McKim lab stocks

Location	Genotype	Comments
G001	*y w; polo[s132408]/TM3*	hypomorph, P allele
G002	*y/y[+]Y; al dp b pr cn c px sp/CyO*	
G003	*w[1118]/Dp(1;Y)y[+]; noc[Sco]/SM6a*	
G004	*al dp b pr cn c px sp/CyO*	
G005	*T(2;3)dp[D], dp[D]/SM5*	outcross
G006	*yw/y[+]Y; mei-1685/TM3*	
G007	*ry[531]/MKRS*	
G008	*y pn cv m f . y[+]/C(1)DX, y w f/Y*	C(1)DX from Hawley G14B
G009	*T(2;3)ltx13*	homo viable
G010	*ry[609]/MKRS*	
G011	*T(3;4)A2/ TM6B*	outcross 886
G012	*y w; T(2;3)B3, CyO: TM6B, Tb/ Pin[88K]*	No Cy
G013	*y w; mast[1]/ TM3*	P allele, S068607
G014	*yw/y[+]Y; mei-217/TM3*	
G015	*mei-217/TM6*	
G016	*yw/y[+]Y; mei-751/TM3*	
G017		
G018	*cn bw*	
G019	*T(1;3)JA29, l(1)1Fc[2]/FM7a*	
G020	*MKRS, ry kar/TM2, ry Ubx*	
G021	*yw/y[+]Y; C(4)RM, ey[R] ci*	
G022	*T(2;3)ltx16/ SM1*	
G023	*mei-1685/TM6*	
G024	*mei-2024/TM6*	

Location	Genotype	Comments
G025	*MKRS, ry kar/TM6B, ry*	
G026	*yw/y[+]Y; mei-2024/TM3*	
G027	*y w [G?7c23]; Gla/SM6a*	Hobo free
G028		
G029	*T(2;3)C202, al/SM1*	
G030	*al dp b pr cn/ SM6a*	
G031	*al[1] dp[ov1] b[1] pr[1] Bl[1] cn[1] c[1] px[1] s*	
G032	*y/y[+]Y; al dp b Sp pr cn bw / CyO*	From G71 with marked X and Y.
G033	*T(3;4)A30/ TM6B*	from 890
G034	*mei-2541/TM6*	
G035	*C(1:Y)1, y v f B:y[+]/C(1)RM, y v; C(4)RM, ci ey[*	= Y[S]X . Y[L], In(1)EN
G036	*Gal4 P{UASP:lacIGFP}; P{lacO}2-2*	nosGal4?, labelled UAS:lacO
G037	*w; mnk[P6]/ SM6, TM6B, Tb*	from M. Brodsky
G038	*y cv v l(1)692-19 f / FM7w/y[+]Y*	FM7 made in Hawley lab with a
G039	*T(2;3)C287/TM6B, Tb*	from 1210
G040	*P{UASP:Incenp[myc]}10/TM3*	N-terminal myc tag
G041	*thr kar ry[606] cv-c*	
G042	*y[1]? w[67c23]?; pch2[EY01788a] e ca*	recombinant rom 15536 to remov
G043	*y w; polo[s025604]/ TM3*	P allele
G044	*mio[1]/ SM6*	from M. Lilly
G045	*nosGal4; P{lacO}2-16/FM7/y[+]Y*	labelled UAS:lacO
G046	*Dp(1;Y)B[S]Yy[+] /C(1)DX, y f/R(1)2, y w f*	Ring
G047	*C(2)EN, b pr*	Hawley B57
G048	*y/y[+]Y; cn[35]; spa[pol]*	Hawley AB12
G049	*T(2;3)S[L], S[L]/In(2L)Cy, In(2R)Cy, Cy[1] E(S)[1*	
G050	*In(1)dl49, y Hw m g/ C(1)DX, y f/ y[+]Y; spa[pol]*	Hawley A71

Location	Genotype	Comments
G051	*+/y[+]YB[S]; C(2)EN, bw sp*	Hawley AB3
G052	*y cv v f car/y[+]; spa[pol]*	Hawley A75R3
G053	*C(3L), st[1]; C(3R), e[s]*	Hawley C4
G054	*FM7,w/y[+]Y*	
G055	*y w/ y[+]Y; Pr, ru h th st cu sr e ca / TM3*	From G59 with marked X and Y
G056	*Bwinscy/ C(1)DX, y f/ y[+]Y*	Hawley A262
G057	*C(2L), b; C(2R), px*	Hawley B23
G058	*+/B[S}Y; st/TM6, Sb*	outcrossed, isogenized
G059	*ru h thr st cu sr e Pr ca/ TM6B, Bsb Tb*	Hawley C45
G060	*y w /y[+]Y; D/TM3, Sb Ser*	Hawley AC21
G061	*Df(2L)f02036-d04880*	Df of cdc14, viable
G062	*P{lacO}2-19; D/TM3*	labelled UAS:lacO
G063	*y Hw w/ C(1)DX/ B[S]Y*	
G064	*w; ru h P{FRT(X97,I-site)RS3r}75CD sr e ca/ TM*	Tandem white with SceI between
G065	*H2lvd::GFP / TM3*	GFP fusion of variant histone
G066	*y(?) w dnl4[169a]*	ligase 4 mutant
G067	*ru h th st cu sr e ca/ TM6B, Tb*	derived from BT 576 and G96, h
G068	*mio[2] b pr/ SM6*	from M. Lilly
G069	*P{UASP:Incenp[myc]}1/TM3*	N-terminal myc tag
G070	*In(2LR)bw[VI], ho dp b pr bw[VI]/ CyO = Plum*	
G071	*al dp Sp b pr cn bw/ CyO*	probably carries a bw mutation
G072	*w; P{UASP:ZFMryA14D}, P{UASP:ZFMryB9B}/C*	Two ry zinc fingers, on 2, homo f
G073	*y/y[+]Y; cn bw/ SM6; ry[531]*	not downstairs
G074	*P{UASP:Incenp[myc]}7/TM3*	N-terminal myc tag
G075	*TM6, Sb/ TM2 Ubx*	
G076	*Sb Ubx/ TM6B*	made from BT588

Location	Genotype	Comments
G077	*y/y[+]Y; b pr c px sp/ CyO*	from b pr c px sp homozygous
G078	*R(1)2, y f/Dp(1;Y)B[S]Yy[+]/C(1)DX, y f*	
G079	*FM7c/y[+]Y/C(1)DX*	
G080	*y/y[+]Y;dp[lv1]b/SM5*	
G081	*P{UASP:Incenp[myc]}6/SM6*	
G085	*yw/y[+]Y; Sb Ubx/ TM6B, Dr*	
G086	*w; orb[F303]/ TM3, Ser*	
G087		
G088	*b BicD[R26] FRT40A / CyO, bw*	BicD null allele with FRT
G089	*w; orb[mel]/ TM3, Ser*	
G090	*BicD[PA66], cn bw/ CyO, bw*	BicD hypomorph - makes 16 cell
G091		
G092	*w; orb[F343]/ TM3, Ser*	
G093	*egl[PV27] cn bw/ CyO*	egl null allele = egl[5]
G094	*yw/y[+]Y; Dr/ TM3, Sb Ser*	
G095	*y/y[+]Y; cn; ry[531]; spa[pol]*	from G48 and G73.
G096	*+/B[S]Y; ru h th st cu sr e ca Pr/TM6B, Tb*	
G097	*st cu sr e ca/ TM6B, Tb Bsb*	from rucuca and G96
G098		
G099		
G100	*y pn cv m f . y[+]/ FM7c*	
G101	*y w/ y[+]Y; CycE[P2}; spa[pol]*	P allele, viable
G102	*cv Sb[sbd-2] gl e*	
G103		
G104	*st[1] Sb[sbd-1] e[s] ro[1] ca[1]*	
G105	*cu[1] kar[1]*	

Location	Genotype	Comments
G106	*w; P{UASP:ZFMryA14C}, P{UASP:ZFMryB1B}*	two ry zinc fingers, on 2
G107	*y w; P{w+mC y+mDint2=EPgy2}CG17509[EY064*	near spo76, got from Bellen
G108	*y[1]; cn[1] bw[1] sp[1]*	
G109	*y w/y[+]Y; CycE[P7}; spa[pol]*	P allele, viable
G110	*w; Sp/Cyo; Sb delta2-3/TM6*	from P36
G111	*w P{UASP:ZFMryA?}*	Zinc finger on X, ID number?
G112	*w P{UASP:ZFMryB2A}*	Zinc finger on X
G113	*Df(2R)BSC39, cn[1] bw[1]/SM6a, bw[k1]*	pds5 Df
G114		
G115	*y[1] Sxl[f2] / FM7c*	
G116	*y[1] cv[1] Sxl[f4] v[1] f[1] / FM0*	=Sxl[fs1]
G117	*y[1] cv[1] Sxl[f5] v[1] f[1] / FM3*	=Sxl[fs2]
G118	*P{lacO}2-13; Gal4 PUASP:lacIGFP}*	labelled UAS:lacO, nosGal4?
L_1	*neb[419]/TM3, Sb*	From Goldberg
L_2	*gamma-Tub37C[1] pr Ddc[ts1]/ CyO*	
L_3	*P{y[+mDint2] w[BR.E.BR]=SUPor-P}polo[16-1]*	
L_4	*Cks30A[RA74] cn[1] bw[1]/CyO, l(2)DTS513[1]*	
L_5	*y w; P{w+mC=lacW}Bub1[k03113]/CyO*	P allele, now BubR1?
L_6	*y/y[+]Y; sub[137D]/SM6*	Excision of EP616
L_7	*capu[1] cn[1] bw[1]/CyO, l(2)DTS513[1]*	
L_8	*twe[1] cn[1] /CyO*	recombinant removed bw and let
L_9	*w; L Pin/ Kr:Gal4 UAS:GFP, CyO*	Green balancer
L10	*Df(1)BK10, r f/Dp(1;Y)W73/C(1)DX*	Df(1)BK10 is a X-ray induced B
L11	*y/y[+]Y;cn sub[202]/SM6*	excision of EP616, recombinant
L12	*y(?)/y[+]Y; sub[131] bw sp/CyO*	excision of EP616, recombinant
L13	*y/y[+]Y; sub[202] bw/SM6*	excision of EP616, recombinant

Location	Genotype	Comments
L14	*trx[1] st/ TM1*	hypomorph, viable
L15	*asl[1]/ TM6C, Sb Tb*	male sterile
L16	*zw[10] FRT18A/ FM7c*	
L17	*Df(1)BK18/FM7; Dp(1;4)rK20/spa[pol]*	
L18	*w; mre11[58S] / CyO, S*	H230Y (CAT - TAT), originally
L19	*y cv v Df(1)815-6 f/FM7/y[+]Y; Dp(1;4)rK20*	Dp derived from Dp(1;4)r[+].
L20	*w[67c23] P{w[+mC]=lacW}ran[G0075]/FM7c*	P-insertion allele
L21	*Df(1)BK28, r f/ C(1)DX/ Dp(1;Y)W73; Dp(1;4)r[+*	copy 2□Dp(1;Y)W73 : Eberl et
L22	*y(?)/y[+]Y; cn sub[131] /CyO*	excision of EP616, recombinant
L23	*"4696" fs from stock center*	
L24	*gamma-Tub37C[3]/ CyO*	
L25	*w; mre11[delta35K1] (GFP)] / TSTL, CyO TM6B*	mre11 null, arm-GFP on chromo
L26	*y w; P{w+mC=lacW}64A pavB200 th st cu sr e ca/*	replaced original TMBB, Tb vers
L27	*d-tacc[stella592]/ TM3*	from J. Raff, d-tacc allele
L28	*al dp incenp[3747] sub[131]/SM6*	
L29	*Df(1)BK16/FM7; Dp(1;4)rK20/spa[pol]*	rk20 a rudimentary deletion of D
L30	*w; atm[wk], Pw+/ TM6B*	atm weak allele
L31	*asl[2]/ TM6C, Sb Tb*	late lethal
L32	*w[1118]; P{w+mC=lacW}ksrj5E2 cu e/TM3, Sb S*	Recombinant from 10212
L33	*r f B; Dp(1;4)fK4/+*	reverts to B+
L34	*y klp3A[1124] cv v f /FM7*	origin Hawley lab T012
L35	*b cn Incenp[QA26] c / SM6*	Incenp hypo, female sterile
L36	*aur[87Ac-3]/TM6B*	outcross from 6188, lethal allele
L37	*P{deltaCT}XC, w; l(3)His2Av[810]/TM6B*	Transgene is deleted for C-term p
L38		
L39	*rD1/C(1)DX/y[+]Y; Df(1;4)fK27/spa[pol]*	

Location	Genotype	Comments
L40	*cn[1] bie[*] bw[1] fs(2)ltoAPV63[APV63]/CyO, l(*	
L41	*Dp(1;Y)B[S]; ru st aur[1] e ca/TM3, Sb*	maternal effect
L42	*sub[131] ; ncd[1] ca[nd1]/T(2;3)B3, CyO, TM6B,*	double mutant lethal, no Cy, fro
L43	*w[*]; alphaTub67C[2] th[1] st[1] cu[1] sr[1] e[s]*	
L44	*w; atm[str], Pw+/ TM6B*	strong atm allele
L45	*w; myt[r3]/ TM6B, Tb Hu e*	viable, sterile?
L46	*al dp incenp[3747] sub[1] bw /CyO*	
L47	*Df(1)rD1, f /C(1)DX/y[+]Y; Dp(1;4)fK7/spa[pol]*	Associated with a T(3;4)
L48	*rD1/C(1)DX/y[+]Y; Dp(1;4;Y)fK24/spa[pol]*	Dp derived from Dp(1;4)r[+].
L49	*mus309[D3]/TM6*	from Sekelsky, Bloom's homolog
L50	*Df(1)BK28, r f/ C(1)DX/ Dp(1;Y)W73; Dp(1;4)r[+*	males only survive with both dup
L51	*Df(1)rD1/C(1)DX/y[+]Y; Df(1;4)fK4/spa[pol]*	
L52	*+; sub[HM26], cn bw/ CyO*	
L53	*msps[MJ208]/TM3*	Hypomorphic female sterile allel
L54	*msps[P]/TM6B*	lethal allele
L55	*mwh[1] Dhc64C[4-19] jv[1] ca[1]/TM6B, Tb[+] c*	
L56	*mwh[1] Dhc64C[6-10] h[1] st[1] p[p] e[s]/TM6B,*	
L57	*+/B[S]Y; sub[1], cn bw/CyO*	from BT5117
L58	*BubR1[k03113] sub[131] bw sp/SM6; ncdGFP*	ncd???
L59	*y klp3A[521] cv v f /FM0*	Hawley lab origin, T011, male le
L60	*P{ry+t7.2=PZ}Klp61F[07012] FRT/TM3, ry[RK]*	lethal, with FRT
L61	*P{ry+t7.2=PZ}Eb1[04524] cn/CyO; ry[506]*	lethal
L62	*b Eb1[2] cn/CyO*	semi-lethal???
L63	*y w; P{w+mC=lacW}cana[k07716]/CyO*	lethal
L64	*P{ry+t7.2=PZ}cmet[04431] cn/CyO; ry[506]*	lethal
L65	*w[1]; wee[ES1] cn[1]/CyO, P{w[+mC]=ActGFP}*	

Location	Genotype	Comments
L66	*w[1]; wee[DS1] cn[1]/CyO, P{w[+mC]=ActGFP}*	
L67	*feo[EA86]/ FM7a*	
L68	*y w; P{w+mC=lacW}CycB3[L6540]/TM3, Ser*	
L69	*y w; P{y+mDint2 wBR.E.BR=SUPor-P}CLIP-190[*	viable
L70	*y; P{y+mDint2 wBR.E.BR=SUPor-P}CLIP-190[K*	no Cy+, lethal?, some ry?
L71	*Bjl[vv179] FRT/ TM3; y w hsFLP*	hsFLP unknown
L72	*y w; P{FRT(w[hs])}G13 par-1[Δ16]/ CyO*	from Jan lab
L73	*w; CycB3[2]/TM3, Sb P{w+mC=35UZ}2*	
L74	*w; P{FRT(w[hs])}G13 incenp[3747]/ SM6*	no germ line clones
L75	*w; P{w+mC=UAS-CycA.W}II.2*	overexpress CycA
L76	*w; CycB[2]/CyO, P{ry+t7.2=ftz/lacB}E3*	lethal, rud ovaries with other alle
L77	*y; P{y+mDint2 wBR.E.BR=SUPor-P}CycB[KG088*	some homo, rud. Ovaries
L78	*Df(2L)Exel7049 sub[1] bw / CyO*	Aur B Df, from BT 7821
L79	*Dp(1;Y)B[S]; ru[1] st[1] polo[1] e[s] ca[1]/TM3,*	female sterile, hypomorph
L80	*w[1118]; P{w[+mC]=EP}Incenp[EP2340]*	P isertion, lethal allele
L81	*CycE[01672]/ CyO*	female sterile hypomorph
L82	*w; P{FRT(w[hs])}G13 px[1] sp[1] pds5[E3]/ SM6*	P-excision allele, lethal, other let
L83	*y w; pds5[E6]/ SM6*	P-excision allele, lethal, other let
L83	*w; P[w+, UASp-sub[L6]Rain/TM3, Sb*	lack of motor domain insertion, "
L84	*Df(3R)XS, Dp(3;3)XS, asp[1] ats[1] p[p]/TM6B, T*	
L85	*sub[1]; ncd[1] ca[nd1] /T(2;3)B3, CyO: TM6B, Tb*	no Cy expression, double mutant
L86		
L87	*w; al dp incenp[3747] c px sp/ CyO; Kr:GFP*	EMS allele, c?
L88	*y w[67c23]; P{w[+mC] y[+mDint2]=EPgy2}Ubc*	rad6, lethal
L89	*cn[1] cnn[HK21] bw[1]/CyO, l(2)DTS513[1]*	
L90	*CycE[712]/ CyO*	A Zuker collection allele, probab

Location	Genotype	Comments
L91	*y w; FRT rad50[CP1]/ CyO*	
M_1	*mei-218[6-7] car/C(1)DX/y[+]Y*	Gone from both collections?????
M_2	*y w mei-218[6] f car/FM7*	
M_3	*yw; mei-2024 th st cu e ca/ TM3*	
M_4	*sub[137D]/ CyO; nosGAL4::VP16*	
M_5	*y cv v mei-218[8] f car/C(1)DX/y[+]Y; spa[pol]*	
M_6	*mei-218[6-7] car/FM7*	
M_7	*mei-218[8] f car/C(1)DX/y[+]Y*	
M_8	*y cv v mei-218[7] f car/C(1)DX/y[+]Y*	
M_9	*y/y[+]Y; rec[2] bx[34e]/MKRS*	from Sekelsky
M10	*y/y[+]Y; ord[10] cn bw sp/SM1*	from Bickel 3/09, no homos?
M100	*c(2)M[EP]/CyO; c(3)G[68] ca/TM3*	
M101	*y/y[+]Y; ncd[1] ca[nd1] / TM3*	from BT1290 with lethal remove
M102	*y/y[+]Y; Sco sub[Dub] / SM1*	"iso Dy"
M103	*y/y[+]Y; pr cn sub[Dub] sp / SM1*	" iso AB"
M104	*w; P{w[+], gc(2)M[myc]}III.5*	from Page, on 2
M105	*al dp c(2)M[EP]/CyO*	
M106	*okr[WS] c(2)M[EP]/CyO*	
M107	*okr[WS] cn bw/CyO*	"V1"
M108	*e ca spn-A[1]/ TM6B*	from M100 recombinant, much h
M109	*c(2)M/ CyO;st spnB[BU] e/TM6*	TM3??
M11	*y cv v mei-218[6] f car/C(1)DX/y[+]Y; spa[pol]*	
M110	*y/y[+]Y; c(2)M[8c]/ SM6*	Doc
M111	*y/y[+]Y; c(2)M[1062]/ SM6*	Doc
M112	*c(2)M[EP]/CyO; spn-A e ca/ TM6*	
M113	*c(2)M[EP]/CyO; spnD e? ca?/ TM6*	

Location	Genotype	Comments
M12	*sub[1794], cn bw/CyO; ncd[1]/ TM3, Sb*	
M13	*sc w[bl] mei-218[hfnd]/C(1)DX/y[+]Y*	
M14	*y mei-38 w/y[+]Y/C(1)DX*	
M15	*mei-218[5] f car/FM7*	
M16	*st mus301[D4]/TM3, Sb Ser*	
M17	*okr[Z0682] dp b pr cn/ SM6*	
M18	*w; P{w[+], gc(2)M[myc]}III.3*	from Page, on 3
M19	*cn sub[1794] bw/ CyO; nosGAL4::VP16*	for sub transgenes
M20	*w; P{UASP:c(2)M[HA.C-term]}*	C-terminal fusion of 3 Has
M21	*y mei-41[D18] f/FM7/B[S]Y; spa[pol]*	recombinant from original y cv v
M22	*y/y[+}Y; rec[1] bx[34e]/ MKRS, Sb*	from Sekelsky, no Sb+
M23	*y cv v mei-218[8] f /FM7*	
M24	*y mei-38/ C(1)DX/ y[+]Y; spa[pol]*	
M25	*P{ry[+t7.2]=neoFRT}82B tefu[atm-8] e[1]/TM3,*	
M26	*y/y[+]Y; net Ercc1[X]/CyO*	
M27	*y mei-218[4] f car/C(1)DX/y[+]Y*	
M28	*y w mei-217[g10]/ FM7c/y[+]Y*	
M29	*w[*]; P{ry[+t7.2]=neoFRT}82B tefu[atm-3] e[1]/*	
M30	*y/y[+]Y; mus-312/TM3*	
M31	*y mei-218[4] f car/FM7*	
M32	*r[?] mei-218[7] f car/C(1)DX/y[+]Y*	
M33	*mei-218[4] f /y[+]Y*	
M34	*y/y[+]Y; c(3)G[68] ca/TM3, Sb*	healthy homos
M35	*w mei-41[D3]/FM7*	female sterile allele.
M36	*y cv v mei-218[6] f/ FM7c*	
M37	*mei-218[1]/FM7/y[+]Y*	

Location	Genotype	Comments
M38	*y mei-9[a] f[36]/ FM7/ B[S]Y*	
M39	*y / y[+]Y; sub[1794]/SM6; spa[pol]*	from Z40
M40	*y nod[a]/C(1)DX/y[+]Y*	
M41	*st mus301[D1]/TM3, Sb Ser*	
M42	*FM7a, nod[4] /Dp(1;Y)y[+] / C(1)DX, y f; svs[pa-*	
M43	*y; bw; mei-S282 / TM2; pol*	Hawley M60
M44	*y mei-218[1]/ C(1)DX, y f/ y[+]Y; spa[pol]*	Hawley M8, Canonical mei-218
M45	*Dp(1;Y)B[S]; ru mus301[D3] st e ca/TM3, Sb*	was spn-C[3], lost
M46	*mei-41[D5] f/ C(1)DX/Y*	Hawley
M47	*st mus311[D1] / TM3, Sb Ser*	
M48	*st mus311[D2]/ TM3, Sb Ser*	
M49	*y mei38[J23]/FM7c*	Probably with an inversion
M50	*y mei-9[A2]/ B[S]Y*	
M51	*y mei-218[g4]/FM7; pol*	X-screen
M52	*y hdm[g6]/FM7; pol*	
M53	*y hdm[g7]/C(1)DX/y[+]Y; pol*	
M54	*y hdm[g8]/FM7; pol*	
M55	*y mei-218[g9]/FM7; pol*	X-screen
M56	*al dp c(2)M[3880]/ SM6*	homo lethal
M57	*y mei-218[j1]/FM7; pol*	X-screen
M58	*y mei-218[j2]/FM7; pol*	X-screen
M59	*y mei-9[j3]/FM7; pol*	also known as mei-9[12], meiosis
M60	*c(2)M[EP2115] pr cn/ CyO*	
M61	*y hdm[g7] m f . y[+]/ FM7*	
M62	*+/ B[S]Y; ru[1] st[1] spn-B[1] e[1] ca[1] /TM3,S*	
M63	*y pn cv hdm[g7]/ FM7c*	

Location	Genotype	Comments
M64	*ru[1] th[1] st[1] ri[1] roe[1] p[p] e[s] spn-D[2] /*	=349
M65	*mei-9[a] v /FM7; ry[531]*	used for gene conversion experi
M66	*y w mei-38/ FM7c*	
M67	*spn-B[BC] st sr e/TM3, Sb*	
M68	*spn-B[BU] st sr e/TM3, Sb*	
M69	*spn-D[150] st e ca/TM3, Ser*	from Schupbach, null?
M70	*spn-D[349] st ri roe p[P] e[s] / TM3, Ser*	from Schupbach,
M71	*mei-9[a] hdm[g7]/ FM7c*	
M72	*okra[RU] cn bw/CyO*	outcrossed with new balancer
M73	*y mei-217[g10] /C(1)DX; spa[pol]*	
M74	*okr[WS], cn bw sp/ CyO, l(2)DTS100*	synonym = okr[W31], outcross w
M75	*y mei-218[g2]/FM7/y[+]Y; pol*	X-screen
M76	*y mei-217[R1]/ C(1)DX, y f/ y[+]Y; pol*	
M77	*y mei-217[r1] . y[+]/ FM7c*	
M78	*y pn cv m mei-217[R1] f/ FM7c*	
M79	*y mei-mei-217[g10] . y[+]/ FM7c*	
M80	*y mei-41[D18] f/ FM7; mei-W68/ CyO*	
M81	*mei-9[a] mei-218[1]/ FM7; ry[531]*	received from J. Sekelsky 11/98
M82	*st c(3)G[1] ca / TM3*	outcross of BT606 originally c(
M83	*y mei-g5/y mei-g5/ y[+]Y; spa[pol]*	Originally FM7 stock, but now h
M84	*y w mei-218[KO] /FM7*	targetted KO, deletions at both e
M85	*yw/ y[+]Y; ru h mei-2024 st/ TM3, Sb*	May carry a lethal near ru deriv
M86	*y pn cv m mei-217[g10] / FM7c*	
M87	*y[1] / Dp(1;Y) y[+] ; cn[1] mei-S332[1] / CyO ! l*	Original stock outcrossed to G4.
M88	*w; mer-3[9]/ TM3*	Excision allele
M89	*y mei-53*	isolate 5-3

Location	Genotype	Comments
M90	*y mei-53/ C(1)DX, y f/ y[+]Y*	isolate 5-3
M91	*y hdm[g7]/y[+]Y; spa[pol]*	
M92	*y hdm[g7] mei-218[1]/ FM7*	
M93	*cn sub[1] bw/ CyO; nos:Gal4:VP16*	
M94	*y mei-41[D18]/FM7; spnB[BU], st sr e/ TM3*	
M95	*+/B[S]Y; st[1] ncd[D] /TM3, Sb Ser*	outcross of BT2243
M96	*w; mer-3[35]/ TM3*	excision allele
M97	*y ; ru h st c(3)G[68] cu ca/ TM3*	
M98	*yw/ y[+]Y;c(2)M[EP2115]/ CyO*	From M99, outcross of original
M99	*y[1] Klp3A[mei-352]/y[+]/C(1)DX, y[1] f[1]; sv[s*	originall mei-352
P_1	*P{hsp83:mei-P22[GAL4-HA]}13/FM7,w*	N-terminal GAL4 fusion plus H
P_2	*y/y[+]Y; mei-P22[206] e ca/TM3*	
P_3	*y w; P{w[+]; hsp83::mei-P22[3XHA]}9-1/+; mei-*	triple-HA tagged mei-P22 transg
P_4	*yw/y[+]Y; mei-P22[1]/TM3, Sb Ser*	Isogenized 8/98
P_5	*yw; P{UASP:mei-P22[HA]}/ CyO; mei-P22[103] t*	
P_6	*P{UASP:EB1[HA]}3/TM3*	HA tag EB1
P_7	*y w; P{y[SceI], Pw+, y[Pst]/ SM6*	Pwfl construct with y[sceiI]
P_8	*P{UASP:EB1[HA]}2/TM3*	HA tag EB1
P_9	*y w; P{w[+]; hsp83::mei-P22[3XHA]}4-2; mei-P2*	transgene on 2
P10	*yw; mei-P22[206]/TM3, Sb*	EMS induced allele, LOST
P11	*y w P{w[+]; hsp83::mei-P22[3XHA]}X-1; mei-P2*	X-linked transgene
P12	*yw; mei-P22[N1]/TM3, Sb*	EMS induced allele. Carries a li
P13	*mei-P22[103] st /TM3*	
P14	*mei-P22[206] spn-A[1] e ca/ TM3*	
P15	*w; P{UASP: mei-P22[3190]}58*	C-terminal deletion of mei-P22
P16	*yw; mei-P22/TM3, Sb*	Rapidly becomes P22/P22

Location	Genotype	Comments
P17	*w; P{UASP: mei-P22[2660]}58*	N-terminal deletion of mei-P22
P18	*FM7/y[+]; nosGal4, P{UASP:lacIGFP}*	
P19	*y w/y[+]Y; P{hsp83::mei-P22[3XHA]}9-1/SM6*	transgene in stock without a mei-
P20	*w; P{UASP:lacIGFP}/SM1; Pr Bsb/ TM3, Sb*	
P21	*FM7/y[+]; P{UASP:LacIGFP}/ TM3*	3rd chrom viable
P22	*mei-P22[103] th st P{w+mC=GAL4::VP16-nos.U*	T36 recombinant
P23	*+/B[S]Y; mei-P22[103] th st cu e ca /TM3, ry Sb*	lethal removed from parental chr
P24		
P25	*+/Y; mei-P22[1] ry[531]/TM3, Sb*	P-element insertion mutation
P26		
P27		
P28	*P{UASP:EB1[HA]}1/TM3*	HA tag EB1
P29	*mei-P22[103] e spn-A ca/ TM3*	
P30	*y pn cv mei-P26 /FM7c*	original P insertion allele
P31	*y w mei-P26/FM7c/y[+]Y; spa[pol]*	
P32	*(y) w; P{w+; UAS: mei-P22[3XHA]} / TM3*	wild-type tagged with HA
P33	*yw/ y[+]Y; Bc Elp/ CyO, H{w+, delta2-3}HOP1*	HOP1 = H{w[+], delta 2-3}
P34	*y w/y[+]Y; Dr/ TMS, Sb P{ry[+], delta2-3}99B*	outcrossed from stock center
P35	*(y) w; P{w+; UAS: mei-P22[3XHA]} / SM6*	wild-type tagged with HA
P36	*Sp/ CyO; delta2-3, Sb/ TM6*	2-4C
P37	*(y) w P{hsp83::mei-P22[3XHA]}2/FM7*	
P38	*yw/ y[+]Y; Sb delta 2-3/ TM6*	
P39	*mei-P22[103] st sr spn-B[BU] e / TM3*	
P40	*mei-P22[103] th st c(3)G[68]/ TM3*	
P41	*y w; mei-P22[N1] th st cu e ca/ TM3, Sb*	
P42	*y w; P{hsp83::mei-P22[3XHA]}6/SM6*	

Location	Genotype	Comments
P43	*y[1] w; H{w+, delta2-3}HOP8*	X-linked transposase
P44	*yw;/ y[+]Y; mei-P22[N1]/ TM3*	outcross of original N1 stock to r
P45	*mei-P22[206] , th st cu e ca/ TM3*	
R_1	*T(2;3)77, ry[531]/ TM6B, Tb*	
R_2	*T(2;3)45, ry[531]/ TM6B, Tb*	
R_3		
R_4	*T(2;3)34, ry[531]/ TM6B, Tb ry*	
R_5	*T(2;3)F, ry[531]/ TM6B, Tb*	
R_6	*T(1;3)Z, ry[531]/ TM6B, Tb*	
R_7	*T(2;3)49, ry[531]/ TM6B, Tb Bsb*	
R_8	*T(2;3)92, ry[531]/ TM6B, Tb*	
R_9	*T(2;3)77, ry[531]/ MKRS, Sb*	
R10	*T(1;3)X, ry[531]/ MKRS, Sb*	
T_1	*y w/y[+]Y; P{w+mC=GAL4::VP16-nos.UTR}MV*	Express HA tagged C(2)M with
T_2	*y w; TM6B, P{w[+], cre}/ MKRS, P{ry[+], hsFLP*	= crew3
T_3	*y w; Gla/SM6, mei-I*	jumped from orginal T6 on norm
T_4	*w P{UASP:AurB[HA]}37-2/FM7*	
T_5	*P{COG-GAL4::VP16-nos.UTR}iso1-4*	germline driver on X
T_6	*w; P{w[+}, mei-I}/ CyO*	homozygous viable
T_7	*w mei-218 f car; P{w[+}, hsp83-FLAG-mei-218}7*	
T_8	*w mei-218 f car; P{w[+}, hsp83-FLAG-mei-218}3*	
T_9	*w; P{w[+}, polo-GFP}*	from Sunkel, original GFP
T10	*p{UASP:CBTPMGFP}35*	non-degradable CycB-GFP, U69
T11	*w; P[w+, UASp-GFPS65C-al-tub84B]#14-6/TM3,*	GFP Tub, NG10, possibly
T12	*w; P{UASP;PAGFP-a-Tub84B}F11.1*	GFP PA - Tub, on 3
T13	*w; P{w[+mc] GAL4-Hsp70,PB}89-2-1*	hsp70-GAL4 on 3

Location	Genotype	Comments
T14	*w; P{UAS:mei-217[3XHA]}3-2*	on 2
T15	*w; P[w+, UASp-GFPS65C-al-tub84B]#14-6/SM6a*	NG9
T16	*w; P{UAS:mei-217[3XHA]}7-3*	on 3
T17	*yw; P{UASP-sub[Pst]}7/TM3*	
T18	*w[1118]; P{70FLP}10*	efficient excision of tandem FLP
T19	*ncd[GFP] 4121/TM3*	M3M1; M9F1
T20	*armGAL4 (2)*	
T21	*armGAL4 (2X3)*	
T22	*P{UASP:mei-38[HA]}RZC-8/TM3*	
T23	*w; P{UASP:AurB[HA]}37-5/TM3*	on 2, original endsout, has only o
T24	*w; P{w+, UASp-sub[ATP-HA]}4/TM3, Sb*	ATP mutant
T25	*w; P{UASP:AurB[HA]}18-3/TM3*	
T26	*P{UASP:mei-38[HA]}RZC-8 P{UASP:nosGAL4:V*	
T27	*y w; P{ry+t7.2=70FLP}23 P{v+t1.8=70I-SceI}4A/*	
T28	*P{ry+t7.2=hsFLP}12, y[1] w; noc[Sco]/CyO ! FL*	
T29	*P{UASP:mei-38[HA]}RZD-10/SM6*	
T30	*y w v(?); P[ry+, 70FLP]4, P[v+, 70I-SceI]2B, Sco*	For gene targetting
T31	*P{UASP:mei-38[HA]}RZA-5L/TM3*	
T32	*cn sub[1] bw/ CyO; P[w+, UASp-GFPS65C-al-tub*	Using GFP-Tub from T11
T33	*P{w+mC=GAL4::VP16-nos.UTR}MVD1 cu c(3)G*	Strong ovary GAL4 driver
T34	*w P[w+, UASp-sub[RFP]X*	FRP tag on X
T35	*P{w[+m*]=Ubi-tacc.GFP}1, y[1] w[*]*	GFP - TACC
T36	*w[1118]; P{w+mC=GAL4::VP16-nos.UTR}MVD1*	high expression in germline. ins
T37	*P[w+, UASp-GFPS65C-al-tub84B]#14-6; P{w+;*	GFP-Tub
T38	*b c(2)M[910], P{hsp83::mei-P22[3XHA]}9/SM6*	
T39	*w; P{UASp,c(2)M[3XHA]}65L/TM3, Sb*	on 3, lethal

Location	Genotype	Comments
T40	*w; P{UASp,c(2)M[3XHA]}28*	on 3
T41	*w; P{w+mC=GAL4-nos.NGT}40*	lower level than nos::VP16, on 2
T42	*w P{w+, UASp-sub[ATP-HA]}1/FM7, w*	ATP mutant
T43	*w; P{w+, UASp-sub[ATP-HA]}2/SM6*	ATP mutant
T44	*y w/ Y; mei-I mei-W68[Z4472] bw/ SM6*	
T45	*w; P[w+, UASp-sub[L6]10/TM3, Sb*	lack of motor domain insertion
T46	*P{ry+t7.2=hsFLP}1, y [w1118]; Dr[Mio]/TM3, ry*	
T47	*y [w1118] P{ry+t7.2=70FLP}3F/Dp(1;Y)y[+]; T*	
T48	*y [w1118] P{ry+t7.2=70FLP}3F/Dp(1;Y)y[+]; no*	
T49	*GFP;ord*	fusion from Bickel, T076
T50	*w; P{UASP:AurB[HA]}18-5/SM6*	
T51	*w; P[w+, UASp-sub[RFP]28/TM3, Sb*	
T52	*P{UASP:mei-38[HA]}REA-1/SM6*	
T53	*w; P[w+, UASp-sub[RFP]1/TM3, Sb*	
T54		
T55		
T56		
T57	*y[1] w[*]; P{w[+mC]=tubP-GAL4}LL7/TM3, Sb[*	ubiquitous GAL4 expression
T58	*w[1118]; P{w[+mC]=GAL4::VP16-nos.UTR}MV*	GFP tubulin with driver, differen
T59	*w[*]; P{w[+mC]=matalpha4-GAL-VP16}V37*	alphaTub67C promoter for GAL
T60	*w[*]; P{w[+mC]=matalpha4-GAL-VP16}V2H*	GAL4-VP16 fusion protein expre
T61	*w; P[w+, UASp-sub[GFP]2/TM3, Sb*	full length sub with amino-GFP,
T62	*w; P[w+, UASp-sub[GFP]35/TM3, Sb*	full length sub with amino-GFP,
T63	*w; P[w+, UASp-sub[GFP]89/SM6*	full length sub with amino-GFP
T64	*w; P[w+, UASp-sub[3XHA]45/TM3, Sb*	full length sub with amino-HA, l
T65	*w; P[w+, UASp-sub[3XHA]21/TM3, Sb*	full length sub with amino-HA

Location	Genotype	Comments
T66	*w; P[w+, UASp-sub[3XHA]31/TM3, Sb*	full length sub with amino-HA, l
T67	*w P[w+, UASp-sub[3XHA]70*	full length sub with amino-HA
T68	*w; P[w+, UASp-sub[S500A-GFP]43/TM3, Sb*	S-A change at amino acid 500 wi
T69	*w; P[w+, UASp-sub[S500A-GFP]5/TM3, Sb*	S-A change at amino acid 500 wi
T70	*w; P[w+, UASp-sub[S500A-GFP]48/TM3, Sb*	S-A change at amino acid 500 wi
T71	*w; P[w+, UASp-sub[1801-GFP]58/TM3, Sb*	amino-term deletion with amino-
T72	*w; P{UASP:AurB[myc]}87/SM6*	
T73	*w P[w+, UASp-sub[1801-GFP]31*	amino-term deletion with amino-
T74	*w; P[w+, UASp-sub[3001-GFP]1/TM3, Sb*	C-terminal domain only with ami
T75	*w; P[w+, UASp-sub[3001-GFP]64/TM3, Sb*	C-terminal domain only with ami
T76	*w; P[w+, UASp-sub[3001-GFP]43/TM3, Sb*	C-terminal domain only with ami
T77	*w; P[w+, UASp-sub[SSAA-GFP]59/TM3, Sb*	Two serines changed near amino
T78	*w; P[w+, UASp-sub[SSAA-GFP]47/TM3, Sb*	Two serines changed near amino
T79	*w P[w+, UASp-sub[SSAA-GFP]74*	Two serines changed near amino
T80	*y w, P[w+, UASp-sub[Pst-GFP]/FM7,w*	N-terminal fragment
T81	*w; P[w+, UASp-sub[RFP]7/SM6*	
T82	*w; P[w+, UASp-sub[L6]13L/TM3, Sb*	lack of motor domain insertion
T84	*w; P[w+, UASp-sub[L6]sun/FM7c*	lack of motor domain insertion, "
T85	*w; P[w+, UASp-sub[L6]shrek/SM6*	lack of motor domain insertion
W_1	*y/y[+]Y; Df(2R)W68-2/SM6a*	originally mei-W68[w-2]. From
W_2	*yw/y[+]Y; Df(2R)LL5/ CyO*	W46 lost Cy, outcrossed to W50
W_3	*y; y[Sce1-1]*	C31 insertion
W_4	*y w; mei-W68[k5603]/ CyO*	P{lacW} inserted in 5' UTR□ori
W_5	*y/y[+]Y; mei-W68[1]/CyO*	
W_6	*al dp b pr cn mei-W68[1] /SM6; y/y[+]Y*	
W_7	*y/B[S]Y; mei-W68[2-0949], cn bw /SM6*	Zuker EMS mutant

Location	Genotype	Comments
W_8	*y/B[S]Y; mei-W68[1] px mr bw sp/SM6a*	
W_9	*y; y[Sce1-2]*	C31 insertion
W10	*B[S]Y/+; mei-W68[L1]/SM6*	
W11	*y w/y[+]Y; dp b pr cn Df(2R)LL4/ SM6*	Homozygous viable, best when c
W12	*cn mei-W68[L1]/SM6*	
W13	*al dp b Sp pr cn mei-W68[L1]/SM6*	
W14	*y; mei-W68[w3]/ SM6*	Lethal excision of EP0899. Also
W15	*y/ B[S]Y; mei-W68[2-5736], cn bw /SM6*	Zuker allele, outcrossed with ne
W16	*okr[WS] cn mei-W68[L1] / CyO*	
W17	*y[1] w[1118]; P{w[+mC]=UAS-y.C}MC1*	wild-type UAS yellow
W18	*P{UASp:mei-W68[3XHA]} mei-W68/SM6*	all Cy+????
W19	*mei-I mei-W68/SM6*	recombinant mei-I and mei-W68
W20	*w mei-41[D3]/FM7; mei-W68/ CyO*	
W21	*mei-W68[L1]/CyO; ry[531]*	
W22	*y w; cn mei-W68[k5603]/ SM6*	from 5603/ cn bw
W23		
W24	*y/y[+]Y; al mei-W68/SM6*	
W25	*cn mei-W68[4572] bw/ CyO; spn-B[BU]/ TM6*	
W26	*w/; P{w[+]; pup:mei-W68[GAL4]}32/ TM3*	mei-W68 GAL4 fusion, with c(2)
W27	*P{w[+]; UAS??} f/ FM7; cn mei-W68 bw/CyO*	
W28	*P{UASp:mei-W68[3XHA]}-2 mei-W68/SM6*	all Cy+?
W29	*+/Y; mei-W68[2-5736]/CyO*	
W30	*al dp b pr cn mei-W68[2-4572]/ SM6*	Zuker allele, recombinant chrom
W31	*y; P{w[+]; pup:mei-W68[GAL4]}7L-9b mei-W68*	mei-W68 GAL4 fusion, with c(2)
W32	*w/+; P{w[+]; pup:mei-W68[GAL4]}21D mei-W6*	mei-W68 GAL4 fusion, with c(2)
W33	*y/y[+]Y; mei-W68/CyO; ry[531]*	

Location	Genotype	Comments
W34	*al dp b pr cn mei-W68[2-1049] bw/ CyO*	
W35	*y w/Y; Df(2R)LL99/SM6*	Excision from EP(2)0899. Hom
W36		
W37		
W38		
W39		
W40	*P{UASp:mei-W68[3XHA]}-3 mei-W68/SM6*	all Cy+????
W41		
W42		
W43		
W44		
W45	*y w/y[+]Y; Df(2R)[LL4], P{lacW}/SM6a*	Derived by mobilization of P{lac
W46		
W47	*al dp b pr cn mei-W68[2-1049]/ SM6*	Zuker allele, recombinant chrom
W48	*w ; l(2)k06821 / CyO*	
W49	*w ; EP(2)1144/CyO*	outcrossed
W50	*y w / y[+]Y ; W68[1] / CyO*	
W51	*w ; EP(2)0899*	
W52		
W53	*w; P{w[+]ovo[D4]:mei-W68[mutY-F]}H*	#3
W54	*w; P{w[+]ovo[D4]:mei-W68[WT]}13A*	#5
W55	*w; P{w[+]ovo[D4]:mei-W68[mutY-F]}K*	#6
W56	*w; P{w[+]ovo[D4]:mei-W68[WT]}B1*	#8
W57	*P{hsp83::mei-P22[3XHA]9 mei-W68[4572] bw/S*	few homozygotes
W58	*w; P{w[+]ovo[D4]:mei-W68[YY-FF]}EV*	#12
W59	*w; P{w[+]ovo[D4]:mei-W68[WT]}ST/ C(1)DX, y*	#16

Location	Genotype	Comments
W60	*w; P{w[+]ovo[D4]:mei-W68[WT]}DD*	#17
W61	*okr[WS] cn mei-W68[1]/SM6*	
W62	*y w/y[+]Y; P{w[+]; pup:mei-W68[GAL4]}7D4 m*	mei-W68 GAL4 fusion, with c(2)
W63	*w; P{w[+]ovo[D4]:mei-W68[WT]}A1*	#20
W64	*y/y[+]Y; P{w[+]; pup:mei-W68[GAL4]}21L mei-*	mei-W68 GAL4 fusion, with c(2)
W65		
W66		
W67	*w; P{w[+}ovo[D4]:mei-W68[mutY-F]}HH ! leth*	#24
Z01	*2-4143 cn bw / CyO*	
Z02	*okr[Z0682] b pr cn c px sp / SM6*	
Z03	*al dp b pr cn sub[1794] bw / SM6*	
Z04	*y; c(2)M[3880] pr c px sp / SM6*	
Z05	*(y); cn dwd[4134] sp/ SM6*	
Z06	*y/y[+]Y; c(2)M[3880] cn bw / SM6*	
Z07	*c(2)M[910] cn c px sp / SM6*	
Z08	*2-6006 cn bw / CyO*	
Z09	*al dp b c(2)M[910] cn bw / SM6*	
Z10	*al dp b cn 2-5106 bw / SM6*	
Z11	*al dp cn bw 1789 / SM6*	
Z12	*al dp 6066 cn bw / SM6*	
Z13	*y/B{S}Y; cn 2-5106 px sp / SM6*	
Z14		
Z15	*mei-W68[2-4472] cn bw / CyO*	Allele of meiW68
Z16	*cn 2-6106 px sp / SM6*	
Z17	*mei-W68[2-0949] cn bw / CyO*	
Z18	*y/B[S]Y; ord[2-3397] cn bw / SM6*	

Location	Genotype	Comments
Z19	*y/y[+]Y; 5105 px sp/ SM6*	
Z20	*2-0824 cn bw / CyO*	
Z21	*2-1655 cn bw / CyO*	
Z22	*2-1689 cn bw / CyO*	
Z23	*y/B[S]Y; 2-6066 cn bw / SM6*	
Z24	*c(2)M[2-3780] cn bw / CyO*	
Z25	*2-2029 cn bw / CyO*	
Z26		
Z27	*2-3397 cn bw / CyO*	
Z28	*2-4134 cn bw / CyO*	
Z29	*y/B[S]Y; 2-2029 cn bw / SM6*	outcross of original stock
Z30	*2-5006 cn bw / CyO*	Zuker vial
Z31	*2-5315 cn bw / CyO*	
Z32	*2-6066 cn bw / CyO*	
Z33	*2-2525 cn bw / CyO*	
Z34	*2-5955 cn bw / CyO*	
Z35	*2-3090 cn bw / CyO*	
Z36	*2-1694 cn bw / CyO*	
Z37	*y / y[+]Y ; mei-W68[2-4472] cn bw / CyO*	
Z38	*y / B[S]Y ; 2-5006 cn bw / SM6a*	
Z39	*y / B[S]Y ; 2-4134 cn bw / SM6a*	
Z40	*y / B[S]Y ; sub[1794] cn bw / SM6a*	Zuker 2-1694
Z41	*y / B[S]Y ; 2-4143 cn bw / SM6a*	
Z42	*y / y[+]Y ; 2-6006 cn bw / CyO*	
Z43	*y / B[S]Y ; c(2)M[2-0810] cn bw / SM6a*	Zuker 2-810, outcross
Z44	*y / B[S]Y ; 2-1655 cn bw / SM6a*	

Location	Genotype	Comments
Z45	*y / y[+] Y ; 2-5955 cn bw / CyO*	
Z46	*y / B[S] Y ; okr[Z0682] cn bw / SM6a*	outcrossed
Z47	*(y); dwd[4134] c bw/ SM6*	
Z48	*y / B[S] Y ; 2-1689 cn bw / SM6a*	
Z49	*yw / B[S] Y ; cn sub[1794] px sp / SM6a*	
Z50	*y ?/ B[S] Y ; 2-1655 c px sp/ SM6a !!low or no N*	
Z51	*y? / B[S] Y ; al dp b 2-5955 cn bw / SM6a*	
Z52	*y? / B[S] Y ; 2-5955 cn px sp / SM6a*	
Z53	*+ / B[S] Y ; al dp cn 2-4134 bw / SM6a*	
Z54	*y? / B[S] Y ; cn 2-4134 sp / SM6a*	