Figure legends

Figure S1. *wg* expression patterns in imaginal discs are conserved between two *Drosophila* species. *In situ* hybridizations with third instar larval imaginal discs. (*A*) *D. melanogaster* wing disc. (*B*) *D. melanogaster* eye-antennal disc. (*C) D. melanogaster* leg disc. (*D*) *D. guttifera* wing disc. (*E*) *D. guttifera* eye-antennal disc. (*F*) *D. guttifera* leg disc. All discs are oriented with anterior to the left and dorsal on top.

Figure S2. *wg* expression in longitudinal vein tips and campaniform sensilla of the pupal wing is unique to *D. guttifera*. *In situ* hybridization for the *wg* gene is shown in pupal wings of various species. (*A*) *Drosophila melanogaster*. (*B*) *D. guttifera*. (*C*) *D. deflecta*. (*D*) *D. nigromaculata*. (*E*) *D. palustris*. (*F*) *D. quinaria*. *D. melanogaster* belongs to *melanogaster* species-group of the subgenus *Sophophora*, while the other species belong to the *quinaria* species group of the subgenus *Drosophila*. All probes are species-specific.

Figure S3. Map of the Wnt region of *D. melanogaster* and *D. guttifera*. Vertical bars connected with black lines indicate sequences of 40bp or longer with 100% sequence conservation between species. Red lines indicate conserved but inverted sequences. Numbered horizontal solid bars indicate DNA fragments tested by transgenic reporter assays. *D. melanogaster* fragments were tested in transgenic *D. melanogaster* using phiC31 integration. *D guttifera* fragments were tested in transgenic *D. guttifera* using the *piggyBac* transposon.

Figure S4. The novel activity of the *D. guttifera* gutCV-T enhancer arose within a conserved enhancer. Sequence alignment of the gutCV-T enhancer and the orthologous defCV enhancer (Serial Cloner v2.6.1, local alignment, and word size=15 were used). Mismatches are marked with a #. Red bars indicate identical sequence matches >10bp with the melCV-core enhancer from *D. melanogaster*. The extent of collinear sequence conservation (with no significant rearrangements) between the two species indicates that a small change or an accumulation of small changes is responsible for the new enhancer activity in the vein tips.

Figure S5. Expression patterns of four *Wnt* genes in pupal wings of *D. guttifera* visualised by *in situ* hybridizations. (*A*) *Wnt4*. (*B*) *wingless*. (*C*) *Wnt6*. (*D*) *Wnt10*.

Figure S6. *wingless* is the predominant *Wnt* gene expressed in the pupal thorax. *Wnt* genes expressed in the pupal thorax (top) and embryo (middle) were detected by RT-PCR and reaction products profiled by gel electrophoresis. Control reactions from genomic DNA are shown at the bottom

Figure S7. *Cis-* and *trans*-regulatory changes are responsible for the novel campaniform sensillum expression of *wg* in *D. guttifera*. (*A*) In the *D. melanogaster* pupal wing, the melCS enhancer shows no restricted expression (EGFP, green). (*B*) In the *D. guttifera* pupal wing, the melCS enhancer shows no expression (DsRed, magenta). (*C*) *D. melanogaster* pupal wing, the gutCS enhancer shows no restricted expression (EGFP, green). (*E*): *D. guttifera* pupal wing, the gutCS enhancer drives reporter expression in the campaniform sensillum (DsRed, magenta).

Figure S8. *Cis-* and *trans*-regulatory changes are responsible for the striped expression of *wg* in the pupal thorax of *D. guttifera*. (*A*) *D. melanogaster* pupal thorax, the melTS fragment shows no stripe expression (EGFP, green). (*B*) *D. guttifera* pupal thorax, the melTS fragment shows no stripe expression (DsRed, Magenta). (*C*) *D. melanogaster* pupal thorax, the guts enhancer is expressed in incomplete stripes (EGFP, green). (*D*) *D. guttifera* pupal thorax, the gutTS enhancer drives full stripe expression which corresponds to the adult pigmentation pattern (DsRed, magenta).

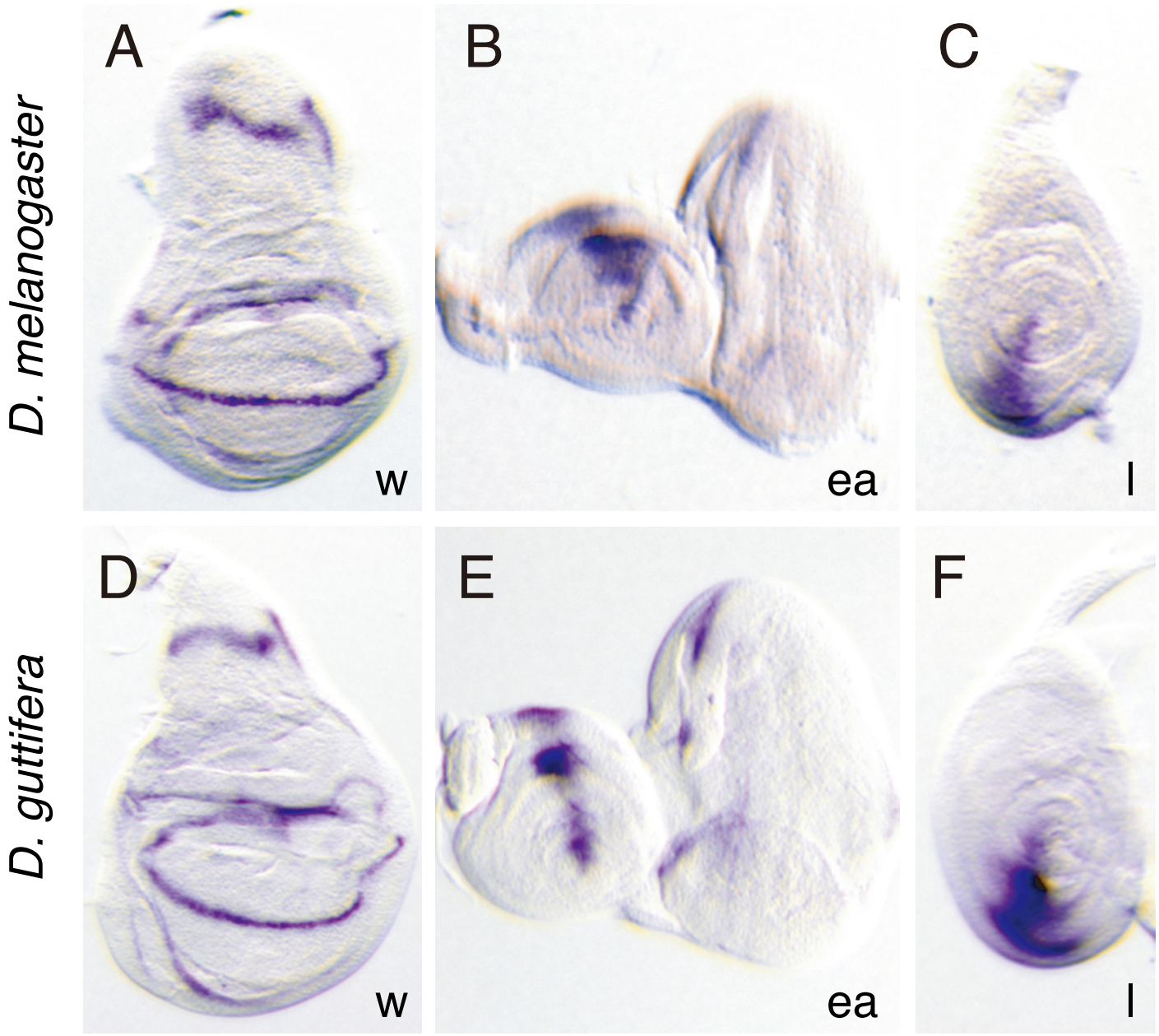


Figure. S1

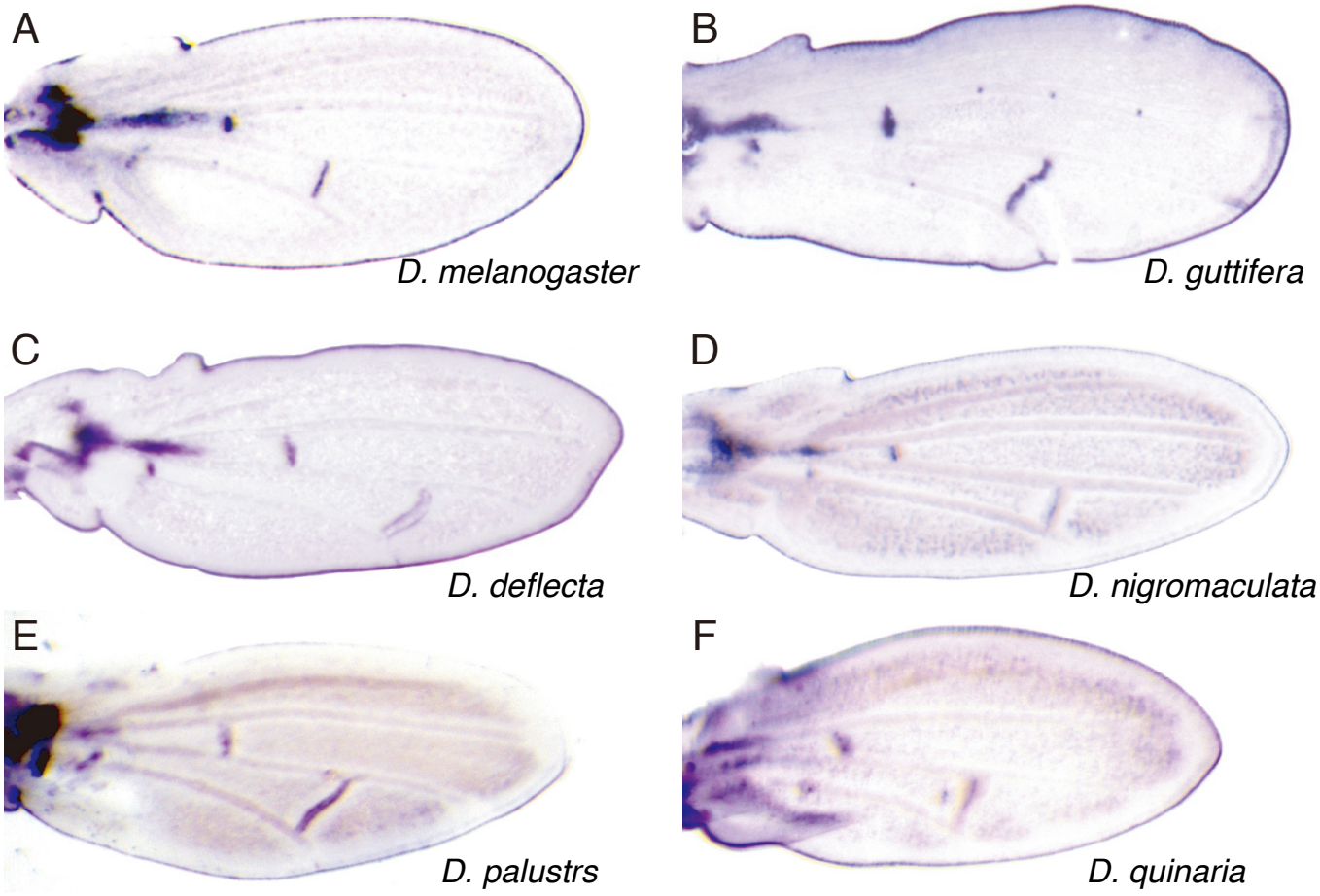

A
D. melanogaster
B
D. guttifera
C
D. deflecta
D
D. nigromaculata
E
D. palustrs
F
D. quinaria

Figure. S2

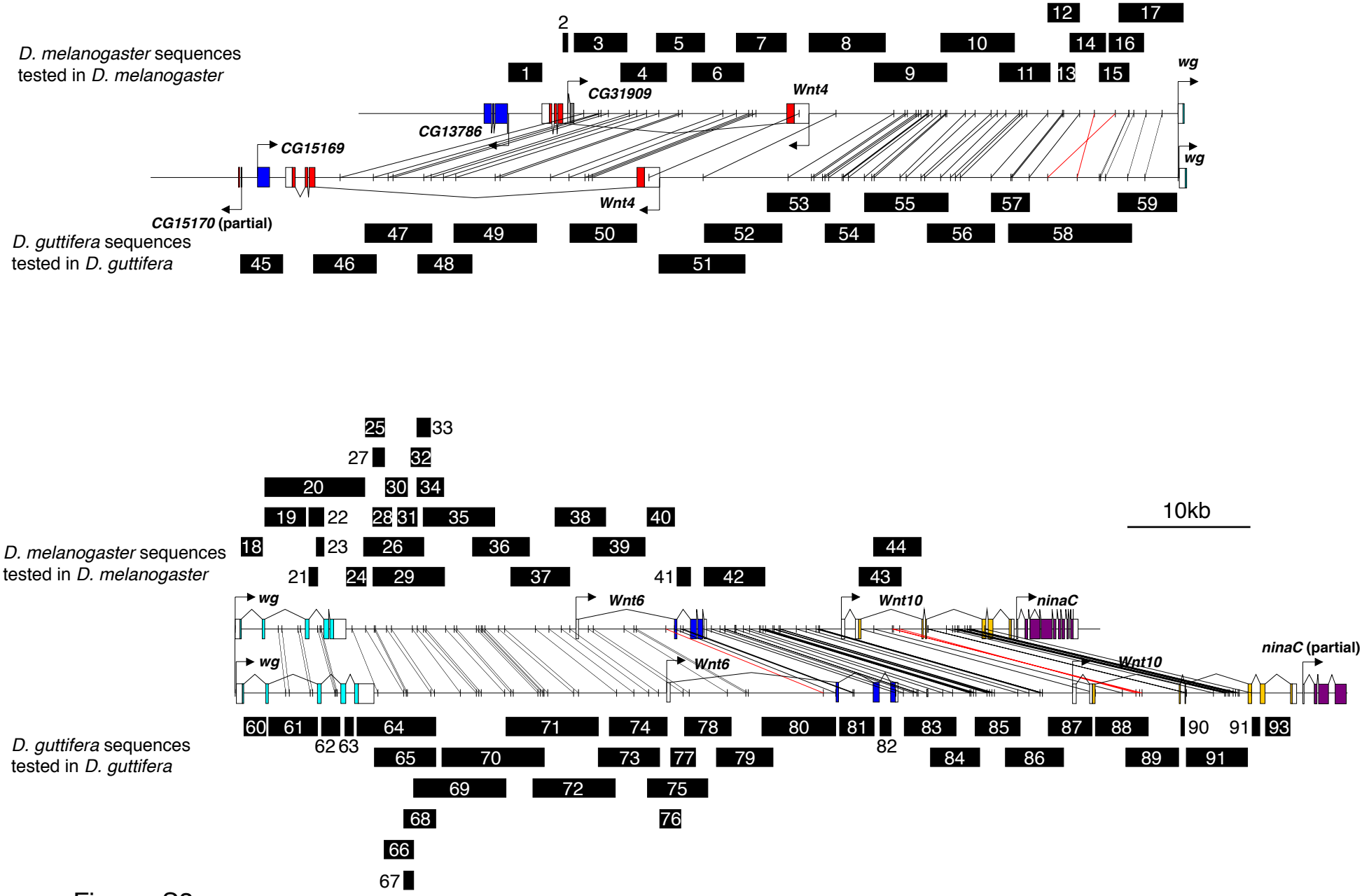
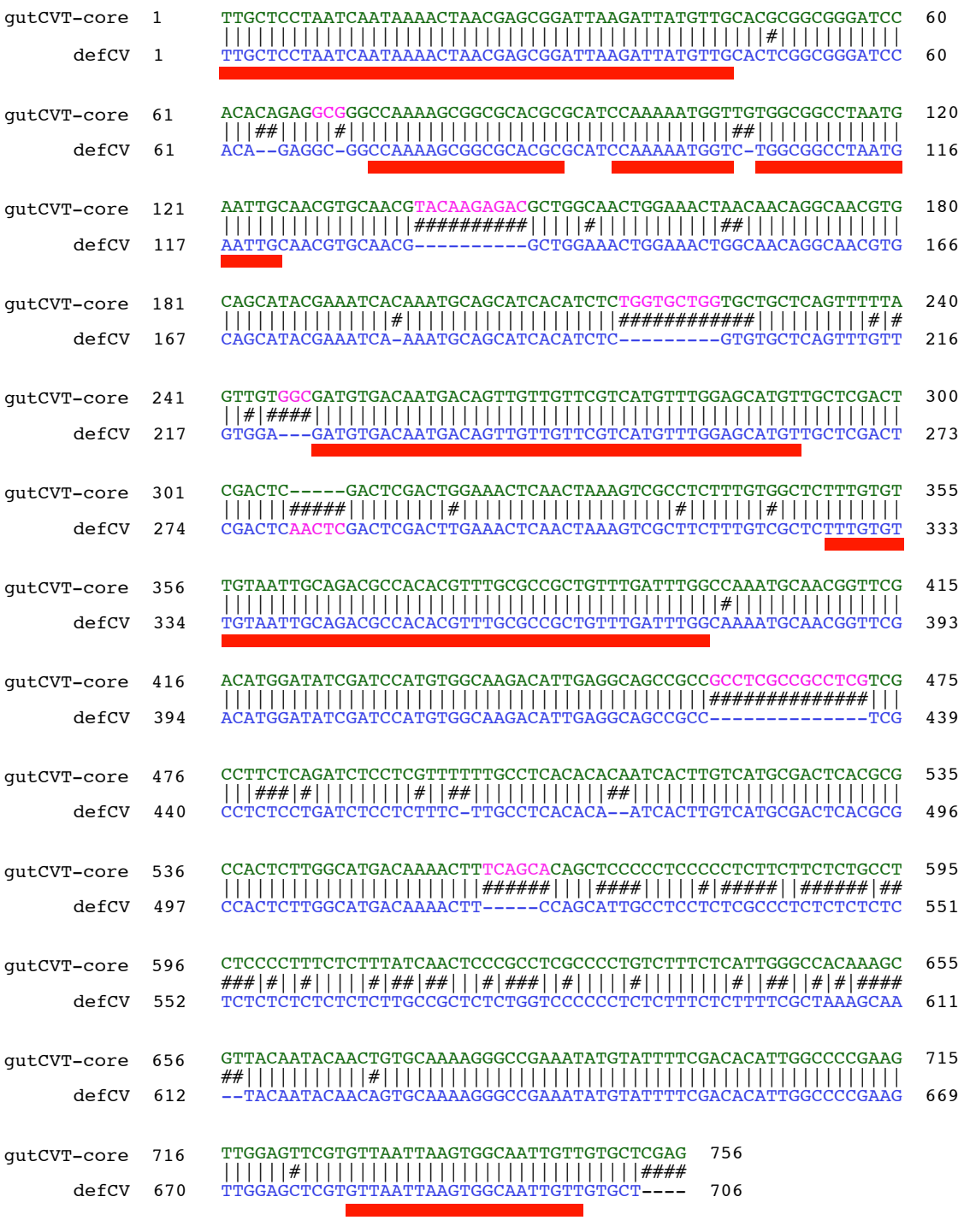

D. melanogaster sequences tested in D. melanogaster
D. guttifera sequences tested in D. guttifera
CG15170 (partial)
CG15169
CG13786
CG31909
Wnt4
wg
10kb
Wnt6
Wnt10
ninaC
ninaC (partial)

Figure. S3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defCV 1 TTGCTCCTAATCAATAAAACTAACGAGCGGATTAAGATTATGTTGCACTCGGCGGGATCC 60
gutCVT-core 61 ACACAGAGGCGGGCCAAAAGCGGCGCACGCGCATCCAAAAATGGTTGTGGCGGCCTAATG 120
defCV 61 ACA--GAGGC-GGCCAAAAGCGGCGCACGCGCATCCAAAAATGGTC-TGGCGGCCTAATG 116
gutCVT-core 121 AATTGCAACGTGCAACGTACAAGAGACGCTGGCAACTGGAAACTAACAACAGGCAACGTG 180
defCV 117 AATTGCAACGTGCAACG----------GCTGGAAACTGGAAACTGGCAACAGGCAACGTG 166
gutCVT-core 181 CAGCATACGAAATCACAAATGCAGCATCACATCTCTGGTGCTGGTGCTGCTCAGTTTTTA 240
defCV 167 CAGCATACGAAATCA-AAATGCAGCATCACATCTC---------GTGTGCTCAGTTTGTT 216
gutCVT-core 241 GTTGTGGCGATGTGACAATGACAGTTGTTGTTCGTCATGTTTGGAGCATGTTGCTCGACT 300
defCV 217 GTGGA---GATGTGACAATGACAGTTGTTGTTCGTCATGTTTGGAGCATGTTGCTCGACT 273
gutCVT-core 301 CGACTC-----GACTCGACTGGAAACTCAACTAAAGTCGCCTCTTTGTGGCTCTTTGTGT 355
defCV 274 CGACTCAACTCGACTCGACTTGAAACTCAACTAAAGTCGCTTCTTTGTCGCTCTTTGTGT 333
gutCVT-core 356 TGTAATTGCAGACGCCACACGTTTGCGCCGCTGTTTGATTTGGCCAAATGCAACGGTTCG 415
defCV 334 TGTAATTGCAGACGCCACACGTTTGCGCCGCTGTTTGATTTGGCAAAATGCAACGGTTCG 393
gutCVT-core 416 ACATGGATATCGATCCATGTGGCAAGACATTGAGGCAGCCGCCGCCTCGCCGCCTCGTCG 475
defCV 394 ACATGGATATCGATCCATGTGGCAAGACATTGAGGCAGCCGCC--------------TCG 439
gutCVT-core 476 CCTTCTCAGATCTCCTCGTTTTTTGCCTCACACACAATCACTTGTCATGCGACTCACGCG 535
defCV 440 CCTCTCCTGATCTCCTCTTTC-TTGCCTCACACA--ATCACTTGTCATGCGACTCACGCG 496
gutCVT-core 536 CCACTCTTGGCATGACAAAACTTTCAGCACAGCTCCCCCTCCCCCTCTTCTTCTCTGCCT 595
defCV 497 CCACTCTTGGCATGACAAAACTT-----CCAGCATTGCCTCCTCTCGCCCTCTCTCTCTC 551
gutCVT-core 596 CTCCCCTTTCTCTTTATCAACTCCCGCCTCGCCCCTGTCTTTCTCATTGGGCCACAAAGC 655
defCV 552 TCTCTCTCTCTCTCTTGCCGCTCTCTGGTCCCCCCTCTCTTTCTCTTTTCGCTAAAGCAA 611
gutCVT-core 656 GTTACAATACAACTGTGCAAAAGGGCCGAAATATGTATTTTCGACACATTGGCCCCGAAG 715
defCV 612 --TACAATACAACAGTGCAAAAGGGCCGAAATATGTATTTTCGACACATTGGCCCCGAAG 669
gutCVT-core 716 TTGGAGTTCGTGTTAATTAAGTGGCAATTGTTGTGCTCGAG 756
defCV 670 TTGGAGCTCGTGTTAATTAAGTGGCAATTGTTGTGCT---- 706
Similarity : 633/756 (83.73 %)

Figure. S4

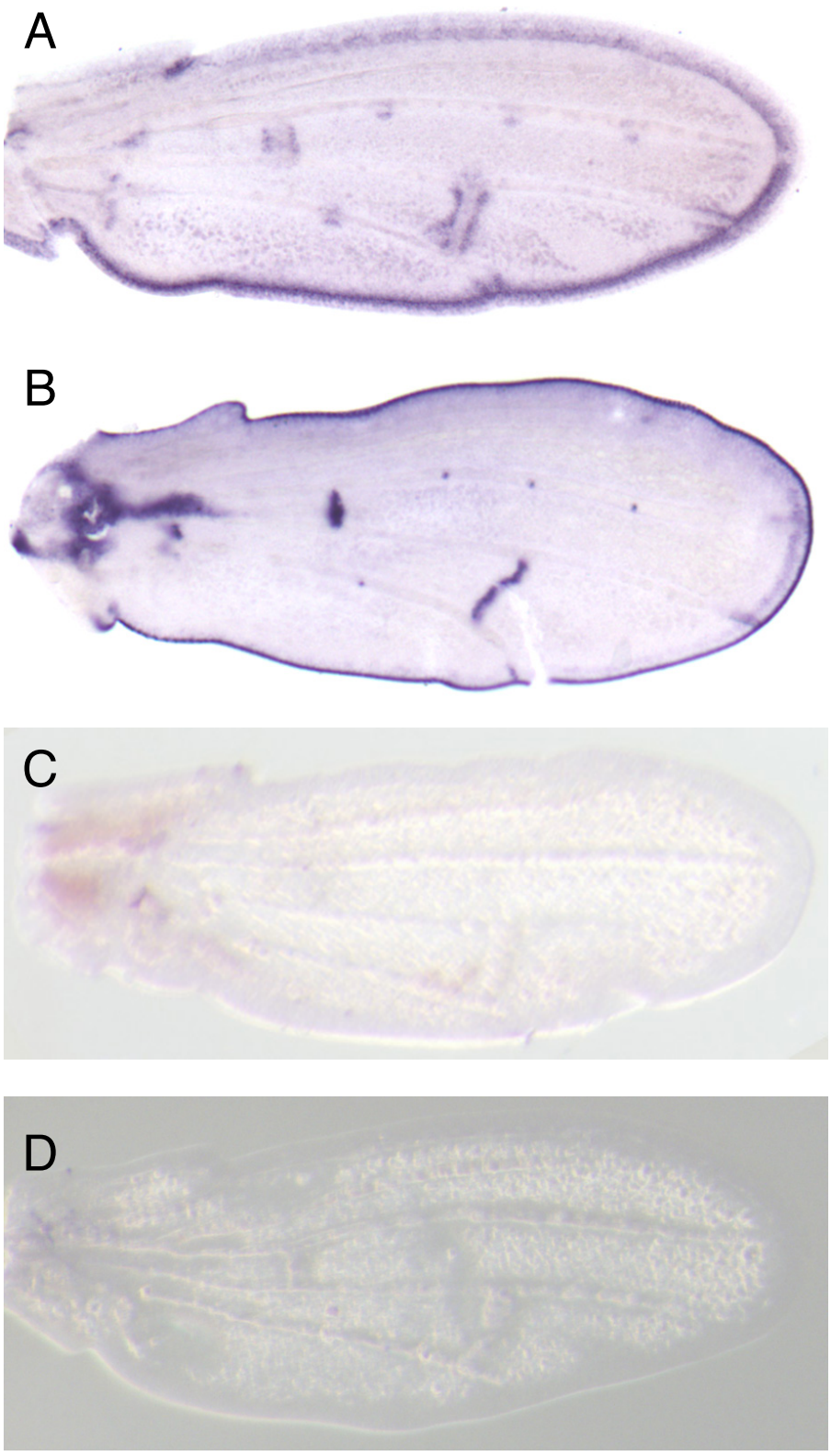

Figure. S5

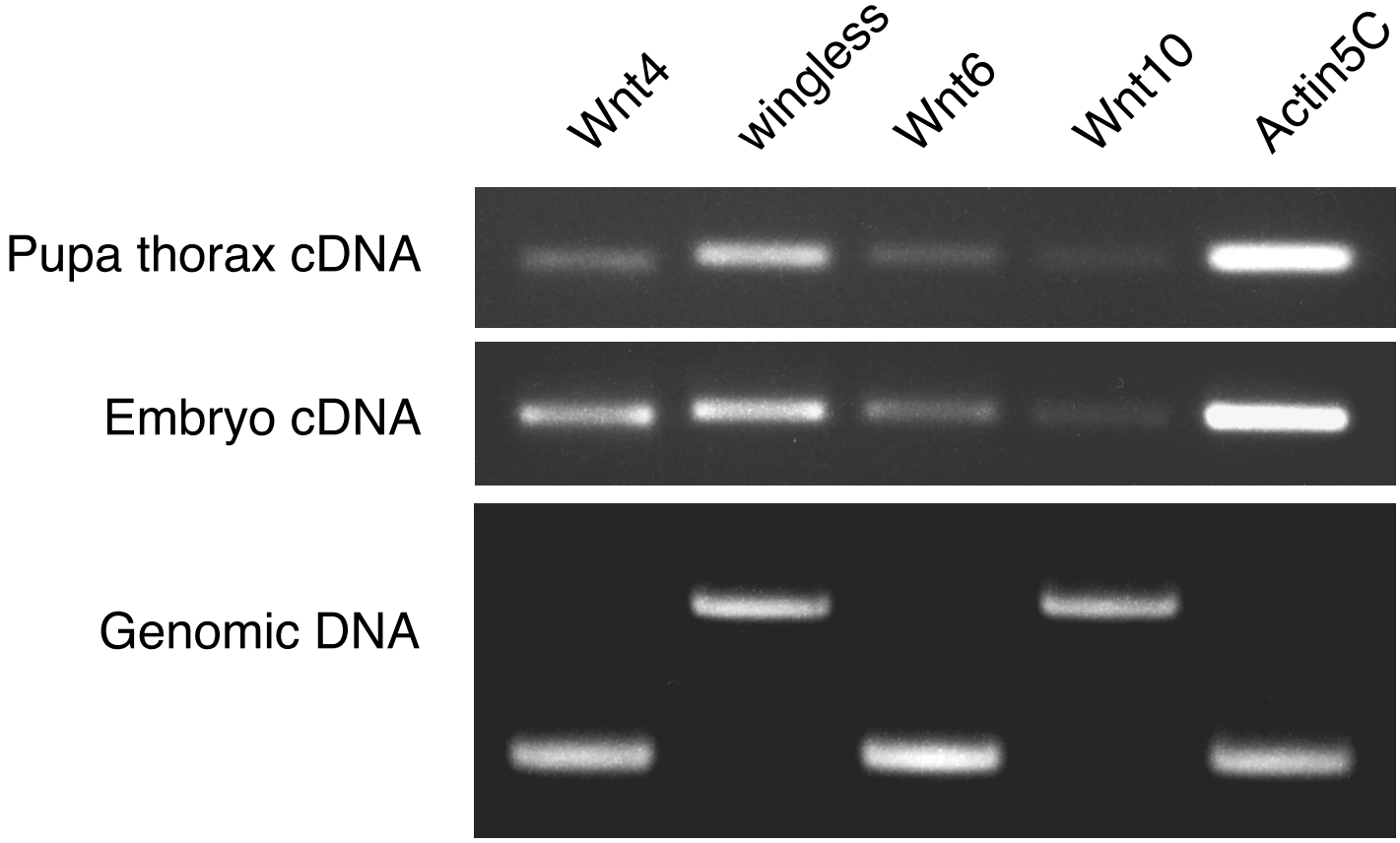
Wnt4
wingless
Wnt6
Wnt10
Actin5C
Pupa thorax cDNA
Embryo cDNA
Genomic DNA

Figure. S6

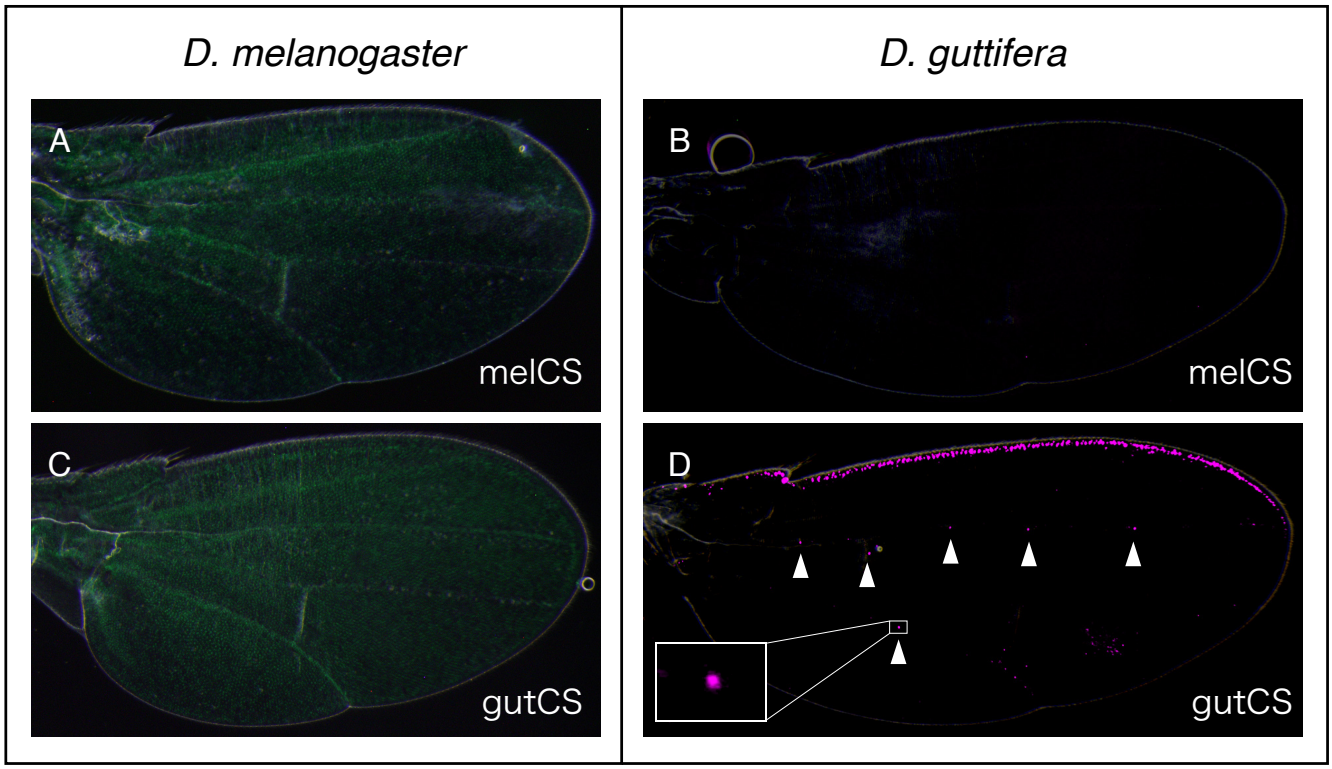

Figure. S7

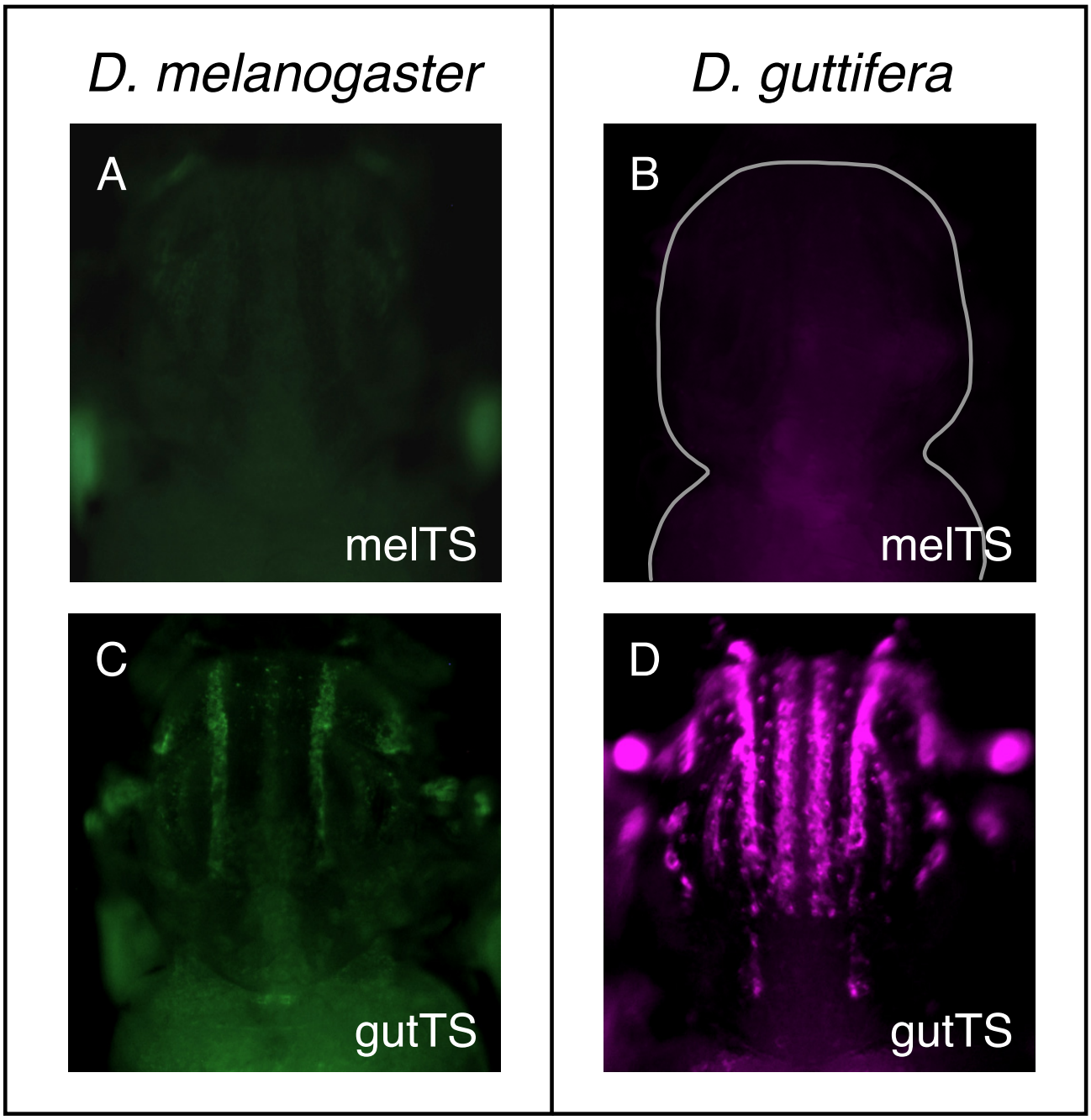

Figure. S8

Table S1. Primers used in the study.

Primers for enhancer screening (Figure S3)	Restriction site	Fragment in FigS3	Host species	Template specie	Cloning ststem
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ATCACCTGCAGGCAAAACTGAAATGAAATATAAAGTTTCAC	SbfI	1	melanogaster	melanogaster	S3aG
TCATGGCGCGCCCTGGCAGACATAAAGTATTCGAAATT	AscI	2	melanogaster	melanogaster	S3aG
ATCACCTGCAGGAGTTTTCTGACTGTAGGGTCTATT	SbfI	2	melanogaster	melanogaster	S3aG
TCATGGCGCGCCGACAGGTGGGATTTGAAAAATGTA	AscI	3	melanogaster	melanogaster	S3aG
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TCATGGCGCGCCCGAAAACCGCAAGACAAACGT	AscI	4	melanogaster	melanogaster	S3aG
AACACCTGCAGGCGTGGCAGCATTTGATTTGTTTAGA	SbfI	4	melanogaster	melanogaster	S3aG
TCATGGCGCGCCGCGAAACGCAATCAACATTAAAAGT	AscI	5	melanogaster	melanogaster	S3aG
AACGCCTGCAGGGGGTATTTTAATTCAATTTGTGGGCT	SbfI	5	melanogaster	melanogaster	S3aG
ACTAGGCGCGCCGTTTGACATTAAACGCGTGTTAATTTC	AscI	6	melanogaster	melanogaster	S3aG
AACGCCTGCAGGCATCGCCATGTGGCACCA	SbfI	6	melanogaster	melanogaster	S3aG
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AACGCCTGCAGGCAAAAGATAAGAGCAGCGCCATA	SbfI	9	melanogaster	melanogaster	S3aG
ACTTGGCGCGCCCGGTAATATGAGCAGTAATAAAATTTGA	AscI	10	melanogaster	melanogaster	S3aG
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AACGCCTGCAGGACAGGTGATAATTATATAATTTGTGAC	SbfI	12	melanogaster	melanogaster	S3aG
TCTTGGCGCGCCGAATTCTCGTGAACCTCCCAGCA	AscI	13	melanogaster	melanogaster	S3aG
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TACACCGCGGTCTAAGAGGTTCTAATTGAGCCAAT	SacII	55	guttifera	guttifera	pBac
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AACGGCTAGCCTAATATGTGTAAACCRCGTTAGTTAC	NheI	57	guttifera	guttifera	pBac
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AACGGCTAGCAGAARGTGAATCTGCACAACAACGAG	NheI	62	guttifera	guttifera	pBac
AACGGGTACCCATGCCRTGGCATTTGCACTC	KpnI	62	guttifera	guttifera	pBac
TCAACCGCGGAAACTTACAATTCAAAACTATTACTTGTTTTA	SacII	63	guttifera	guttifera	pBac

AACGGCTAGCGTAAAGAAGAAAATGAAATAGAAAGTGGA	NheI	63	guttifera	guttifera	pBac
TCAACCGCGGAAGATTATACACACGTGTCTGTAAG	SacII	64	guttifera	guttifera	pBac
AACGGCTAGCGCGAACGCTGCATAATGGAA	NheI	64	guttifera	guttifera	pBac
TCAACCGCGGAATAAGAATTGCGAATGCAATGCACA	SacII	65	guttifera	guttifera	pBac
AACGGCTAGCGCGAACGCTGCATAATGGAA	NheI	65	guttifera	guttifera	pBac
TCAACCGCGGAAAAAAAAGTGATCGTGCTACATGTGT	SacII	66	guttifera	guttifera	pBac
AACGGCTAGCAGCACAACAATTGCCACTTAATTAACA	NheI	66	guttifera	guttifera	pBac
TCAACCGCGGGACATTGCTCCTAATCAATAAAACTAA	SacII	67	guttifera	guttifera	pBac
AACGGCTAGCGCGAACGCTGCATAATGGAA	NheI	67	guttifera	guttifera	pBac
TCAACCGCGGGACATTGCTCCTAATCAATAAAACTAA	SacII	68	guttifera	guttifera	pBac
AACGGCTAGCAGCACAACAATTGCCACTTAATTAACA	NheI	68	guttifera	guttifera	pBac
ATATCTCGAGGACATTGCTCCTAATCAATAAAACTAA	XhoI	69	guttifera	guttifera	pBac
TCTGGCTAGCTGCCAATTTATCGATCAACACGCT	NheI	69	guttifera	guttifera	pBac
TCAACCGCGGTGCGCCAACTTTGTTTAACTTTG	SacII	70	guttifera	guttifera	pBac
AACGGCTAGCGCGATTCATAATATCATATTTCACTTCCT	NheI	70	guttifera	guttifera	pBac
TCAACCGCGGCGGCGAACAAAATTGCGAATGAA	SacII	71	guttifera	guttifera	pBac
AACGGCTAGCTTTTTTTAAAGATTTCTTTGATGAAAATTAGTAAG	NheI	71	guttifera	guttifera	pBac
TCAACCGCGGGCTGACGTTTAGTCATAAAATATTCCA	SacII	72	guttifera	guttifera	pBac
AACGGCTAGCGGCGCTGATTCAACAAATAAACAAA	NheI	72	guttifera	guttifera	pBac
TCAACCGCGGTGTTTAGAAATTTTAAGTTTCTCGATAAATG	SacII	73	guttifera	guttifera	pBac
AACGGCTAGCAAAATGAAATTAAACGCGCGCTTTAATCA	NheI	73	guttifera	guttifera	pBac
TCAACCGCGGATTTTATGACCCATTGATAGTGCGAA	SacII	74	guttifera	guttifera	pBac
AACGGCTAGCGACGCGCGAATTAATCAAGCG	NheI	74	guttifera	guttifera	pBac
TCAACCGCGGTCTACAGATACATTAGAAAATATCTCAAA	SacII	75	guttifera	guttifera	pBac
AACGGCTAGCTGGAATTTCCATTCATTTAACGGCAC	NheI	75	guttifera	guttifera	pBac
TCAACCGCGGGCTTTGATTAAAGCGCGCGTTTA	SacII	76	guttifera	guttifera	pBac
AACGGCTAGCGTTTGTAATGGGTTTTTTGGCATTGAA	NheI	76	guttifera	guttifera	pBac
TCAACCGCGGGAGTTCACAGTTAAAGTTCGAGC	SacII	77	guttifera	guttifera	pBac
ATCGGCTAGCCCTTGAACGGCACAGATGAAG	NheI	77	guttifera	guttifera	pBac
TCAACCGCGGGACCTTCAAAATGTGACGTTTGATTTA	SacII	78	guttifera	guttifera	pBac
AACGGCTAGCCCGCGACGTATCTTTATTTTGC	NheI	78	guttifera	guttifera	pBac
ATATCTCGAGTCGGCCAATTGCCAAAAATTAATGCA	XhoI	79	guttifera	guttifera	pBac
TCAACCGCGGGAGATTATGACAACTTAATAGCTACAGA	SacII	79	guttifera	guttifera	pBac
TCAACCGCGGTCCATTGAGGCCTATAACGACA	SacII	80	guttifera	guttifera	pBac
AACGGCTAGCAGAATATATAATTGTAAGAATAAGACTTTTAGA	NheI	80	guttifera	guttifera	pBac
TCAACCGCGGAAGTGCAACTAAATATGTAACTACTACAA	SacII	81	guttifera	guttifera	pBac
AACGGCTAGCGTGGGAGGCAGAAAGGATAAC	NheI	81	guttifera	guttifera	pBac
ATATCTCGAGAAGTAGGCAAAGAAAGAAGAAATCCT	XhoI	82	guttifera	guttifera	pBac
AACGGCTAGCGGAAAATGGAGCAAAAAGAATGCTT	NheI	82	guttifera	guttifera	pBac
TCAACCGCGGATGATTAAGCGTAATTTAATGAAGACAACA	SacII	83	guttifera	guttifera	pBac
AACGGCTAGCCAATTAAACAATTCATTTATCTTAATTTGTCTG	NheI	83	guttifera	guttifera	pBac
TCAACCGCGGTGACTTTCCCATAAATTAACACAATTTTATTGT	SacII	84	guttifera	guttifera	pBac
AACGGCTAGCGACTGCATTAAAAATCAACTTAATTTCA	NheI	84	guttifera	guttifera	pBac
TCAACCGCGGGTTAATAAAAAACACATAATTGCGTATGTT	SacII	85	guttifera	guttifera	pBac
AACGGCTAGCGTTTACAAACCACAGCACGCA	NheI	85	guttifera	guttifera	pBac
TCAACCGCGGAGCGCCACATCAACGTCATAA	SacII	86	guttifera	guttifera	pBac
AACGGCTAGCACACTTGTTACACTTCAAAGGACTT	NheI	86	guttifera	guttifera	pBac
TCAACCGCGGGTGATGGTGATTCATGGCAATTG	SacII	87	guttifera	guttifera	pBac
AACGGCTAGCAAACATTTTGTAGAACACATTGAAGAAAT	NheI	87	guttifera	guttifera	pBac
TCAACCGCGGAAGTGAGGCGAGCGTATCTTATA	SacII	88	guttifera	guttifera	pBac
AACGGCTAGCTGCCTAACTGGCTTAGATAGCTAA	NheI	88	guttifera	guttifera	pBac
TCAACCGCGGGCCTAGTAGTTGCAGCTTGTTA	SacII	89	guttifera	guttifera	pBac
AACGGCTAGCGGCATATAAATAGACAGTTTGAATTTATTA	NheI	89	guttifera	guttifera	pBac
TCAACCGCGGAAATAAATCTGCTTCTAATGCGAAAATG	SacII	90	guttifera	guttifera	pBac
AACGGCTAGCACATCAGCATAAAAAGCATAAATAAAGAAAG	NheI	90	guttifera	guttifera	pBac
AACGGGTACCGAGTTCCCCAGGTTTCCAC	KpnI	91	guttifera	guttifera	pBac
AACGGCTAGCTTACAGGAGGTAGGAAGAATGAGAA	NheI	91	guttifera	guttifera	pBac
TCAACCGCGGAAGCGAAGAAAAATACTAGTTTCAATTACA	SacII	92	guttifera	guttifera	pBac
AACGGCTAGCAGAAGTCGAAATGGGCAAGATATTAG	NheI	92	guttifera	guttifera	pBac
TCAACCGCGGGAGTGGGAGTTTTACGAATTGAATGT	SacII	93	guttifera	guttifera	pBac
AACGGCTAGCAAAAATTTAACAAAGAATGTGGGGTTAGATA	NheI	93	guttifera	guttifera	pBac

Primers for cis/trans test (Figure 3, 4, 5, S7, S8)		Fragment name	Host species	Template specie	Cloning system
TCAACCGCGGAAAAAAAGTGATCGTGCTACATGTGT	SacII	gutCV-T	guttifera	guttifera	pBac

AACGGCTAGCAGCACAACAATTGCCACTTAATTAACA	NheI	gutCV−T	guttifera	guttifera	pBac
AACGCCTGCAGGAAAAAAAGTGATCGTGCTACATGTGT	SbfI	melCV	melanogaster	melanogaster	S3aG
ACTAGGCGCGCCCTCACGCCTCGAAACAATTGC	AscI	melCV	melanogaster	melanogaster	S3aG
AACGCCTGCAGGTAAAAAAGTGATCGTGCTACATGTGT	SbfI	gutCV−T	melanogaster	guttifera	S3aG
TCTTGGCGCGCCCTCGAGCACAACAATTGCCACT	AscI	gutCV−T	melanogaster	guttifera	S3aG
TCAACCGCGGATAAAAAGTGATCGTGCTACATGTGT	SacII	defCV	guttifera	deflecta	pBac
ATATCTCGAGGACATTGCTCCTAATCAATAAAACTAA	XhoI	defCV	guttifera	deflecta	pBac
TCTGGCTAGCTGCCAATTTATCGATCAACACGCT	NheI	gutME	guttifera	guttifera	pBac
ATATCTCGAGGACATTGCTCCTAATCAATAAAACTAA	XhoI	gutME	guttifera	guttifera	pBac
AACGCCTGCAGGAAGTATTTAGTTTCAATTTGTTGCTTTG	SbfI	melME	melanogaster	melanogaster	S3aG
ACTTggcgcgccGCGAATGAAGACGTTTCGTGA	AscI	melME	melanogaster	melanogaster	S3aG
AACGCCTGCAGGAAAAAAAGTGATCGTGCTACATGTGT	SbfI	gutCVT5	melanogaster	guttifera	S3aG
TCTTGGCGCGCCAGACCGCGACGATGCGAT	AscI	gutCVT5	melanogaster	guttifera	S3aG
AACGCCTGCAGGGACATTGCTCCTAATCAATAAAACTAA	SbfI	gutCVT−core	melanogaster	guttifera	S3aG
TCTTGGCGCGCCCTCGAGCACAACAATTGCCACT	AscI	gutCVT−core	melanogaster	guttifera	S3aG
TCAACCGCGGGACATTGCTCCTAATCAATAAAACTAA	SacII	gutCVT−core	guttifera	guttifera	pBac
AACGGCTAGCAGCACAACAATTGCCACTTAATTAACA	NheI	gutCVT−core	guttifera	guttifera	pBac
ATCACCTGCAGGGCCTAGTAGTTGCAGCTTGTTA	SbfI	melCS	melanogaster	melanogaster	S3aG
ACTTGGCGCGCCGCAACGAAATGGGGTACAGTATTA	AscI	melCS	melanogaster	melanogaster	S3aG
ATATCTCGAGGCCTAGTAGTTGCAGCTTGTTA	XhoI	melCS	guttifera	melanogaster	pBac
TCAACCGCGGGCAACGAAATGGGGTACAGTATTA	SacII	melCS	guttifera	melanogaster	pBac
ATCACCTGCAGGGCCTAGTAGTTGCAGCTTGTTA	SbfI	gutCS	melanogaster	guttifera	S3aG
ACTTGGCGCGCCGGCATATAAATAGACAGTTTGAATTTATTA	AscI	gutCS	melanogaster	guttifera	S3aG
TCAACCGCGGGCCTAGTAGTTGCAGCTTGTTA	SacII	gutCS	guttifera	guttifera	pBac
AACGGCTAGCGGCATATAAATAGACAGTTTGAATTTATTA	NheI	gutCS	guttifera	guttifera	pBac
ATCACCTGCAGGGAGTCTCTCATCTATCCTAAGAC	SbfI	melTS	melanogaster	melanogaster	S3aG
ACTTGGCGCGCCGGCAAAATGCATTTTAATTGGCTGA	AscI	melTS	melanogaster	melanogaster	S3aG
ATATCTCGAGGAGTCTCTCATCTATCCTAAGAC	XhoI	melTS	guttifera	melanogaster	pBac
TCAACCGCGGGCAACGAAATGGGGTACAGTATTA	SacII	melTS	guttifera	melanogaster	pBac
ATCACCTGCAGGAAGTGAGGCGAGCGTATCTTATA	SbfI	gutTS	melanogaster	guttifera	S3aG
ACTTGGCGCGCCGGCATATAAATAGACAGTTTGAATTTATTA	AscI	gutTS	melanogaster	guttifera	S3aG
TCAACCGCGGAAGTGAGGCGAGCGTATCTTATA	SacII	gutTS	guttifera	guttifera	pBac
AACGGCTAGCTGCCTAACTGGCTTAGATAGCTAA	NheI	gutTS	guttifera	guttifera	pBac

Primers for in situ hybridization	Gene	Target species
CACGTCCAAGCGGAGATGCG	wg	melanogaster
GGCGACGGCATGTTCGGGTG	wg	melanogaster
CACGTTCAGGCGGAGATGCG	wg	guttifera, deflecta, nigromaculata, palstris, quinaria
GGCGATGGCATATTGGGATGATG	wg	guttifera, deflecta, nigromaculata, palstris, quinaria
CGAACACTTTATATCGGAGCA	Wnt4	guttifera
GAGTCATGTCGCAATATTTCGG	Wnt4	guttifera
GCCATTCGCGATGCGATG	Wnt6	guttifera
CTAGAGGCATGTGTTGACCTC	Wnt6	guttifera
GCCGTGTCCAATAACATGGAGT	Wnt10	guttifera
CCTGTATATCCGCTCCTAGAT	Wnt10	guttifera

Primers for RT−PCR	Gene	Template species
GAGCAGCAACTGTTGCTGTC	Wnt4	guttifera
GCCAATCCTTTGTTCACATTGATTC	Wnt4	guttifera
GAGTGCAAATGCCACGGCAT	wg	guttifera
GGCTCCAGATAGACAATATCCTT	wg	guttifera
GCCATTCGCGATGCGATG	Wnt6	guttifera
AATTATGTTCATGACTCTGCCGAG	Wnt6	guttifera
GTTATCGGGAAAGTGCTTTTGC	Wnt10	guttifera
CTTCAGCACTTTGCCAACAATGT	Wnt10	guttifera
ATGTGTGACGAAGAAGTTGCT	Act5C	guttifera
TAGATGGGCACAGTGTGG	Act5C	guttifera