











A MULTI - MODEL APPROACH TO DESCRIBE THE **CHIKUNGUNYA EPIDEMIC IN ITALY DURING SUMMER 2007**

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AIM OF THE STUDY: investigate the transmission potential of CHIKV in Italy, to provide insight into the possible impact of future outbreaks in temperate climate regions, and the effectiveness of the interventions performed during the outbreak for the epidemic control.

EPIDEMIC: Chikungunya virus (CHIKV), arthropod-borne virus transmitted to humans by Aedes mosquitoes (Reiter et al., 2006); VECTOR: Aedes albopictus;

EPIDEMIC HOTBED: Castiglione di Cervia and Castiglione di Ravenna

(Emilia Romagna region), 3968 inhabitants in a built-up area of about 70 ha

CONFIRMED CASES: 161 laboratory confirmed cases (Rezza et al., 2007);

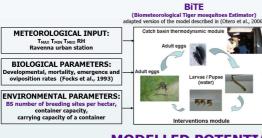
PREVALENCE: estimated 10.2% (Moro et al., 2010), La Reunion Island '05 -'06 epidemic

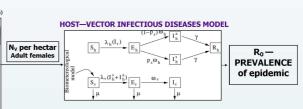
38.2%(Gerardin et al., 2008) and in Mayotte, Indian Ocean, 37.2%(Sissoko et al., 2008):

INDEX CASE: foreigner arrived in Italy from India on June 21, recorded on June 23;

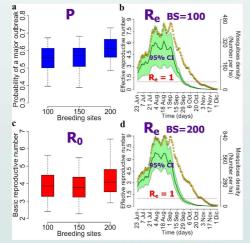
INTERVENTIONS: (Rezza et al., 2007) started at August 23. Breeding sites removal in the entire area, eggs removal (1 day), larvicides and adults removal (3 days).







MODELLED POTENTIAL TRANSMISSION OF CHIKV



BITING RATE K = 0,22 day-1 95% CI 0,15-0,32

fitted using notification data before the interventions and by assuming a number of breeding sites equal to 200 (0,25 day reported in Fonteneille et al., 2007)

MAX RATIO OF MOSQUITOES TO HUMANS $N_V/N_H = 10 - 15$

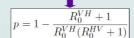
obtained with the mean number Nv estimated through the biometeorological model over a 70 ha area

(20 reported in Fonteneille et al., 2007)



Table 1: Epidemiological and entomological para

 $R_0 = R_0^{HV} R_0^{VH} = k^2 \frac{\overline{N_v}}{N_v} \frac{\chi_h \chi_v}{\chi_h \chi_v} \underline{\quad \omega_v}$ $R_0 = 4.3 95\% \text{ CI } 2.9-6.5$ obtained with the mean number of Nv estimated through the biometeorological $N_h \gamma \mu \omega_v + \mu$



PROBABILITY OF MAJOR OUTBREAK p = 0,61 95% CI 0,49-0,73

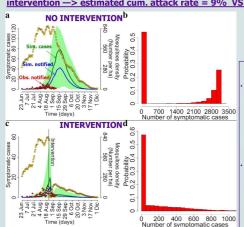
obtained with BS = 200 (almost constant with BS variation)

major outbreaks are possible only for large value of Nv/Nh!!!

MODEL VALIDATION

Cumulative attack rate = symptomatic cases / population at the end of epidemic Prevalence = cum. attack rate / $P_{\rm S}$ symptomatic ratio

-> estimated cum. attack rate = 739 intervention —> estimated cum. attack rate = 9% VS 8,4% (Moro et al., 2010)



192 sim. VS 161 obs. notified cases (Rezza et al., 2007)

DISCUSSION:

Transmission potential of CHIKV in a Italy was similar to the one observed in tropical regions where Chikungunya fever is widespread confirming the high risk of vector borne diseases as a consequence of globalization and global warming .

This analysis strongly support the efficacy of the disinfestation strategy performed during the Italian outbreak, which drastically contributed to reduce the cumulative attack rate (of about 88%).

EFFECTIVENESS OF INTERVENTIONS

INTERVENTIONS (Rezza et al., 2007)	BS removal	Eggs removal	Larvicide	Adulticide
Reference scenario (estimated effects)	-40%	-40%	-90%	-95% for 3 days

Sensitivity analysis through the variation of each single intervention (modelled with BiTE biometeorlogical model), all togheter or separately: A combined strategy can result in a drastic reduction of the epidemic impact!

Adulticides are the most effective single intervention since mosquitoes are in a decaying phase at the time of the

