

Protein Alignment Summary Table (Vpx)

Name(s)	Accession No.	Region	First Author	Reference
ROD	M15390	complete genome	Guyader M	<i>Nature</i> 326 (6114), 662–669 (1987)
NIHZ	J03654	complete genome	Zagury JF	<i>PNAS</i> 85 5941–5945 (1988)
ISY	J04498	complete genome	Franchini G	<i>PNAS</i> 86 2433–2437 (1989)
ST	M31113	complete genome	Kumar P	<i>J Virol</i> 64 890–901 (1990)
BEN	M30502	complete genome	Kirchhoff F	<i>Virology</i> 177 305–311 (1990)
CAM2	D00835	complete genome	Tristem MF	<i>J Gen Virol</i> 72 721 (1991)
D194	J04542	complete genome	Kuehnel H	<i>PNAS</i> 86 2383–2387 (1989)
GHI	M30895	complete genome	Hasegawa A	<i>ARHR</i> 5 593–604 (1989)
KR	U22047	complete genome	Kraus GK	<i>PNAS</i> 90 (9), 4226–4230 (1993)
MDS	Z48731	gag, pol, vif, tat, rev, nef, env	Becker M	Unpublished (1995)
UC2	U38293	complete genome	Barnett SW	Unpublished (1997)
UC1	L07625	complete genome	Barnett SW	<i>J Virol</i> 67 1006–1014, (1993)
D205	X61240	complete genome	Dietrich U	<i>ARHR</i> 8 1619 (1992)
EHQA	U27200	complete genome	Rey-Cuille MA	<i>Virology</i> 202 (1), 471–476 (1994)
MM251	M19499	complete genome	Franchini G	<i>Nature</i> 328 539–543 (1987)
MM1A11	M76764	complete genome	Luciw PA	<i>ARHR</i> 8 395–402 (1992)
MM32H	D01065	complete genome	Rud EW	<i>J Gen Virol</i> 75 529 (1994)
MM316S	L22790	vif, vpx, vpr	Kodama T	<i>J Virol</i> 67 6522–6534 (1993)
MM239	M33262	complete genome	Regier DA	<i>ARHR</i> 6 1221–1231 (1990)
MM142	M16403	complete genome	Chakrabarti L	<i>Nature</i> 328 543–547 (1987)
MNE	M32741	complete genome	Benveniste RE	Unpublished (1990)
SMMPBJ14.441A	M31325	complete genome	Dewhurst S	<i>Nature</i> 345 636–640 (1990)
SMMH9	M80194	complete genome	Cournaud V	<i>J Virol</i> 66 414–419 (1992)
SMMH4	X14307	complete genome	Hirsch VM	<i>Nature</i> 339 389–391 (1989)
SM62A	U04982	partial env, nef, tat, rev	Hirsch VM	<i>J Virol</i> 68 2649–2661 (1994)
STM	M83293	complete genome	Novembre FJ	<i>Virology</i> 186 783–787 (1992)

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