Vector Hazard Report: Mosquito-Borne Diseases of The Middle East



Information gathered from products of The Walter Reed Biosystematics Unit (WRBU):

Systematic Catalogue of the Culicidae VectorMap



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Table of Contents

1. Background

- General background
- <u>Countries of the Middle East, WHO profiles</u>
- Climate of the Middle East
- Monthly Climate Maps

2. Malaria

Disease risk maps: Plasmodium falciparum Plasmodium vivax Infectious days EIR: P. falciparum Prevalence: P. vivax **Dominant Malaria Vectors** Vector Habitat Suitability Models: Anopheles culicifacies s.l. Anopheles fluviatilis Anopheles sacharovi Anopheles stephensi Anopheles superpictus Malaria Vector Bionomics: Anopheles culicifacies Anopheles claviger



Malaria Vector Bionomics con't: <u>Anopheles fluviatilis</u> <u>Anopheles messeae</u> <u>Anopheles multicolor</u> <u>Anopheles pulcherrimus</u> <u>Anopheles sacharovi</u> <u>Anopheles sergentii</u> <u>Anopheles stephensi</u> <u>Anopheles superpictus</u>

3. Dengue Fever and Chikungunya virus

Disease risk maps Dengue virus prediction model CHIKV country of occurence Vector Habitat Suitability Models: Ae. albopictus **Vector Bionomics** 4. West Nile Virus Disease risk map: WNV country of occurrence **Vector Bionomics** 5. References

Background

Vector-borne disease hazards in the middle east include malaria, dengue fever, chikungunya virus, West Nile virus, leishmaniasis and Crimean-Congo hemorrhagic fever. This brief identifies factors that can impact disease risk including climate anomalies, vector distributions and host densities. All maps are available for download on VectorMap.org. More information on mosquito and sand fly vectors can be found at wrbu.org.



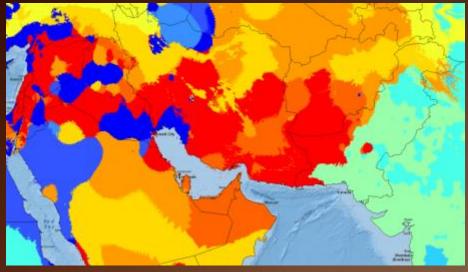
Countries of the Middle East

Background from the World Health Organization (WHO)

> Afghanistan Bahrain Iran Iraq Israel Jordan Lebanon Oman Pakistan Qatar Saudi Arabia United Arab Emirates



Climate of the Middle East









11. Nov 12. Dec VRAIR

1. Jan

2. Feb

3. Mar

4. Apr

5. May

6. Jun

7. Jul

8. Aug

9. Sep

10. Oct

Back to Table of Contents

Jan

Feb

Mar

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May

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Jul

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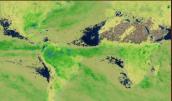
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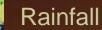
Dec

Monthly Climate Maps Click here to view the maps described below









This map shows the accumulated rainfall for the past month. Updated monthly. -NASA Earth Observations

Consistent Above and Below Average Precipitation

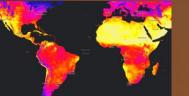
Areas with consistent above average monthly rainfall over the past 3 months may indicate increased mosquito breeding sites which may lead to increased mosquito-borne disease transmission. Areas with consistent below average rainfall may also indicate increased water storage or ponding which can provide additional habitat for mosquito species that lay eggs in human containers, protected micro environments, or long lasting pools. Updated monthly. -NASA Earth Observations.

Drought Breaking Rain

Areas receiving above average rainfall for the past month and below average rainfall for the previous 12 months. Drought breaking rain may indicate recent suitable conditions for vectors and diseases in a stressed environment or human population. Updated monthly. -WorldClim, Giovanni online data system NASA GES DISC, Tropical Rainfall Measuring Mission (TRMM).

Temperature anomaly

This map shows where earth's temperatures were warmer or cooler in the daytime for the past month than the average temperatures for the same month from 2001-2010. Updated monthly. -NASA Earth Observations



Land Surface Temperature

This map shows the temperature of the earth's lands during the daytime. Updated monthly. -NASA Earth Observations



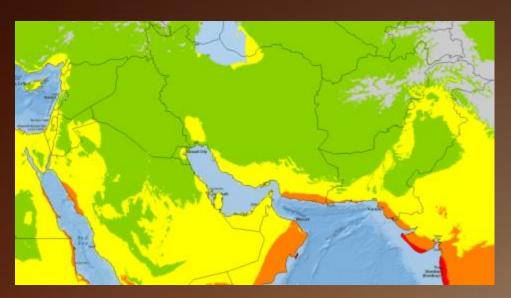
Malaria

Malaria is a mosquito borne disease caused by parasites of the genus Plasmodium. It is primarily spread through the bite of mosquitoes of the genus Anopheles. Symptoms of malaria are flu-like including high fever and chills. According to the CDC there were an estimate 219 million cases of malaria world-wide in 2010 with approximately 91% of infections occurring in Africa (CDC, 2014).

The World Health Organization (WHO) reports that about half the world's population is at risk of malaria infection. Although there is no current vaccine for malaria, the disease is preventable and treatable (WHO, 2014).



Plasmodium falciparum

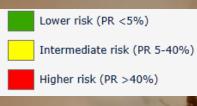


The normalized Z(T) index of temperature suitability that incorporates the duration and degree of suitability across an average year -Gething et al. 2011



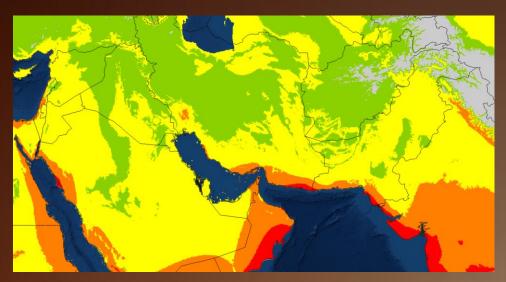


Stratified estimate proportion of 2-10 year olds in the general population that are infected with P. falciparum at any one time averaged over the 12 months of 2010. -Malaria Atlas Project





Plasmodium vivax



The normalized Z(T) index of temperature suitability that incorporates the duration and degree of suitability across an average year -Gething et al. 2011





Stratified estimate proportion of the general population that are infected with P. vivax at any one time averaged over the 12 months of 2010. -Malaria Atlas Project

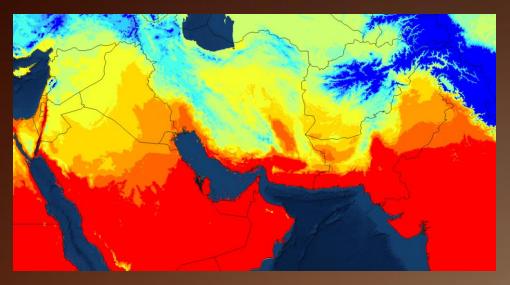


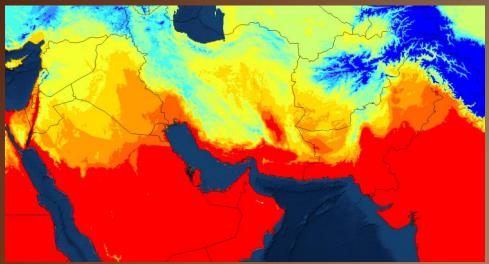
Intermediate risk (PR 1-5%)

Higher risk (PR >5%)



Infectious Days





The number of days in an average year (grouped by month), in which the annual temperature regime could support potentially infectious vectors of *P. falciparum*. -Gething et al. 2011



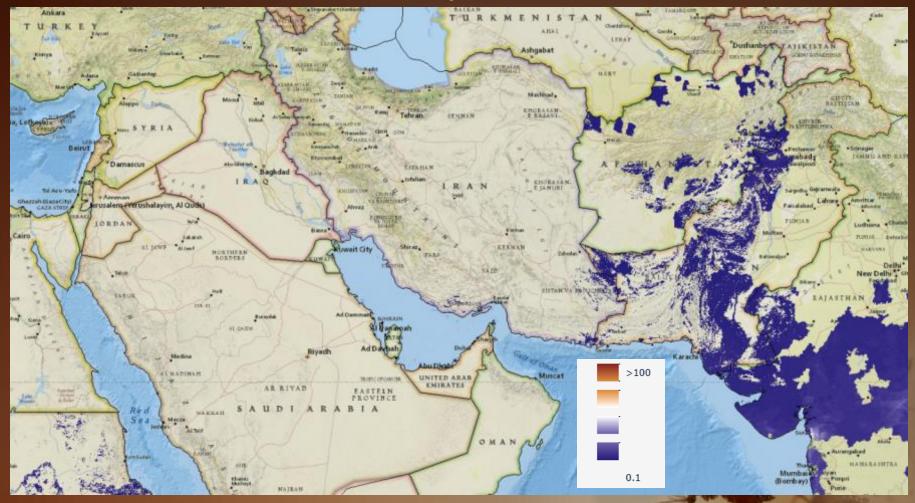
The number of days in an average year (grouped by month), in which the annual temperature regime could support potentially infectious vectors of *P. vivax*. -Gething et al. 2011





Malaria (Plasmodium falciparum) Entomological Inoculation Rate, 2010

Number of expected bites from infected mosquitoes per person, per year. -Gething et al. 2011







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Malaria (*Plasmodium vivax*) Estimated proportion of population infected with P. vivax, 2010

-Malaria Atlas Project



Malaria Vectors: Habitat Suitability Models

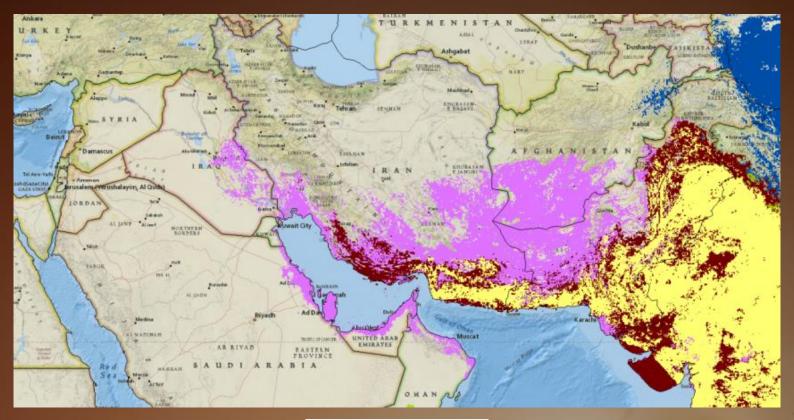




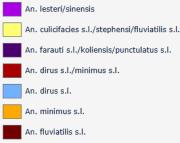


Dominant Malaria Vectors

-Malaria Atlas Project



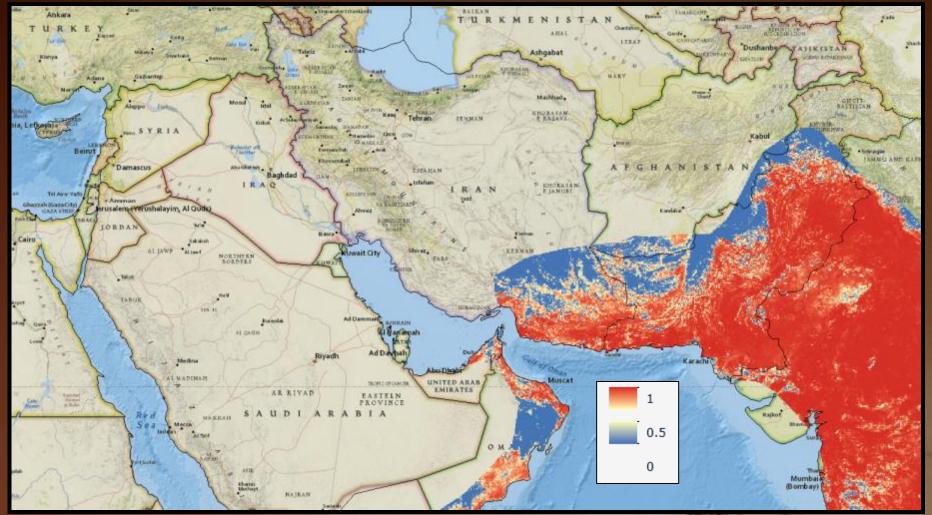




Back to Table of Content

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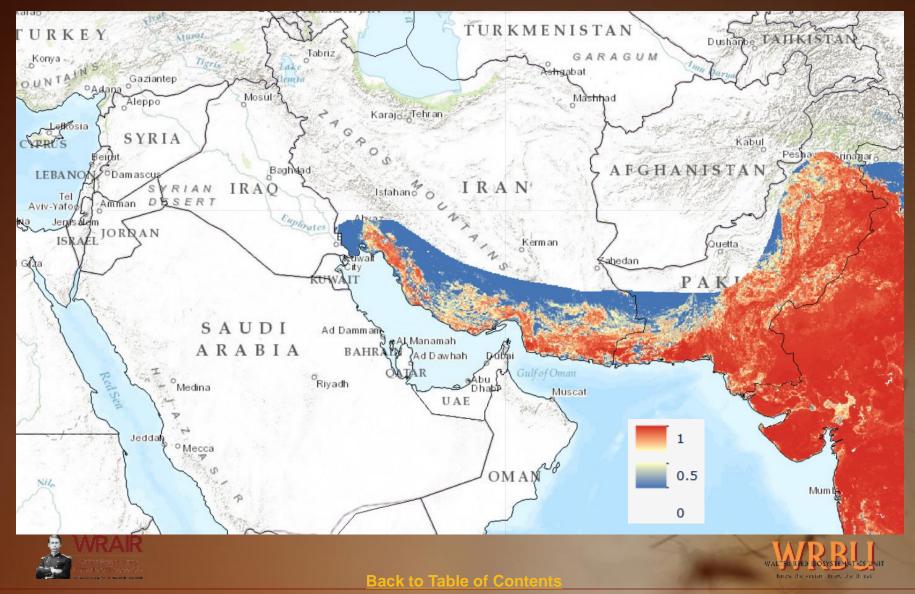
Habitat Suitability Models: Anopheles (Cel.) culicifacies s.l.







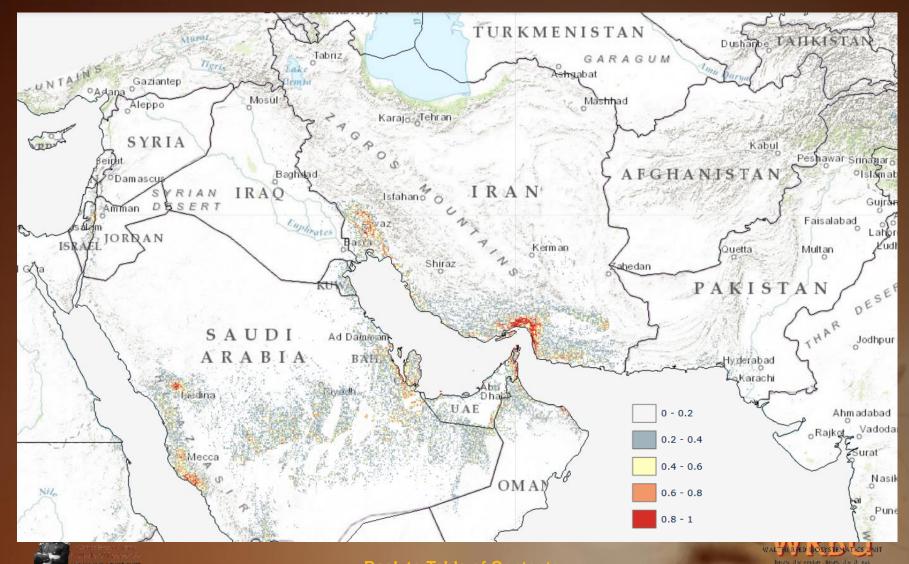
Habitat Suitability Model: An. (Cel.) fluviatilis



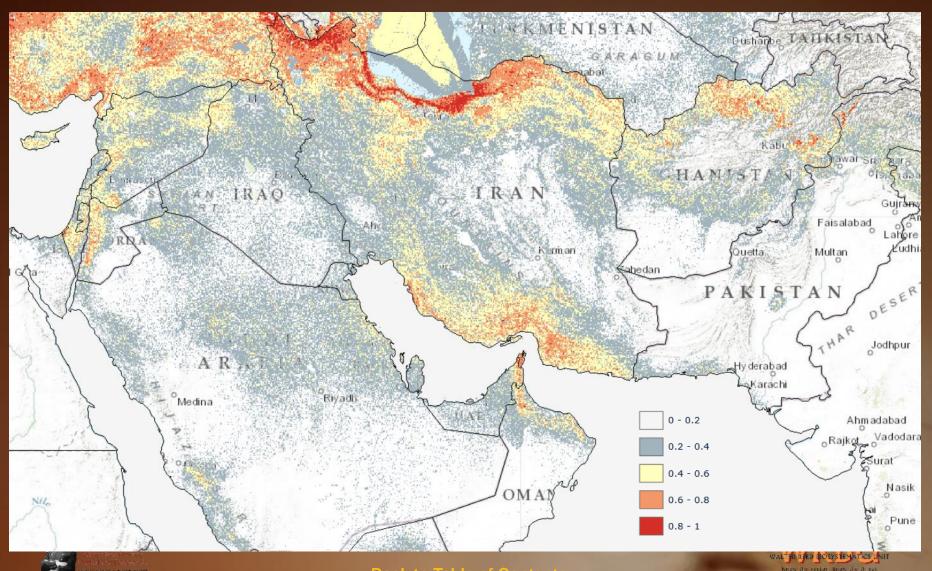
Habitat Suitability Model: Anopheles (Ano.) sacharovi



Habitat Suitability Model: Anopheles (Cel.) stephensi



Habitat Suitability Model: Anopheles (Cel.) superpictus



Anopheles (Cel.) culicifacies Giles, 1901

Bionomics:

Larvae are found in fresh water irrigation ditches, rain pools, pools in riverbeds, freshly dug pits or holes and wells. Females avoid oviposition site with emergent vegetation. Larvae found between 35 and 960m. in Thailand; in Vietnam only over 914 and in Pakistan usually between 1524 - 1829 but also up to 2286m (Harrison 1980).

Medical Importance:

Primary malaria vector (Harrison 1980).

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Anopheles (Ano.) claviger Meigan 1804

Bionomics

In higher latitudes the larvae are found in permanent pond and lake margins especially where shade is present. In southern parts of its range the species is restricted to cool spring-fed pools and similar habitats (Bates in Boyd 1949).

Medical Importance

Malaria vector in Middle East but not considered a vector in Europe (Bates in Boyd 1949).

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Anopheles (Ano.) fluviatilis James 1902

Bionomics:

Not usually found below 1000 ft. Larvae are found in the grassy edges of slow moving streams, springs, irrigation channels, sometimes in the edges of swamps and lakes. Females readily bite man, feeding generally before midnight (Christophers 1933)

Medical Importance:

Primary malaria vector (Christophers 1933)

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Anopheles (Ano.) messeae Falleroni 1926

Bionomics

Larvae are characteristically found in great inland river valleys and large marshes. This species rarely bites man and is associated with malaria only in a few places where a rather dense human population lives in close association with large marshy areas (Bates in Boyd 1949)

Medical Importance

The species rarely bites man and is associated with malaria only in a few places where a rather dense human population lives in close association with large marshy areas (parts of Hungary, and Lake Malik in Albania) (Bates in Boyd 1949).

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Anopheles (Cel.) multicolor Cambouliu 1902

Bionomics

An inland and coastal breeder of semi-arid regions in pans, oases and collections of brackish water. Breeds in fresh water, cesspools and highly saline pools. Females readily enter houses and bite man (Gillies and deMeillon 1968).

Medical Importance

Secondary malaria vector (Gillies and deMeillon 1968).

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Anopheles (Cel.) pulcherrimus Theobald 1902

Bionomics

Larvae are found in warm, sunny, stagnant habitats with abundant submerged vegetation. Where sufficiently warm, rice fields are used. Females readily bite man and animals (Beklemischev in Boyd 1949).

Medical Importance

Secondary malaria vector (Beklemischev and Boyd 1949).

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Anopheles (Ano.) sacharovi Favre 1903

Bionomics

In the Mediterranean area it breeds typically in large brackish marshes though larvae may at times be found in a wide variety of habitats. Adults are anthropophagous (Bates in Boyd 1949).

Medical Importance

Primary malaria vector (Bates in Boyd 1949).

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Anopheles (Cel.) sergentii Theobald 1907

Bionomics

Larvae occur in oases and irrigated areas in many types of water, shaded and unshaded, with and without vegetation. Females enter houses and readily bite man, (Gillies and deMeillon 1968).

Medical Importance

Primary malaria vector, (Gillies and deMeillon 1968).

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Anopheles (Cel.) stephensi Liston 1901

Bionomics

In urban areas the larvae are found in a wide variety of artifical containers, cisterns, wells, tubs and fountains. In the wild they are found in stream pools, stream margins, seepages, irrigation channels and springs. Females avidly bite man, (Puri in Boyd 1949).

Medical Importance

Primary malaria vector, (Christophers 1933)

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Anopheles (Cel.) superpictus Grassi 1899

Bionomics

Larvae are found in flowing water, including streams, rivers and irrigation channels in hilly and mountainous areas. Adults readily enter houses to bite man (Barraud 1933)

Medical Importance

Primary malaria vector, (Beklemischev in Boyd 1949)

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Dengue Fever and Chikungunya virus

According to the CDC, dengue fever is the leading cause of illness in the tropics and sub-tropics. Dengue fever is caused by one of a group of four viruses. Symptoms of Dengue include high fever, severe headache and joint pain (CDC, 2014).

The World Health Organization estimates that approximately 2.5 billion people are at risk from dengue with as many as 50 to 100 million annual infections. Currently it is now endemic in over 100 countries in Africa, the Americas, the Eastern Mediterranean, South-east Asia and the Western Pacific (<u>WHO, 2014</u>).

Chikungunya (pronounced: \chik-en-gun-ye) virus (CHIKV) is transmitted to humans by mosquitoes of the genus *Aedes*. The most common symptoms of chikungunya virus infection are fever and joint pain. Other symptoms may include headache, muscle pain, joint swelling, or rash (<u>CDC, 2014</u>).

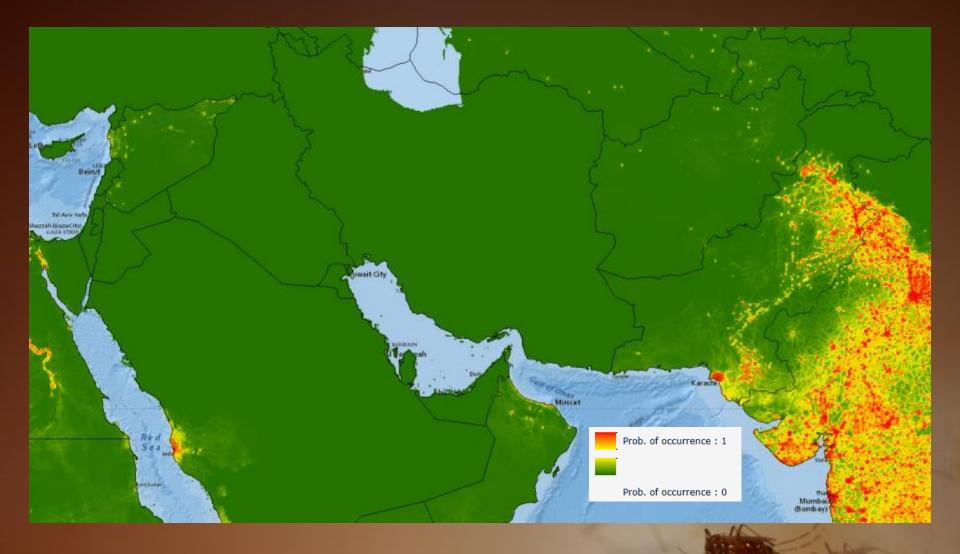
According to the World Health Organization (WHO), CHIKV has been reported in over 40 countries in Asia, Africa, Europe and also in the Americas (WHO, 2014).

Currently, there is no vaccine to prevent or medicine to treat dengue and chikungunya viruses.



Dengue Virus Prediction Model

-Bhatt, S. et al. 2013





Back to Table of Contents

Countries reporting CHIKV cases to CDC as of March 2015





Back to Table of Contents

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Habitat Suitability Model: Aedes aegypti



Habitat Suitability Model: Aedes albopictus



Aedes (Stg.) aegypti (Linnaeus, 1762) "Yellow Fever Mosquito"

Bionomics:

In association with man, aegypti will use any and all natural and artificial containers. Away from urban areas the species tends to favor pools in river beds, tree stumps, tree holes and natural containers. Females are primarily day biters and readily enter buildings to feed. They have also been taken in lesser numbers at night (Christophers 1960).

Medical Importance:

This species is a primary vector of dengue, chikungunya virus and yellow fever (Christophers 1960).

WRBU Catalog species page







Aedes (Stg.) albopictus (Skuse, 1894) "Asian Tiger Mosquito"

Bionomics:

Also known as the "Asian Tiger Mosquito", the immatures are found in natural containers, including treeholes, bamboo stumps, coconut shells, rockholes, palm fronds, and leaf axils. They are also found in all varieties of artificial containers and will breed indoors. The females readily bite man (Huang 1972).

Medical Importance:

This species is a known vector of dengue and yellow fever in the wild. Under laboratory conditions it has also been shown to vector bird malarias, Eastern and Western equine encephalitis, West Nile, chikungunya and Japanese encephalitis viruses (Huang, 1972).

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West Nile Virus

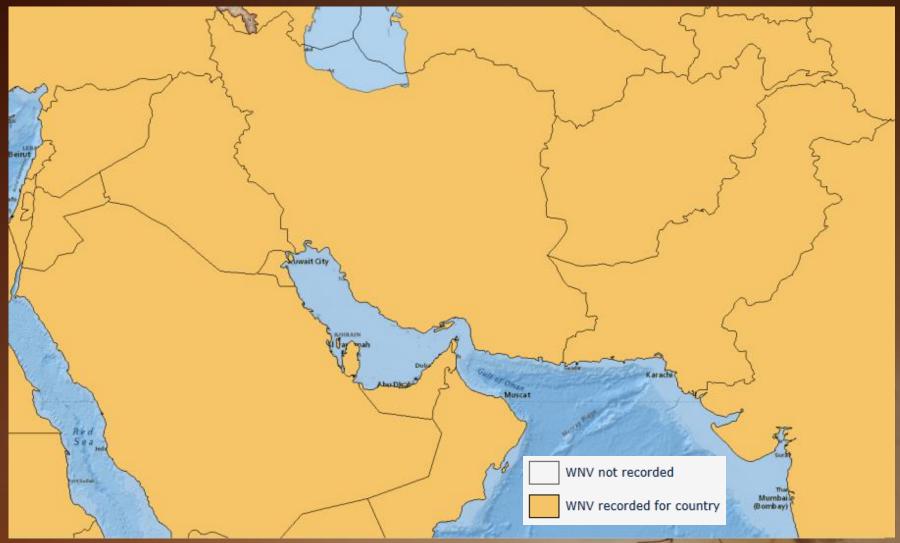
West Nile Virus is a mosquito-borne illness that primarily cycles between mosquitoes and birds. If enough of the virus is present in an infected bird and a mosquito takes a blood meal, that mosquito can then transmit the virus to a humans and other mammals. Mammals are a 'dead-end' host to WNV as there is typically not enough virus within a mammal to infect mosquitoes. (CDC, 2014)

According to the CDC, most people infected with WNV do not show symptoms. Approximately 1 in 5 infected people will show symptoms of fever and other flu-like symptoms. Fewer than 1% of humans infected with WNV will develop a serious, sometimes fatal, neurologic illness.

(<u>CDC, 2014</u>)



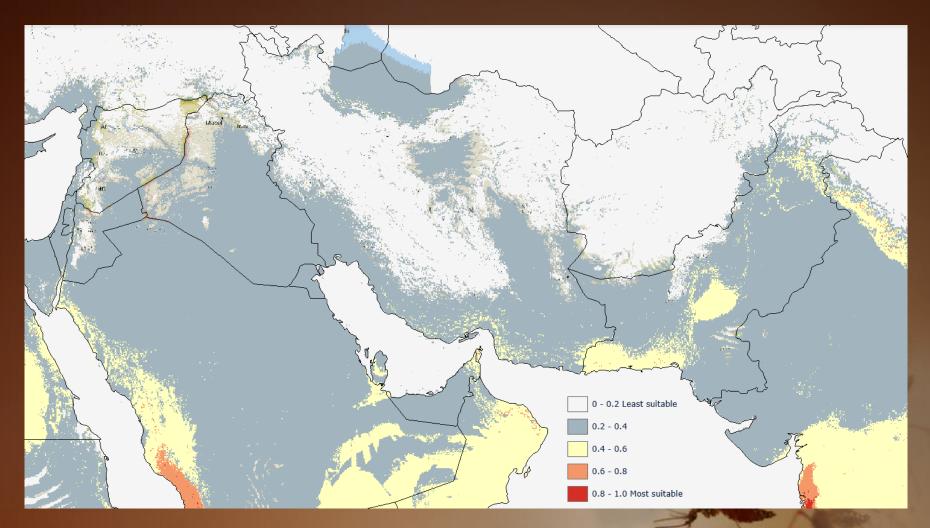
Countries Reporting West Nile Virus







Habitat Suitability Model: Aedes vexans







Habitat Suitability Model: *Culex (Cux.) pipiens*



Back to Table of Contents

Know the worker, know the threat

Aedes (Adm.) vexans (Meigan, 1830)

Bionomics

Immatures are found in unshaded fresh water flood pools in secondary scrub, but have also been collected in ditches, swamps, rice fields, and elephant foot prints. Habitats usually have little aquatic vegetation or algae. Females are night biters and readily feed on man and cattle (Reinert 1973).

Medical Importance

Capable of transmitting Eastern equine encephalitis virus (EEE), Western equine encephalitis virus (WEE), SLE, West Nile Virus (WNV) (Turell et al. 2005:60). It is also a vector of dog heartworm (Reinert 1973).

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Aedes (Och.) dorsalis (Meigan, 1830)

Bionomics

The larvae of *Ae. dorsalis* occur in a variety of habitats including both brackish and fresh water. They are often found in large numbers in tidal marshes but also occurs in fresh-water marshes and in overflow from artesian wells and irrigation ditches. (Carpenter and LaCasse 1955:176)

Medical Importance

Ae. dorsalis is considered a vector of Western Equine Encephalitis (WEE) and West Nile Virus (WNV) (Turell et al. 2005:60).

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Culex (Cux.) pipiens Linnaeus, 1758

Bionomics

Larvae are found in numerous and variable breeding places ranging from highly polluted cesspits to clear water pools and containers. This species usually breeds in stagnant water in either shaded or unshaded situations. Females readily attack man both indoors and outdoors, (Harbach 1988)

Medical Importance

It has been found naturally infected with Sindbis virus and West Nile viruses in Israel, West Nile and Rift Valley Fever in Egypt, and is a primary vector of periodic Bancroftian filariasis, (Harbach 1988)

WRBU Catalog Species Page





References

Maxent model of *Ae. aegypti* habitat suitability. A. Nyari. GlobalUniqueIdentifier: USNM:MosquitoModeI:Ae_aegypti_Nyari_1 Maxent model of *Ae. vexans* habitat suitability. A. Nyari. GlobalUniqueIdentifier: USNM:MosquitoModeI:Ae_vexans_Nyari_1 Maxent model of *Ae. albopictus* habitat suitability. GlobalUniqueIdentifier: USNM:MosquitoModeI:Ae_albopictus_Dornak_1 Maxent model of *Ae. aegypti* habitat suitability Maxent model of *Cx. pipiens* habitat suitability, Dornak, L. Maxent model of *An. superpictus* habitat suitability, Dornak, L. Maxent model of *An. stephensi* habitat suitability, Dornak, L. *An. culicifacies* species complex. Sinka et al. 2011. Malaria Atlas Project. *An. sacharovi* species complex. Sinka et al. 2011. Malaria Atlas Project.

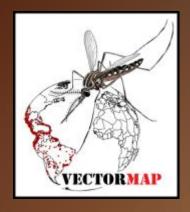
- CDC. Countries where West Nile Virus is present in year 2000. http://www.cdc.gov/ncidod/dvbid/westnile/map.htm
- CDC. Countries with reported current or previous local transmission of chikungunya virus (as of Oct 21, 2014). http://www.cdc.gov/chikungunya/map/index.html
- Plasmodium falciparum, Temp suitability index. The normalized Z(T) index of temperature suitability that incorporates not just the duration but also the degree of suitability across an average year. Source: Gething et al. Parasites & Vectors 2011, 4:92.
- Plasmodium vivax, Temp suitability index. The normalized Z(T) index of temperature suitability that incorporates not just the duration but also the degree of suitability across an average year. Source: Gething et al. Parasites & Vectors 2011, 4:92.
- Plasmodium falciparum, Infectious Days. The number of days in an average year in which the annual temperature regime could support potentially infectious vectors. Source: Gething et al. Parasites & Vectors 2011, 4:92.
- Plasmodium vivax, Infectious Days. The number of days in an average year in which the annual temperature regime could support potentially infectious vectors. Source: Gething et al. Parasites & Vectors 2011, 4:92
- Estimated proportion of the general population that are infected with P. vivax at any one time, averaged over the 12 months of 2010. Malaria Atlas Project http://www.map.ox.ac.uk/browse-resources/endemicity/Pv_mean/world/
- Bhatt, S. et al. 2013. The Global Distribution and Burden of Dengue. Natrure, 496: 504-507.

Mosquito images courtesy of J. Stoffer, Walter Reed Biosystematics Unit





The Walter Reed Biosystematics Unit is part of the Walter Reed Army Institute of Research and is based at the Smithsonian Institution Museum Support Center. To access taxonomic keys, the Systematic Catalog of Culicidae or to learn more about WRBU visit www.wrbu.org.



VectorMap is only as good as the data you provide. If you have collection records, models or pathogen testing results please contact the VectorMap team to learn how to contribute data at mosquitomap@si.edu.





Back to Table of Contents







Vector Photos Courtesy of Judith Stoffer, Walter Reed Biosystematics Unit

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