

Overexpressed genes				
GO term	Genes	Log2fold Change		
		24°C	28°C	30°C
Drug metabolic process (GO:0017144)	ATP synthase F1 subunit gamma protein (LbrM.33.0830)	1.681	1.487	1.559
	ATP synthase. epsilon chain (LbrM.30.3630)	1.801	1.968	2.224
	Ubiquinol-cytochrome c reductase complex 14kd subunit (LbrM.34.0140)	1.237	1.144	1.316
	ATP synthase F1. alpha subunit (LbrM.05.0510)	1.559	1.057	1.423
	ATP synthase delta (OSCP) subunit (LbrM.35.7110)	1.721	1.570	1.796
Cellular response to external stimulus (GO:0071496)	Eukaryotic translation initiation factor 2 subunit alpha (LbrM.03.0890)	1.145	1.513	1.511
Cellular response to extracellular stimulus (GO:0031668)				
Cellular response to nutrient levels (GO:0031669)				
Cellular response to starvation (GO:0009267)				
Response to extracellular stimulus (GO:0009991)				
Response to nutrient levels (GO:0031667)				
Response to starvation (GO:0042594)				
Cellular response to heat (GO:0034605)	ALBA-domain protein 1 (LbrM.13.0270)	1.958	1.271	1.561
Response to heat (GO:0009408)				