





# Final stakeholder meeting of the







# IDAMS and DENFREE Consortium in collaboration with WHO-TDR at Institut Pasteur, Paris 21-22 November 2016 Amphi Duclaux

Free Registration deadline: November 15, 2017

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The IDAMS and DENFREE consortium in collaboration with WHO-TDR are pleased to welcome you to our final stakeholder meeting, which will take place at the Institut Pasteur, Paris on the 21-22 November 2016. This joint meeting is part of a common dissemination strategy of progress made in our comprehension of dengue transmission, protective immunity and disease pathogenesis, and in the development of innovative tools to improve disease surveillance, prediction of severe disease, vaccines and mosquito control.

### **BACKGROUND**

DENFREE: Our main focus was on finding key factors determining dengue transmission and dynamics in order to develop new tools and strategies for controlling dengue transmission. We estimated the risk of spreading DENV to uninfected areas, especially in Southern Europe where susceptible vectors exist. The major tools generated are predictive models that enable specific interventions, whether concerning the environment, mosquito or human, to be made and that can undermine an epidemic. We aimed to develop an easy-to-use point of care diagnostic tool that is sensitive to detect virus in both human and mosquito samples, which improved DENV surveillance. Inherent in this approach was the belief that improved surveillance and diagnosis of the asymptomatic dengue carriers contributed to effective intervention, especially during early stages of pathogen invasion into a naïve region.

IDAMS: The overall concept of the IDAMS research project was to assemble a consortium of international experts working together to develop new and innovative tools to be applied to the control of dengue in a global context. The core of the work focussed on:

- improving diagnosis and clinical management of dengue through two linked work packages designed **a**) to identify readily available clinical and laboratory parameters and/or viral and immunological markers, that differentiate between dengue and other common febrile illness within 3 days of fever onset, and **b**) to identify any of the available markers that are predictive of the likelihood of evolving to a more severe disease course
- assessing the risk of dengue spread though linked work packages focused on **a**) mapping and modelling techniques to define the current extent of dengue disease globally and to evaluate possible scenarios of spread or risk to previously uninfected regions in the future, and **b**) developing effective and affordable early warning and outbreak response systems.

## OVERALL OBJECTIVE (AIM) AND SPECIFIC OBJECTIVES OF THE MEETING

### 1. Aim:

To identify best practices in dengue risk assessment, management and surveillance as a contribution to the Global Strategy for Dengue Prevention and Control

# 2. Objective:

- 1. To review, collate and summarize outputs of IDAMS and DENFREE leading to:
- Evidence supporting policy to be passed onto global & country policy-makers
- Relevant implementation/operational research needs to facilitate the deployment and scaleup of evidence-supported measures
- Gap in knowledge that have not been filled or have been highlighted by the 2 projects
- 2. To discuss the evidence generated through the IDAMS and DENFREE Consortia regarding: a) Burden of disease; b) Dengue surveillance including dengue alert for outbreaks and early response; c) Clinical and laboratory warning signs for severe dengue; d) Viral and immunological aspects; e) Vector surveillance and control.
- 3. To take notice of the current spread of other viral diseases transmitted by the same vector and identify further priority areas for research on dengue and Zika.
- 4. To develop, in a participatory approach with key stakeholders (particularly Ministries of Health, International Organizations such as WHO, RCCC and national dengue control services), recommendations for the adaptation of the global dengue guidelines according to the new evidence generated.

DAY 1 21st November		Speaker / Chair	
8.30-9.00	Registration		
9.00-9.30	Welcome, Presentation of participants Opening: Purpose of the meeting	DENFREE, IDAMS, TDR, EU officer, IP director	
Dengue burden IDAMS WP4, DENFREE WP1			
9.30-9.50	Role of evaluating and monitoring dengue burden in countries and WHO Regions	R. Velayudhan, WHO	
9.50-10.10	Global dengue burden re-assessed	S. Hay / O. Brady	
10.10-10.30	Coffee Break		
10.30-10.50	Asymptomatic infection epidemiology	C. Fortas	
10.50-11.10	Economic assessment of dengue burden	A. Farlow	
11.10-11.30	Wrap up discussion and use of data		
Dengue case definition, dengue diagnostics, prediction of severe disease IDAMS WP1 & WP2			
11.30-12.00	Overall and diagnostic profile of the IDAMS observational prospective study	T. Jaenisch, B. Wills, C. Simmons, M. Guzman	
12.00-12.30	Clinical case definition for dengue & differentiation from other febrile diseases	B. Wills, T. Jaenisch, P. Lam & K. Rosenberger	
12:30-13:30	Lunch Break		
13:30-14.00	Warning signs for severe dengue – rationale and findings	T. Jänisch, B. Wills	
14.00-14.20	Virological and genetic predictors of severe dengue	C. Simmons, M. Guzman	
14.20-14.50	Wrap up discussion and use of data		
14.50-15.20	Coffee break		
Virology, Immunology and genetics DENFREE WP 2,6,7,8, IDAMS WP2			
15.20-15.40	Human genetic susceptibility to dengue	B. Sierra / L. Pereira	
15.40-16.00	Viral population study in asymptomatic vs symptomatic dengue infection	V. Doung, M. Vignuzzi	
16.00-16.20	Antibody responses to dengue viruses	F. Sallusto and C. Simmons	
16.20-16.40	Highly potent broad spectrum anti-dengue antibodies	A. Rouvinski	
16.40-17.00	Serotyping Dengue virus infections	M. Schreiber	
17.00-17.20	Immunological profile of asymptomatic dengue	T.Cantaert /A.Sakuntabhai	
17.20-17.40	Wrap up discussion and use of data		
19.00	Reception		

Day 2 22 <sup>nd</sup> November		Speaker / Chair	
Mathematical modelling DENFREE WP3,4			
9.00-9.15	Forecast/Prediction dengue epidemic models	Nico Stollenwerk	
9.15-9.30	Zika R0 in South Pacific	Bernard Cazelles	
9.30-9.45	Successful & Unsuccessful examples in dengue prediction: Lessons for chikungunya and Zika outbreaks	Xavier Rodo	
9.45-10.00	Agent based model in Bangkok	E. Daudé – A. Vaguet	
10.00-10.15	Modeling Dengvaxia implementation	M. Aguiar	
10.15-10.30	Dengue in Periodic Environments	J. Rocklov	
10.30-10.45	Wrap up uptake and use of data		
10.45-11.00	Coffee Break		
Outbreak response IDAMS WP3 and WP4			
11.00-11.15	The expansion of dengue vectors	M. Kraemer, O. Brady	
11.15-11.30	Alarm signals for dengue outbreaks and early response	L. Bowman, M. Petzold, A. Kroeger	
11.30-11.45	Country experiences with dengue alarm and early response: Brazil, Mexico, Malaysia,	G. Coelho, G. Tejeda, L. Hakim	
11.45-12.00	Handbook for dengue outbreak detection & response	S. Runge Ranzinger	
12.00-12.15	Wrap up uptake and use of data	3 3	
12.15-13.30	Lunch Break		
Vector control: IDAMS WP4, DENFREE WP 5 and DengueTools			
13.30-13.45	A new tool for community involvement in dengue control	F. Monasso	
13.45-14.00	Implementation of auto-dissemination in Il de Madeira	G. Devine	
14.00-14.15	European vector competence and viral adaptation	P.S. YEN/L. Lambrechts	
14.15-15.00	DengueTools presentation 2	P. Reiter	
15.00-15.15	Wrap up uptake and use of data		
15.15-15.40	Coffee Break		
	Dengue Futures & Zika		
15.40-16.00	Zika	NN, WHO	
16.00-16.20	Dengue Futures and Zika risk mapping	J. Messina	
16.20-16.40	Zika in Latin American IDAMS sites	E. Marques, P. Brasil, A. Tami, NN El Salvador	
16.40-17.00	Harmonisation of tools & protocols for Zika research	Thomas Jaenisch	
17.00–17.20	Zika-Dengue Interactions	J. Mongkolsapaya	
17.20-17.40	Diagnostic challenges of Zika	R. Peeling & C. Simmons	
17.40-18.00	Wrap up discussion, use of data and conclusion		