

**Table S1.** Primers used for PCR genotyping of *dcNDP2* alleles and PCR product details.

<b>name</b>	<b>Primer sequence (5'-&gt;3')</b>	<b>Amplicon</b>
gntp-1	ggagattgccttctcaataaagtgg	no product from wild-type flies
gntp-2	gtacagagagggagagtcac	2672 bp from homozygous $\Delta$ <i>CNDP2</i> <sup>(mw)</sup> mutant flies
gntp-3	aaactcgacaccggtataacttc	no product from wild-type flies
gntp-4	cgttgactctgagaacacttctctg	2800 bp from homozygous $\Delta$ <i>CNDP2</i> <sup>(mw)</sup> mutant flies
gntp-5	ttttggcaggtaaacaataaaaacg	5457 bp from wild-type flies
gntp-6	athtagctgtttccagacgatag	4332 bp from homozygous $\Delta$ <i>CNDP2</i> <sup>(mw)</sup> mutant flies
		1271 bp from homozygous $\Delta$ <i>dcNDP2</i> mutant flies
gntp-5	ttttggcaggtaaacaataaaaacg	2382 bp from wild-type flies
gntp-7	tccattccaccatacgaccg	2467 bp from homozygous $\Delta$ <i>dcNDP2</i> <sup>(rescue+mw)</sup> flies
		2467 bp from homozygous $\Delta$ <i>dcNDP2</i> <sup>(rescue)</sup> flies
gntp-8	gacgaacaaatgaaaatctgctgtg	1354 bp from wild-type flies
gntp-6	athtagctgtttccagacgatag	7551 bp from homozygous $\Delta$ <i>dcNDP2</i> <sup>(rescue+mw)</sup> flies
		1435 bp from homozygous $\Delta$ <i>dcNDP2</i> <sup>(rescue)</sup> flies
gntp-9	catgatgaaataacataaggtgtcc	no product from wild-type flies
gntp-6	athtagctgtttccagacgatag	992 bp from homozygous $\Delta$ <i>dcNDP2</i> <sup>(rescue+mw)</sup> flies