

Supplementary Table 3: Genes differentially expressed at the basal level (± 2 -fold) in *mutS* mutant *E. coli* compared to wildtype. Only genes for which ANOVA $p \leq 0.05$ are listed.

Amino acid transport and metabolism				
Gene	Blattner Number	Signal Log Ratio	ANOVA p-value	Gene Product and Function
gadB	b1493	1.10	0.000	glutamate decarboxylase isozyme
trpA	b1260	1.71	0.048	tryptophan synthase, alpha protein
trpB	b1261	1.65	0.027	tryptophan synthase, beta protein
trpC	b1262	1.52	0.013	N-(5-phosphoribosyl)anthranilate isomerase and indole-3-glycerolphosphate synthetase
trpD	b1263	1.49	0.016	anthranilate synthase component II, glutamine amidotransferase and phosphoribosylanthrani
trpE	b1264	1.13	0.023	anthranilate synthase component I
Carbohydrate transport and metabolism				
Gene	Blattner Number	Signal Log Ratio	ANOVA p-value	Gene Product and Function
eno	b2779	1.12	0.052	enolase
fba	b2925	1.12	0.009	fructose-bisphosphate aldolase, class II
malE	b4034	1.60	0.017	periplasmic maltose-binding protein; substrate recognition for transport and chemotaxis
malK	b4035	1.35	0.048	ATP-binding component of transport system for maltose
Cell motility				
Gene	Blattner Number	Signal Log Ratio	ANOVA p-value	Gene Product and Function
flgE	b1076	1.25	0.027	flagellar biosynthesis, hook protein
fliC	b1923	1.26	0.045	flagellar biosynthesis; flagellin, filament structural protein
Energy production and conversion				
Gene	Blattner Number	Signal Log Ratio	ANOVA p-value	Gene Product and Function
ppa	b4226	1.13	0.033	inorganic pyrophosphatase
Replication, recombination and repair				
Gene	Blattner Number	Signal Log Ratio	ANOVA p-value	Gene Product and Function
dps	b0812	1.02	0.029	global regulator, starvation conditions
Translation				
Gene	Blattner Number	Signal Log Ratio	ANOVA p-value	Gene Product and Function
rplO	b3301	1.42	0.052	50S ribosomal subunit protein L15
Unclassified, unknown or general function prediction only				
Gene	Blattner Number	Signal Log Ratio	ANOVA p-value	Gene Product and Function
lamB	b4036	2.00	0.039	phage lambda receptor protein; maltose high-affinity receptor
yfiD	b2579	1.20	0.009	putative formate acetyltransferase