

TABLE S3 Summary of genes up-regulated in CF74ΔT6SS

gene_id	gene	product	direction	log2 (fold_change)	p_value
CFCDC_1139	<i>malK</i>	maltose/maltodextrin transporter ATP-binding protein	+	-3.7477	0.0007
CFCDC_2642	<i>csgC</i>	autoagglutination protein	+	-11.0695	0.0028
CFCDC_0517	<i>dcuD</i>	C4-dicarboxylate transporter	-	-3.2288	0.0029
CFCDC_0476	<i>nanK</i>	N-acetylmannosamine kinase	+	-3.2815	0.0173
CFCDC_1137	<i>malF</i>	maltose transporter membrane protein	-	-2.3432	0.0177
CFCDC_1044	<i>evgA</i>	DNA-binding transcriptional activator EvgA	+	-3.9065	0.0182
CFCDC_1749	<i>sbp</i>	sulfate-binding protein (sulfate starvation-induced protein 2)	+	-2.2109	0.0219
CFCDC_1140	<i>lamB</i>	maltoporin	+	-2.4846	0.0255
CFCDC_0153	<i>malS</i>	periplasmic alpha-amylase precursor	-	-2.1120	0.0277
CFCDC_3321	<i>yeeN</i>	hypothetical protein	-	-2.1912	0.0284
CFCDC_2753	<i>potD</i>	spermidine/putrescine ABC transporter periplasmic substrate-binding protein	-	-2.1112	0.0288
CFCDC_1138	<i>malE</i>	maltose ABC transporter periplasmic protein	-	-2.2756	0.0347
CFCDC_1255	<i>ecnR</i>	transcriptional regulatory protein	-	-2.0784	0.0412
CFCDC_1829	<i>zraP</i>	zinc resistance protein	-	-11.1286	0.0443
CFCDC_2975	<i>lcrS</i>	LcrS	-	-10.4444	0.0443