

Zika Virus Phage Display Library

Virus	Strain or Antigen	GenBank Accession No.
ZIKV	H/PF/2013	KJ776791
	MR 766 Uganda	AY632535
	ArD128000	KF383117
	ArD157995	KF383118
	PRVABC59	KU501215
	Peptide #1 identified in peptide array	https://zika.labkey.com/announcements/OConnor/thread.view?rowId=101
	Peptide #2 identified in peptide array	
DENV-1	Myanmar 1996	AY722802
	Western Pacific	U88536
	DENV-1/BR/BID-V2401/2008	FJ850093
DENV-2	Burkina Faso 1983	EU056810
	16803	GU289914
	NewGuineaC	AF038403
	DENV-2/BR/BID-V3653/2008	GU131885
DENV-3	Mozambique 1985	FJ882575
	DENV-3/BR/BID-V2403/2008	FJ850094
DENV-4	P73-1120	JF262780
	H781363, 2011	JQ513345 (genotype I)
	BR/SJRP/850/2013	KP188566 (genotype II)
WNV	NY99	DQ211652
	WN02	KR868734
YFV	Uganda1948a	AY968065
	BeH655417, 2002	JF912190
JEV	Laos, 2009	KC196115
TBEV	MDJ-03	JF316708
CHIKV	KPA15	HQ456254 (structural proteins were translated from sequence and included in library)
	TR206/H804187, 2014	KP164572
ONNV	SG650	AF079456
RVFV	Kenya 9800523	DQ380169.1 for S DQ380196.1 for M DQ375400.1 for L
Malaria	HRP-II, Kenya	FJ871238
HIV-1	V3	Group M consensus
	MPER	Group M consensus
	Gp41 immunodominant	Group M consensus
	C5 immunodominant	Group M consensus
	Full-length HIV	Consensus Clade A, B, C, D (LANL)
	Full-length Q23	AF004885.1 & LANL Alignments
	BF520.W14M.C2	KX168094

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	BG505.W6.C2	DQ208458
	QA013 initial Env	FJ866134
	QA013 SI Env	FJ396015
	QB850 intial Env	Val's sequence info
	QB850 SI Env	Val's sequence info
	Q461.d1 Env	AF407155
	QC406.F3 Env	FJ866133
Rhinovirus B	Xu et al Rhinovirus B epitope (Table S2)	