CLUSTAL O(1.2.4) multiple sequence alignment

PB1_NCBI PB1_pHW2000	MDVNPTLLFLKVPAQNAISTTFPYTGDPPYSHGTGTGYTMDTVNRTHQYSEK SESRQTI*MDVNPTLLFLKVPAQNAISTTFPYTGDPPYSHGTGTGYTMDTVNRTHQYSEK ************************************	52 59
PB1 NCBI	GKWTTNTETGAPQLNPIDGPLPEDNEPSGYAQTDCVLEAMAFLEESHPGIFENSCLETME	112
PB1 pHW2000	GKWTTNTETGAPQLNPIDGPLPEDNEPSGYAQTDCVLEAMAFLEESHPGIFENSCLETME	119
	****************	
PB1_NCBI	VVQQTRVDKLTQGRQTYDWTLNRNQPAATALANTIEVFRSNGLTANESGRLIDFLKDVME	172
PB1_pHW2000	VVQQTRVDKLTQGRQTYDWTLNRNQPAATALANTIEVFRSNGLTANESGRLIDFLKDVME ************************************	179
PB1_NCBI	SMDKEEMEITTHFQRKRRVRDNMTKKMVTQRTIGKKKQRVNKRGYLIRALTLNTMTKDAE	232
PB1_pHW2000	SMDKEEMEITTHFQRKRRVRDNMTKKMVTQRTIGKKKQRVNKRGYLIRALTLNTMTKDAE ************************************	239
PB1_NCBI	RGKLKRRAIATPGMQIRGFVYFVETLARSICEKLEQSGLPVGGNEKKAKLANVVRKMMTN	292
PB1_pHW2000	RGKLKRRAIATPGMQIRGFVYFVETLARSICEKLEQSGLPVGGNEKKAKLANVVRKMMTN ***********************************	299
PB1_NCBI	SQDTELSFTITGDNTKWNENQNPRMFLAMITYITKNQPEWFRNILSIAPIMFSNKMARLG	352
PB1_pHW2000	SQDTELSFTITGDNTKWNENQNPRMFLAMITYITKNQPEWFRNILSIAPIMFSNKMARLG ************************************	359
PB1_NCBI	KGYMFESKRMKLRTQIPAEMLASIDLKYFNESTRKKIEKIRPLLIDGTASLSPGMMMGMF	412
PB1_pHW2000	KGYMFESKRMKLRTQIPAEMLASIDLKYFNESTRKKIEKIRPLLIDGTASLSPGMMMGMF ********************************	419
PB1_NCBI	NMLSTVLGVSILNLGQKKYTKTTYWWDGLQSSDDFALIVNAPNHEGIQAGVDRFYRTCKL	472
PB1_pHW2000	NMLSTVLGVSILNLGQKKYTKTTYWWDGLQSSDDFALIVNAPNHEGIQAGVDRFYRTCKL ************************************	479
PB1_NCBI	VGINMSKKKSYINKTGTFEFTSFFYRYGFVANFSMELPSFGVSGINESADMSIGVTVIKN	532
PB1_pHW2000	VGINMSKKKSYINKTGTFEFTSFFYRYGFVANFSMELPSFGVSGINESADMSIGVTVIKN ************************************	539
PB1_NCBI	NMINNDLGPATAQMALQLFIKDYRYTYRCHRGDTQIQTRRSFELKKLWDQTQSRAGLLVS	592
PB1_pHW2000	NMINNDLGPATAQMALQLFIKDYRYTYRCHRGDTQIQTRRSFELKKLWDQTQSRAGLLVS ***********************************	599
PB1_NCBI	DGGPNLYNIRNLHIPEVCLKWELMDEDYRGRLCNPLNPFVSHKEIESVNNAVVMPAHGPA	652
PB1_pHW2000	DGGPNLYNIRNLHIPEVCLKWELMDEDYRGRLCNPLNPFVSHKEIESVNNAVVMPAHGPA ************************************	659
PB1_NCBI	KSMEYDAVATTHSWIPKRNRSILNTSQRGILEDEQMYQKCCNLFEKFFPSSSYRRPVGIS	712
PB1_pHW2000	KSMEYDAVATTHSWIPKRNRSILNTSQRGILEDEQMYQKCCNLFEKFFPSSSYRRPVGIS ************************************	719
PB1_NCBI	SMVEAMVSRARIDARIDFESGRIKKEEFSEIMKICSTIEELRRQK	757
PB1_pHW2000	SMVEAMVSRARIDARIDFESGRIKKEEFSEIMKICSTIEELRRQK**I*LVLHEKMPCFY *********	776