Supplementary table 1. Detailed summary of all diversification events summarized in Figure 1 as estimated from the BEAST analysis including 84 partial and complete Asian ZIKV genomes

	Thom the DL/151 analysis merading 64 p	<u></u>		8
Node	Diversification event	tMRCA	95% HPD	
A	Malaysia 1966 / other ZIKV	1954–04	1948–08	1959–05
В	India / other ZIKV	1956-10	1949-09	1964-05
C	Philippine–Micronesia / other ZIKV	1997–01	1994-09	1999-03
D	Cambodia / other ZIKV	2001-01	1999-02	2003-12
E	Vietnam / other ZIKV	2004-06	2002-06	2006-04
F	Thailand–Bangladesh / other ZIKV	2005-01	2003-11	2009-05
G	Indonesia / other ZIKV	2006-03	2004-09	2007-06
H	Thailand / other ZIKV	2006-09	2005-05	2007-01
I	Thailand / other ZIKV	2007-10	2006-06	2009-01
J	Thailand–Singapore / other ZIKV	2009-01	2007-12	2010-02
K	Vietnam / other ZIKV	2010-02	2009-01	2011-03
L	Vietnam / other ZIKV	2011–05	2010-01	2012-07
M1	Pacific islands / Pi–American ZIKV	2012-06	2011–12	2013-01
M2	Pacific islands / Pi–American ZIKV	2012-09	2012-01	2013-02
M3	Pacific islands / Pi–American ZIKV	2012-11	2012-07	2013-03
M4	Pacific islands / American ZIKV	2013-01	2012-09	2013-05
N	American ZIKV	2013-03	2012-11	2013-06
	Mean evolutionary rate	7.26x10 ⁻⁴		
	95% HPD evolutionary rate	$6.28 - 8.19 \times 10^{-4}$		