Table S1: Supplementary information on genes in the three gene sets

Walser et al. 2006
Heat-shock promoters: targets for evolution by *P* transposable elements in *Drosophila*

Sources used for the tables and for further information:

(1) Drosophila developmental gene expression timecourse
http://genome.med.yale.edu/Lifecycle/

Drosophila Developmental Gene Expression Timecourse (http://genome.med.yale.edu/Lifecycle/).

Color coded time course for gene list: 

Yellow color represents high relative levels of expression while blue represents low levels. The brightest color is three-fold or greater differential from the reference black

(2) FlyBase - a database of the Drosophila Genome
http://flybase.bio.indiana.edu/

(3) FlyGRID - a database of genetic and physical interactions developed
http://biodata.mshri.on.ca:80/fly_grid/servlet/SearchPage

(4) MitoDrome – a database with the aim to annotate the complete set of *Drosophila melanogaster* nuclear genes encoding for mitochondrial proteins in order to contribute to their functional characterization.
http://www2.ba.itb.cnr.it/MitoDrome/

(5) BDGP Gene Expression Summary
http://www.fruitfly.org/cgi-bin/ex/bquery.pl?qpage=entry&qtype=summary

Gene Set I. Heat-inducible genes, *hsp* genes, chaperone genes. ID: FlyBase ID; SYM This is the standard abbreviation (gene symbol) for the name of the gene.; Location: Chromosome arm and Location of the gene on the cytological (polytene chromosome) map. Developmental Gene Expression Timecourse (for color code see above)

					Metamorphosis		Adult		
ID	SYM	Location	Embryo	Larvae	Prepupae	Pupae	□	□	Ref.
CG4460	*Hsp22*	3L-67B2							1-5
CG4463	*Hsp23*	3L-67B3							1,5
CG4183	*Hsp26*	3L-67B2							1,5
CG4466	*Hsp27*	3L-67B3							1,5
CG5436	*Hsp68*	3R-95D11							5
CG18743	*Hsp70Ab*	3R-87A3							1,5,6
CG1242	*Hsp83*	3L-63B11							5
CR31400	*Hrsω*	3R-93D4							7
CG10578	*DnaJ-1*	3L-64E5							8
CG12240	*DnaJ-60*	2R-60C1							9
CG5504	*DnaJ*	2R-59F6							10
CG9920		3R-88A8							
CG12101	*Hsp60*	X-10A4							11
CG2830	*Hsp60b*	2L-21E2							12
CG7756	*Hsc70-2*	3R-87D10							13-15
CG4147	*Hsc70-3*	X-10E3-4							14,15
CG4264	*Hsc70-4*	3R-88E4							13-15
CG8542	*Hsc70-5*	2R-50E6							14,15
CG4167	*Hsp67Ba*	3L-67B21							5

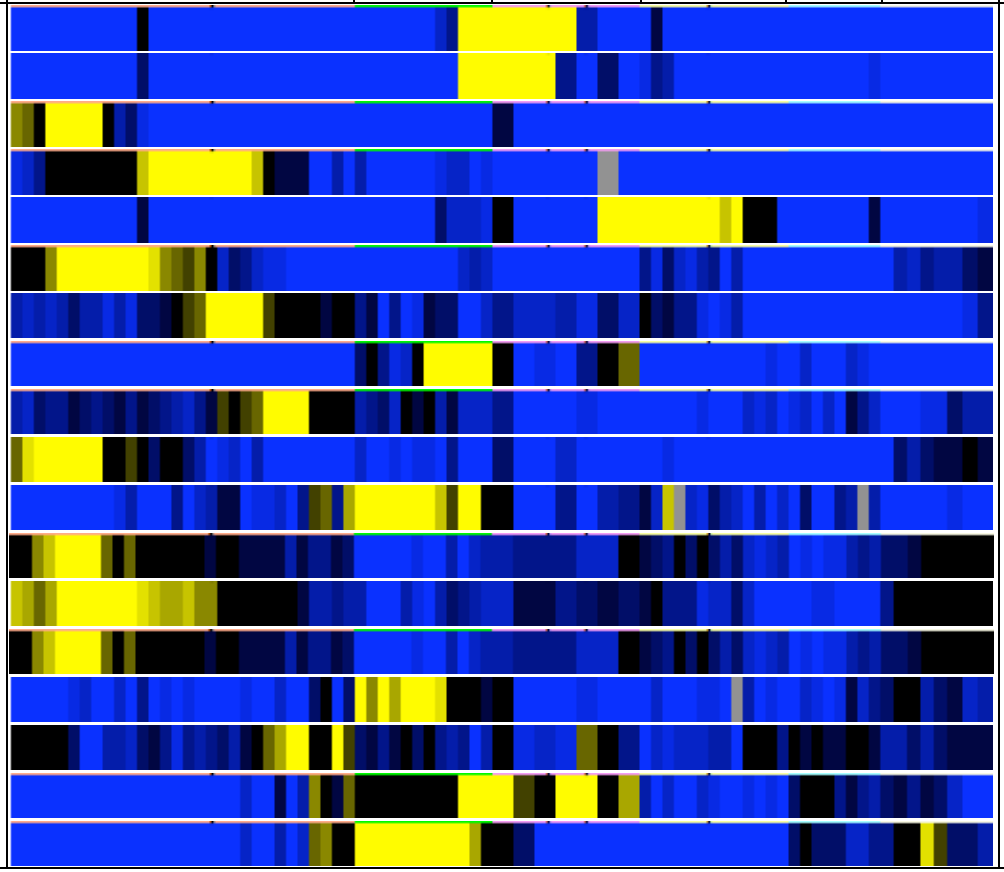
Red: Location in areas known for high recombination (Charlesworth B (1996) Background selection and patterns of genetic diversity in Drosophila melanogaster. Genetical Research 68: 131-149)

Gene Set II. Genes resembling *hsp70* in regulation by polymerase pausing or DNAse hypersensitivity. ID: FlyBase ID; SYM This is the standard abbreviation (gene symbol) for the name of the gene; Location: Chromosome arm and Location of the gene on the cytological (polytene chromosome) map. Genes resembling *hsp70* in regulation by A) polymerase pausing, B) Dnase hypersensitivity, C) high germline transcriptional activity [16], D) dublicated genes. Developmental Gene Expression Timecourse (for color code see above) and Reference.

				Metamorphosis		Adult		
ID	SYM	Location		Embryo	Larvae	Prepupae	Pupae	Ref.
CG12055	*Gapdh1*	2R-43E17	A					17
CG8893	*Gapdh2*	X-13F17	A					18-21
CG9277	*βTub56D*	2R-56D7-8	A					18
CG10798	*dm*	X-3D2	A					22-25
CG3481	*adh*	(2L)-35B3	A					
CG12181	*Sgs4*	X-3C10	B					26,27
CG6222	*su(s)*	X-1B13	B					28
CG4027	*Act5C*	X-5C7	B					18,29,30
CG4550	*ninaE*	3R-92B4	B					31
CG7939	*RpL32*	3R-99D3	B					32
CG6382	*Elf*	2L-33E4-5	C					
CG8040	*CG32061*	3L-67E3	C					
CG13340		2R-50C5	C					
CG4750		2R-53C7	C					
CG2512	*αTub84D*	3R-84D9	D					33-36
CG1913	*αTub84B*	3R-84R4	D					33-36
CG2238	*Ef2b*	2L-39Ef	D					37
CG11181	*cup*	X-26F5-6	D					38

Red: areas known for high recombination (Charlesworth B (1996) Background selection and patterns of genetic diversity in Drosophila melanogaster. Genetical Research 68: 131-149)

Gene Set III. Non-*hsp* genes with limited transcription during the life cycle and negligible germ-line expression (features expected to minimize transposition and its vertical transmission). ID: FlyBase ID; SYM This is the standard abbreviation (gene symbol) for the name of the gene.; Location: Chromosome arm and Location of the gene on the cytological (polytene chromosome) map. Developmental Gene Expression Timecourse (for color code see above) and Reference.

					Metamorphosis		Adult		
ID	SYM	Location	Embryo	Larvae	Prepupae	Pupae	□	□	Ref.
CG7906		3L-70D4							
CG7924		3L-70D4							
CG15634		2L-25A3							
CG12011		3L-62B1							
CG4818		3L-72E2							
CG2916	*Sep5*	2R-43F8							
CG10591		3L-64D5							
CG11650	*Lcp1*	2R-44CR							
CG2961	*Ipod*	X-9B4							
CG9506	*slam*	2L-26C4							
CG7252		3L-68E3							
CG4602	*Srp54*	2L-30E4							
CG9984	*TH1*	X-14C2							
CG14025	*Bsg25D*	2L-25D2-3							
CG10154		3L-70B1							
CG7724		3L-73E5							
CG2070		2R-43E8							
CG6296		3R-97D13							

Red: Location in areas known for high recombination (Charlesworth B (1996) Background selection and patterns of genetic diversity in Drosophila melanogaster. Genetical Research 68: 131-149)

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