





CD030_RS16270:ABC transporter ATP-binding protein/permease; protein_CD030_RS07580:purtl, formyltetrahydrofolate deformylase; protein_id=WPCD030_RS03630:GGDEF domain-containing phosphodiesterase; protein_CD030_RS03630:GGDEF domain-containing phosphodiesterase; protein_

CD030_RS03630:GGDEF domain–containing phosphodiesterase; protein, CD030_RS05195:mscL, large conductance mechanosensitive channel pro CD030_RS04135:MBL fold metallo–hydrolase; protein_id=WP_015445556 CD030_RS15555:cobN, cobaltochelatase subunit CobN; protein_id=WP_0

CD030_RS06970:TonB-dependent receptor; protein_id=WP_010968430.1 CD030_RS06960:MFS transporter; protein_id=WP_010968432.1 CD030_RS04545:flhA, flagellar biosynthesis protein FlhA; protein_id=WP_CD030_RS05080:SDR family NAD(P)-dependent oxidoreductase (pseudo CD030_RS00270:A2C family ABC transporter permease; protein_id=WP_

CD030_RS02435:rkpf, UDP_glucose 6-dehydrogenase; protein_id=WP_CD030_RS02435:rkpf, UDP_glucose 6-dehydrogenase; protein_id=WP_CD030_RS08438:autoinducer binding domain—containing protein (pseudogenase). CD030_RS04400:XRE family transcriptional regulator; protein_id=WP_010CD030_RS01240prA, DNA-processing protein DprA; protein_id=WP_010D030_RS01240prA, DNA-processing protein protein_protein_id=WP_010908_CD030_RS0125;sppA, signal peptide peptidase SppA; protein_id=WP_010908_CD030_RS01253:winged helix—turn—helix domain—containing terratricopeption_CD030_RS01735:winged helix—turn—helix domain—containing terratricopeption_CD030_RS01730:winged_terratricoper_domain_containing_terratricoper_domain_conta

CD030_RS00140:exbD, TonB system transport protein ExbD; protein_id=W CD030_RS02865:HAMP domain-containing histidine kinase; protein_id=W CD030_RS16895:BCCT family transporter; protein_id=WP_010969412.1 CD030_RS0525:Rrl2 family transcriptional regulator; protein_id=WP_081876498.1 CD028_RS01640:hypothetical protein_jrotein_id=WP_088195498.1

CD030_RS16855-in/pothetical protein; protein_id=WP_015445609.1
CD030_RS16125-LIA synthase family protein; protein id=WP_003534093.
CD030_RS01435:NADH ubiquinone oxidoreductase; protein_id=WP_0109
CD030_RS01610:hypothetical protein; protein_id=WP_013844967.1
CD030_RS07580:purU, formyltetrahydrotolate deformylase; protein_id=WP_013844967.3

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CD030_RS16263.cytochrome cf; protein_id=WP_01999491.1
CD030_RS11905.electron transfer flavoprotein subunit beta/FixA family proc CD030_RS16025.0mpA family protein; protein_id=WP_04170153.1
CD030_RS06705.GIY_YIG nuclease family protein; protein_id=WP_01096

CD030_RS06705:GIY_YIG nuclease family protein; protein_id=WP_01096 CD030_RS17140:molybdopterin molybdotransferase MoeA; protein_id=WP CD030_RS02085:pdxA, 4-hydroxythreonine-4-phosphate dehydrogenase CD030_RS0200:hybrid sensor histidine kinase/response regulator; protein

CD030_RS05203_UAX, 4—Iyoux/inteninie—phiospirate entrylinogeriase CD030_RS05200-hybrid sensor histidine kinase/response regulator; proteii CD030_RS16380:recA, recombinase RecA; protein_id=WP_010969478.1 CD030_RS02735-PAS-domain containing protein; protein_id=WP_013844 CD030_RS12270:glutamine synthetase family protein; protein_id=WP_013