

Table 3: Effect of Low Fe Growth on Transcription of Genes in the Pigment Synthesis Pathway

Enzyme	Gene Name	Gene Designation	Regulation in LoFe (+) ↑ or (-) ↓)	p-value
<i>Chl Biosynthesis</i>				
Glu-tRNA synthetase	<i>gltx</i>	s110179	--	
Glu-tRNA reductase	<i>hemA</i>	slr1808		
Glutamate-1-semialdehyde aminotransferase	<i>hemL/gsa</i>	s110117	--	
PBG synthase	<i>hemB</i>	s111994	--	
Porphobilinogen deaminase	<i>hemC</i>	slr1887	--	
Uroporphyrinogen III synthase	<i>hemD</i>	s110166	--	
Uroporphyrinogen III decarboxylase	<i>hemE</i>	slr0536	--	
Coproporphyrinogen III oxidase	<i>hemF</i> <i>O₂-indep.</i> <i>hemN</i>	s111185 s111876 s111917	-1.6X (12h) -4.0X (12h) -1.5X (12h)	0.0015 2 x 10 ⁻⁸ 0.0026
Protoporphyrinogen IX oxidase	<i>hemK</i>	s111237		
Ferrochelatase	<i>hemH</i>	slr0839		
Magnesium chelatase	<i>chlI</i> <i>chlH</i> <i>chlD</i>	slr1030 slr1055 slr1777	-- -- -1.8X	
Mg-protoporphyrin IX methyl-transferase	<i>ChlM</i>	slr0525	-1.5X	0.02
Cyclase	<i>BchE</i>	slr0905	-3.4X	2 x 10 ⁻⁸
Protochlorophyllide reductase	<i>chlN</i> <i>chlB</i> <i>chlL</i>	slr0750 slr0772 slr0749	-6.3X (12h) ---	6 x 10 ⁻⁸
Light-induced, Fe-subunit	<i>bchB</i> (or <i>bchK</i>)	ssr2049	-1.4X	0.04
Protochlorophyllide oxidoreductase	<i>pcr</i>	slr0506	+1.5X	0.014
Chlorophyll synthetase	<i>chlG</i>	slr0056	+1.7X	0.01
<i>Phycobilin Synthesis</i>				
Heme oxygenase	<i>ho1</i> <i>ho2</i>	sllll84 slll875	-1.6X	0.02
Biliverdin reductase	<i>bvdR</i>	slr1784		