



# Transcriptomic Analyses Reveal Differential Gene Expression of Immune and Cell Death Pathways in the Brains of Mice Infected with West Nile Virus and Chikungunya Virus

Stephanie M. Lim<sup>1</sup>, Henk-Jan van den Ham<sup>2</sup>, Minoushka Oduber<sup>2</sup>, Eurydice Martina<sup>1</sup>, Fatiha Zaaraoui-Boutahar<sup>2</sup>, Jeroen M. Roose<sup>1</sup>, Wilfred F. J. van IJcken<sup>3</sup>, Albert D. M. E. Osterhaus<sup>1,4</sup>, Arno C. Andeweg<sup>2</sup>, Penelope Koraka<sup>2</sup> and Byron E. E. Martina<sup>1,2\*</sup>

## OPEN ACCESS

### Edited by:

Dirk Dittmer,  
University of North Carolina at Chapel  
Hill, United States

### Reviewed by:

Helen Lazear,  
University of North Carolina at Chapel  
Hill, United States  
Sara Louise Cosby,  
Queen's University Belfast, Ireland

### \*Correspondence:

Byron E. E. Martina  
b.martina@artemisononehealth.com

### Specialty section:

This article was submitted to  
Virology,  
a section of the journal  
Frontiers in Microbiology

**Received:** 27 April 2017

**Accepted:** 02 August 2017

**Published:** 17 August 2017

### Citation:

Lim SM, van den Ham H-J, Oduber M,  
Martina E, Zaaraoui-Boutahar F,  
Roose JM, van IJcken WFJ,  
Osterhaus ADME, Andeweg AC,  
Koraka P and Martina BEE (2017)  
Transcriptomic Analyses Reveal  
Differential Gene Expression of  
Immune and Cell Death Pathways in  
the Brains of Mice Infected with West  
Nile Virus and Chikungunya Virus.  
*Front. Microbiol.* 8:1556.  
doi: 10.3389/fmicb.2017.01556

<sup>1</sup> Artemis One Health Research Foundation, Delft, Netherlands, <sup>2</sup> Department of Viroscience, Erasmus Medical Center, Rotterdam, Netherlands, <sup>3</sup> Center for Biomix, Erasmus Medical Center, Rotterdam, Netherlands, <sup>4</sup> Research Center for Emerging Infections and Zoonoses (RIZ), University of Veterinary Medicine, Hannover, Germany

West Nile virus (WNV) and chikungunya virus (CHIKV) are arboviruses that are constantly (re-)emerging and expanding their territory. Both viruses often cause a mild form of disease, but severe forms of the disease can consist of neurological symptoms, most often observed in the elderly and young children, respectively, for which the mechanisms are poorly understood. To further elucidate the mechanisms responsible for end-stage WNV and CHIKV neuroinvasive disease, we used transcriptomics to compare the induction of effector pathways in the brain during the early and late stage of disease in young mice. In addition to the more commonly described cell death pathways such as apoptosis and autophagy, we also found evidence for the differential expression of pyroptosis and necroptosis cell death markers during both WNV and CHIKV neuroinvasive disease. In contrast, no evidence of cell dysfunction was observed, indicating that cell death may be the most important mechanism of disease. Interestingly, there was overlap when comparing immune markers involved in neuroinvasive disease to those seen in neurodegenerative diseases. Nonetheless, further validation studies are needed to determine the activation and involvement of these effector pathways at the end stage of disease. Furthermore, evidence for a strong inflammatory response was found in mice infected with WNV and CHIKV. The transcriptomics profile measured in mice with WNV and CHIKV neuroinvasive disease in our study showed strong overlap with the mRNA profile described in the literature for other viral neuroinvasive diseases. More studies are warranted to decipher the role of cell inflammation and cell death in viral neuroinvasive disease and whether common mechanisms are active in both neurodegenerative and brain infectious diseases.

**Keywords:** transcriptomics, Genomics, West Nile virus, chikungunya virus, neuroinvasive disease, cell death mechanisms, immune response