



Complete Mitochondrial Genomes of *Lygus lineolaris* and *Lygus hesperus*

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The mitochondrial genome of *Lygus lineolaris* is 17027-bp. The *Lygus hesperus* mt genome is 17747-bp.

Both genomes contain 13 protein coding regions, 22 tRNA genes, and two ribosomal RNA genes.

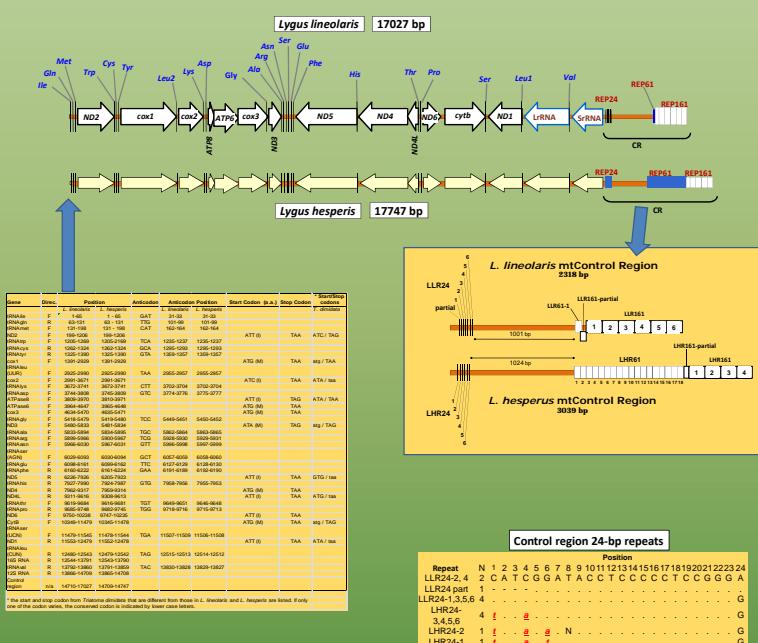
The gene arrangement corresponds to the common order found among insect mtDNAs.

Excluding the control region, the difference between the two genomes (14709 bp) is 3.3%.

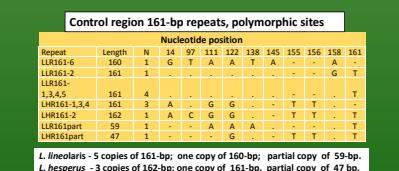
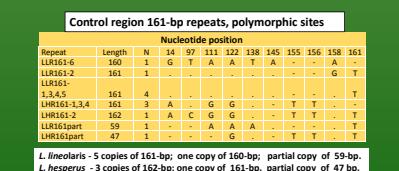
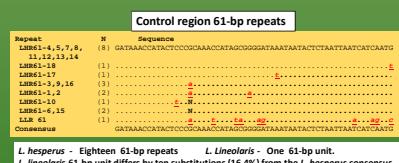
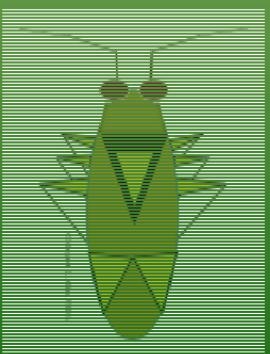
If only the ORFs (7450 bp) are compared, the sequence divergence is 1.4%.

The net 721 bp difference in genome lengths is completely accounted for by differences in the control regions.

The control regions contain an array of repetitive sequences which could be a source of species diagnostic sequences.



* One start and stop codon from Tenebrionidae that are different from those in *L. lineolaris* and *L. hesperus* are listed. Only one of the codon series, the converted codon is indicated by lower case letters.



Gene	Lugus A+T content of coding regions and RNAs											
	1st Position	2nd Position	3rd Position	Total	Size	%A+T	%G	%C	%T	%G	%C	%A+T
nd2	52	51	51	51	120	50.0	25.0	11.7	12.5	1.7	0.0	51.7
nd3	53	41	37	34	108	28.5	31.5	11.1	29.2	4.5	0.0	47.6
nd4	74	73	63	69	146	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5	74	73	63	67	140	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6	83	82	73	76	160	50.6	14.0	11.3	24.0	1.3	1.3	50.6
cyt b	74	73	63	67	140	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd1	84	84	85	85	168	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd4s	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5s	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6s	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5a	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6a	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5b	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6b	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5c	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6c	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5d	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6d	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5e	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6e	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5f	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6f	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5g	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6g	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5h	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6h	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5i	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6i	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5j	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6j	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5k	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6k	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5l	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6l	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5m	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6m	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5n	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6n	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5o	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6o	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5p	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6p	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5q	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6q	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5r	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6r	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5s	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6s	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5t	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6t	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5u	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6u	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5v	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6v	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5w	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6w	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5x	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6x	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5y	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6y	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5z	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6z	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5aa	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6aa	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5ab	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6ab	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5ac	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6ac	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5ad	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6ad	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5ae	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6ae	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd5af	74	73	63	66	132	50.0	14.0	11.1	24.4	1.4	1.4	50.0
nd6af	74	73										