Search

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VectorBase

Bioinformatics Resource for Invertebrate Vectors of Human Pathogens

What is search and how is it organized?

Search

Filter Results

Domain (Reset Filter)	Hits
Genome	4297312

Sub-domain	Hits
Genomic sequence assembly	3665236
Gene	584805
Mitochondrial genome	47271

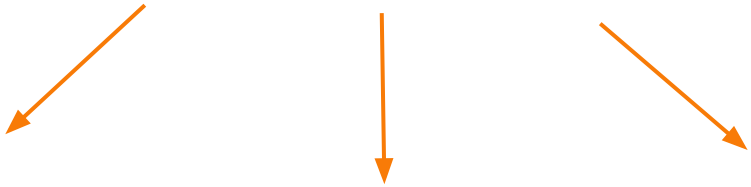
Species	Hits
Ixodes scapularis	960900
Biomphalaria glabrata	754144
Aedes albopictus	527435

Where is the Search box located?

Demo: Try Search using the keyword the wild card or asterisk *

Table 1
List of genes analyzed in the study. They are 1:1:1 orthologous genes that are predicted to code for perfectly conserved proteins among the three mosquitoes.

Gene name	Ortholog trios (Agam/Aaeg/Cqui)
Actin	AGAP005095/AAEL001673/CPIJ016462



AGAP******

Anopheles gambiae
PEST

AAEL******

Aedes aegypti
Liverpool

CPIJ******

Culex quinquefasciatus
Johannesburg

Rodriguez et al. 2012.
Infect Genet Evol. 12 (7):1535-42

Demo: Try Search using as keywords the gene name and the gene IDs.

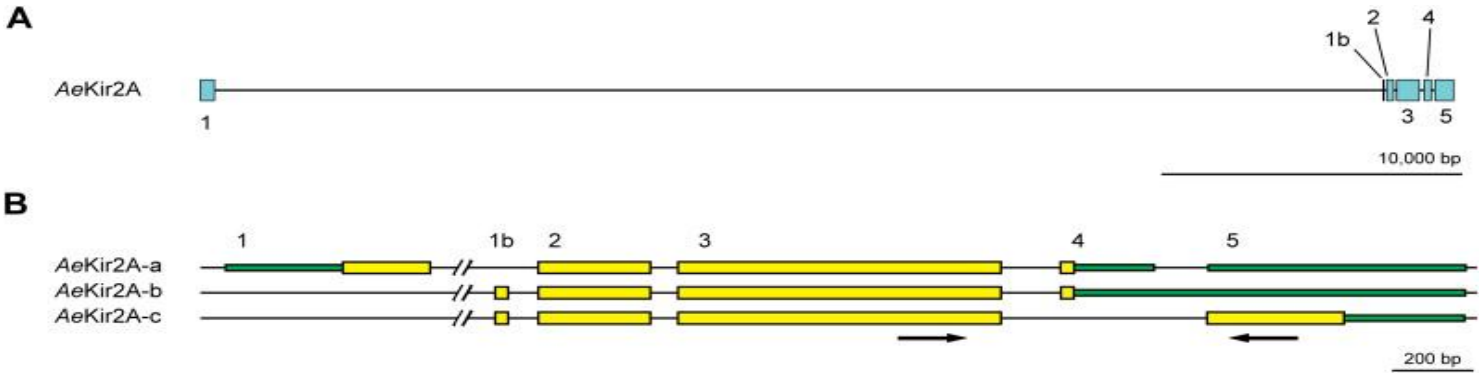


Fig. 2.

Genomic structure of the *Aedes aegypti* Kir2A gene: (A) Representation of the *Ae*Kir2A gene (to scale) on supercontig1.358 of the Liverpool strain of *Aedes aegypti*. Exons are indicated by blue boxes. Horizontal black bars represent introns. (B) Representation of the exon composition of the *Ae*Kir2A cDNAs (to scale) cloned in the present study. Exons are indicated by the colored bars. Green shading indicates a 5′ or 3′ untranslated region; yellow shading indicates protein coding sequence. Arrows signify the location of the splice-form specific RT-PCR primers (Supplemental Fig. 1). (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

Rouhier et al. 2014.
Insect Biochem Mol Biol 48: 91-9

Demo: Try Search using the as keywords the gene name

Keywords to look for a gene

- VectorBase gene ID
- Gene metadata (name and description) - Not recommended
- Terms from external databases such as InterPro or Gene Ontology (GO)
- Links from the archival databases, such as GenBank, to VectorBase

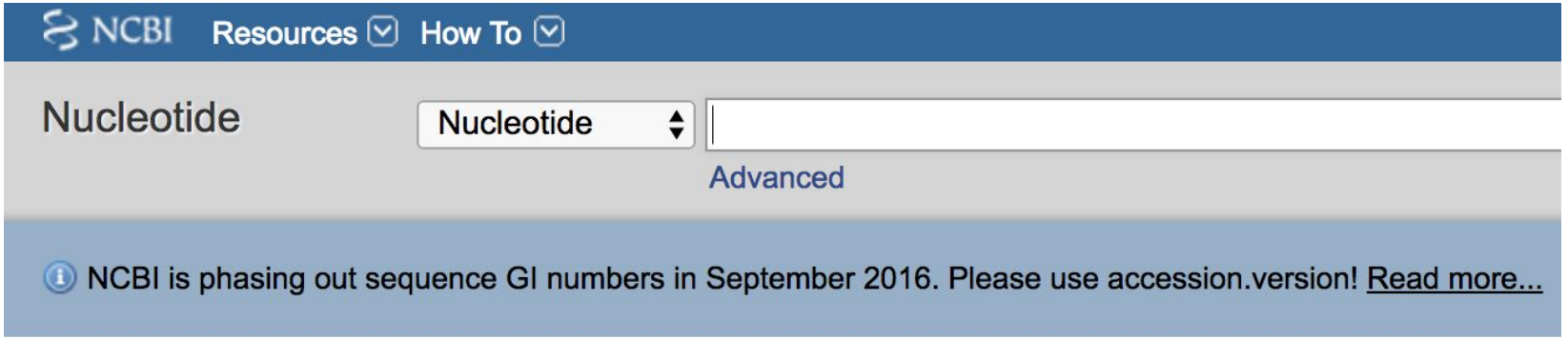
Abstract

Background

Pyrethroid insecticides, especially permethrin and deltamethrin, have been used extensively worldwide for mosquito control. However, insecticide resistance can spread through a population very rapidly under strong selection pressure from insecticide use. The upregulation of aldehyde dehydrogenase (*ALDH*) has been reported upon pyrethroid treatment. In *Aedes aegypti*, the increase in ALDH activity against the hydrolytic product of pyrethroid has been observed in DDT/permethrin-resistant strains. The objective of this study was to identify the role of individual ALDHs involved in pyrethroid metabolism.

Lumjuan et al. 2014.
PLoS ONE 9(7): e102746

Demo: Try Search using the as keywords the InterPro ID and gene description (with and without quotation marks)



GenBank

Aedes aegypti AAEL010392-RA partial mRNA

NCBI Reference Sequence: XM_001654436.1

FASTA Graphics

Go to:

```
LOCUS       XM_001654436            1999 bp    mRNA    linear   INV 05-MAR-2015
DEFINITION  Aedes aegypti AAEL010392-RA partial mRNA.
ACCESSION   XM_001654436
VERSION     XM_001654436.1  GI:157125998
```

Demo: Navigate from GenBank to VectorBase, following the gene ID link

In summary

How to search for more information or help?

E-mail us at
info@vectorbase.org

Thank you!