

Sc Pol5	1	MTGKVNRLDFKIASDLREERLHAAVALIKDLSALDLPDDAEWWSYVLNRLIKGLSSDRN	60
Sp pol5	1	MATKTQIELEFKITSNDKAIRLSSAACLIDSL-----NEELKYSLNRLTKGLSSGRE	54
Sc Pol5	61	SARLGFSLCLTEVINLAVNMPPGQRFKGLESTNEFLSTLSTILNVNVNEGTKKSMKQKDE	120
Sp pol5	55	SARIGFAVALTHELL-----T-----RTKEIRAT----HVIDLLVKHNTASG--N-IKQODE	98
Sc Pol5	121	RGILFGKLPGLKSLNNEPLFSEIFVKDLEKGNTEFIRFTECLIDLALKKMWIKFPCFFT	180
Sp pol5	99	RDFYFGLPGLCSIV---YSGILTHKES--TIELEQVVDILLQLSGKKNWLQDVCFYV	152
		TV	
Sc Pol5	181	LFQTMKMLLPFMDES-SAEKILLIYDKYDLTLTNEGLSTYLLIKYEGEESLIPSVLDLKN	239
Sp pol5	153	IYKLVCEIPEISFSSNAFLAVNKLLQTPAVSKSTXGVGLFLCLTRVFN-VKSEEVAMAN	211
		II	
		pol5-3	
		R	
Sc Pol5	240	PGMKDNDPLARGNPLLLTFVLRNSSVIFDANGGLKETKKQKNTNWNFRLHFVMSVLLPLF	299
Sp pol5	212	--WEPAPLHKSNLVTLISKIMRQA----DAS----ETGGQ-NSAWKCKIPMVWKYIFEEY	260
Sc Pol5	300	GNGKLENTSHISKRRKKTNNKQVNSICFPFVKMAVDESFFNEKASSERKYLGFLLIIDA	359
Sp pol5	261	-----QRTYSGLAFPHDFWAVVVDGIFSSSTSSLERKFWGFCIMEL	302
		M	
Sc Pol5	360	AFKAVPGSYIGFCFSCNVMRTLINOSIISORVLNKISQLTLDIVKACREDSANRLVPCL	419
Sp pol5	303	ALDYVSSDNIGDIFSKNFLHCLINHLSEEDRYLYRAAKRVTISKLEXSKQ---NPTLVYI	359
		VI	
		pol5-2	
		S	
Sc Pol5	420	NAMLFGFHGSINFTKLTKSGVSKLIAIK-ELPSTVLAQLLDVFFLQLQ-----EFKGVL	473
Sp pol5	360	AIHLLGERGSLNFTRVINTKIVEHILPLADEQGILQLFOLLLSYVKRCPEDIASETKAVE	419
		pol5-1	
		E	
Sp pol5	420	WRRQWATDTMLSILRSKRSIKQBPWVREITRNIHCLMATLRYQQSEEVIPKPFSEGTQNMF	479
Sc Pol5	530	KERLYSILGELTINKEIRCK-DPEIN--SWQYLTIKLIDDIENSHVGLINPLDENLENI	586
Sp pol5	480	RLRLMSALSYLSSSAFQOSOTDHOLOGDNWPYVALNVILLELEKSPKNNILISMDESIVIEI	539
		III	
		pol5-2	
		S	
Sc Pol5	587	KNEAISCLSKVCR-----SRTASWGI STLISMCLVCIYAGD TDSISYIHELCE-----	634
Sp pol5	540	VOKSLSVLHKVTKKIDKKAQHLQLNAFQLLYSLVLLCVYAGD TDSILVLEDICNCYSKV	599
		I	
		pol5-1	
Sc Pol5	635	---FSKHENNS-----MVGITEILLSLIAQKKALLRKLSLIIWQCFIEEVGLIHELQILLD	687
Sp pol5	600	FNKKSREKRESTSNEPTAMEIITEVMSLSISRPSILLRKLVEMLFTSFSEDMNRRESIHLICD	659
Sc Pol5	698	ILKARENKQGFACIFEGEEFPERIKEENDASSEKSTGSESESESESDSDDADEKDERDE	747
Sp pol5	660	VLKAKESVKDSEGMFAGEEVEEDAFGETEMDEIDFEBIDTCEIEECSDWEMISNCDASD-	718
Sc Pol5	748	ANEDILNIDKEATSALVVMINLPDNIIVNDKGEVDLDQLEGLSDDGGDDEESMCDKMM	807
Sp pol5	719	---NE---ELERKLDKVIEDADAKVKD---E-----ESSEBELMNDQML	754
		V	
Sc Pol5	808	ELDDCISEIFKRRKEALSSISTGNQRFKFEVKQSRENVISFKHRVVDMLAVVVKYCEKLT	867
Sc Pol5	868	ANKSEHSNNLGGSLSKLVYFIIIPMLKCVNETLDRFLADKISKLLKGIKFKIKVTAFKDMN	927
Sp pol5	806	-N-----NGLG-----PEFLIPLLEMILKTKHKVLEEKQAVFRNRLSKLWTEEKPS	853
Sc Pol5	928	KDIELMDLLKXTHKMLTSPKPGQHAAVFYSMCSTSSLFSLKLYVEIGGNDKIDELIDLYT	987
Sp pol5	854	K--NVLEALKKVHVLGK-----AS-LGSTGSSISQLLIKLLADTP---YLKEGVEVYL	902
Sc Pol5	998	ATTKEAMQ---KQKCGPNIFIDFINWLSK-----KCTVMDKE-----	1022
Sp pol5	903	KSFLLIQEPSKSHYNANIFHDFINWCAQORLKHQQTSTAASSPKTGHH	