Table S1. Insertion position and loss- or gain-of-function analysis of targeted genes

Gene	EP line	Position into targeted gene	Loss-or gain-of function analysis towards GG17
A. Ras, Rac and Rho signalling proteins			
MESR4	EP(2)386	-575 bp from mRNA start	Behaves like UAS-MESR4 (Huang and Rubin, 2000).
AKAP200 */MESR2*	EP(2)2072 EP(2)2254	-43 bp from mRNA start -114 bp from mRNA start	Behaves like UAS-MESR2 (Huang and Rubin, 2000).
DC3G	EP(1)1613*	In first intron, -475 bp from AUG start codon	UAS-DC3G lines (Ishimaru, 1999) similarly enhanced GG17 (see text).
CG31012	EP(3)3700	-14 bp from mRNA start	*P{SUP-or-P}CG31012P[KG05741]* has no effect
pbl	EP(3)3415*	5'UTR, -457bp from AUG start codon	Described as gof in (Prokopenko et al., 1999). *pbl[5]* has no effect (see text).
GEF(64C)	EP(3)3035 EP(3)3322*	In first untranslated exon (+26bp) -469 from mRNA start	UAS-GEF (64C) line (Bashaw, 2001) enhanced GG17. *GEF* mutants have no effect (see text).
grp	EP(2)587	In first intron of transcript CG17161-RA, - 16 kb from AUG start codon	Described as gof (Abdelilah-Seyfried et al., 2000). *grp[06034]* has no effect.
CG7097	EP(2)2445	-44 bp from mRNA start	*Df(2R)P34* (55E-56C) has no effect.
SNF4Aγ	EP(3)648	-40 bp from mRNA start of *SNF4Aγ* transcript CG17299-RH	*P{SUPor-P}SNF4Aγ[P{KG00325}]* has no effect.
B. Cytoskeletal control and membrane trafficking			
Myo31DF	EP(2)2491	-28 bp from mRNA start	*P{lacZ}l(2)k09116* has no effect
Beach1	EP(2)2299*	In first intron, + 194 bp downstream of AUG start codon: may generate a Nter truncated protein.	Described as gof in (Abdelilah-et al, 2000; Kraut et al., 2001). *Df(2L)cl7* (25E-26A) has no effect.
C. Chromatin remodelling factors			
kis	EP(2)474	In first intron of *kis* CG3696-RA transcript, - 5 kb from CTG start codon	Described as gof in (Kraut et al., 2001; Pena-Rangel et al., 2002). *kis[1]* has no effect.
	EP(2)563	- 1kb from start of *kis* CG3696-RB transcript	
dom	EP(2)2371	-23 bp from mRNA start	*dom[P{K08108}]* has no effect.
Dp1	EP(2)2422	In first intron, -1152 kb from AUG start codon	*P{GT1}Dp1[BG01405b]* and *P{GT1}Dp1[BG02288]* have no effect.
D. Spermatogenesis			
poe	EP(2)349 EP(2)737	-18 bp from mRNA start -18 bp from mRNA start	EP737 described as gof in (Kraut et al., 2001). *poe[O3420]* has no effect.
TMS1d	EP(3)807	-24 bp from mRNA start	*Df(3L)st-e4* (72D-73A) enhanced GG17 (see text).
cdi	EP(3)3319	In first non coding exon	Described as gof in (Kraut et al., 2001) *cdi[r47]* has no effect.
E. Others			
Traf2	EP(1)325	- 257 bp from mRNA start	*Df(1)GE202* (7D12-7E3) has no effect
wun2	EP(2)652	-4,6kb from mRNA start may affect *wun* UAS+/-	May affect *wunen* . *wun[EMS4]* and *P{lacW}wun[K10201]* have no effect.
CG14959	EP(2)714 EP(2)3139	Both EPs inserted -7,6 kb from mRNA start	Described as gof (Tseng and Hariharan, 2002; Pena-Rangel et al., 2002). *P{SUP-orP}KG0365* has no effect
CG5261	EP(2)816	-3kb from mRNA start	Described as gof (Abdelilah-Seyfried et al., 2000; Pena-Rangel et al., 2002).
		May affect *chameau (chm)* (3' to *chm*)	*P{GT1}chm[BG02254]* has no effect
CG3624	EP(2)827	- 40 bp from mRNA start	*P{SUPor-P}CG3624[KG05061]* has no effect
rho-6	EP(2)2023	-958 bp from mRNA start	*P{SUPor-P}rho-6[KG09603]* has no effect
CG6701	EP(2)2054	5'UTR, -251 bp from AUG	Described as gof (Tseng and Hariharan, 2002; Pena-Rangel et al., 2002). *P{SUPor-P}CG6701[KG00917]* has no effect
CG8740	EP(2)2233	-36 pb from mRNA (GH05582) start	/
numb	EP(2)2455	- 520 bp from *numb* transcript CG3779-RA	*numb[1]* has no effect
cpo	EP(3)3395 EP(3)3608	Both in second intron of *cpo* transcript, -2,2 kb from CUC start codon	*cpo[01432]* has no effect
CG2017	EP(3)3503	5' UTR, - 40 bp from AUG start codon	Described as gof (Kraut et al 2001). *Df(3R)Tpl10,DpDfd[rv1]* has no effect

Note that Gal4-dependent regulation can be functional at considerable distance of up to10 kb (Nicolai et al., 2003).

References
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