





**EFSA's risk assessments  
on vector-borne diseases:  
an example of assessing the risk  
of mosquito-borne diseases**

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**Outline**

- Vector-borne diseases in EFSA
- Fever and Mintrisk
- Example of risk assessment of mosquito-borne diseases in the EU
- Future challenges


2

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# ■ VECTOR-BORNE DISEASES IN EFSA

3

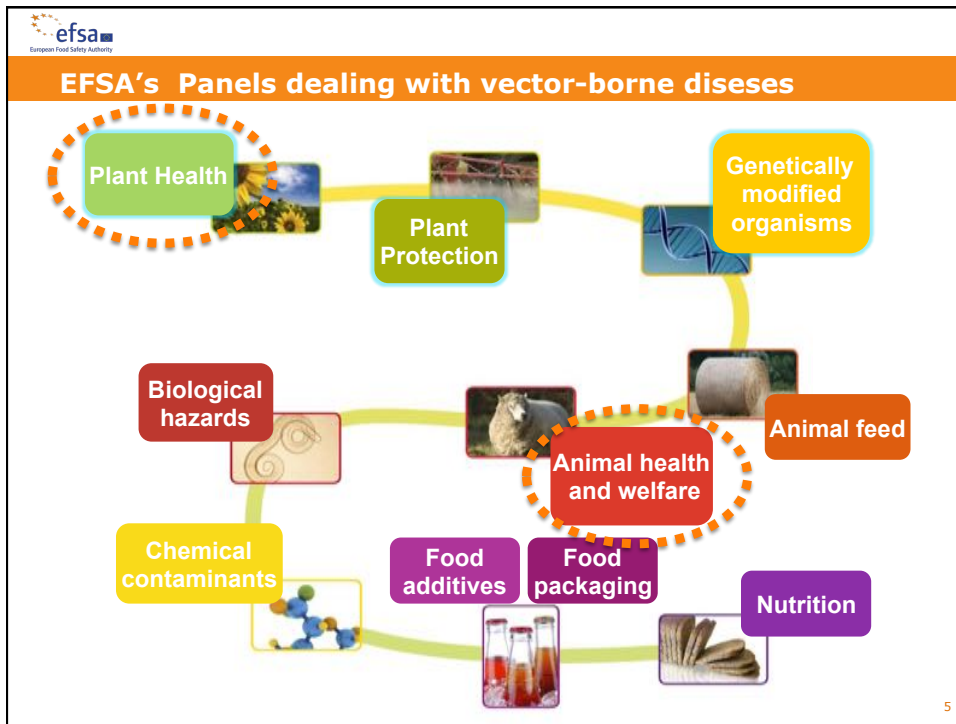
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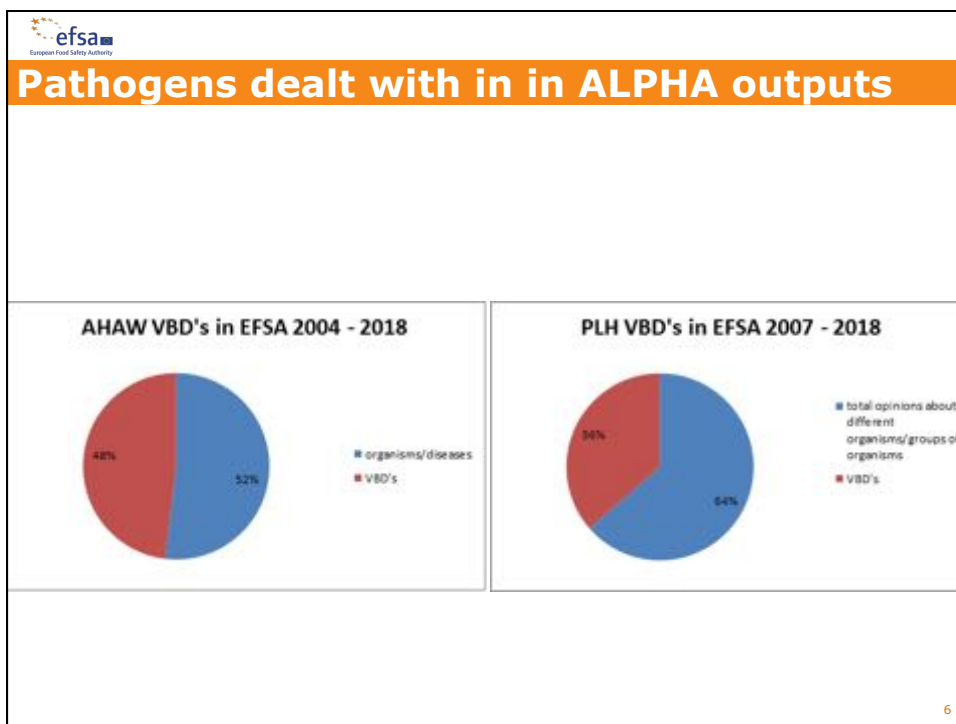
## How does it work?



4



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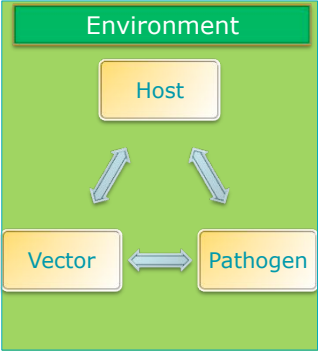
**ALPHA: One-health approach**

- Climate change, social and political instability, changes in landscape : potential drivers for their introduction
- Complex interaction between host, pathogen and environment > complex diseases to prevent and control

-**Human VBD's**: 17% of the estimated global burden of all infectious diseases

-**Animals VBD**: bluetongue virus: ~1.4B \$USD in economic losses in France in 2007 alone (Tabachnik et al. 2008)

-**Plant VBD**: citrus tristeza virus (an aphid-borne disease) has killed tens of millions of citrus trees worldwide, and currently threatens California orange crops



The diagram illustrates the One-Health approach. It features a green background with a dark green bar at the top labeled 'Environment'. Below this, a yellow box labeled 'Host' is positioned at the top. At the bottom, two yellow boxes labeled 'Vector' and 'Pathogen' are connected by a double-headed arrow. Two double-headed arrows also connect the 'Host' box to the 'Vector' and 'Pathogen' boxes, indicating complex interactions between all three components within the environment.

**AGENTS/PATHOGENS**

ANIMALS AND/OR HUMANS	PLANTS
<p><b>Viruses:</b> e.g: Dengue fever virus, Rift Valley fever virus, Japanese encephalitis virus, West Nile virus, Crimean-Congo hemorrhagic fever virus, etc</p> <p><b>Bacteria:</b> e.g.: <i>Borrelia burgdorferi</i> (bacteria causing Lyme disease) or <i>Yersinia pestis</i> (causing plague)</p> <p><b>Protozoa:</b> e.g.: <i>Trypanosoma brucei gambiense</i> (sleeping disease) or <i>L. infantum</i> (leishmaniasis), <i>Plasmodium spp.</i> (malaria)</p>	<p><b>Viruses:</b> e.g: Banana bunchy top virus BBTV, Arabis mosaic virus, Raspberry ringspot virus (more than 200 plant viruses transmitted by a single whitefly, <i>Bemisia tabaci</i>)</p> <p><b>Viroids</b> (eg Pospiviroids transmission by aphids and bumble bees)</p> <p><b>Bacteria:</b> e.g.: <i>Xylella fastidiosa</i> causing Pierce's disease by xylem-sap feeder insects</p> <p><b>Phytoplasma eg.</b> eg grapevine Flavescence dorée by leafhoppers</p> <p><b>Nematodes:</b> e.g.: <i>Bursaphelenchus xylophilus</i> (vectored by <i>Monochamus spp.</i> insect)</p> <p><b>Fungi:</b> <i>Gibberella circinata</i>, <i>Monilinia fructicola</i>, <i>Bretziella fagacearum</i> by beetles</p>


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## VECTORS


**ANIMALS AND/OR HUMANS**

Vector species: the largest part are blood-sucking **arthropods**


Insects (6 legs)

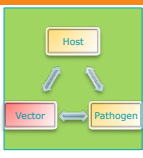


Arachnids (8 legs)




Other creepy stuff






Insects (6 legs)




Snail (1 leg)

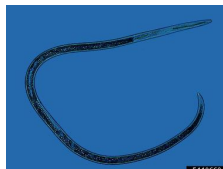


**PLANTS**

Arachnids (8 legs)



Nematodes (no legs...)




And...some fungi transmitting viruses  
(like *Synchytrium endobioticum*)

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## INSECT VECTORS

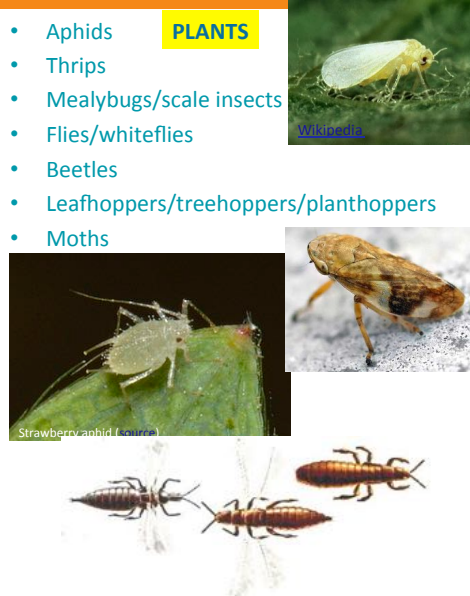
**ANIMALS AND/OR HUMANS**

- Mosquitoes (*Culicidae*)
- Culicoids (*Culicoides* spp.)
- Sandflies (*Phlebotomus* spp.)
- Black flies (*Simuliidae*)



**PLANTS**

- Aphids
- Thrips
- Mealybugs/scale insects
- Flies/whiteflies
- Beetles
- Leafhoppers/treehoppers/planthoppers
- Moths





Strawberry aphid (source)

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## ARACHNID VECTORS

**ANIMALS AND/OR HUMANS**


- Mites
- Ticks

Hyalomma Dermacentor Rhipicephalus Ixodes

**PLANTS**

- Mites



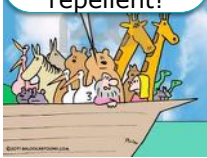
120 µm

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## HOSTS

**ANIMALS AND/OR HUMANS**

When this is over I am going to need a lot of mosquito repellent!




- Large range of animals (pets, livestock, wildlife, e.g. rodents, birds...)
- Humans
- Both animals and humans (=zoonotic VBD)

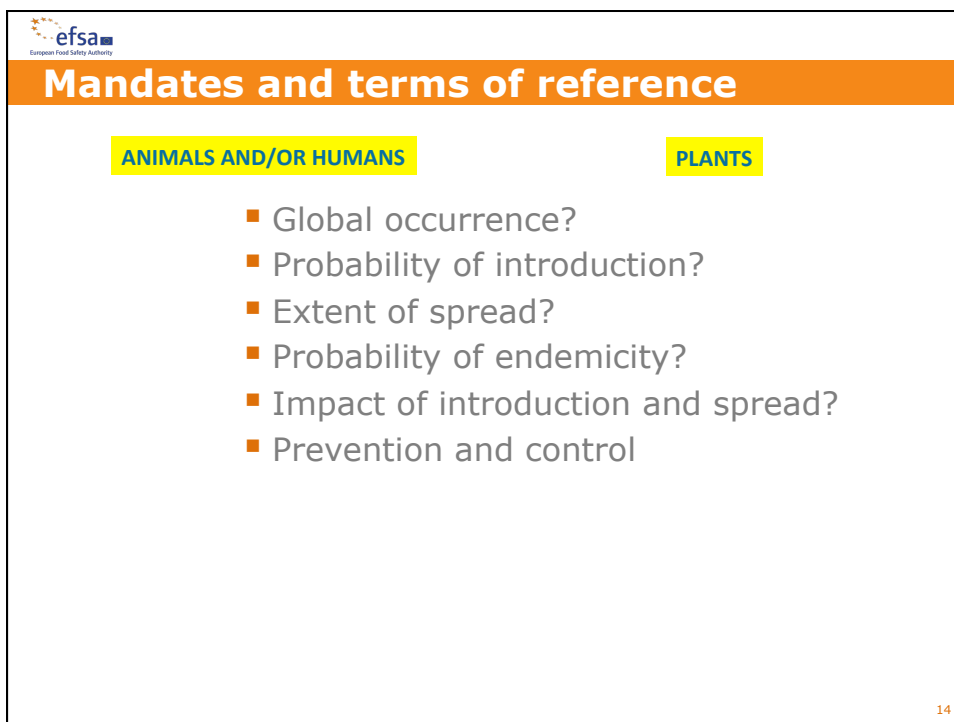
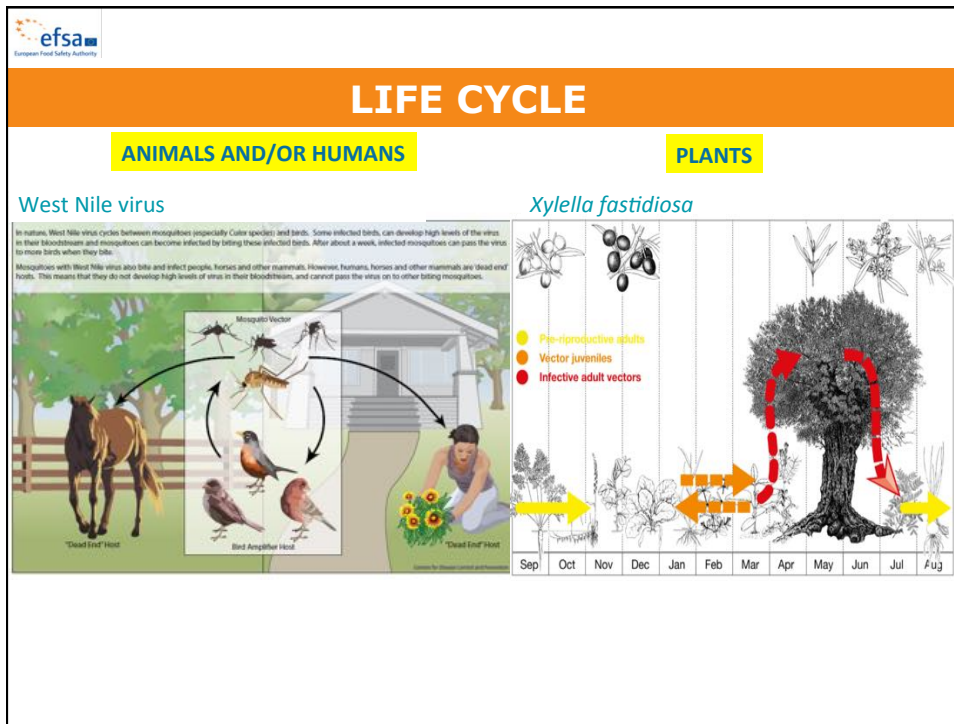
**PLANTS**

Tree of Plant Evolution

- Very large range of plant species: (ca. 300 000 species):
- Herbaceous crops (cereals, legumes, forage, vegetables ...)
- Fruit trees and bushes
- Forest trees
- Wild plants in natural habitats
- Wild crop relatives
- Ornamentals (cut flowers, pot plants, landscape and amenity trees)



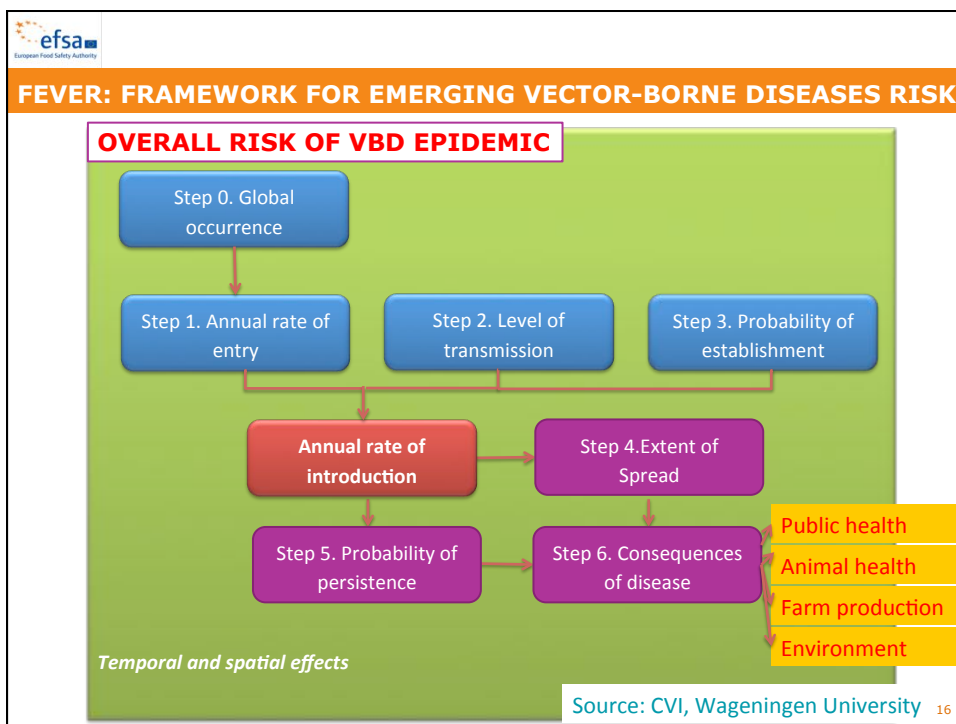
ALL OTHER LIFE




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## ■ FEVER and MINTRISK

15






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**METHOD FOR INTEGRATED RISK ASSESSMENTS**

<https://www.wecr.wur.nl/mintrisk/>

Qualitative or quantitative data input and output



17

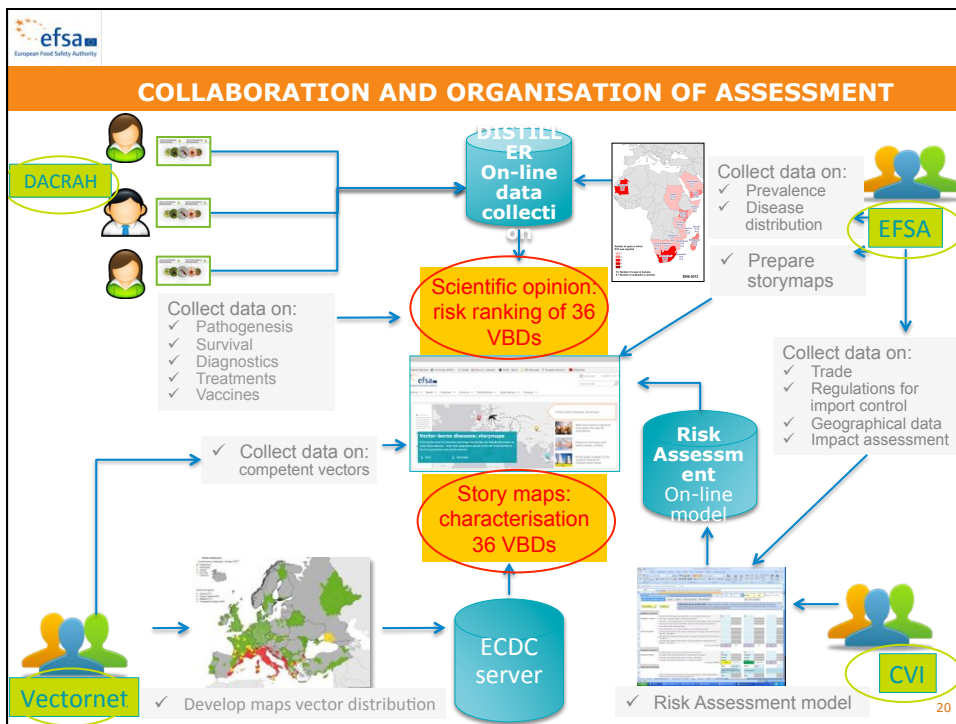
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**■ EXAMPLE: Mosquito borne diseases**

18

**Characterisation of mosquito-borne diseases**

MBD's	Species	Pathogens	Hosts											
			OIE notifiable	Deer	Wild boar	Cattle	Sheep	Goats	Pigs	Horse	Dogs	Birds	Humans	
1	CVV	Bunyamwera virus virus	No	X		x	X	x	x	X				
2	EEEV	Eastern equine encephalitis virus	Yes							X				x
3	GETV	Getah virus	No		X				x	X				
4	HJV	Highlands J. virus	No							X				
5	JEV	Japanese encephalitis virus	Yes		X				x	x				x
6	MIDV	Middelburg virus	No							x				
7	PHSV	Peruvian horse sickness virus	No							x				
8	RVFB	Rift Valley fever virus	Yes	X		x	X	x						x
9	SLEV	St. Louis encephalitis virus	No							x				
10	VEE	Venezuelan equine encephalitis virus	Yes							x				x
11	VSV	Vesicular stomatitis virus*	No	X		x	X	x	x	x				
12	WEEV	Western equine encephalitis virus	Yes							x				X
13	WSLV	Wesselsbron virus	No	X		x	X	x						
14	YUOV	Yunnan orbivirus	No					X	x	x				



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## The project – expected outcome

**Common vector distribution database**

**Support to risk assessments and opinions**

**Filling gaps**

**Improved human and animal health in Europe**


21

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## STORY MAPS OF VBD-MANDATE (2016 – 2017)


Vector-borne disease map journals

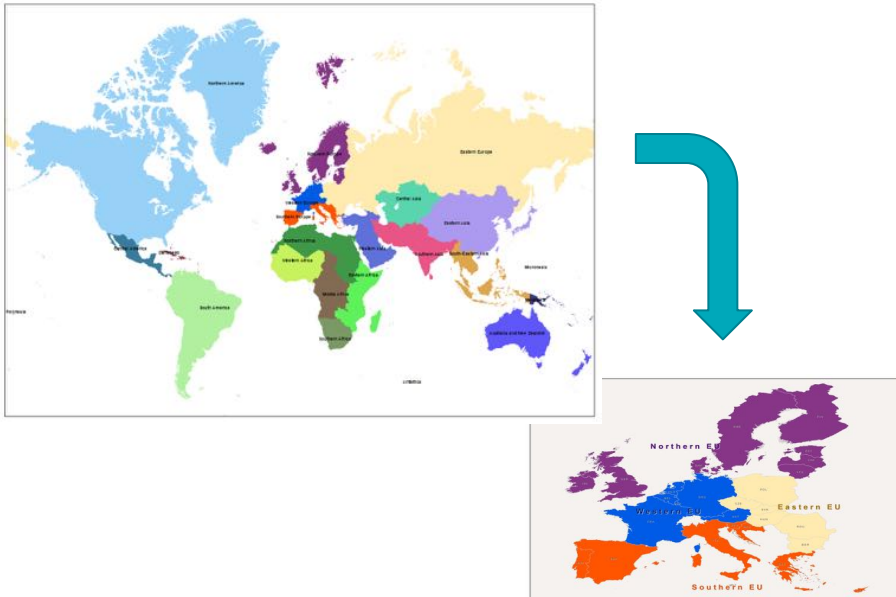
EFSA's vector-borne disease map journals

 **EXAMPLES OF THE STORY MAP ONLINE**

**Japanese encephalitis (JEV)**



 **Global occurrence**



The world map shows JEV occurrence in North America, Europe, Asia, and Australia. The zoomed-in map of Europe is divided into Northern EU, Western EU, Eastern EU, and Southern EU.

24

**Global occurrence**

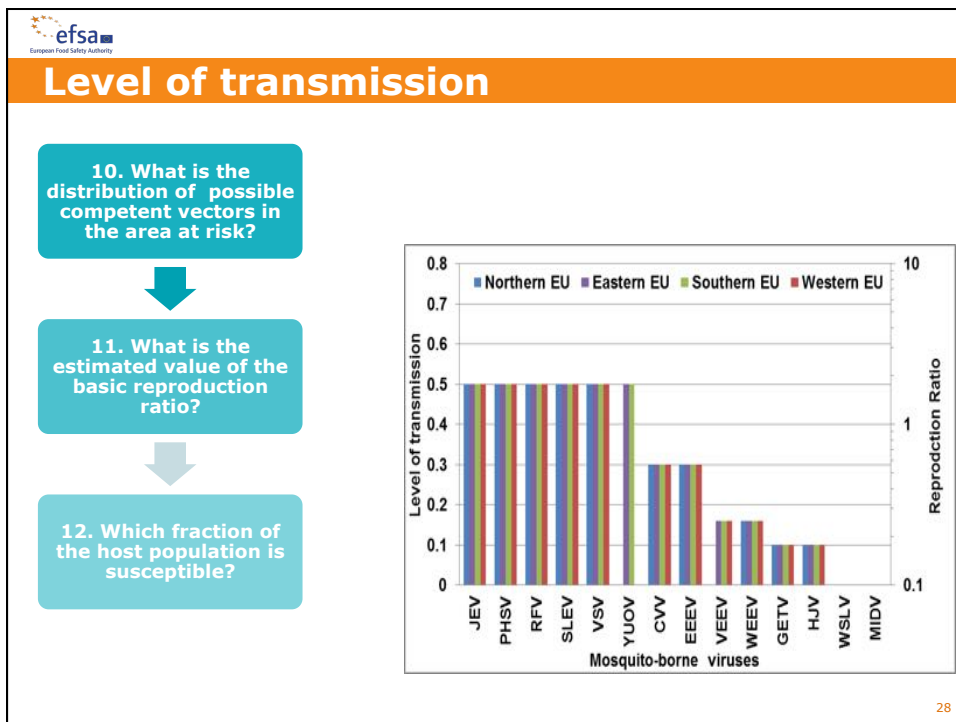
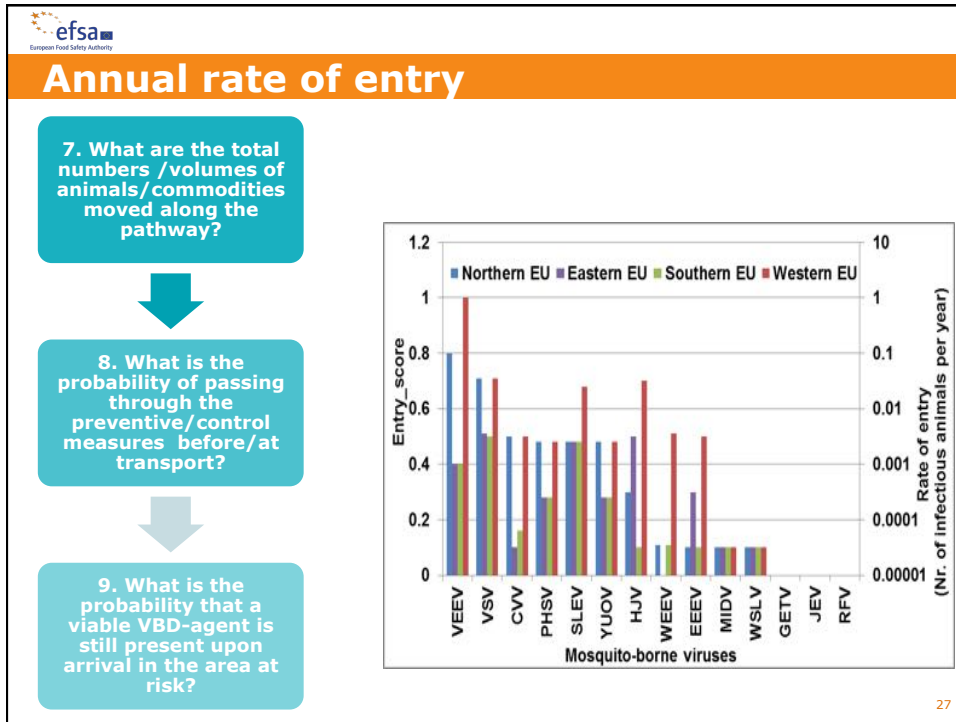
1. What is the geographic distribution of the VBD in the world?
2. How likely is it that the VBD is notified to the OIE?
3. What is the relative size of the infected area?
4. What is the duration of undetected spread?
5. What is the frequency of the epidemics in the infected area?
6. What is the prevalence?

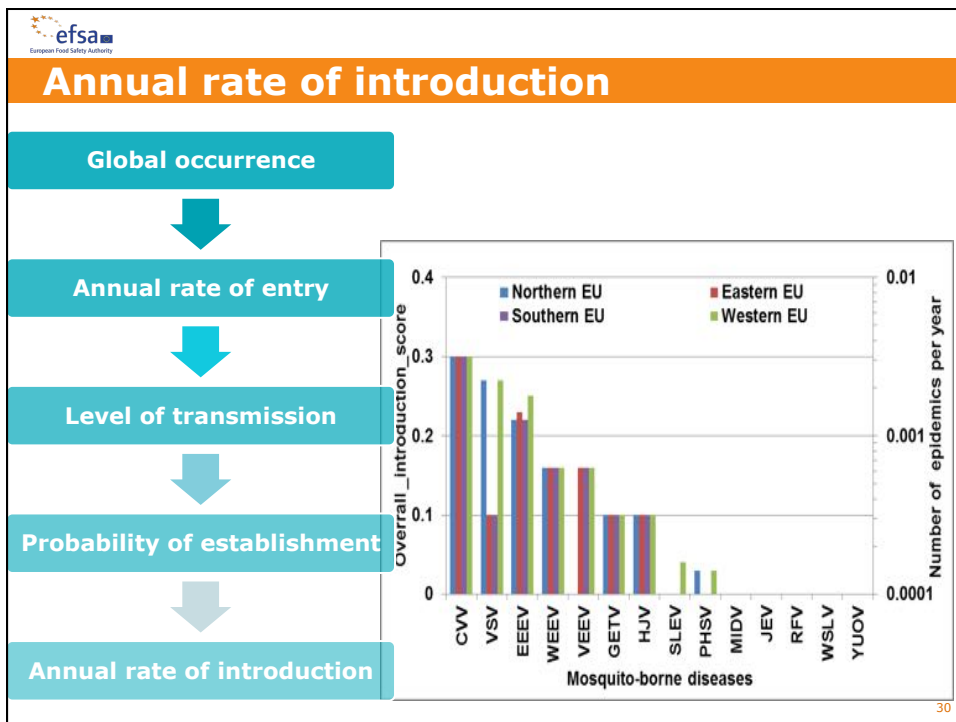
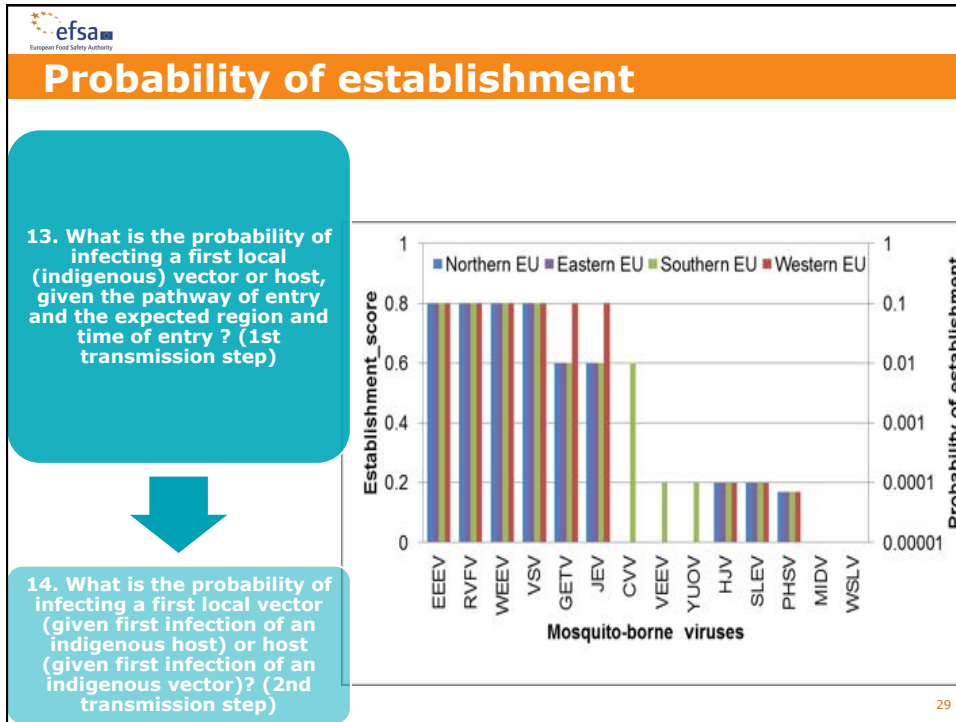
25


**Global occurrence**

Mosquito borne virus	Abbreviation	Australia and New Zealand	Northern America	Central America	Caribbean	South America	Northern Africa	Eastern Africa	Middle Africa	Western Africa	Southern Africa	Western Asia	Eastern Asia	South-Eastern Asia	Southern Asia
Bunyamwera virus	CVV		X	X	X	X									
Eastern equine encephalitis virus	EEEV		X	X		X						X			
Getah virus	GETV												X	X	X
Highlands J. virus	HJV		X												
Japanese encephalitis virus	JEV												X	X	
Middelburg virus	MIDV							X	X	X	X				
Peruvian horse sickness virus	PHSV	X				X									
Rift Valley fever virus	RVSV						X	X	X	X	X	X			
St. Louis encephalitis virus	SLEV		X	X	X	X									
Venezuelan equine encephalitis virus	VEE			X		X									
Vesicular stomatitis virus	VSV		X			X									
Western equine encephalitis virus	WEEV		X	X	X	X									
Wesselsbron virus	WSLV						X	X	X	X				X	
Yunnan orbivirus	YUOV	X			X								X		X

26







  
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## ■ Challenges?

31


  
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## Irreconcilable differences?

<b>Risk managers:</b> <ul style="list-style-type: none"> <li>• No time</li> <li>• Urgent response</li> </ul>	<b>Risk assessment:</b> <ul style="list-style-type: none"> <li>• Need more time</li> <li>• Prevention</li> </ul>
<b>Complex methods</b> <ul style="list-style-type: none"> <li>• Epidemiological models</li> <li>• Systematic reviews</li> <li>• Expert elicitation processes</li> </ul>	<b>Simple methods</b> <ul style="list-style-type: none"> <li>• Rapid tools</li> <li>• Expert opinion</li> </ul>
<b>Big data</b> <ul style="list-style-type: none"> <li>• Animal movements</li> <li>• Populations</li> </ul>	<b>No data</b> <ul style="list-style-type: none"> <li>• Crucial epidemiological parameters (vector capacity?)</li> </ul>

**Preparedness!!**

**Keep assessments up to date: living reports/living opinions**

**Collaboration**

32



 **COLLABORATION: CALL FOR VECTORNET 2 ON-LINE**

**A collaborative approach to data collection activities on vectors and pathogens they transmit**

1. To collect existing published or unpublished information on the geographical distribution of priority vectors;
2. To further develop/maintain Network of medical, veterinary entomologists and public health professionals working in the field of vectors or vector-borne diseases;
3. To deliver ad-hoc scientific advice to support ECDC and EFSA;
4. To carry out targeted entomological surveillance.


<https://etendering.ted.europa.eu/cft/cft-display.html?cftid=4079::> **DEADLINE: 23/10/2018 16:00**

 **Vector control : With what- where -how?**


- Approved products?
- Approved applications?
- Resistance
- Priorities?

**Collaborative one-health approach in EU**







34


 **ACKNOWLEDGEMENTS**

- **AHAW Panel:** More S. (chair), Bicout D., Bøtner A., Butterworth A., Calistri P., Depner K., Edwards S., Garin-Bastuji B., Good M., Schmidt C.G. , Michel V., Miranda M.A., Saxmose N.S., Raj M., Sihvonen L., Spoolder H., Stegeman A., Thulke H.H., Velarde A., Willeberg P., Winckler C.
- **VBD working group:** Stegeman A. (chair), Bicout D., Aline de Koeijer, Miranda M.A., Thulke H.H.
- **EFSA staff:** Dhollander S. (coordinator), Bau A., Beck B.B., Carnesecchi E., Georgiadis M., Casier P. Czwienczek E., Gogin A., Lima E., Matteucci F., Pasinato L. , Richardson J., Riolo F., Rossi G. and Watts M.

35

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36