHAPOLFL_00030)
sn-glycerol-3-phosphate import ATP-binding protein UgpC (EHAPOLFL_00033)
Trehalose-recycling ABC transporter (VF0842) (EHAPOLFL_00036)
Trehalose-recycling ABC transporter (VF0842) (EHAPOLFL_00035)
IS3 family transposase ISPmar2 (EHAPOLFL_00032)
IS3 family transposase ISNha4 (EHAPOLFL_00031)
Acetyl-CoA:oxalate CoA-transferase (EHAPOLFL_00029)
34
IMGPR_plasmid_3300042205_000099 18440:31170(+)
Diacetylchitobiose uptake system ATP-binding protein MsiK (BCMNNOKJ_00008)
sn-glycerol-3-phosphate import ATP-binding protein UgpC (BCMNNOKJ_00005)
Trehalose-recycling ABC transporter (VF0842) (BCMNNOKJ_00002)
Trehalose-recycling ABC transporter (VF0842) (BCMNNOKJ_00003)
IS3 family transposase ISPmar2 (BCMNNOKJ_00006)
IS3 family transposase ISNha4 (BCMNNOKJ_00007)
Acetyl-CoA:oxalate CoA-transferase (BCMNNOKJ_00009)
4
HTH-type transcriptional regulator LutR (BCMNNOKJ_00001)
IMGPR_plasmid_3300048599_000377 132:13677(-)
Diacetylchitobiose uptake system ATP-binding protein MsiK (NDJIFMNN_00008)
sn-glycerol-3-phosphate import ATP-binding protein UgpC (NDJIFMNN_00005)
Trehalose-recycling ABC transporter (VF0842) (NDJIFMNN_00002)
Trehalose-recycling ABC transporter (VF0842) (NDJIFMNN_00003)
IS3 family transposase ISPmar2 (NDJIFMNN_00006)
IS3 family transposase ISNha4 (NDJIFMNN_00007)
Acetyl-CoA:oxalate CoA-transferase (NDJIFMNN_00009)
4
HTH-type transcriptional regulator LutR (NDJIFMNN_00001)
IMGPR_plasmid_3300049784_000010 77:13439(-)
Diacetylchitobiose uptake system ATP-binding protein MsiK (JPMDBKKM_00006)
sn-glycerol-3-phosphate import ATP-binding protein UgpC (JPMDBKKM_00003)
Trehalose-recycling ABC transporter (VF0842) (JPMDBKKM_00001)
IS3 family transposase ISPmar2 (JPMDBKKM_00004)
IS3 family transposase ISNha4 (JPMDBKKM_00005)
Acetyl-CoA:oxalate CoA-transferase (JPMDBKKM_00007)
2
IMGPR_plasmid_3300049822_000313 664:12411(-)
Diacetylchitobiose uptake system ATP-binding protein MsiK (AGCHBCKE_00010)
sn-glycerol-3-phosphate import ATP-binding protein UgpC (AGCHBCKE_00007)
Trehalose-recycling ABC transporter (VF0842) (AGCHBCKE_00004)
Trehalose-recycling ABC transporter (VF0842) (AGCHBCKE_00005)
IS3 family transposase ISPmar2 (AGCHBCKE_00008)
IS3 family transposase ISNha4 (AGCHBCKE_00009)
Putative alkyl/aryl-sulfatase YjcS (AGCHBCKE_00001)
Acetyl-CoA:oxalate CoA-transferase (AGCHBCKE_00011)
6
3
2
IMGPR_plasmid_3300042204_000188 149:16159(-)
6000 nt
Virulence