

These tables includes the taxonomically significant peptides from Vero cell cultured VACV samples purified by sucrose cushion, two sucrose gradients with and without CsCl gradient.

The peptide, uniprot ID and protein name from one of the homologous proteins is provided.

The species specificity is provided along with the frequency of observation.

Peptides observed in more than replicate data set are included.

**Supplemental Table 2B**  
**Sucrose cushion and two sucrose gradient purified, n = 13**

Peptide	Uniport ID	Protein Description	Organism Specificity	Freq.
YGLIPEEFFQFLYPK	F7EHQ1	ATP synthase subunit b, mitochondrial	All Primates	0.38
LATQSNEITIPVTFESR	F6XTF7	Heat shock protein beta-1	All Primates	0.31
LPEEWSQWLGGSSWPGYVR		Heat shock protein beta-1		
PLPPAAIESPAVAAPAYSRSR	F6XTF7	ATP synthase subunit alpha	All Primates	0.15
FEN AFLSHVISQHQALLGTIR	F7ETD0	Complement component 1 Q	Primates, no great apes	0.62
GVDNTFADELVELSTALEHQ		subcomponent-binding		
EYISFLEDLK	Q9MZE0	protein, mitochondrial	Primates, no great apes	0.62
		Uncharacterized protein (A-kinase anchor protein 8		
GEKDDEDDEVK	F7DSP9	(Homo sapiens )	Primates, no great apes	0.15
TLSSIATSTDAAASVVHSTDLV		Hydroxyacyl-coenzyme A		
VEAIVENLK	F7DIX8	dehydrogenase	Primates, no great apes	0.31
		Superoxide dismutase		
GDSPVQGTINFEQK	Q8HXQ0	[Cu-Zn]	Primates, no great apes	0.23
		Superoxide dismutase	Primates, no great apes, Mus musculus, Sus scrofa	
HVGDLGNVTAGK	Q8HXQ0	[Cu-Zn]		0.77

**Supplemental Table 2C**

**Sucrose cushion, two sucrose gradient and CsCl gradient purified, n = 35**

Peptide	Uniport ID	Protein Description	Organism Specificity	Freq.
LPEEWSQWLGGSSWPGYVR		Heat shock protein beta-		
PLPPAAIESPAVAAPAYSR	F6XTF7	1	All Primates	0.06
		Superoxide dismutase		
GDSPVQGTINFEQK	P62938	[Cu-Zn]	Primates, no great apes	0.09
		Superoxide dismutase	Primates, no great apes, Mus musculus,	
HVGDLGNVTAGK	P62938	[Cu-Zn]	Sus scrofa	0.63