

EBNA2	(452)	ADLDES	W	DY	IF	ETTE	(466)
VP16	(469)	DMADFE	F	EQ	MF	TDAL	(483)
P53	(44)	MLSPDD	I	EQ	WF	TEDP	(58)
			Φ	XX	$\Phi\Phi$		

Supplementary Figure S2. Tfb1/p62 binding regions of acidic TADs.

Sequence alignment of the regions from the TADs of p53, VP16 and EBNA2 (that form α helices when bound to the Tfb1/p62 subunit of TFIID). Key hydrophobic residues (Φ) of the Φ XX $\Phi\Phi$ motif that directly interact with Tfb1PH/p62PH are highlighted in black. Numbers of residues shown on each side of the sequence are inclusive.