		_		
	3'	dilp6	5'	dilp7
No. sites				
Intronic	n.a.	870	n.a.	170
Synonymous	n.a.	75.8	n.a.	112.1
Silent	766	1312.8	29	334.1
Non-synonymous	n.a.	245.2	n.a.	364.9
Total	766	1558	29	669
S				
Intronic	n.a.	18 (10)	n.a.	5 (0)
Synonymous	n.a.	0	n.a.	6 (4)
Silent	4(1)	22 (11)	1(1)	13 (5)
Non-synonymous	n. a.	1 (1)	n. a.	1 (1)
Total	4(1)	23 (12)	1(1)	14 (6)
$\pi$	. ,			. ,
Intronic	n.a.	0.006	n.a.	0.01
Synonymous	n.a.	0	n.a.	0.012
Silent	0.002	0.005	0.006	0.011
Non-synonymous	n.a.	0.0007	n.a.	0.0005
Total	0.002	0.004	0.006	0.005
h	6	11	2	9
Hd	0.85	0.98	0.17	0.91
K				
Intronic	n.a.	0.062	n.a.	0.166
Synonymous	n.a.	0.177	n.a.	0.118
Silent	0.043	0.071	n.a.	0.164
Non-synonymous	n.a.	0.021	n.a.	0.010
Total	0.043	0.062	n.a.	0.073

S, number of segregating sites (number of singletons in parentheses);  $\pi$ , nucleotide diversity; h, number of haplotypes; Hd, haplotype diversity; K, nucleotide divergence; n. a., not applicable.