

# 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

→ <b>Domain (Reset Filter)</b>	<b>Hits</b> ▼
Genome	4297312
→ <b>Sub-domain</b>	<b>Hits</b> ▼
Genomic sequence assembly	3665236
Gene	584805
Mitochondrial genome	47271
→ <b>Species</b>	<b>Hits</b> ▼
<i>Ixodes scapularis</i>	960900
<i>Biomphalaria glabrata</i>	754144
<i>Aedes albopictus</i>	527435

# Where is the Search box located?



The screenshot shows the top section of the VectorBase website. On the left is a circular logo featuring a mosquito. To its right, the text reads "VectorBase" in a large blue font, followed by "Bioinformatics Resource for Invertebrate Vectors of Human Pathogens" in a smaller black font. In the top right corner, there is a search bar with the placeholder text "Enter search terms" and a blue "GO" button. Below the search bar, the text "Advanced Search" is visible. To the right of the search bar is a green "LOGIN" button. Below these elements is a horizontal navigation menu with buttons for "ABOUT", "ORGANISMS", "DOWNLOADS", "TOOLS", "DATA", "HELP", "COMMUNITY", and "CONTACT US". Below the navigation menu, there is a "Welcome to VectorBase!" section with a brief description of the site's purpose. To the right of this section, there is a box with the text "Want to see your BLAST, ClustalW and HMMer jobs? Log in or Register here."

**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.

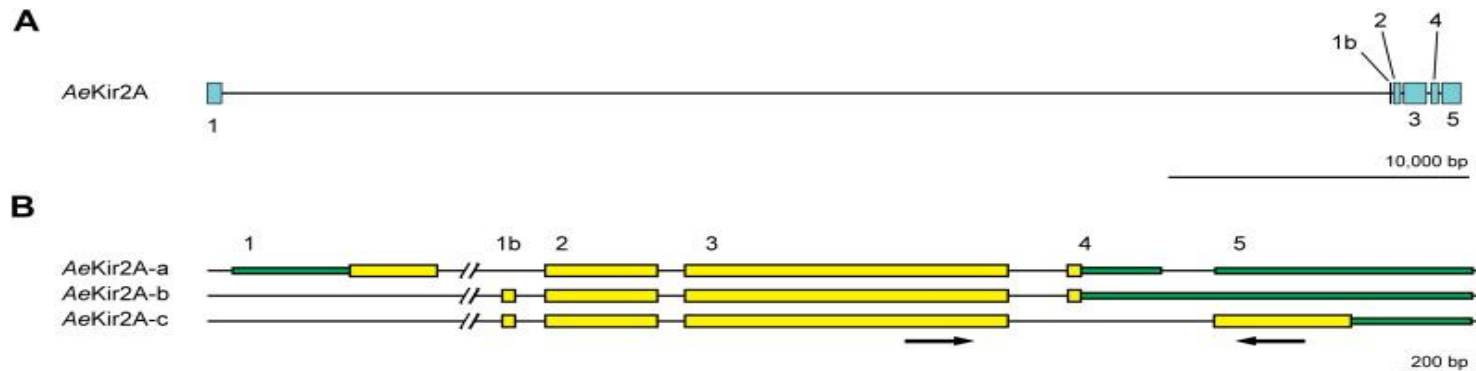






Fig. 2.

Genomic structure of the *Aedes aegypti* Kir2A gene: (A) Representation of the AeKir2A 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 AeKir2A 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)

Nucleotide

Nucleotide [Advanced](#)

 NCBI is phasing out sequence GI numbers in September 2016. Please use accession.version! [Read more...](#)

[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](mailto:info@vectorbase.org)

**Thank you!**