

# A Dominant EV71-specific CD4 T cell epitope is conserved among enteroviruses

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# Hand, foot, and mouth disease (HFMD) and EV71









# **HFMD : Severe complications**

### Cardiorespiratory manifestations :

- Pneumonia
- Pulmonary hemorrage
- Pulmonary oedema
- Myocarditis

### > Neurological manifestations :

- Brainstem encephalitis (58%)
- Aseptic meningitis (36%)
- Encephalitis
- ANS dysregulation (4%)

### Systemic manifestations : Liver injury, myositis, thrombocytopenia

Since 2008, nearly 6 millions HFMD cases have been reported, and over 2000 children have been dead in China.



# Cellular rather than humoral immunity is correlated with clinical outcome of HFMD



Yang C, et al. Virology J 2011

Chang LY, et al. Pediatr 2008



## Three Human EV71 CD4+ T cell Epitopes Had Been discovered

Only EV71 VP1antigen alone



Foo DG et al. Viral Immunol. 2008 Jun;21(2):215-24.



Q	
X	

SP2	E VV PQ L LQ YM FV P P G
EV71/GDFS/3/2008	www.weither.com.weither.com.weither.com.weither.com.weither.com.weither.com.weither.com.weither.com.weither.com
FY23	
EV71/Fuyang Anhui P.R.C/17.08/3	
EV71/Lanzhou01	
EV71/HENAN/DC/2010	
SZ/HK08-6	
SZ/HK08-5	
EV71/Jiang su P.R.C/07.08/10	
BJ06-SJS06	
SHZH03	
EV71-Hubei-09-China	
BrCr	PP

#### LVVRIYMRMKHVRAW

L--------

---R-----

---R-----Q--P-----

EV71/GDFS/3/2008	
FY23	
EV71/Fuyang Anhui P.R.C/17.08/3	
EV71/Lanzhou01	
EV71/HENAN/DC/2010	
SZ/HK08-6	
SZ/HK08-5	
EV71/Jiang su P.R.C/07.08/10	
BJ06-SJS06	
SHZH03	
FY23-K14	
FY23-K12	
28/SHENZHEN/08	
EV71/Zhejiang08	
121/SHENZHEN/08/	
EV71-Hubei-09-China	I
BrCr	I

FY23-K14

FY23-K12

SP3

28/SHENZHEN/08

121/SHENZHEN/08/

EV71/Zhejiang08





### EpiMatirx Algorithm is applied to predict new EV71 CD4+ T cells epitopes



Anne De Groot

University of Rhode Island.

EpiMatrix has been successfully applied to the analysis of previously published epitopes, and in the prospective selection of epitopes from HIV, Mycobacterium tuberculosis, Tularemia to vaccinia virus.

FY573 polyprotein sequence was subjected to analysis by EpiMatirx Algorithm.





11 predicted epitopes are located in structural viral protein regions, and the rest are located in none-structural protein regions.

# All tested 6 adults respond to A3 eiptope, and to various degree to other epitopes

Peptide	Sequence	Cluster Address	Proportion of Responding Donor (%)
A1	NAQFHYLYRSGFCIHVQ	164 - 180	16.7
A2	ADGFELQHPYVLDAGISISQL	224 - 244	33.3
A3	PHQWINLRTNNCATII	248 - 263	100.0
A4	HCNFGLLVVPISPLD	278 - 292	66.7
A6	TGSFMATGKMLIAYTPPGGPLP	445 - 466	50.0
A7	IWDFGLQSSVTLVIPWISNTH	479 - 499	16.7
A8	NTAYIIALAAAQKNFTMKL	533 - 551	83.3
A10	PLVVRIYMRMKHVRAWIP	811 - 828	16.7
A11	NQNYLFKANPNYAGNSI	833 - 849	33.3
A13	HYPVSFSKPSLIYVE	933 - 947	33.3
A14	SREVEALKNYFIGSE	1034 - 1048	50.0
A19	GLEWVSNKISKFIDWL	1128 - 1143	33.3
A23	GVSFTSKFVIASTNASNIIVP	1321 - 1341	16.7
A24	VSELIREYNNRSAIGNTIE	1418 - 1436	16.7
A27	RLEVDFEQALFSKYVGNTLYEP	1780 - 1801	16.7
A28	EAALHYANQLKQLEINTSQMSMEE	1807 - 1830	16.7
A30	DSVYLRMAFGHLYETF	1914 - 1929	66.7
A31	GHLYETFHANPGTITGSA	1923 - 1940	16.7
A33	SPVWFRALELVLREIGYSE	1973 - 1991	16.7
A34	THHVYRNKTYCVLGGMPSGCS	2003 - 2023	16.7
SP1	IETRCVLNSHSTAET	66-67	16.7
SP2	EVVPQLLQYMFVPPG	145-159	33.3





### A3 and A8 are two dominant EV71 CD4+ T cell epitopes







### A3 and A8 epitopes are conservative among EV71 isolates, and A3 is a common CD4+ T cell epitope in EV71, Coxsackie viruses A4, A6 and A16.

	A3	A8	A14	SP2
	PHQWINLRTNNCATII	NTAYIIALAA AQKNFT MKL	SREVE ALK NYFIGSE	EVVPQLLQYMFVPPG
FY23			8-1	
FY23-K12			H I	
FY23-K14				
EV71/GDFS/3/2008				
EV71-Hubei-09-China				
EV71/HENAN/DC/2010			!	
EV71/Lanzhou01				
SHZH03				
SZ/HK08-6				
SZ/HK08-5				PP
121/SHENZHEN/08				L
28/SHENZHEN/08				LP
EV71/Fuyang.Anhui.P.R.C/17.08	/3			R
EV71/Jiangsu.P.R.C/07.08/10				R
BJ06-SJS06				QP

В

А

#### A3

#### A8

40	BUOWLNI BTNNCATLL	NTAVILALAAAOVNETUVI
A3	PHQWINLRINNCALLI	NIATIIALAAAQKNFIMKL
CAV A4		TVADN
CAV A6		-E-NMG-GNL
CAVA16		SE-NGENL
CAV A9	sv	SSCMCFVS-CND-SVR-
CAV B3	sv	SDCK-LCFVS-CND-SVRM
CAV B5	sv	SNCS-LCFVS-CND-SVRM
CAV B6	sv	T-ST-LCFVS-CND-SVR-
Echovirus 30	sv	-SSS-MCFAS-CND-SVRM
Echovirus 15	sv	TDCIVLCFVS-CND-SVRM
Echovirus 5	V	TQCD-LGFVS-CND-SVRM
Poliovirus 1	I L V	REMD-LGFVS-CND-SVR-
Poliovirus 2	I LV	RKMD-LGFVS-CND-SVR-
Poliovirus 3	I S V	KSM SMLGEVS-CND-SVR-











### A3 can stimulate PBMCs derived from previous EV71 infected healthy adults PBMCs to produce multiple cytokines







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