

Table S8 Results for one-way ANOVAs with Dunnett's tests comparing pre-stress relative transcript abundances with high temperature (38.5°) and recovery (25C°). LSMean= least squares mean, significant terms are bolded.

Gene	Transcript	time	LSMean	Pvalue
CG10264	mRNA	0	-16.7583361	
		0.25	-15.9483352	0.9714
		0.53	-16.3533329	0.9995
		4.315	-15.5166648	0.8255
		8.315	-15.4900004	0.8124
		12.315	-15.0916692	0.5932
		36.315	-15.9966668	0.9791
		48.315	-16.2116654	0.9967
		Pre-mRNA	0	-16.4033326
	0.25		-12.4600007	0.0001
	0.53		-12.0166658	0.0000
	4.315		-11.9433335	0.0000
	8.315		-15.4550002	0.5240
	12.315		-16.8533346	0.9638
	36.315		-17.7083332	0.2303
	48.315		-16.7416690	0.9918
	CG12267	mRNA	0	-7.4566667
0.25			-7.1916666	0.9822
0.53			-6.6366666	0.2932
4.315			-7.4450001	1.0000
8.315			-6.9816666	0.7843
12.315			-6.9500000	0.7369
36.315			-7.0916666	0.9188
48.315			-7.5283333	1.0000
pre-mRNA			0	-15.3516682
		0.25	-11.1433333	0.0001
		0.53	-8.1049998	0.0000
		4.315	-8.8616665	0.0000
		8.315	-9.5366668	0.0000
		12.315	-10.0699998	0.0000
		36.315	-9.8750003	0.0000
		48.315	-11.4766660	0.0002
CG32187		mRNA	0	-16.8366663
	0.25		-16.9783335	0.9999
	0.53		-16.3133328	0.8981
	4.315		-13.4783311	0.0002
	8.315		-12.8400027	0.0001
	12.315		-15.5800008	0.1967
	36.315		-15.5066658	0.1585
	48.315		-16.1916670	0.7838
	pre-mRNA		0	-16.2349999
		0.25	-15.3650017	0.4116
		0.53	-15.8116684	0.9310
		4.315	-13.7866662	0.0014
		8.315	-13.9674991	0.0068
		12.315	-16.0000006	0.9968
		36.315	-15.9383329	0.9879
		48.315	-16.6700025	0.9225

<i>CG5608</i>	mRNA	0	-8.3016668	
		0.25	-8.0700000	0.9986
		0.53	-7.6583333	0.7849
		4.315	-8.4150002	1.0000
		8.315	-7.4016666	0.4928
		12.315	-7.8383334	0.9388
		36.315	-7.0016667	0.1721
		48.315	-7.9349999	0.9802
		pre-mRNA	0	-14.3666667
	0.25		-9.8333336	0.0000
	0.53		-6.6416667	0.0000
	4.315		-7.9500001	0.0000
	8.315		-9.2449999	0.0000
	12.315		-11.7933334	0.0019
	<i>Hsp83</i>	mRNA	0	-2.2916667
0.25			-1.6900000	0.7346
0.53			-1.3366667	0.3080
4.315			0.5733333	0.0002
8.315			0.2375000	0.0024
12.315			-2.4100000	1.0000
36.315			-1.4116667	0.3833
48.315			-1.3033333	0.2782
pre-mRNA			0	-14.8999997
		0.25	-7.6866666	0.0000
		0.53	-6.5233334	0.0000
		4.315	-5.7883333	0.0000
		8.315	-7.0400001	0.0000
		12.315	-8.7466664	0.0000
<i>aur</i>		mRNA	0	-4.3550000
	0.25		-3.8966667	0.9939
	0.53		-3.9383333	0.9965
	4.315		-5.2000000	0.8809
	8.315		-3.3883333	0.8069
	12.315		-3.7900000	0.9810
	36.315		-5.3866667	0.7621
	48.315		-3.9850000	0.9983
	pre-mRNA		0	-17.2433333
		0.25	-9.8766670	0.0000
		0.53	-7.1400001	0.0000
		4.315	-7.9633332	0.0000
		8.315	-7.1616666	0.0000
		12.315	-8.1666669	0.0000
	<i>Gr85a</i>	mRNA	0	-18.4216659
0.25			-18.1483327	0.9992
0.53			-18.4624992	1.0000
4.315			-17.1700001	0.4185
8.315			-16.7466676	0.1727
12.315			-17.0500007	0.3321

	36.315	-18.2216673	0.9999
	48.315	-17.7816679	0.9174
pre-mRNA	0	-17.4133322	
	0.25	-12.1449989	0.0000
	0.53	-11.9566662	0.0000
	4.315	-12.3683326	0.0000
	8.315	-13.8200010	0.0005
	12.315	-15.0516677	0.0189
	36.315	-15.2083341	0.0296
	48.315	-15.5583342	0.0789
