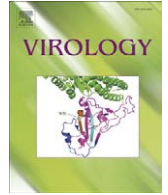




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Corrigendum

Corrigendum to “Identification of structural protein–protein interactions of herpes simplex virus type 1”
[Virology 378 (2008) 347–354]

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Table 2 was labeled incorrectly. The terms “Bait” and “Target” should be swapped. For the reader's convenience, the corrected table is reproduced here along with its legend.

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Table 2
Yeast two-hybrid matrix screen for binary interactions of HSV-1 structural proteins^a

Target \ Bait	Capsid						Tegument														Envelope															
	pUL6	pUL18	pUL19 UD	pUL25	pUL35	pUL38	pUL4	pUL11	pUL13	pUL14	pUL16	pUL17	pUL21	pUL36 N1	pUL37	pUL41	pUL46	pUL47	pUL48	pUL49	pUL51	pUL5	pUS2	pUS3	pUS10	pUS11	gM CT	gB CT	gC CT	pUL45 CT	gD CT	gI CT	gE CT	pUS9 CT		
Capsid																																				
pUL6	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUL18 (VP23)	-	+	-	-	-	+++	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE
pUL19 UD (VP5)	-	-	-	-	-	-	-	-	-	-	++	-	+++	-	-	-	-	-	-	+	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE
pUL25	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	
pUL35 (VP26)	-	+++	+	+	-	++	-	+	-	++	++	-	++	-	+++	-	-	-	++	-	+++	-	-	+	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUL38 (VP19C)	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	
Tegument																																				
pUL4	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUL11	-	-	-	-	-	-	-	-	-	-	+++	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE
pUL13	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	NE	
pUL14	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	
pUL16	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	
pUL17	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	
pUL21	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	
pUL36 N (VP1/2)	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+++	-	-	-	+++	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUL37	-	-	-	-	-	+	-	-	-	-	-	-	++	+++	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUL41	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUL46 (VP11/12)	-	+++	+	+	-	+	-	-	-	-	-	+	-	+++	-	-	-	+++	+	-	-	-	-	+++	+	-	-	++	-	NE	-	NE	-	NE	NE	
pUL47 (VP13/14)	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUL48-AD (VP16)	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA		
pUL49 (VP22)	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+++	+++	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUL51	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUL56	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUS2	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
pUS3	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	
pUS10	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	
pUS11	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+++	-	-	NE	-	NE	-	NE	NE	
Envelope																																				
gpUL10 CT (gM)	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	
gpUL27 CT (gB)	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	+++	NE	-	NE	-	NE	NE	
gpUL44 CT (gC)	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
gpUS6 CT (gD)	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
gpUS7 CT (gI)	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	
gpUS8 CT (gE)	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	
pUS9 CT	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	NE	-	NE	-	NE	NE	

Positive interactions (shaded) were observed as blue colonies with + indicating lightest blue and +++ the darkest blue. No growth (-) indicates a negative interaction.

AA, autoactivation; NE, not expressed; N, amino acids 1–1874; N1, amino acids 1–767; CT, cytoplasmic domain; -AD, minus activation domain (amino acids 412–490); UD, upper domain (amino acids 451–1054).

^a Interactions were assessed using a qualitative in vivo plate assay (at day 3) with readout being expression of the reporter genes *LEU2* and *lacZ*.