

<https://helda.helsinki.fi>

Experimental transmission of Zika virus by mosquitoes from central Europe

Heitmann, A.

2017-01-12

Heitmann , A , Jansen , S , Luehken , R , Leggewie , M , Badusche , M , Pluskota , B , Becker , N , Vapalahti , O , Schmidt-Chanasit , J & Tannich , E 2017 , ' Experimental transmission of Zika virus by mosquitoes from central Europe ' , Eurosurveillance , vol. 22 , no. 2 , 30437 , pp. 4-6 . <https://doi.org/10.2807/1560-7917.ES.2017.22.2.30437>

<http://hdl.handle.net/10138/177038>

<https://doi.org/10.2807/1560-7917.ES.2017.22.2.30437>

cc_by

publishedVersion

Downloaded from Helda, University of Helsinki institutional repository.

This is an electronic reprint of the original article.

This reprint may differ from the original in pagination and typographic detail.

Please cite the original version.

Experimental transmission of Zika virus by mosquitoes from central Europe

A Heitmann^{1,2}, S Jansen^{1,2,3}, R Lühken¹, M Leggewie^{1,3}, M Badusche¹, B Pluskota⁴, N Becker^{4,5}, O Vapalahti⁶, J Schmidt-Chanasit^{1,3}, E Tannich^{1,3}

1. Bernhard Nocht Institute for Tropical Medicine, Hamburg, Germany

2. These authors contributed equally to this work

3. German Centre for Infection Research (DZIF), partner site Hamburg-Luebeck-Borstel, Hamburg, Germany

4. Institute for Dipterology (IfD), Speyer, Germany

5. University of Heidelberg, Heidelberg, Germany

6. University of Helsinki and Helsinki University Hospital, Helsinki, Finland

Correspondence: Renke Lühken (renkeluhken@gmail.com)

Citation style for this article:

Heitmann A, Jansen S, Lühken R, Leggewie M, Badusche M, Pluskota B, Becker N, Vapalahti O, Schmidt-Chanasit J, Tannich E. Experimental transmission of Zika virus by mosquitoes from central Europe. *Euro Surveill.* 2017;22(2):pii=30437. DOI: <http://dx.doi.org/10.2807/1560-7917.ES.2017.22.2.30437>

Article submitted on 15 December 2016 / accepted on 10 January 2017 / published on 12 January 2017

Mosquitoes collected in Germany in 2016, including *Culex pipiens pipiens* biotype *pipiens*, *Culex torrentium* and *Aedes albopictus*, as well as *Culex pipiens pipiens* biotype *molestus* (in colony since 2011) were experimentally infected with Zika virus (ZIKV) at 18°C or 27°C. None of the *Culex* taxa showed vector competence for ZIKV. In contrast, *Aedes albopictus* were susceptible for ZIKV but only at 27°C, with transmission rates similar to an *Aedes aegypti* laboratory colony tested in parallel.

In 2015, Zika virus (ZIKV) emerged in Columbia and Brazil and spread rapidly across the American continent and the Caribbean, causing an epidemic with notable numbers of associated clinical cases of microcephaly and Guillain–Barré syndrome [1]. Mosquitoes of the species *Aedes aegypti* and *Ae. albopictus* are considered the primary and secondary vectors of ZIKV [2]. However, with transmission rates below 50%, their vector competence for ZIKV in the laboratory is low [3]. The question therefore remains whether other common mosquito species such as *Culex* spp. play a role in the transmission cycle of ZIKV. The few studies performed so far have provided inconclusive results and suggested that at least *Culex quinquefasciatus* might be able to transmit ZIKV [4–9]. In addition, for an assessment of the risk of possible spread to regions with temperate climate such as central Europe, information is lacking on ZIKV vector competence of mosquitoes under reduced temperature conditions (< 20°C).

This study aimed to evaluate the vector competence of central European mosquito species for ZIKV. Therefore, German populations of *Culex pipiens pipiens* biotype *pipiens* (*Cx. p. pipiens*), *Culex pipiens pipiens* biotype *molestus* (*Cx. p. molestus*), *Culex torrentium* and *Ae. albopictus* (*Ae. albopictus*, GER) were experimentally

infected with ZIKV, using *Ae. aegypti* and an Italian *Ae. albopictus* (*Ae. albopictus*, ITA) as positive controls.

Experimental infection of mosquitoes

Two long-established laboratory strains (*Ae. aegypti* (Bayer company) and *Cx. p. molestus* (in colony since 2011, collected in Heidelberg, Germany)) and four species collected in summer 2016 (*Cx. p. pipiens* Fo (collected in Hamburg, Germany), *Culex torrentium* Fo (collected in Hamburg, Germany), *Ae. albopictus* F7 (collected in Freiburg, Germany) and *Ae. albopictus* F7 (collected in Calabria, Italy)) were analysed and maintained as previously described [10,11]. All colonies tested negative in pan-flavivirus PCRs [12].

Between 150 and 200 female mosquitoes 4–14 days-old were starved for 24 h before application of infectious blood meals containing ZIKV (strain ZIKV_FB-GWUH-2016, GenBank KU870645, fifth passage) [13] at a final concentration of 107 plaque-forming units (PFU)/mL. Artificial feeding was performed using a Hemotek Feeder (*Aedes* spp.) or by cotton sticks (*Culex* spp.). Engorged females were incubated at 80% humidity at either 18°C or 27°C. Analyses for ZIKV were done 14 and 21 days post infection (dpi) for approximately 35 randomly selected females and twice the number for *Ae. aegypti* at 27°C. For salivation, mosquitoes were anaesthetised and the proboscises were inserted into cropped 10 µL filter tips containing 10 µL phosphate-buffered saline (PBS). After 30 min, tips were removed and saliva-containing PBS was analysed for the presence of infectious virus particles by measuring its cytopathic effect (CPE) on Vero cells within the following 8 days. ZIKV in the supernatant of cytopathic cells was confirmed by qRT-PCR using Real Star Zika Virus RT-PCR Kit (Altona diagnostics, Hamburg, Germany). In addition, bodies of all challenged mosquitoes,

TABLE

Susceptibility and transmission rates of mosquitoes experimentally infected with Zika virus (n = 856)

Mosquito taxa	T in °C	14 days post infection			21 days post infection		
		IR ^a (%)	Mean (SD) log ₁₀ RNA copies/specimen ^b	TR ^c (%)	IR ^a (%)	Mean (SD) log ₁₀ RNA copies/specimen	TR (%)
<i>Aedes aegypti</i>	18	17/31 (55)	4.70 (0.86)	0/17	18/33 (55)	4.33 (0.63)	0/18
	27	31/63 (49)	8.69 (1.60)	14/31 (45)	36/50 (72)	6.82 (1.75)	11/36 (31)
<i>Aedes albopictus</i> , ITA	18	19/30 (63)	4.05 (0.59)	0/19	14/39 (36)	5.52 (0.87)	0/14
	27	22/31 (71)	6.34 (2.14)	4/22 (18)	15/29 (52)	7.41 (2.22)	2/15 (13)
<i>Aedes albopictus</i> , GER	18	4/32 (13)	6.22 (1.25)	0/4	11/32 (34)	6.36 (1.39)	0/11
	27	20/31 (65)	6.78 (2.41)	4/20 (20)	18/34 (53)	8.61 (1.82)	6/18 (33)
<i>Culex p. molestus</i>	18	12/41 (29)	3.40 (0.38)	0/12	2/32 (6)	2.48 (0.29)	0/2
	27	7/29 (24)	3.73 (0.38)	0/7	12/38 (32)	4.02 (0.44)	0/12
<i>Culex p. pipiens</i>	18	16/34 (47)	3.38 (0.40)	0/16	3/32 (9)	3.88 (0.43)	0/3
	27	3/37 (8)	3.13 (0.45)	0/3	0/35 (0)	NA ^d	NA ^d
<i>Culex torrentium</i>	18	11/35 (31)	3.15 (0.47)	0/11	1/38 (3)	3.31 (NA)	0/1
	27	4/36 (11)	3.80 (1.79)	0/4	0/34 (0)	NA ^d	NA ^d

GER: from Germany; IR: infection rate; ITA: from Italy; NA: not available; SD: standard deviation; T: temperature; TR: transmission rate.

^a Infection rate: number of ZIKV-positive mosquito bodies per number of fed females.

^b RNA copies were averaged over all ZIKV-positive mosquito bodies excluding the zeros of ZIKV-negative mosquito bodies.

^c Transmission rate: number of mosquitoes with ZIKV-positive saliva per number of ZIKV-positive mosquito bodies.

^d Not available: Mean viral RNA copies and transmission rate could not be calculated for the species–temperature combinations with no ZIKV-positive bodies.

excluding legs and wings, were analysed for ZIKV RNA by qRT-PCR.

Results

At 14 or 21 dpi, ZIKV RNA was detected in the bodies of all challenged mosquito taxa, with infection rates ranging between 3 and 72% in the species–temperature combinations with ZIKV-positive bodies. Infection rates and virus titres were substantially higher in *Aedes* species, with viral RNA copies ranging from 10² to 10⁴ in *Culex* spp. and from 10⁴ to 10⁹ in *Aedes* spp. (Table).

Virus load was generally higher at elevated incubation temperature (27°C vs 18°C). However, transmission of infectious virus particles as measured by CPE of Vero cells incubated with mosquito saliva was not detected in any of the *Culex* taxa. In contrast, saliva was positive for infectious virus particles in all *Aedes* species, but only at 27°C incubation temperature. Interestingly, transmission rates at 21 dpi were similar in *Ae. aegypti* and *Ae. albopictus* from Germany but were substantially lower in *Ae. albopictus* from southern Italy (30% vs 13%).

Discussion

Culex species from central Europe are known as established vectors, able to transmit numerous viruses including West Nile, Sindbis and Usutu virus [14,15]. The results presented here indicate that the three most common *Culex* taxa in central Europe (*Cx. p. pipiens*, *Cx. p. molestus* and *Cx. torrentium*) do not have vector competence for ZIKV. This is in agreement with results from other parts of the world including Italy [4-7,9], which all showed a low degree of competence of the *Cx. pipiens* complex for ZIKV transmission.

The invasive mosquito *Ae. albopictus* is established in large parts around the Mediterranean Sea and is considered to be the main vector in Europe for autochthonous human infections with chikungunya and dengue virus [16]. *Aedes albopictus* are regularly introduced into Germany as accidental cargo via road traffic from southern Europe [17]. In the winter 2015/16, successful overwintering of the species was observed for the first time in southern Germany [18]. The results presented here indicate that specimens of this overwintering population have considerable susceptibility to ZIKV, although only at elevated temperature of 27°C.

Moreover, the transmission rate in this overwintering population was substantially higher than in *Ae. albopictus* from the Calabrian region in southern Italy. Whether the difference in virus susceptibility between German and Italian *Ae. albopictus* populations is due to an ongoing process of adaptation to a new environment or to experimental conditions remains to be determined. Nevertheless, the susceptibility of European *Ae. albopictus* to ZIKV demonstrates the risk of arbovirus transmission associated with the establishment and ongoing spread of this invasive mosquito species in Europe. Of note, none of the tested *Aedes* populations were susceptible to ZIKV at 18°C, which may limit the spread of ZIKV in central Europe to short summer periods with high temperatures. However, for a comprehensive risk assessment of ZIKV transmission in central Europe, further infection studies are needed at intermediate temperatures (e.g. 21°C and 24°C) as well as with other common *Aedes* species such as *Ae. vexans* or the newly established *Ae. japonicus* [19].

Acknowledgements

We thank Ella Weinert and Michelle Helms for excellent technical assistance and Jessica Börstler for analyses of Culex mosquitoes. ML was supported by the Leibniz Association, grant number SAW-2014-SGN-3. This work was financially supported by the German Federal Ministry of Food and Agriculture (BMEL) through the Federal Office for Agriculture and Food (BLE) with the grant number 28-1-91.048-15.

Conflict of interest

None declared.

Authors' contributions

Conceived and designed the study: AH, SJ, RL, JSC, ET. Performed the data collection: AH, SJ, ML. Analysed the data: AH, SJ, RL, JSC, ET. Provided the ZIKA virus strain: OV. Provided mosquito specimens: MB, BP, NB. Wrote the paper: AH, SJ, RL, ET. Contributed to the manuscript drafting: ML. All authors read and approved the final version of the manuscript.

References

- Musso D, Gubler DJ. Zika Virus. *Clin Microbiol Rev.* 2016;29(3):487-524. DOI: 10.1128/CMR.00072-15 PMID: 27029595
- World Health Organization Regional Office for Europe (WHO/Europe). Zika virus technical report. Interim risk assessment WHO European Region. Copenhagen: WHO/Europe; 2016. Available from: <http://www.euro.who.int/en/health-topics/emergencies/zika-virus/technical-reports-and-guidelines-on-zika-virus/zika-virus-technical-report.-interim-risk-assessment-for-who-european-region>
- Chouin-Carneiro T, Vega-Rua A, Vazeille M, Yebakima A, Girod R, Goindin D, et al. Differential susceptibilities of *Aedes aegypti* and *Aedes albopictus* from the Americas to Zika virus. *PLoS Negl Trop Dis.* 2016;10(3):e0004543.
- Aliota MT, Peinado SA, Osorio JE, Bartholomay LC. *Culex pipiens* and *Aedes triseriatus* Mosquito susceptibility to Zika virus. *Emerg Infect Dis.* 2016;22(10):1857-9. DOI: 10.3201/eid2210.161082 PMID: 27434194
- Amraoui F, Atyame-Nten C, Vega-Rúa A, Lourenço-de-Oliveira R, Vazeille M, Failloux AB. *Culex* mosquitoes are experimentally

- unable to transmit Zika virus. *Euro Surveill.* 2016;21(35):30333. DOI: 10.2807/1560-7917.ES.2016.21.35.30333 PMID: 27605199
- Boccolini D, Toma L, Di Luca M, Severini F, Romi R, Remoli ME, et al. Experimental investigation of the susceptibility of Italian *Culex pipiens* mosquitoes to Zika virus infection. *Euro Surveill.* 2016;21(35):30328. DOI: 10.2807/1560-7917.ES.2016.21.35.30328 PMID: 27605056
 - Fernandes RS, Campos SS, Ferreira-de-Brito A, de Miranda RM, Barbosa da Silva KA, de Castro MG, et al. *Culex quinquefasciatus* from Rio de Janeiro Is Not Competent to Transmit the Local Zika Virus. *PLoS Negl Trop Dis.* 2016;10(9):e0004993.
 - Guo XX, Li CX, Deng YQ, Xing D, Liu QM, Wu Q, et al. *Culex pipiens quinquefasciatus*: a potential vector to transmit Zika virus. *Emerg Microbes Infect.* 2016;5(9):e102. DOI: 10.1038/emi.2016.102 PMID: 27599470
 - Huang Y-JS, Ayers VB, Lyons AC, Unlu I, Alto BW, Cohnstaedt LW, et al. *Culex* species mosquitoes and Zika virus. *Vector Borne Zoonotic Dis.* 2016;16(10):673-6. DOI: 10.1089/vbz.2016.2058 PMID: 27556838
 - Vogels CBF, van de Peppel LJJ, van Vliet AJH, Westenberg M, Ibañez-Justicia A, Stroo A, et al. Winter activity and aboveground hybridization between the two biotypes of the West Nile virus vector *Culex pipiens*. *Vector Borne Zoonotic Dis.* 2015;15(10):619-26. DOI: 10.1089/vbz.2015.1820 PMID: 26394124
 - Leggewie M, Badusche M, Rudolf M, Jansen S, Börstler J, Krumkamp R, et al. *Culex pipiens* and *Culex torrentium* populations from Central Europe are susceptible to West Nile virus infection. *One Health.* 2016;2:88-94. DOI: 10.1016/j.onehlt.2016.04.001
 - Chao D-Y, Davis BS, Chang G-JJ. Development of multiplex real-time reverse transcriptase PCR assays for detecting eight medically important flaviviruses in mosquitoes. *J Clin Microbiol.* 2007;45(2):584-9. DOI: 10.1128/JCM.00842-06 PMID: 17108075
 - Driggers RW, Ho C-Y, Korhonen EM, Kuivanen S, Jääskeläinen AJ, Smura T, et al. Zika virus infection with prolonged maternal viremia and fetal brain abnormalities. *N Engl J Med.* 2016;374(22):2142-51. DOI: 10.1056/NEJMoa1601824 PMID: 27028667
 - Jöst H, Bürck-Kammerer S, Hütter G, Lattwein E, Lederer S, Litzba N, et al. Medical importance of Sindbis virus in south-west Germany. *J Clin Virol.* 2011;52(3):278-9. DOI: 10.1016/j.jcv.2011.08.002 PMID: 21893428
 - Nikolay B. A review of West Nile and Usutu virus co-circulation in Europe: how much do transmission cycles overlap? *Trans R Soc Trop Med Hyg.* 2015;109(10):609-18. DOI: 10.1093/trstmh/trv066 PMID: 26286946
 - Medlock JM, Hansford KM, Schaffner F, Versteirt V, Hendrickx G, Zeller H, et al. A review of the invasive mosquitoes in Europe: ecology, public health risks, and control options. *Vector Borne Zoonotic Dis.* 2012;12(6):435-47. DOI: 10.1089/vbz.2011.0814 PMID: 22448724
 - Becker N, Geier M, Balczun C, Bradersen U, Huber K, Kiel E, et al. Repeated introduction of *Aedes albopictus* into Germany, July to October 2012. *Parasitol Res.* 2013;112(4):1787-90. DOI: 10.1007/s00436-012-3230-1 PMID: 23242268
 - Pluskota B, Jöst A, Augsten X, Stelzner L, Ferstl I, Becker N. Successful overwintering of *Aedes albopictus* in Germany. *Parasitol Res.* 2016;115(8):3245-7. DOI: 10.1007/s00436-016-5078-2 PMID: 27112761
 - Huber K, Schuldt K, Rudolf M, Marklewitz M, Fonseca DM, Kaufmann C, et al. Distribution and genetic structure of *Aedes japonicus japonicus* populations (Diptera: Culicidae) in Germany. *Parasitol Res.* 2014;113(9):3201-10. 25056941 DOI: 10.1007/s00436-014-4000-z

License and copyright

This is an open-access article distributed under the terms of the Creative Commons Attribution (CC BY 4.0) Licence. You may share and adapt the material, but must give appropriate credit to the source, provide a link to the licence, and indicate if changes were made.

This article is copyright of the authors, 2017.