



SURE v1.0 nucleotide sequence unanimity alignments for HIV-1 Vif encoding region (created Jun 23, 2015; reformatted Nov 19, 2018, v6)  
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Reference	1	10	20	30	35
	M	Q	D	H	K
	ATG	CAG	GAC	CAC	AAA
Reference	ATG	CAG	GAC	CAC	AAA
MX033	---	---	---	---	---
MX050	---	---	---	---	---
MX066	---	---	---	---	---
MX071	---	---	---	---	---
MX071_2	---	---	---	---	---
MX077	---	---	---	---	---
MX078	---	---	---	---	---
MX081	---	---	---	---	---
MX086	---	---	---	---	---
MX089	---	---	---	---	---
MX091	---	---	---	---	---
MX101	---	---	---	---	---
MX109	---	---	---	---	---
MX110	---	---	---	---	---
MX110_2	---	---	---	---	---
MX111	---	---	---	---	---
MX112	---	---	---	---	---
MX118	---	---	---	---	---
MX119	---	---	---	---	---
MX122	---	---	---	---	---
MX123	---	---	---	---	---
MX125	---	---	---	---	---
MX131	---	---	---	---	---
MX133	---	---	---	---	---
MX158	---	---	---	---	---
MX160	---	---	---	---	---
MX166	---	---	---	---	---
MX168	---	---	---	---	---
MX172	---	---	---	---	---
MX176	---	---	---	---	---
MX186	---	---	---	---	---
MX190	---	---	---	---	---
MX196	---	---	---	---	---
MX207	---	---	---	---	---
MX207_2	---	---	---	---	---
MX230	---	---	---	---	---
MX313	---	---	---	---	---
MX324	---	---	---	---	---
MX327	---	---	---	---	---
MX350	---	---	---	---	---
MX350_2	---	---	---	---	---
MX350_3	---	---	---	---	---
MX351	---	---	---	---	---
MX354	---	---	---	---	---
MX355	---	---	---	---	---
MX370	---	---	---	---	---
MX375	---	---	---	---	---
MX376	---	---	---	---	---
MX381	---	---	---	---	---
MX381_2	---	---	---	---	---
MX385	---	---	---	---	---
MX385_2	---	---	---	---	---
MX390	---	---	---	---	---
MX399	---	---	---	---	---
MX399_2	---	---	---	---	---
MX401	---	---	---	---	---
MX403	---	---	---	---	---
MX409	---	---	---	---	---
MX450	---	---	---	---	---
MX463	---	---	---	---	---
MX464	---	---	---	---	---
MX469	---	---	---	---	---
MX470	---	---	---	---	---
MX473	---	---	---	---	---
MX474	---	---	---	---	---
MX540	---	---	---	---	---
MX542	---	---	---	---	---
MX546	---	---	---	---	---
MX554	---	---	---	---	---
Consensus_D	---	---	---	---	---
MX141	---	---	---	---	---
MX210	---	---	---	---	---
MX352	---	---	---	---	---
MX395	---	---	---	---	---
MX396	---	---	---	---	---
Consensus_A	---	---	---	---	---
Consensus_G	---	---	---	---	---
MX384	---	---	---	---	---
MX384_2	---	---	---	---	---
MX384_3	---	---	---	---	---













## Key to annotations

The following Vif protein motifs descriptions are based on criteria established by Bell *et al*, 2007.

- Tryptophans ( $W^5$ ,  $W^{11}$ ,  $W^{21}$ ,  $W^{38}$  &  $W^{79}$ ) involved in APOBEC3G recognition and suppression are shown in cyan highlight).
- APOBEC3 binding motifs are shown in dark blue highlight; the C-terminal APOBEC3F-binding motif ( $^{171}EDRWN^{175}$ ) is not highlighted for simplicity.
- The  $^{88}EW^{89}$  motif located in central hydrophilic region known to be involved in enhancing steady state expression is shown in yellow highlight and red font.
- The nuclear localization inhibitory signal ( $^{90}RLRR^{93}$ ) is shown in red highlight. Sequences not having the consensus RLRR motif but having residues with conservative physicochemical properties are also shown in highlight.
- The  $^{95}ST^{96}$  CKII and p44/42 Mitogen-Activated Protein Kinase (MAPK) phosphorylation sites are shown in magenta highlighting. Residues other than ST that can also be phosphorylated are also shown in highlight.
- In any case, clinical isolate or subtype consensus sequences not having the aforementioned functional sites or motifs are shown without highlight in the corresponding region (i.e.  $W^{11}$  which in MX071\_2 is represented by an R<sup>11</sup>).
- The zinc-binding motif ( $H^{108}C^{114}C^{133}H^{139}$ ) is shown in black highlight and white font.
- Viral BC Box ( $^{144}SLQYLALAALITPK..K^{158}$ ) is shown in green highlight and based on the criteria suggested by Bizinoto *et al*, 2013.
- The protease processing site ( $L^{150}$ ) is also shown in black highlight.
- Threonine phosphorylation sites ( $T^{155}$  and  $T^{170}$ ) are shown in yellow highlight.
- The Cullin-binding box ( $^{159}KPPLPSVTKLTEDR^{173}$ ) is shown in grey highlight.

## References

*Molecular characterization of the HIV type 1 subtype C accessory genes vif, vpr, and vpu.* Bell CM, Connell BJ, Capovilla A, Venter WD, Stevens WS, Papathanasopoulos MA. AIDS Res Hum Retroviruses. 2007 Feb;23(2):322-30.

*Codon pairs of the HIV-1 vif gene correlate with CD4+ T cell count.* Bizinoto MC, Yabe S, Leal É, Kishino H, Martins Lde O, de Lima ML, Morais ER, Diaz RS, Janini LM. BMC Infect Dis. 2013 Apr 11;13:173.