

Supplementary Figure Legends

Fig. S1. Alignment of *dPLZF*, *PLZF* and *EOR-1*.

(a) The BTB/POZ domain of *dPLZF* is similar to the domain of human *PLZF*, *Drosophila kelch* and *C. elegans EOR-1*. Conserved regions are boxed. Shaded areas indicate identical residues. (b) Alignment of zinc finger motifs regions of *dPLZF*, *PLZF* and *EOR-1*. The C₂H₂ zinc-finger motifs are underlined. Cysteines and histidines are indicated with dots.

Fig. S2. High level of *dPLZF* mRNA expression can be detected in embryo, larva and pupa stages but low in adult stage.

(a) RT-PCR analysis of *dPLZF* expression in wild-type flies showed a single band of 1 kb transcript. Total RNA from various developmental stages of *Drosophila* was reverse transcribed, and PCR amplification was performed for 35 cycles with specific primers for *dPLZF* or ribosomal protein (*rp49*). *rp49* was used as a loading control. (b) Real-time PCR analysis of *dPLZF* expression in various developmental stages. Adult was used as a standard of relative expression.

Fig. S3. Genetic phenotypes by double expressed *dPLZF* induced the extra wing veins under *en-Gal4* or *MS1096-Gal4*.

(a) Double expression of *dPLZF* in *en-Gal4* flies produced several extra wing veins. (b) Double expression of *dPLZF* in *MS1096-Gal4* flies produced the strong extra wing veins. We checked individually more than 200 *Drosophila* wings. Flies were grown at 25 °C.

Fig. S4. Genetic phenotypes by Draf/ERK reduced the interaction with dPLZF RNAi or dPLZF haplodeficiency.

(a) Wing of wild type fly (b) Overexpression of dPLZF in MS1096-Gal4 flies produced several extra wing veins. (c) dPLZF overexpression phenotype was mostly suppressed by dPLZF RNAi. (d) Quantification of flies with extra wing vein phenotype for indicated genotypes (from (a) to (c)). We checked more than 200 *Drosophila* wings for each genotype. (e) Wing of Draf overexpression fly using *vg*-Gal4. (f, g) Draf activity was completely suppressed by *dPLZF* haplodeficiency. (h) Wing of ERK^s overexpression fly using *en*-Gal4. (i, j) ERK^s activity was completely suppressed by *dPLZF* haplodeficiency. (k) Quantification of flies with extra wing vein phenotype for indicated genotypes (from (e) to (j)). We checked more than 200 *Drosophila* wings for each genotype. Flies were grown at 25°C.

Fig. S1.**a**

dPLZF	28	QRRTGQFCDLLLELDSDDSLSSVHFCVLAQSSQFINSNQKQQQFSIH-----NP-
PLZF	27	MRLAGTLCDDVIMVDS---QEFHAHRTVLACTSKMFEILFH-----R----N-S
EOR-1	35	QRKTGRFCDFEIVVQN---KSFAAHRNIIAAHSPYFDAIFK-----YCKIT--K
kelch	180	MRKQKQLCDVILVADD---VEIHAHRMVLASCSPPYFYAMFT-----SFEESR-Q
dPLZF	78	LKITYIRNFSCCTQCLHTIVDFYEDLVSVSKEHELHFRELAQILAVTEL
PLZF	68	QHYTL--DFLSPKTFQQILEYATATLQAKAEDLDDLLYAAEILEIEYL
EOR-1	79	EQLTI--NSKSPQVFELFLNYMYSQTVIIDRSSVELLRFANNFLIVKL
kelch	225	ARITL--QSVDARALELLIDYVYATATVEVNEDNVQVLLTAAANLQLTDV

b

dPLZF	212	LIQSLCEVGFLDWREYDSSLRRHSGDLRK-----
PLZF	404	EQQSVCGVELPDNEAVEQHRKLSG---MKT-YGCELCGKRFLDSLRLRMHLLAHSAGAKAFVCD
EOR-1	424	STCIFCGLRTANEELLEKHKARHN---RNTYYMCHLCEFETNWSKQFYLHCAEHWTEIPIYRCE
dPLZF	241	-----PFFCQQGGIRFNTRAAALLVHQPKHST--ETPHICPHCGKGFK
PLZF	465	QCGAQFSKEDALETHRQHTGTDMAVFCLLCGKRFAQQSALQQMEVHAG--VRSYICSEONRTFP
EOR-1	485	TCPFTSNEIQEFLTHRLQHTDER--FFKQGEQAWKGRTRSQIFAHERMISVLDDRSLHCEECGRGEQ
dPLZF	281	WKQGLSNHILVHNPEKQMLCDVCGYSTTHMKALKSHKLLHTGEF--FACTVSGCKHRANRKENLKLH
PLZF	529	SHTALKRHLRSHTGDHPYECFCGSCFRDESTLKSHKRIHTGEKPYEC--NGCGKKFSLKHQLETH
EOR-1	550	QHSTLDHHVASHNPPRYICEDCGFATKTADHLSLHRRQHTGDN--FSCHIAAGCDYSSTKKSQLAAH
dPLZF	346	IETHKQGRDFICEVCGCKFSQSKNLKRHALKHTENGNRYKQQLCGFSSHRSDKMKEHVQRVH
PLZF	593	YRVHTGEKPFECKLCHQRSRDYSAMIKHLRTHNGASP--YQCTICTEYCPSLSSMQKHM-KGH
EOR-1	615	LRTHMAVRAHLCKICGRGFIEKSHLVRHERIHLLEEK--FKCQCEYASSRRDKLKEHIIKHH

Fig. S2.

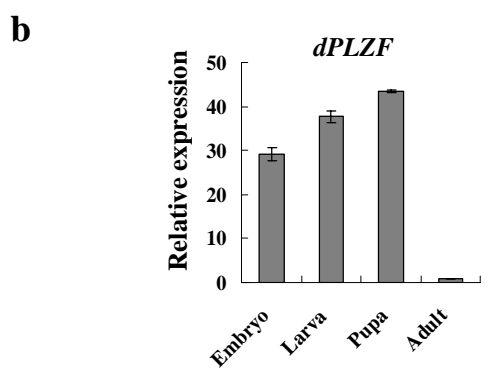
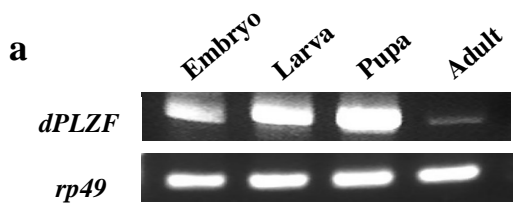
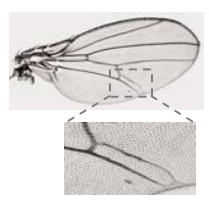


Fig. S3.

a *en-Gal4>2X dPLZF*



b *MS1096-Gal4>2X dPLZF*

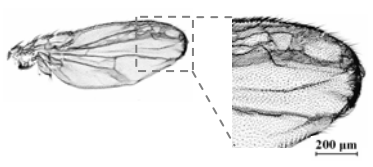


Fig. S4.

