Strain	Copy #a	Mean	CV	Chr.	Trans-	Construct Details	Method of	Transgene
		Exp. ^b	(%) °		gene Allele		Integration	Configuration
TJ375	530	381	22	IV	gpls1	P _{hsp-16.2} ::gfp::unc-54	Gamma	Simple Array
TJ2735 ^d	450	409	26	V	zls2735	P _{hsp-16.2} ::gfp::unc-54	UV	Simple Array
TJ2732	500	73	24	V	zls2732	P _{hsp-16.2} ::dsRed::let-858	UV	Simple Array
TJ2733	940	90	23	IV	zls2733	P _{hsp-16.2} ::dsRed::let-858	UV	Simple Array
CL2070	20	70	24	V	dvls70	P _{hsp-16.2} ::gfp::unc-54;	Gamma	Complex Array
						rol-6(su1006)		
CL2071	22	76	26	III	dvls71	P _{hsp-16.2} ::gfp::unc-54;	Gamma	Complex Array
						rol-6(su1006)		
CL2074	21	103	25	٧	dvls74	P _{hsp-16.2} ::gfp::unc-54;	Gamma	Complex Array
						rol-6(su1006)		
TJ3000 ^d	1	29	26	II	zSi3000	P _{hsp-16.2} ::gfp::unc-54;	MosSCI	Single Copy
						cb-unc-119(+)		
TJ3001	1	32	23	II	zSi3001	P _{hsp-16.2} ::gfp::unc-54;	MosSCI	Single Copy
						cb-unc-119(+)		

a - Copy number was determined via qPCR, using ama-1 as a control.

b – Mean expression level for each reporter derived from several measurements of population mean. Each value was the measured average photomultiplier tube counts.

c - No CV for any *hsp-16.2* reporter was found to be significantly different from any other *hsp-16.2* reporter.

d - These strains were measured thrice; they were not measured as many times as the other reporters. All other strains were measured in together in a series of five or ten experiments (See Materials and Methods). Data shown for TJ375 is from the five experiments with other multicopy reporters in experimental series one. Mean from the ten experiments with TJ3001 is shown in Table 2. Mean expression values for the single copy reporters are not significantly different from one another.