

Mexican influenza virus A/H1N1/2009 pdm segment 7 (M2) protein sequences (n=21, length 97 aa).

	1	30	60	90	97
A/California/04/2009					
A/Mexico/UASLP-004/2009	*****TEVETPTRSEWECRCSDDSDPLVIAANIIGILHLILWITDRLFFKCIYRRFKYGLKRGPESTEGVPESMREEYQQEQQSAVDVDDGHFVNIELE				
A/Mexico/UASLP-018/2009	*****-----*****				
A/Mexico/UASLP-019/2009	*****-----*****				
A/Mexico/UASLP-021/2009	*****-----*****				
A/Mexico/UASLP-022/2009	*****-----*****				
A/Mexico/UASLP-023/2009	*****-----*****				
A/Mexico/UASLP-024/2009	*****-----*****				
A/Mexico/UASLP-026/2009	*****-----*****				
A/Mexico/UASLP-029/2009	***L-----				
A/Mexico/UASLP-030/2010	**LL-----				
A/Mexico/4108/2009	MSLL-----N-----				
A/Mexico/LaGloria-3/2009	MSLL-----				
A/Mexico/4486/2009	MSLL-----F-----				
A/MexicoCity/011/2009	MSLL-----H-----				
A/MexicoCity/007/2009	MSLL-----R-----				
A/MexicoCity/CIA2/2009	MSLL-----G-----				
A/MexicoCity/WR1697T/2009	MSQL-----				
A/MexicoCity/WR1706T/2009	MSQL-----K-----				
A/MexicoCity/WR1090N/2009	MSQL-----G-----				
A/Mexico/InDRE3740/2011	MSLL-----N-----				

Hyphens indicate unanimity to reference, asterisks indicate unknown sequence, full-stops indicate insertions, X indicates pre-mature stop codon.
 Unanimity reformatting algorithm written by James Robinson, Anthony Nolan Research Institute, HLA Informatics Group. Last modified 11/02/03.