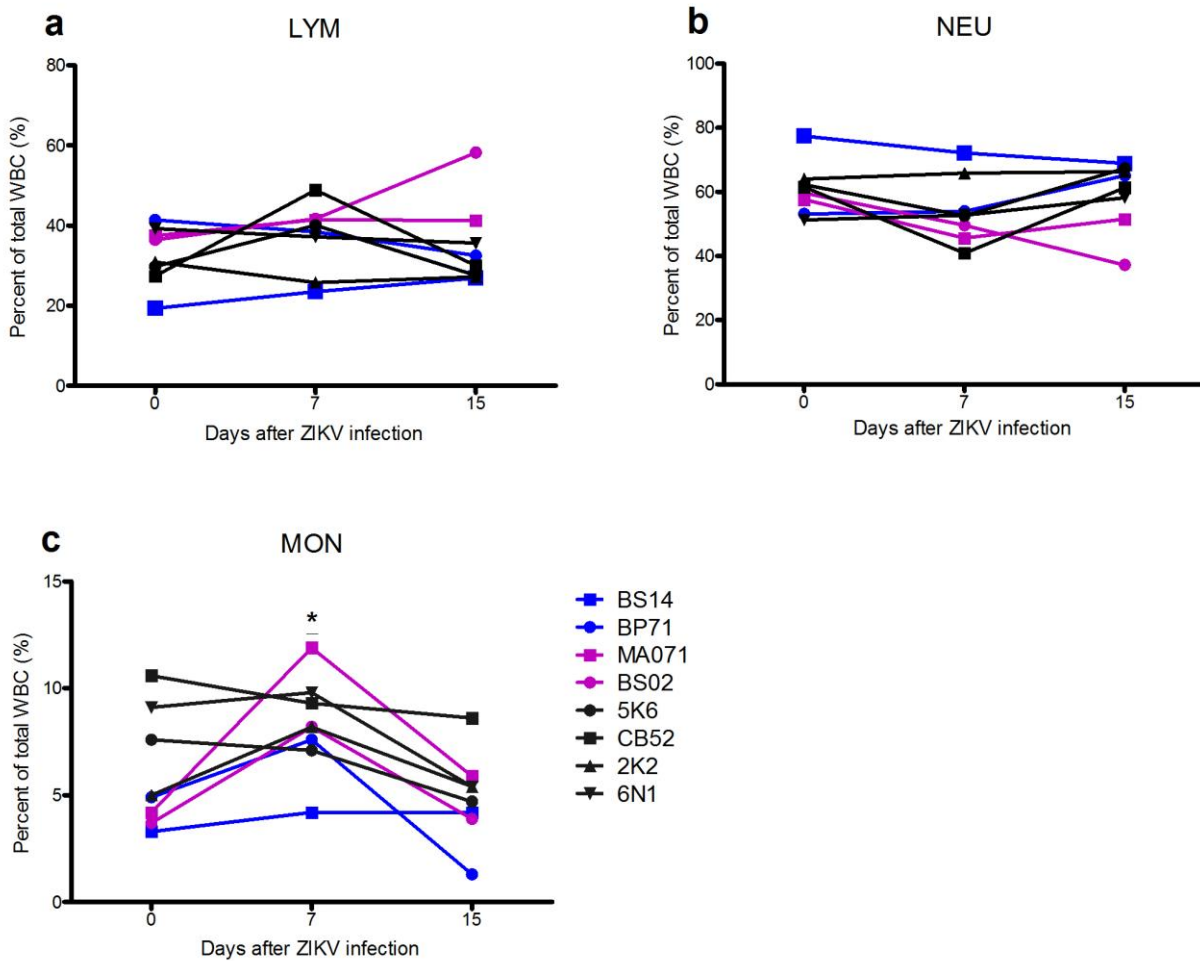
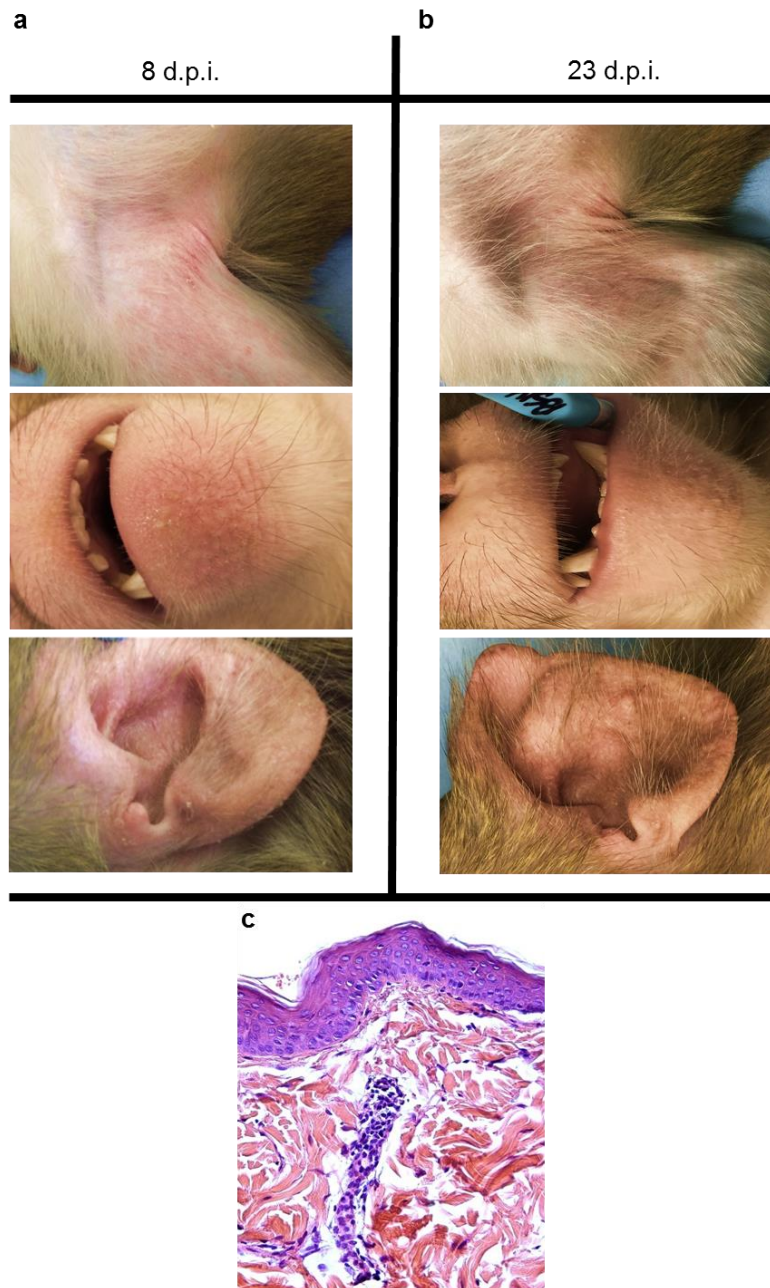


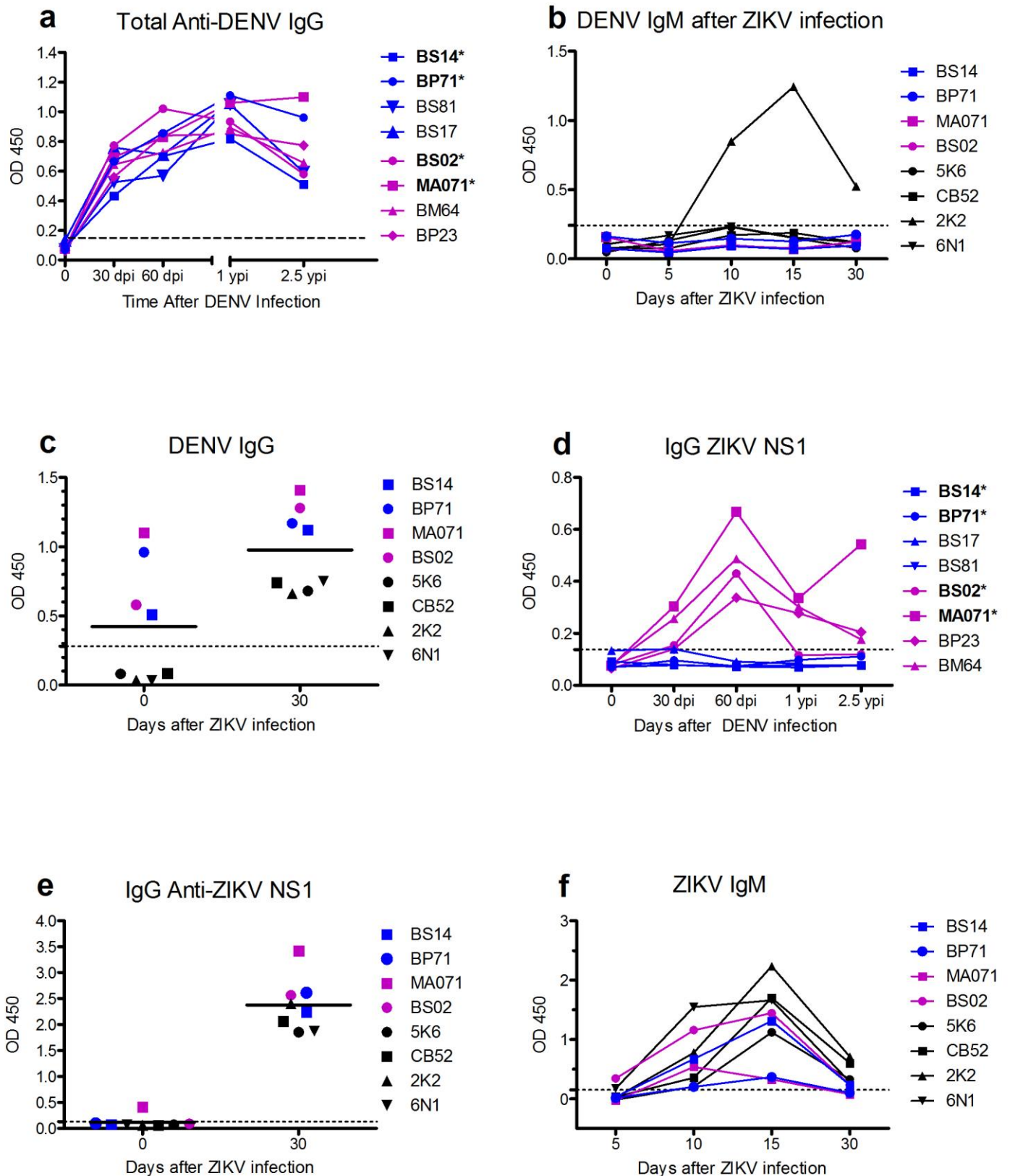
Supplementary Figures



Supplementary Figure 1. Kinetics of cell subsets: (a-c) Cell subsets (LYM: Lymphocytes, NEU: Neutrophils, MON: Monocytes) kinetics obtained from Complete Blood Cell (CBC) counts (% of total WBC) performed at baseline and on days 7 and 15 p.i. are shown in percentage of total cells. Comparison of percentage of MON within cohorts on day 7 p.i. related to their own baseline values was performed using a two-tailed unpaired t test with Sidak-Bonferroni correction ($p < 0.05$). In all panels, DENV-1-pre-exposed macaques are in blue, DENV-2 pre-exposed ones in magenta, and naïve ones in black.

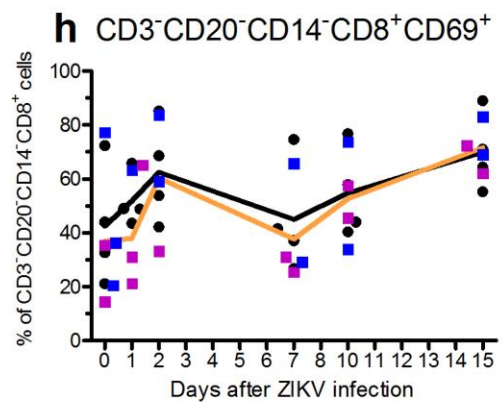
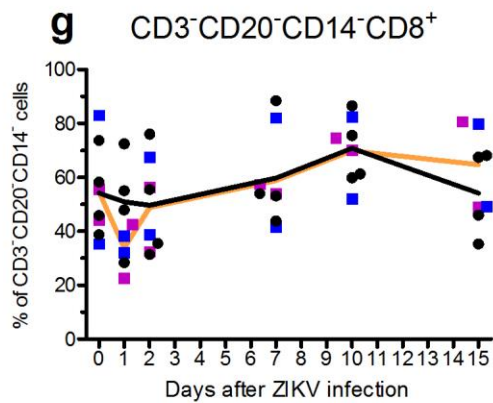
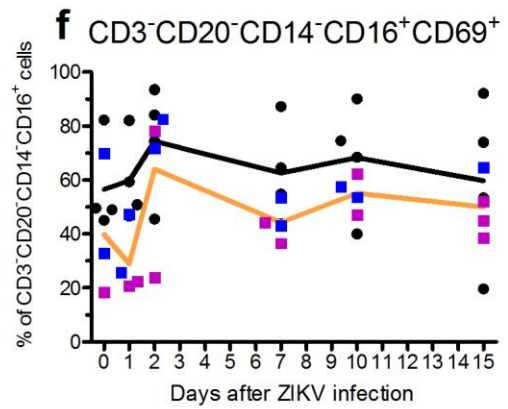
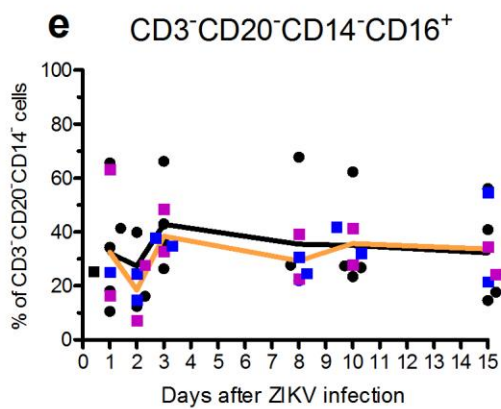
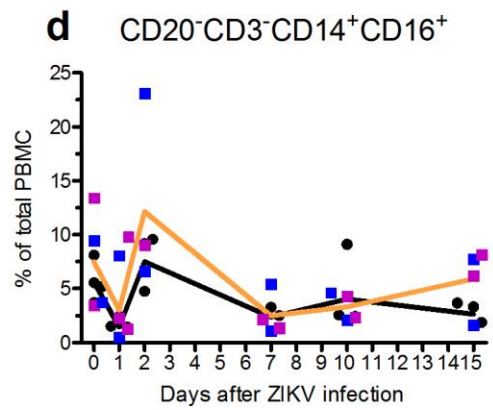
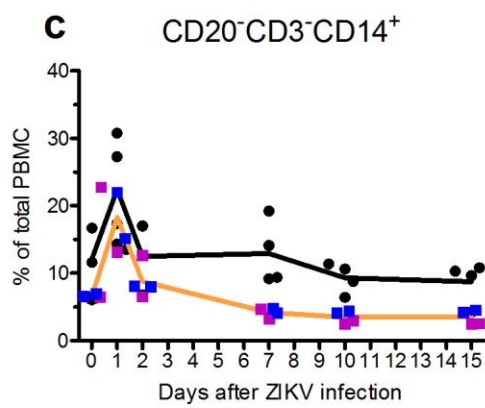
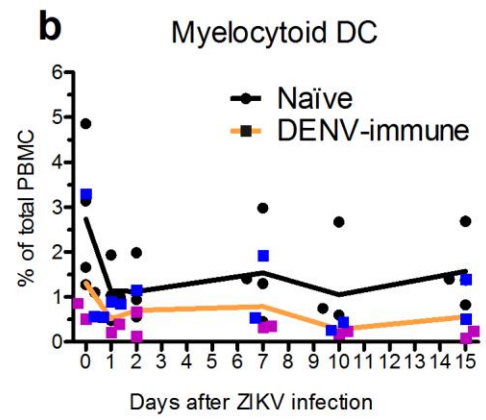
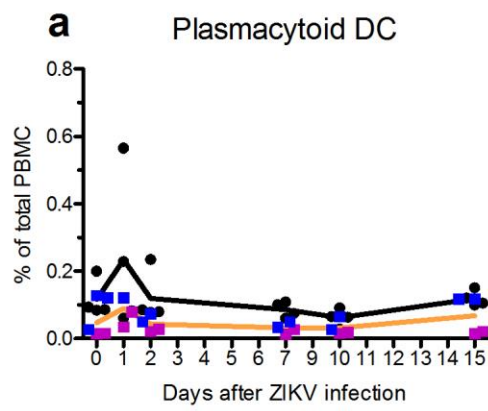


Supplementary Figure 2: Non-pruritic skin rash developed by macaque BS14. BS14 macaque from cohort 1 that was pre-exposed to DENV-1 shows, a) on day 8 p.i. of ZIKV, skin rash in several areas of the body such as axillaries, chin, and ears, and (b) by day 23 p.i., most of the rash was partially resolved in these areas. (c) Hematoxylin and eosin light micrograph of the haired skin of this macaque examined at 40x magnification. Few superficial dermal capillaries exhibit low numbers of perivascular mononuclear inflammatory infiltrate with a predominant population of lymphocytes but lesser plasma cells and histiocytes. Endothelial lining of capillaries is slightly hypertrophied and few dermal capillaries are surrounded by increased space with subtle separation of dermal collagen bundles adjacent to the vessels (edema).

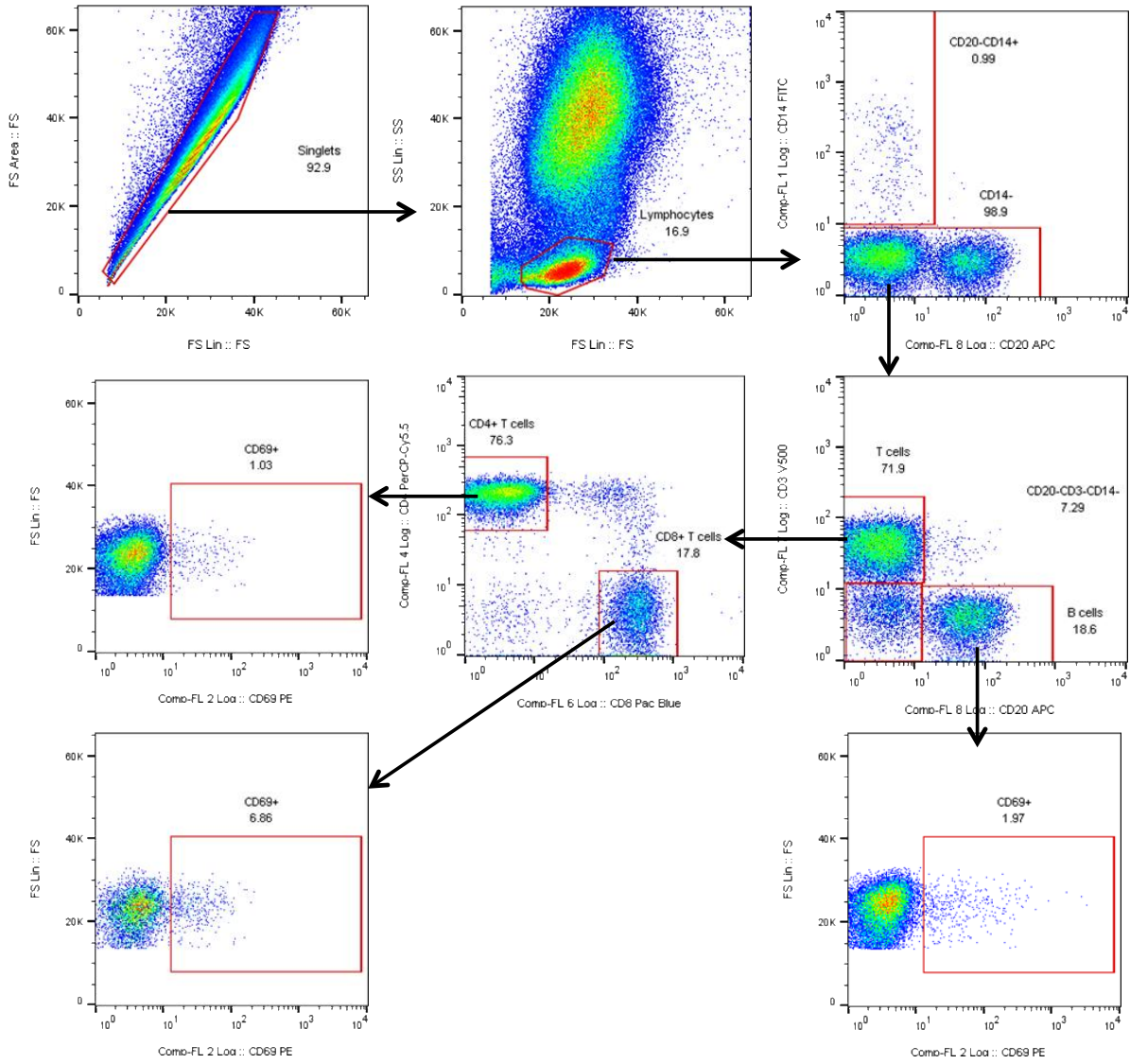


Supplementary Figure 3: Serological profiles of the two cohorts of macaques before and after ZIKV infection. (a) Original cohort of 8 macaques exposed to either DENV-1 (blue, n=4) or to DENV-2 (magenta, n=4) in the year 2013. All eight macaques showed an increase of DENV-IgG with high titers up to 2.5 y.p.i of DENV prior to ZIKV challenge. Two macaques per serotype

(marked with * in panel a) were selected to be challenged with ZIKV. (b) Both naïve and selected DENV-pre-exposed macaques were negative for DENV-IgM previous to ZIKV infection. As shown only one naïve macaque (2K2) developed cross-reacting IgM against DENV, as early as 10 d.p.i. of ZIKV. (c) DENV-naïve macaques were negative for DENV-IgG at baseline and both cohorts developed cross-reacting DENV-IgG after ZIKV infection. (d) Only macaques pre-exposed to DENV-2, but not those exposed to DENV-1, developed cross-reacting IgG to ZIKV-NS1 prior to ZIKV infection. (e) After ZIKV infection both cohorts have detectable IgG to ZIKV NS1 protein. (f) All macaques developed IgM to ZIKV at least by day 5 p.i. of ZIKV. By day 15 p.i., DENV-naïve macaques trend to have higher ZIKV-IgM levels compared to DENV-pre-exposed macaques. By day 30 p.i., IgM levels were back to basal levels.



Supplementary Figure 4: ZIKV does not induce significant changes in the plasmacytoid, myelocytoid dendritic cells, monocytes, and NK cell subsets. (a) Plasmacytoid and (b) Myelocytoid cells were defined as HLA-DR⁺CD3⁻CD14⁻CD20⁻CD66⁻CD123⁺ and CD11c⁺, respectively in PBMCs. No statistical differences were detected. (c) CD20⁻CD3⁻CD14⁺ monocytes, (d) CD20⁻CD3⁻CD14⁺CD16⁺ monocytes, (e) NKCD16⁺ cell subset and (f) its frequency of activation (CD69⁺), and (g) NKCD8⁺ cells subset and (h) its frequency of activation (CD69⁺). Mean values are represented in orange and black lines for DENV-pre-exposed and naïve macaques, respectively. Black circles represent individual naïve macaques, and blue and magenta squares represent individuals previously exposed to either DENV-1 or DENV-2, respectively.



Supplementary Figure 5: Gating strategy for B and T cells and defining subsets. Characterization of T and B cells is shown. Gating for activation marker CD69 is also shown.

Supplementary Tables

Supplementary Table 1. ZIKV RNA detection in urine through 30 days post-infection

RM ID	History	ZIKV RNA in Urine (Days after infection)											Total Days/ Group	
		1	3	5	7	9	15	17	19	21	23	25		30
	Cohort 1													10
BS14	1° DENV1	+++	-	-	+	+	-	-	+	-	-	-	-	
BP71	1° DENV1	+++	-	+	-	-	-	-	-	-	-	-	-	
MA071	1° DENV2	-	-	-	+	+++	-	-	-	-	-	-	-	
BS02	1° DENV2	-	-	-	+	+	-	-	-	-	-	-	-	
	Cohort 2													9
5K6	Naïve	-	+	-	-	-	-	-	-	-	-	-	-	
CB52	Naïve	+	-	-	-	-	-	-	-	-	-	+	-	
2K2	Naïve	-	-	+	-	+++	-	-	+	-	-	-	-	
6N1	Naïve	-	+	+	+++	-	-	-	-	-	-	-	-	

+++ Positive

+ Likely-Positive

- Negative

ZIKV RNA positive urine samples (+++; Ct: ≤ 40.64) using qRT-PCR were detected in both groups within the first 9 d.p.i. Same total of likely-positive samples (+) were detected in both groups (7 samples per group); these samples showed positive-like amplification curves over assay threshold, but their vRNA detection (Ct range: 40.75-41.65) was under the detection limit of the assay (Limited Ct: ≤ 40.64). Negative urine samples (-, no amplification).

Supplementary Table 2 Percent positive stained of unstimulated controls

RM ID	History	Intracellular Staining					
		CD8 ⁽¹⁾ IFN- γ	CD8 TNF- α	CD8 CD107a	CD4 ⁽²⁾ IFN- γ	CD4 TNF- α	CD4 CD107A
Cohort 1							
BS14	1° DENV-1 Day 30	0.20	0.48	1.27	0.11	0.11	0.46
BP71	1° DENV-1 Day 30	0.68	0.62	1.87	0.24	0.46	0.71
MA071	1° DENV-2 Day 30	0.38	0.49	0.34	0.18	0.16	0.21
BS02	1° DENV-2 Day 30	0.28	0.47	0.62	0.10	0.19	0.28
Cohort 2							
5K6	Naïve Day 30	0.19	0.73	0.89	0.10	0.23	1.01
CB52	Naïve Day 30	0.11	0.58	2.10	0.05	0.28	0.72
2K2	Naïve Day 30	0.33	1.40	0.34	0.13	0.69	0.53
6N1	Naïve Day 30	0.20	0.65	0.15	0.12	0.62	0.35
RM ID	History	Intracellular Staining					
		CD8 IFN-γ	CD8 TNF- α	CD8 CD107a	CD4 IFN-γ	CD4 TNF- α	CD4 CD107A
Cohort 1							
BS14	1° DENV-1 Day 60	0.11	0.59	1.17	0.06	0.17	0.33
BP71	1° DENV-1 Day 60	0.23	0.26	1.31	0.01	0.12	0.22
MA071	1° DENV-2 Day 60	0.19	0.29	0.76	0.01	0.19	0.45
BS02	1° DENV-2 Day 60	0.15	0.17	1.09	0.04	0.06	0.12
Cohort 2							
5K6	Naïve Day 60	0.07	0.38	0.76	0.05	0.25	0.53
CB52	Naïve Day 60	0.02	0.11	0.46	0.02	0.04	0.18
2K2	Naïve Day 60	0.06	0.12	0.51	0.02	0.22	0.13
6N1	Naïve Day 60	-	-	-	-	-	-

¹CD8⁺ refers to CD3⁺CD20⁻CD8⁺

²CD4⁺ refers to CD3⁺CD20⁻CD4⁺

Measurement of functional effector response of CD4+ and CD8+ T cells post ZIKV infection'. Antigen-specific CD4+ and CD8+ T cell effector responses were measured 30 d.p.i. of ZIKV to determine if prior DENV exposure impacted the ZIKV-specific functional response from these cells, and measured again after 60 d.p.i. to assess the stability of the functional effector response

Supplementary Table 3 Dengue viremia post challenge in Cohort 1 animals

RM ID	DENV Exposure History	Viremia ^a Days Post-Dengue Challenge (log ₁₀ FFU/ml)										CV ^b log ₁₀ FFU/ml	Average Duration (Days)	
		1	2	3	4	5	6	7	8	9	10			
BS14*	DENV-1	-	2.3	1.2	-	-	-	-	-	-	-	-	3.5	1.8
BP71*	DENV-1	-	2.4	1.4	-	-	-	-	-	-	-	-	3.8	
BS17	DENV-1	-	-	0.9	-	-	-	-	-	-	-	-	0.9	
BS81	DENV-1	-	1.8	1.4	-	-	-	-	-	-	-	-	3.2	
MA071*	DENV-2	1.2	1.8	1.5	1.5	-	-	-	-	-	-	-	6.1	5
BS02*	DENV-2	0.9	1.5	-	0.9	1.2	2	1.8	1.2	-	-	-	9.6	
BP23	DENV-2	-	1.9	2.2	2.5	1.4	-	-	-	-	-	-	7.9	
BM64	DENV-2	-	2.1	1.4	1.5	2.2	1.6	-	-	-	-	-	8.8	

*Animals included in Cohort 1 of this work.

a) Viremia was quantified by immunofocus assay, which measured infectious virus in Vero cells. (-) Indicated below the level of detection of 8 PFU/ml.

b) CV (Cumulative viremia) is defined as the total daily viremia for each animal

Supplementary Table 4

Dengue Viremia post Zika infection

RM ID	DENV Exposure History	Viremia (Qualitative) Days Post-ZIKV Infection									
		1	2	3	4	5	6	7	8	9	10
Cohort 1											
BS14	1° DENV-1	-	-	-	-	-	-	-	-	-	-
BP71	1° DENV-1	-	-	-	-	-	-	-	-	-	-
MA071	1° DENV-2	-	-	-	-	-	-	-	-	-	-
BS02	1° DENV-2	-	-	-	-	-	-	-	-	-	-

For DENV1, DENV-1WP74 RNA was used as a positive control which amplified at a threshold cycle (Ct)= 11.77.

For DENV2, DENV-2 NGC RNA was used as a positive control which amplified at Ct=13.94.
Samples are considered positive when amplification occurs at Ct<36.

Supplementary Table 5 Primers and Probe for Zika RT-PCR

Primer/Probe	Sequence
ZIKV 1086f	5' CCG CTG CCC AAC ACA AG 3'
ZIKV 1162c	5' CCA CTA ACG TTC TTT TGC AGA CAT 3'
ZIKV 1107	FAM 5' AGC CTA CCT TGA CAA GCA GTC AGA CAC TCA A 3' (BHQ-1)

Supplementary Table 6 Antibodies for Cell Phenotyping

Phenotype/activation	Company	Catalog number	Dilution Total ul in 160 ul final volume
CD14 FITC	Beckman-Coulter	6603262	4
CD69 PE	DAKO	R7173	5
CD4 PerCP-Cy5.5	BD-Biosciences	552838	5
CD8 Pac Blue	Fisher-Invitrogen	MHCD0828	2.5
CD3 V500	BD-Biosciences	560770	3
CD20 APC	Biolegend	302310	20
CD16 Alexa700	Biolegend	560713	4
DC control			
CD14 FITC	Beckman-Coulter	6603262	4
CD66abcd	Miltenyi	130-093-132	5
CD20 FITC	Beckman-Coulter	6602381	1
CD8 FITC	Fisher-Invitrogen	MHCD0801	2.5
CD3 FITC	BD-Biosciences	556611	7
Isotype PE	Fisher-Invitrogen	MG104	1.5
HLA-DR PerCP-Cy5.5	BD-Biosciences	552764	5
Isotype APC	BD-Biosciences	554681	1.5
DC tube			
CD14 FITC	Beckman-Coulter	6603262	4
CD66abcd	Miltenyi	130-093-132	5
CD20 FITC	Beckman-Coulter	6602381	1
CD8 FITC	Fisher-Invitrogen	MHCD0801	2.5
CD3 FITC	BD-Biosciences	556611	7
CD11c PE	Biolegend	301606	7.5
HLA-DR PerCP-Cy5.5	BD-Biosciences	552764	5
CD123 APC	BD-Biosciences	560087	10
NK markers			
CD14 FITC	Beckman-Coulter	6603262	4
CD20 FITC	Beckman-Coulter	6602381	1
CD159a (NKG2A) PE	Beckman-Coulter	IM3291U	8
CD337 (NKp30) PC5	Beckman-Coulter	PN A66904	5
CD335 (NKp46) PC7	Beckman-Coulter	PN B38703	3
CD159c (NKG2C) APC*	R&D Systems	MAB1381-100	5
CD16 Alx700	Biolegend	560713	4
CD8 Pacific Blue	Fisher-Invitrogen	MHCD0828	2.5
CD3 V500	BD-Biosciences	560770	5

*In house labeled

Supplementary Table 7 Characterization of fluorochromes antibody panels and cytometer configurations for Natural Killer cells

	Pac Blue	V500	FITC	PE	PerCP-Cy5.5	PE-Cy7	APC	AF700/APC-Cy7
Phenotyping	CD8	CD3	CD14	CD69	CD4	-	CD20	CD16
NK markers	CD8	CD3	CD14, CD20	NKG2A	NKp30	NKp46	NKG2C	CD16
DC control	-	-	CD3, CD8, CD14, CD20, CD66abcd	IgG	HLA-DR	-	IgG	-
DC	-	-	CD3, CD8, CD14, CD20, CD66abcd	CD11c	HLA-DR	-	CD123	-

Cells were defined as CD3⁺CD20⁻CD14⁻ and analyzed for the expression of NK cell markers CD16, CD8, NKG2A, NKG2C, NKp30 and NKp46.

Supplementary Table 8 Antibodies used for cell-immune response assessment

Marker	Stain	Clone	Catalog Number	Vendor	Dilution
CD4	PerCP-Cy-5.5	SK3	566316	BD Biosciences	1:25
CD8 β	PE	ECD	6607123	Beckman-Coulter	1:20
CD3	Pacific Blue	SP34-2	558124	BD Biosciences	1:30
CD20	BV605	2H7	563783	BD Biosciences	1:30
CD107a	FITC	H4A3	555800	BD Biosciences	1:10
CD28	PE-Cy-5	CD28.2	555730	BD Biosciences	1:10
CD95	BV510	DX2	305640	Biolegend	1:30
IFN- γ	APC	B27	554702	BD Biosciences	1:30
TNF- α	PE-Cy-7	Mab11	557647	BD Biosciences	1:30

Supplementary Table 9 Peptides sequences used to stimulate PBMCs 30 and 60 days after Zika infection

Dengue Virus Type 1 Peptides

Peptide	Amino Acid Sequence	Peptide	Amino Acid Sequence	Peptide	Amino Acid Sequence
1	MRCVGIGNRDFV EGLSG	29	TTATITPQA PTSEIQLT	57	KYEGTDAPCKIPFSSQD
2	GNRDFV EGLSGA TWV DV	30	PQAPTSEIQL TDY GALT	58	APCKIPFSSQDEKGVTVQ
3	VEGLSGA TWVDV VLEHG	31	SEQLTDY GALT DCSPR	59	FSSQDEKGVTVQNGRLITA
4	ATWVDV VLEHGSCVTTM	32	YGALTD CSPRTGLDFN	60	KGVTQNGRLITANPIVTD
5	VLEHGSCVTTMAKDKPT	33	DCSPRTGLDFNEMVLLT	61	RLITANPIVTDKEKPVN
6	CVTTMAKDKPTLDIELL	34	GLDFNEMVLLTMEKKS	62	PIVTDKEKPVNIEAE
7	KDKPTLDIELLKTEVTN	35	MVLLTMEKKS WL VHKQW	63	DKEKPVNIEA EPPFGE
8	DIELLKTEV TNPAVLRK	36	EKKS WL VHKQWFLDLPL	64	VNIEA EPPFGESY IVVG
9	TEVTNPAVLRKLCIEAK	37	VHKQWFLDLPLPWTSGA	65	PPFGESY IVVGAGEKAL
10	AVLRKLCIEAKISNTTT	38	LDLPLPWTSGASTSQET	66	YIVVGAGEKALKLSWFK
11	CIEAKISNTTTDSRCPT	39	WTSGASTSQETWNRQDL	67	GEKALKLSWFKKGSSIG
12	SNTTTDSRCPTQGEATL	40	TSQETWNRQDLLVTFKT	68	KLSWFKKGSSIGKMFEA
13	SRCPTQGEATLVEEQDT	41	NRQDLLVTFKTAHAKKQ	69	KGSSIGKMFEA TARGAR
14	GEATLVEEQDTNFVCR	42	VTFKTAHAKKQEVVVLG	70	KMFEA TARGARRMAILG
15	VEEQDTNFVCRRTFVDR	43	HAKKQEVVVLGSQEGAM	71	ARGARRMAILGDTAWDF
16	NFVCRRTFVDRGWGNG	44	VVVLGSQEGAMHTALTG	72	MAILGDTAWDFGSIGGV
17	RTFVDRGWGNGCGLFGK	45	SQEGAMHTALTGATEIQ	73	TAWDFGSIGGVFTSVGK
18	GWGNGCGLFGKGLIT	46	HTALTGATEIQTSGTTT	74	SIGGVFTSVGKLIHQIF
19	CGLFGKGLITCAKFK	47	ATEIQTSGTTTIFAGHL	75	TSVGKLIHQIFGTAYGV
20	KGSLITCAKFKCVTKLE	48	SGTTTIFAGHLKCRKLM	76	IHQIFGTAYGVLFSGV
21	CAKFKCVTKLEKIVQY	49	FAGHLKCRKMDKLT	77	GTAYGVLFSGVSWTMKI
22	VTKLEKIVQYENLKY	50	CRLKMDKLT	78	LFSGVSWTMKIGIGILL
23	GKIVQYENLKYSVIVTV	51	KLTLKGMSYVMCTGSFK	79	WTMKIGIGILLTWLGLN
24	YENLKYSVIVTVHTGDQ	52	MSYVMCTGSFKEKEVA	80	IGILLTWLGLNSRSTSL
25	SVIVTVHTGDQHQVGN	53	TGSFKEKEVA ETQHGT	81	WLGLNSRSTSLSMTCIA
26	HTGDQHQVGNETTEHGT	54	LEKEVA ETQHGT VLVQV	82	RSTSLSMTCIAVGMVTL
27	HQVGNETTEHGTATIT	55	AETQHGT VLVQV KYEGT	83	MTCIAVGMVTLYLGMV
28	TTEHGTATITPQA PT	56	TVLVQV KYEGTDA PCKI	84	GMVTLYLGMV/QA

Supplementary Table 9 Cont

Dengue Virus Type 2 Peptides

Peptide	Amino Acid Sequence	Peptide	Amino Acid Sequence	Peptide	Amino Acid Sequence
1	MRCIGISNRDFV EGV	29	AWLVHRQWFLDLPLPWL	57	MRGAKRMAILGDTA WDF
2	ISNRDFV EGVSGGSWV DI	30	WFLDLPLPWLPGADTQGSNW	58	AILGDTAWDFGSLGGVF
3	GVSGGSWVDIVLEHGSCV	31	PGADTQGSNWQKETLV	59	WDFGSLGGVFTSIGKAL H
4	DIVLEHGSCVTTMAKNK	32	SNWIQKETLVTFKNPHAK	60	VFTSIGKALHQVFGA Y
5	SCVTTMAKNKPTLDFELI	33	LVTFKNPHAKKQDVVVL	61	ALHQVFGA YGAAAFSGV
6	NKPTLDFELIETEAQPA	34	HAKKQDVVVLGSQEGAMH	62	AIYGAAFSGVSWIMKIL I
7	LIETEAQQPATLRKYCI	35	VLGSQEGAMHTAL TGA	63	GVSWIMKILIGV IITWI
8	KQPATLRKYCIEAKL	36	GAMHTALTGATEIQM	64	ILIGV IITWIGMNSR
9	LRKYCIEAKL TTTTDSR	37	ALTGATEIQMSSGNLLF	65	IITWIGMNSRSTSLSVSL
10	KL TTTTDSRCP TQGEPSL	38	IQMSSGNLLFTGHLKCR L	66	SRSTSLSVSLVLVGVVTL
11	RCPTQGEPSLNEEQDKRF	39	LFTGHLKCR L RMDKLQLK	67	SLVLVGVV TLYLGVMVQ A
12	SLNEEQDKRFVCKHSMV	40	RLRMDKLQLK GMSYSM		
13	KRFVCKHSMVDRGWGNG CGL	41	LQLKGMSYSMCTGKFKV V		
14	DRGWGNGCGLFGKGGIV	42	SMCTGKFKV VKEIA ETQH		
15	CGLFGKGGIVTCA MFTCK	43	VVKEIAETQHGTIVIRV		
16	IVTCAMFTCKKNMKGKVV	44	TQHGTIVIRV QY EGDGSPCK		
17	CKKNMKGKVVQPENLEY	45	VQYEGDGSPCKIPFEIM		
18	KVVQPENLEY TIVITPH	46	SPCKIPFEIMDLEKRHVL		
19	LEYTIVITPHSGEEHAV	47	IMDLEKRHVLGRLITV		
20	TPHSGEEHAVGNDTGKH	48	RHVLGRLITV NPV TEK		
21	HAVGNDTGKHGKEIKI	49	ITVNPVTEKDSPVNIEA		
22	TGKHGKEIKITPQSSI	50	EKDSPV NIEA EPPFGDSY		
23	EIKITPQSSITEA ELTGY	51	EA EPPFGDSY IIIIGV		
24	SITEA ELTGYGTVTM	52	FGDSY IIIIGV EPGQLKL		
25	ELTGYGTVTMECSPRTGL	53	IGVEPGQLKL NWFKK		
26	TMECSPRTGLDFNEMVLL	54	GQLKL NWFKKGSSIGQMI		
27	GLDFNEMVLLQMENKAWL	55	KKGSSIGQMIETTMRGAK		
28	LLQMENKAWLVHRQWFL	56	MIETTMRGAKRMA IL		

Supplementary Table 9 Cont

Zika Virus Envelope Peptides

Peptide	Amino Acid Sequence	Peptide	Amino Acid Sequence	Peptide	Amino Acid Sequence
ZIKV59	IRCIQVSNRDFVEGM	ZIKV87	LSVHGSQHSQMIVND	ZIKV115	KGRLSSGHLKCRCLKM
ZIKV60	VSNRDFVEGMSGGTW	ZIKV88	SQHSQMIVNDTGHET	ZIKV116	SGHLKCRCLKMDKRLRL
ZIKV61	FVEGMSGGTWVDVVL	ZIKV89	MIVNDTGHETDENRA	ZIKV117	CRLKMDKLRLLKGVSY
ZIKV62	SGGTWVDVLEHGGC	ZIKV90	TGHETDENRAKVEIT	ZIKV118	DKLRLKGVSYSLCTA
ZIKV63	VVDVLEHGGCVTVMA	ZIKV91	DENRAKVEITPNSPR	ZIKV119	KGVSYSLCTAAFTFT
ZIKV64	EHGGCVTVMAQDKPT	ZIKV92	KVEITPNSPRAEATL	ZIKV120	SLCTAAFTFTKIPAE
ZIKV65	VTVMAQDKPTVDIEL	ZIKV93	PNSPRAEATLGGFGS	ZIKV121	AFTFTKIPAE TLHGT
ZIKV66	QDKPTVDIELVTTTV	ZIKV94	AEATLGGFGSLGLDC	ZIKV122	KIPAE TLHGT V TVEV
ZIKV67	VDIELVTTTVSNMAE	ZIKV95	GGFGSLGLDCEPRTG	ZIKV123	TLHGT V TVEV QYAGT
ZIKV68	VTTTVSNMAEVRSYC	ZIKV96	LGLDCEPRTGLDFSD	ZIKV124	VTVEVQYAGTDGPCK
ZIKV69	SNMAEVRSYCYEASI	ZIKV97	EPRTGLDFSDLYYLT	ZIKV125	QYAGTDGPCKVPAQM
ZIKV70	VRSYCYEASISDMAS	ZIKV98	LDFSDLYYLT MN NKH	ZIKV126	DGPCKVPAQMAVDMQ
ZIKV71	YEASISDMASDSRCP	ZIKV99	LYYLT MN NKH WLVHK	ZIKV127	VPAQMAVDMQ TLTPV
ZIKV72	SDMASDSRCP TQGEA	ZIKV100	MN NKH WLV HKEWFHD	ZIKV128	AVDMQ TLTPV GRLIT
ZIKV73	DSRCP TQGEAYLDKQ	ZIKV101	WLV HKEWFHD I PLPW	ZIKV129	TLTPV GRLIT ANPVI
ZIKV74	TQGEAYLDKQSDTQY	ZIKV102	EWFDI PLPW HAGA D	ZIKV130	GRLIT ANPVI TESTE
ZIKV75	YLDKQSDTQYVCKRT	ZIKV103	I PLPW HAGA DTGTPH	ZIKV131	ANPVI TESTENSKMM
ZIKV76	SDTQYVCKRTLVD RG	ZIKV104	HAGA DTGTPH WNNKE	ZIKV132	TESTENSKMMLELDP
ZIKV77	VCKRTLVD RGVGNGC	ZIKV105	TGTPH WNNKEALV EF	ZIKV133	NSKMMLELDP PFGDS
ZIKV78	LVDRGVGNGCGLFGK	ZIKV106	WNNKEALV EFKDAHA	ZIKV134	LELDPPFGDSYIVIG
ZIKV79	WGNGCGLFGK GSLVT	ZIKV107	ALVEFKDAHAKRQTV	ZIKV135	PFGDSYIVIGVGEKK
ZIKV80	GLFGK GSLV TCAKFA	ZIKV108	KDAHAKRQTVVVLGS	ZIKV136	YIVIGVGEKKITHHW
ZIKV81	GSLV TCAKFA CSK KM	ZIKV109	KRQTVVVLGSQEGAV	ZIKV137	VGEKKITHHWHRSGS
ZIKV82	CAKFA CSK MTGKSI	ZIKV110	VVLGSQEGAVHTALA	ZIKV138	ITHHWHRSGSTIGKA
ZIKV83	CSK MTGKSI QPENL	ZIKV111	QEGAVHTALAGALEA	ZIKV139	HRSGSTIGKA FEATV
ZIKV84	TGKSI QPENLE YRIM	ZIKV112	HTALAGALEAEMDGA	ZIKV140	TIGKA FEATV RGA KR
ZIKV85	QPENLE YRIMLSVHG	ZIKV113	GALEAEMDGAKGRSL	ZIKV141	FEATV RGA KRMAVLG
ZIKV86	EYRIMLSVHGSQHSG	ZIKV114	EMDGAKGRSLSSGHLK	ZIKV142	RGA KRMAVLGDTAWD

Supplementary Table 9 Cont

Peptide	Amino Acid Sequence
ZIKV143	MAVLGDTAWDFGSVG
ZIKV144	DTAWDFGSVGGALNS
ZIKV145	FGSVGGALNSLGKGI
ZIKV146	GALNSLGKGIHQIFG
ZIKV147	LGKGIHQIFGAAFKS
ZIKV148	HQIFGAAFKSLFGGM
ZIKV149	AAFKSLFGMSWFSQ
ZIKV150	LFGMSWFSQILIGT
ZIKV151	SWFSQILIGTLLMWL
ZIKV152	ILIGTLLMWLGLNTK
ZIKV153	LLMWLGLNTKNGSIS
ZIKV154	GLNTKNGSISLMCLA
ZIKV155	NGSISLMCLALGGVL
ZIKV156	LMCLALGGVLIFLST
ZIKV157	LGGVLIFLSTAVSAD
ZIKV158	IFLSTAVSADV GCSV
ZIKV159	AVSADV GCSVDFSKK

Supplementary Table 9 Cont

Zika Virus NS1 Peptides

Peptide	Amino Acid Sequence	Peptide	Amino Acid Sequence	Peptide	Amino Acid Sequence
ZIKV160	VGCSVDFSKKETRCG	ZIKV188	ECPLKHRAWNSFLVE	ZIKV216	GTKVHVEETCGTRGP
ZIKV161	DFSKKETRCGTGVFV	ZIKV189	HRAWNSFLVEDHGFG	ZIKV217	VEETCGTRGPSLRST
ZIKV162	ETRCGTGVFVYNDVE	ZIKV190	SFLVEDHGFGVFHTS	ZIKV218	GTRGPSLRSTTASGR
ZIKV163	TGVFVYNDVEAWRDR	ZIKV191	DHGFGVFHTSVWLKV	ZIKV219	SLRSTTASGRVIEEW
ZIKV164	YNDVEAWRDRYKYHP	ZIKV192	VFHTSVWLKVREDYS	ZIKV220	TASGRVIEEWCCREC
ZIKV165	AWRDRYKYHPDSPRR	ZIKV193	VWLKVREDYSLECDP	ZIKV221	VIEEWCCRECTMPPL
ZIKV166	YKYHPDSPRRLAAAV	ZIKV194	REDYSLECDPAVIGT	ZIKV222	CCRECTMPPLSFRAK
ZIKV167	DSPRRLAAAVKQAWWE	ZIKV195	LECDPAVIGTAVKGK	ZIKV223	TMPPLSFRAKDGCWY
ZIKV168	LAAAVKQAWEDGICG	ZIKV196	AVIGTAVKGKEAVHS	ZIKV224	SFRAKDGCWYGMEIR
ZIKV169	KQAWEDGICGISSVS	ZIKV197	AVKGKEAVHSDLGYW	ZIKV225	DGCWYGMEIRPRKEP
ZIKV170	DGICGISSVSRMENI	ZIKV198	EAVHSDLGYWIESEK	ZIKV226	GMEIRPRKEPESNLV
ZIKV171	ISSVSRMENIMWRSV	ZIKV199	DLGYWIESEKNDTWR	ZIKV227	PRKEPESNLVRSMVT
ZIKV172	RMENIMWRSVEGELN	ZIKV200	IESEKNDTWRLKRAH	ZIKV228	ESNLVRSMVTA GSTD
ZIKV173	MWRSVEGELNAILEE	ZIKV201	NDTWRLKRAHLIEMK	ZIKV229	RSMVTA GSTDHMDHF
ZIKV174	EGELNAILEENGVQL	ZIKV202	LKRAHLIEMKTCEWP		
ZIKV175	AILEENGVQLTVVVG	ZIKV203	LIEMKTCEWPKSHTL		
ZIKV176	NGVQLTVVVGSVKNP	ZIKV204	TCEWPKSHTLWTDGI		
ZIKV177	TVVVGSVKNPMWRGP	ZIKV205	KSHTLWTDGIEESDL		
ZIKV178	SVKNPMWRGPQRLPV	ZIKV206	WTDGIEESDLIIPKS		
ZIKV179	MWRGPQRLPVVNEL	ZIKV207	EESDLIIPKSLAGPL		
ZIKV180	QRLPVVNELPHGWK	ZIKV208	IIPKSLAGPLSHHNT		
ZIKV181	PVNELPHGWKAWGKS	ZIKV209	LAGPLSHHNTREGYR		
ZIKV182	PHGWKAWGKSYFVRA	ZIKV210	SHHNTREGYRTQMKG		
ZIKV183	AWGKSYFVRAAKTNN	ZIKV211	REGYRTQMKG PWHSE		
ZIKV184	YFVRAAKTNN SFVVD	ZIKV212	TQMKG PWHSE ELEIR		
ZIKV185	AKTNN SFVVDGDTLK	ZIKV213	PWHSE ELEIR FE ECP		
ZIKV186	SFVVDGDTLKE CPLK	ZIKV214	ELEIR FE ECPG TKVH		
ZIKV187	GDTLKE CPLKHRAWN	ZIKV215	FE ECPG TKVHVEETC		

Supplementary Note 1

DENV-immune macaques (BS14) developed a nonpruritic skin rash that were centered on lightly haired skin areas (chin, axillary, ears, and inguinal areas) on day 8 p.i. (Supplementary Fig. 2a). This animal did not have a known history of allergy or skin condition. By day 23 p.i. most areas had self-resolved themselves except in the axillary and inguinal areas (Supplementary Fig. 2b). We performed diagnostic testing on affected areas by skin scraping for parasites, skin swabs for bacterial culture, samples for fungal cultures, and impression smears for cytology. We also examined skin biopsies from axillary and inguinal areas by qRT-PCR. Fungal cultures were negative and bacterial cultures showed only normal skin flora. Cytology indicated there was a neutrophilic and plasmacytic dermatitis with hyperkeratosis (Supplementary Fig. 2c). Histopathology of haired skin had a few number of lymphocytes, plasma cells, and lesser macrophages within the superficial dermis with predominant perivascular distribution. There was mild to moderate diffuse orthokeratotic hyperkeratosis (Supplementary Fig. 2d). No other abnormal clinical sign was noted in this animal during this period. qRT-PCR for ZIKV RNA from skin biopsy samples also did not show any specific amplification. To rule out the unlikely event of latent dengue reactivation as cause of this rash we tested the serum samples from the first ten days after ZIKV infection for the presence of dengue viremia in all four cohort 1 animals. Dengue RNA was not detected at any time after ZIKV challenge (Supplementary Table 4 and Supplementary Method)

Supplementary Methods

Dengue RT-PCR

Viral RNA from serum samples was extracted using QIAmp Viral RNA mini kit (Qiagen, Valencia, CA) according to manufacturer instructions. Real-time RT-PCR (TaqMan) singleplex assay-Specific primers and probes for DENV1: Forward (5' CAA AAG GAA GTC GTG CAA TA 3'), Reverse (5' CTG AGT GAA TTC TCT CTA CTG AAC C 3'), Probe (5' CAT GTG GTT GGG AGC ACG C 3' FAM/BHQ1) and DENV2: Forward (5' CAG GTT ATG GCA CTG TCA CGA T 3'), Reverse (5' CCA TCT GCA GCA ACA CCA TCT C 3') and Probe (5' CTC TCC GAG AAC AGG CCT CGA CTT CAA 3' HEX/BHQ1) (Sigma-Aldrich) were used. RNA from known DENV1 or DENV2 was included as positive control. For the reaction mixture, 10uL of RNA was combined with 100uM primers and probes in a 50uL total volume using qScript™ One-Step qRTPCR master mix kit (Quanta Biosciences™) according to manufacturers instructions. Thermocycling parameters were as follows: Reverse transcription for 30min at 50°C and reverse transcriptase inactivation for 12:30min. at 95°C was followed by 45 cycles of 95°C for 15 seconds and annealing at 60°C for 1 min using the iCycler IQ5 Real Time Detection System (Optical System software version 2.1; Bio Rad, CA) ¹.

Interpretation: Cycle threshold (Ct) is placed above background signal, usually intersecting the initial exponential phase of the amplification curve for each sample. Curves with treshold >36 render erratically and are difficult to determine as Ct values increases. Results may be unreliable, hence considered negative ^{1,2}.

Dengue viremia by immunofocus assay

Animals in cohort 1 were previously challenge with DENV1 or DENV2 in October 2013. After the challenge infectious virus in the blood was determined in serum samples within 10 days of challenge. As previously described ³ monolayer of Vero-81 cells were seeded in 24-well plates and inoculated with 100ul of challenge diluted serum samples (1:2.5, 1:5, 1:10, 1:20, 1:40) in triplicate wells. Cells were incubated for 1hr at 37C, then overlaid with 1ml of 1% methylcellulose Optic-MEM (Gibco) supplemented with 2% FBS (Cellgro) and antibiotics. After 4-5 days in a 37C incubator, cells were washed and fixed in 80% methanol. Immunocyto289 staining of DENV infected cells were assessed using anti-flavivirus monoclonal antibody 4G2 followed by a secondary antibody, HRP-conjugated goat anti-mouse Ab (Sigma, IL). Virus foci were visualized by the addition of TrueBlue HRP substrate (KPL, MD). Foci were counted and viral titers were calculated by standard methods.

Supplementary References

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- 3 White, L. J. *et al.* An alphavirus vector-based tetravalent dengue vaccine induces a rapid and protective immune response in macaques that differs qualitatively from immunity induced by live virus infection. *J Virol* 87, 3409-3424, doi:10.1128/JVI.02298-12 (2013).