ACTION OF THE PROPERTY OF THE		DD_3	LL ₃	Diel ₃	DD ₆	LL ₆	Diel ₆	
Company								SAR116_0006 hypothetical protein
Section of the control of the contro	K09013 sufC; Fe-S cluster assembly ATP-binding protein							SAR116_0015 FeS assembly ATPase SufC
### ### ### #### #####################								SAR116_0048 transcriptional regulatory protein, Lacl family, putative
SOURCE CONTROL	K01808 rpiB; ribose 5-phosphate isomerase B [EC:5.3.1.6]							SAR116_0160 sugar–phosphate isomerase, RpiB/LacA/LacB family
### 1970 Common	K03499 trkA; trk/ktr system potassium uptake protein							SAR116_0219 K+ transport systems, NAD-binding component
FORTER AND CONTROL CON								SAR116_0259 hypothetical protein
### PATE OF THE CONTRIBUTION OF THE PATE OF THE PATE OF THE CONTRIBUTION OF THE PATE OF TH	K00010 iolG; myo-inositol 2-dehydrogenase / D-chiro-inositol 1-dehydrogenase [EC:1.1.1.18 1.1.1.369]							SAR116_0266 myo–inositol dehydrogenase (idhA)
EXTRACTOR COLOR CO	K03116 tatA; sec-independent protein translocase protein TatA	. –						SAR116_0415 twin–arginine translocation protein, TatA/E family subunit
Application and continues that is written under a second color of the property of the color of t	K00817 hisC; histidinol–phosphate aminotransferase [EC:2.6.1.9]							SAR116_0442 histidinol–phosphate aminotransferase
EXECUTION CONTRIBUTION CONTRIBU								SAR116_0444 hypothetical protein
TRIBUTION CONTROL OF THE CONTROL OF	K05568 mnhD; multicomponent Na+:H+ antiporter subunit D							SAR116_0460 NADH dehydrogenase (quinone)
Month Color Table Tabl	K02234 cobW; cobalamin biosynthesis protein CobW							SAR116_0572 cobalamin biosynthesis protein CobW
ACCIDITION OF ANY OF CONTROL PROCESSOR AND ACCIDITION OF ACCIDITI								SAR116_0573 hydrogenase accessory protein
### ACTIVE AND Company of Active transport of Active Company of Ac								SAR116_0878 sulfate permease family protein
ACTION ON THE ANALYSIS AND COLOR OF CALLES AND	K01686 uxuA; mannonate dehydratase [EC:4.2.1.8]							SAR116_0904 mannonate hydrolase
ACCEPTION CONTRACTOR C								SAR116_0933 putative major facilitator family (MFS) transporter
ACCEPTING Section Control control section and extended the Control of Acceptance of Control of Cont	K02160 accB; acetyl-CoA carboxylase biotin carboxyl carrier protein							
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