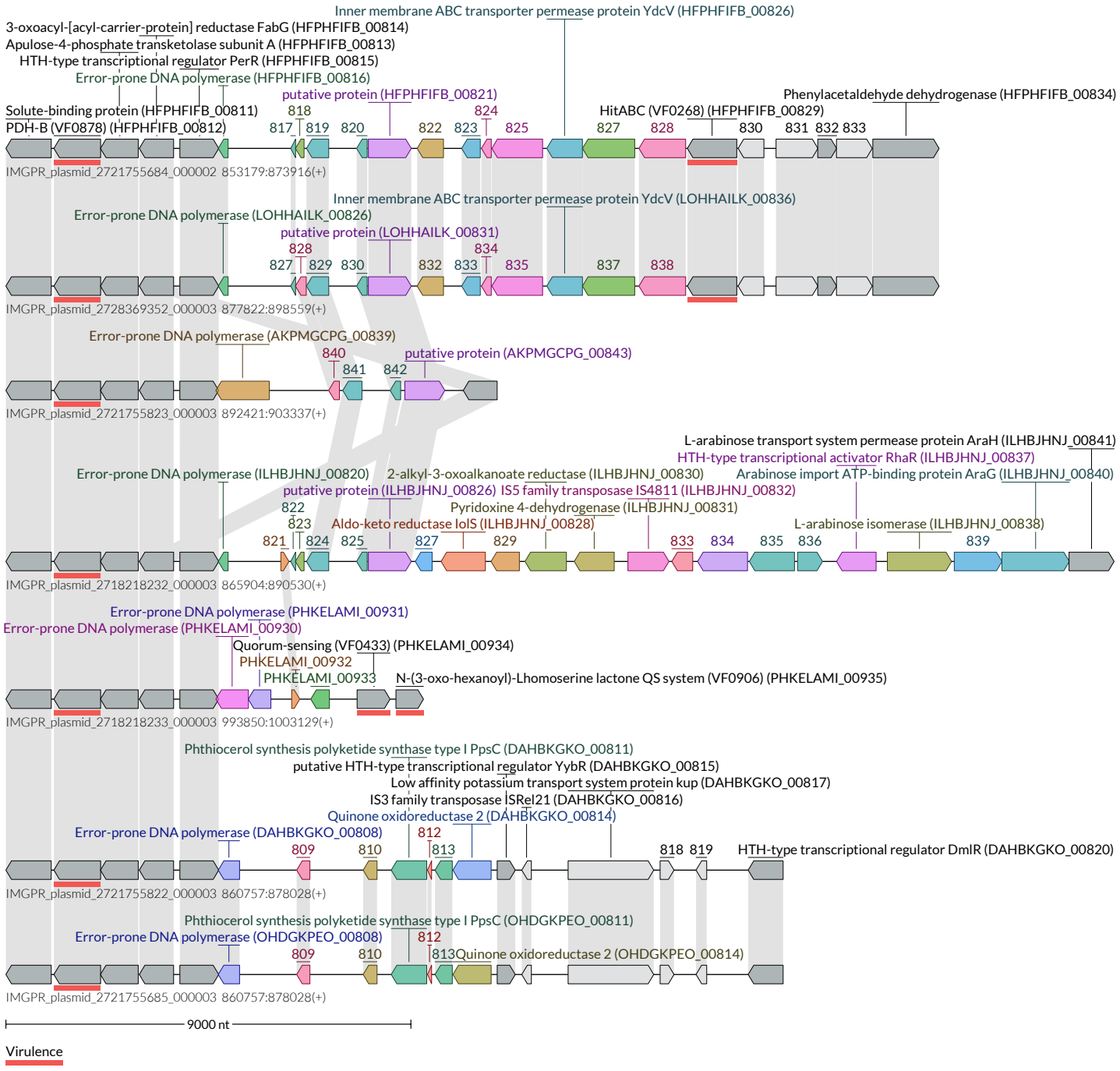
Inner membrane ABC transporter permease protein YdcV (HFPHFIFB_00826)
3-oxoacyl-[acyl-carrier-protein] reductase FabG (HFPHFIFB_00814)
Apulose-4-phosphate transketolase subunit A (HFPHFIFB_00813)
HTH-type transcriptional regulator PerR (HFPHFIFB_00815)
Error-prone DNA polymerase (HFPHFIFB_00816)
putative protein (HFPHFIFB_00821)
Phenylacetaldehyde dehydrogenase (HFPHFIFB_00834)
Solute-binding protein (HFPHFIFB_00811)
PDH-B (VF0878) (HFPHFIFB_00812)
HitABC (VF0268) (HFPHFIFB_00829)
818
824
817 819 820 822 823 825 827 828 830 831 832 833
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Inner membrane ABC transporter permease protein YdcV (LOHHAILK_00836)
Error-prone DNA polymerase (LOHHAILK_00826)
putative protein (LOHHAILK_00831)
828
834
827 829 830 832 833 835 837 838
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Error-prone DNA polymerase (AKPMGCPG_00839)
840
putative protein (AKPMGCPG_00843)
841 842
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L-arabinose transport system permease protein AraH (ILHBJHNJ_00841)
HTH-type transcriptional activator RhaR (ILHBJHNJ_00837)
Error-prone DNA polymerase (ILHBJHNJ_00820)
2-alkyl-3-oxoalkanoate reductase (ILHBJHNJ_00830)
Arabinose import ATP-binding protein AraG (ILHBJHNJ_00840)
putative protein (ILHBJHNJ_00826)
IS5 family transposase IS4811 (ILHBJHNJ_00832)
822
Pyridoxine 4-dehydrogenase (ILHBJHNJ_00831)
823
Aldo-keto reductase IolS (ILHBJHNJ_00828)
L-arabinose isomerase (ILHBJHNJ_00838)
821 824 825 827 829 833 834 835 836 839
IMGPR_plasmid_2718218232_000003 865904:890530(+)
Error-prone DNA polymerase (PHKELAMI_00931)
Error-prone DNA polymerase (PHKELAMI_00930)
Quorum-sensing (VF0433) (PHKELAMI_00934)
PHKELAMI_00932
PHKELAMI_00933
N-(3-oxo-hexanoyl)-Lhomoserine lactone QS system (VF0906) (PHKELAMI_00935)
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Phthiocerol synthesis polyketide synthase type I PpsC (DAHBKGKO_00811)
putative HTH-type transcriptional regulator YybR (DAHBKGKO_00815)
Low affinity potassium transport system protein kup (DAHBKGKO_00817)
IS3 family transposase ISRel21 (DAHBKGKO_00816)
Quinone oxidoreductase 2 (DAHBKGKO_00814)
Error-prone DNA polymerase (DAHBKGKO_00808)
812
809 810 813 818 819
HTH-type transcriptional regulator DmlR (DAHBKGKO_00820)
IMGPR_plasmid_2721755822_000003 860757:878028(+)
Phthiocerol synthesis polyketide synthase type I PpsC (OHDGKPEO_00811)
Error-prone DNA polymerase (OHDGKPEO_00808)
812
809 810 813
Quinone oxidoreductase 2 (OHDGKPEO_00814)
IMGPR_plasmid_2721755685_000003 860757:878028(+)
9000 nt
Virulence