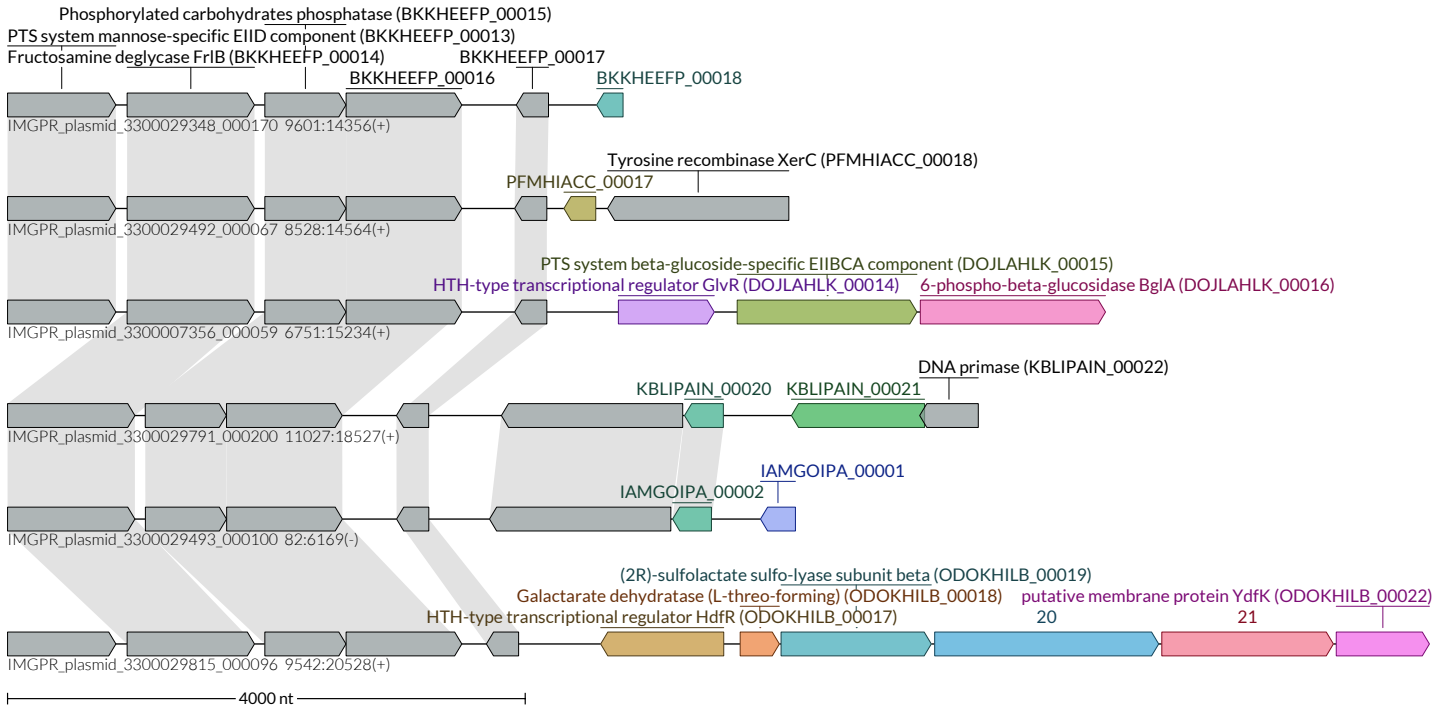
Phosphorylated carbohydrates phosphatase (BKKHEEFP_00015)
PTS system mannose-specific EIID component (BKKHEEFP_00013)
Fructosamine deglycase FrlB (BKKHEEFP_00014)
BKKHEEFP_00017
BKKHEEFP_00016
BKKHEEFP_00018
IMGPR_plasmid_3300029348_000170 9601:14356(+)
Tyrosine recombinase XerC (PFMHIACC_00018)
PFMHIACC_00017
IMGPR_plasmid_3300029492_000067 8528:14564(+)
PTS system beta-glucoside-specific EIIBCA component (DOJLAHLK_00015)
HTH-type transcriptional regulator GlvR (DOJLAHLK_00014)
6-phospho-beta-glucosidase BglA (DOJLAHLK_00016)
IMGPR_plasmid_3300007356_000059 6751:15234(+)
DNA primase (KBLIPAIN_00022)
KBLIPAIN_00020
KBLIPAIN_00021
IMGPR_plasmid_3300029791_000200 11027:18527(+)
IAMGOIPA_00001
IAMGOIPA_00002
IMGPR_plasmid_3300029493_000100 82:6169(-)
(2R)-sulfolactate sulfo-lyase subunit beta (ODOKHILB_00019)
Galactarate dehydratase (L-threo-forming) (ODOKHILB_00018)
putative membrane protein YdfK (ODOKHILB_00022)
HTH-type transcriptional regulator HdfR (ODOKHILB_00017)
20
21
IMGPR_plasmid_3300029815_000096 9542:20528(+)
4000 nt