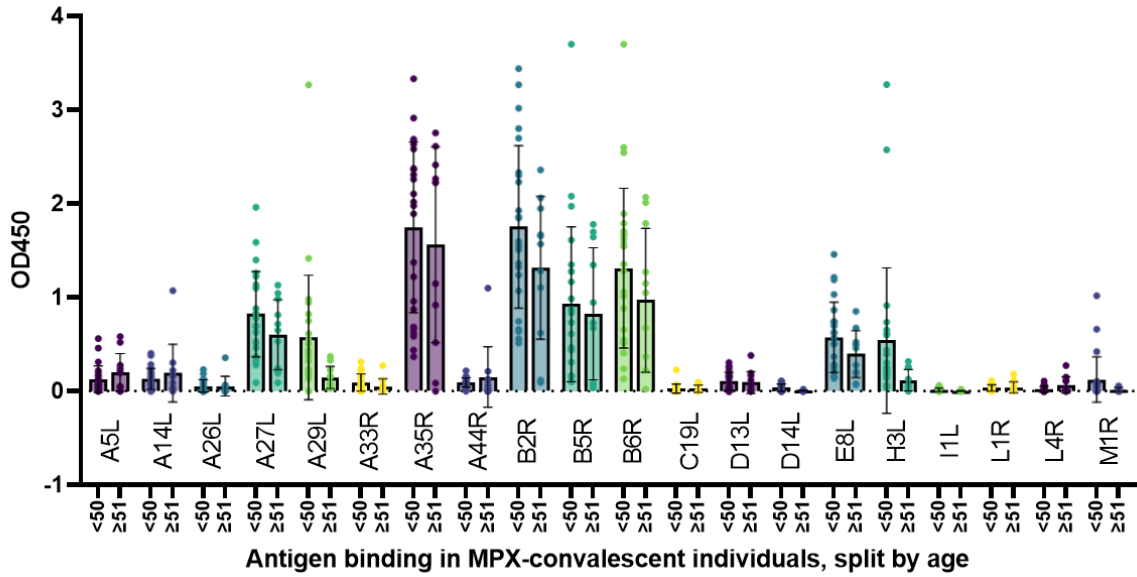


1 **Supplementary figures**

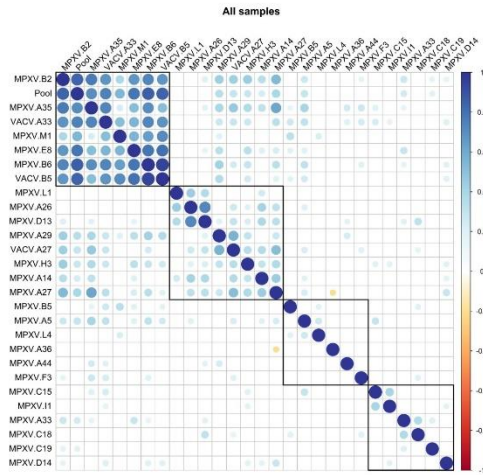
2
3
4



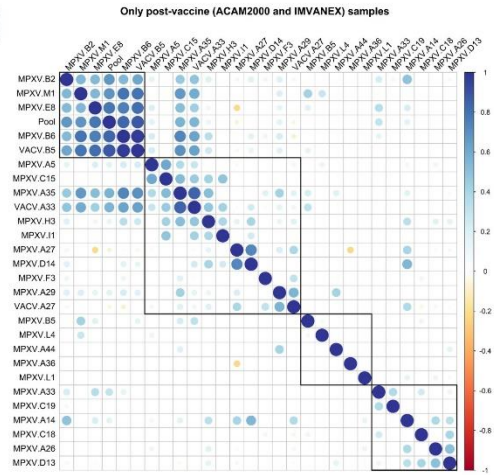
5
6
7
8
9
10

Supplementary Figure 1: Antigen binding in MPX individuals to different MPXV/VACV antigens, split according to age group (<50, ≥51). Points represent samples from singular individuals (<50 n=24, ≥50 n=11). Error bars represent mean ± standard deviation.

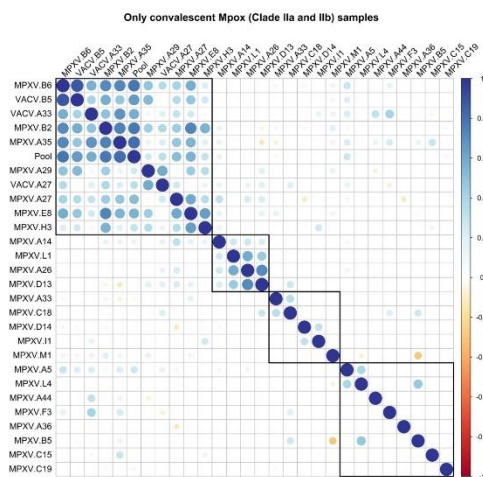
A)



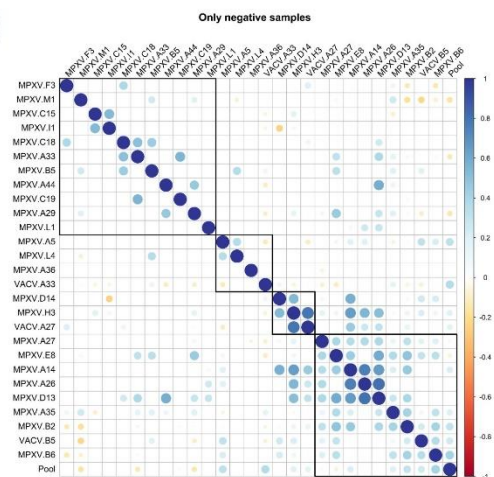
B)



C)



D)

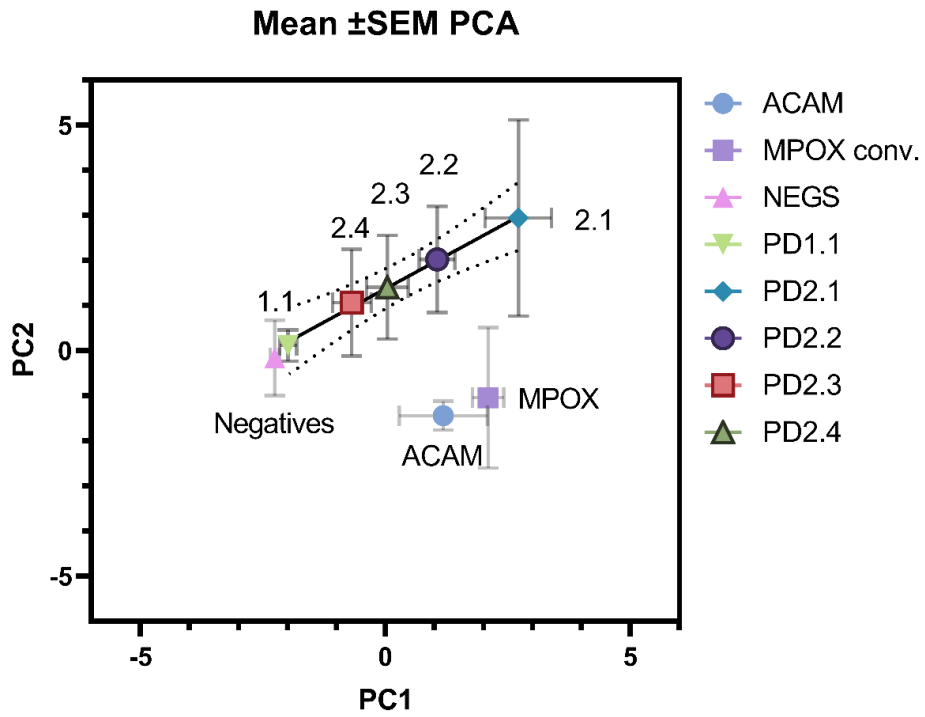


11

12

13 **Supplementary Figure 2:** Pearson correlation plots for **A)** All samples, **B)** post-vaccination
 14 samples only, **C)** Mpox-infected only (Clade IIa and IIb), and **D)** negative (pre-vaccination)
 15 samples. Only significant correlations are shown ($p > 0.05$), with boxes identifying groups
 16 through hierarchical clustering using Ward's method.

17



19

20 **Supplementary Figure 3:** Principal component analysis, with mean PCA \pm SEM for each
21 group. Simple linear regression (solid black line) was applied for all post-Imvanex vaccination
22 samples.
23

23

24 **Supplementary tables**

25 **Supplementary table 1:** Antigens used in this study, the viral source, company, expression
 26 host, expressed fragment and sequence provided by the manufacturer.

27

MPXV/VACV Protein	Virus	Company	Expression Vector	Expressed Fragment	Sequence
A5	MPXV	Native Antigen Company	E. coli	Total protein	MDFFNKFSQG LAESSTPKSS IYYSEEKDPD TKKDEAIEIG LKSQESYYQR QLREQLARDN MMTASRQPTQ PLQPTIHITP QPVPPTTPAP ILLPSSTAPV LKPRQQTNTS SDMSNLFDWL STDTDAPAST LLPALTPSNT VQDIISKFNK DQKMTTPPST QPSQTLPTTT CTQQSDGSIS CTTPTVTPLQ PPIVATVCTP TPTGGTVCTT AQQNPNPGAA SQQNLDDMTL KDLMSSVEKD MRQLQAETND LVTNVYDARE YTRRAIDQIL QLVKGFERFQ K
A14	MPXV	ProteoGenix	E. coli	Asn24-Ala70	-
A26	MPXV	ProteoGenix	E. coli	Asp2-Glu75	-
A27	MPXV	ProteoGenix	E. coli	Leu428-Thr695	-
A27	VACV	SinoBiological	E. coli	Met1-Glu110	-
A29	MPXV	ProteoGenix	Mammalian cells	Asp2-Glu110	-
A33	MPXV	SinoBiological	E. coli	-	-
A33	VACV	SinoBiological	E. coli	Val57-Asn185	-
A35	MPXV	SinoBiological	Mammalian cells (HEK293 Cells)	-	-
A36	MPXV	ProteoGenix	E. coli	Ile29-Lys168	-
A44	MPXV	ProetoGenix	E. coli	Asp2-Thr74	-
B2	MPXV	Abbexa	Mammalian cells	Ser19-Asp274	SP QTSKKIGDDA TISCSRNNTN YYVVMSAWYK EPNSIILLAA KSDVLYFDNY TKDKISYDSP YDDLVTITIT KSLTAGDAGT YICAFFMTST TNDTDKVDYE EYSIELIVNT DSESTIDIIL SGSTPETISE KPEDIDNSNC SSVFEIATPE PITDNVEDHT DTVTYTSDSI NTVNASSGES TTDETPPEIT DKEEDHTVTD TVSYTTVSTS SGIVTTKSTT DDADLYDTYN DNDTVPPPTTV GGSTTSISNY KTKD
B5	MPXV	ProteoGenix	E. coli	Asn26-Thr371	-
B5	VACV	SinoBiological	Mammalian cells (HEK293 Cells)	Tyr18-His279	-
B6	MPXV	ProteoGenix	Mammalian cells	Thr20-His279	-
C15	MPXV	ProteoGenix	Mammalian cells	Met1-Pro176	-
C18	MPXV	ProteoGenix	E. coli	Met1-Leu635	-
C19	MPXV	Abbexa	E. coli	Met1-Ile372	MWPFASVPAG AKCRLVETLP ENMDFRSDDL TTFECFNEII TLAKKIYIYA SFCCNPLSTT RGALIFDKLK EVSEKGIKII

					VLLDERGKRN LGELQSHSPD INFITVNIDK KNNVGLLLGC FWVSDDERCY VGNASFTGGS IHTIKTLGVY SDYPPPLATDL RRRFDTFKAF NSAKNSWLNL CSAACCLPVS TAYHIKNPIG GVFFTDSPFH LLGYSRDLDT DVVIDKLKSA KTSIDIEHLA IVPTTRVDGN SYYWPDYNS IIEAAINRGV KIRLLVGNWD KNDVYSMATA RSLDALCVQN DLSVKVFTIQ NNTKLLIVDD EYVHITSANF DGTHYQNHGF VSFNSIDKQL VSEAKKIFER DWVSSHKSLSL KI
D13	MPXV	ProteoGenix	E. coli	Met1-Glu315	-
D14	MPXV	ProteoGenix	Mammalian cells	Tyr20-Ala216	-
E8	MPXV	ProteoGenix	Mammalian cells	Pro2-Lys274	-
F3	MPXV	ProteoGenix	E. coli	Met1-Phe153	-
H3	MPXV	SinoBiological	Mammalian cells (HEK293 Cells)	-	-
I1	MPXV	SinoBiological	E. coli	-	-
L1	MPXV	ProteoGenix	E. coli	Met1-Gln152	-
L4	MPXV	Native Antigen Company	E. coli	-	MSLLENLIE EDTIFFAGSI SEYDDLQPMVI AGAKSKFPRS MLSIFNIVPR TMSKYELELI HNENITGAMF TTMYNIRNNL GLGDDKLTIE AIENYFLDPN NEVMPLIINN TDMTTVIPKK SGRRKNKNMV IFRQGSSPIL CIFETRKKIN IYKENMESVS TEYTPIGDNK ALISKYAGIN ILNVYSPSTS MRLNAIYGFT NKNKLEKLST NKELESYSSS PLQEPRLND FLGLLECVKK NIPLTDIPTK D
M1	MPXV	SinoBiological	Mammalian cells (HEK293 Cells)	-	-

28

29