

FIGURA 7

sp O60449 LY75_HUMAN	MRTGWATPRRPAGLLMLLFWFFDLAEPFSGRAANDPFTIVHGNTKGCIKIPVYGMIVADDCD	60
MS-Peptides_Elutela	-----CIKIPVYGMIVADDCD	
MS-Peptides_Elutela1b	-----CIKIPVYGMIVADDCD	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	ETEDKILWKWVSQHRLFHLHSQKCLGLDITKSVNELRMFSCDSSAMLWWKCEHHSLYGAAR	120
MS-Peptides_Elutela	ETEDKILWK-----CEHHSLYGAAR	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	YRLALKDGHGTAISNASDVWKKGGSEESLCDQPYHEIYTRDGNYSYGRPCPEFPFLIDGTWH	180
MS-Peptides_Elutela	-----DGHGTAISNASDVWKKGGSEESLCDQPYHEIYTR-----	
MS-Peptides_Elutela1b	-----KGGSEESLCDQPYHEIYTR-----	
MS-Peptides_Elute2a	-----GGSEESLCDQPYHEIYTR-----	
MS-Peptides_Elute2b	-----GGSEESLCDQPYHEIYTR-----	
LC-Peptides	-----DGHGTAISNASDVWK-----	
sp O60449 LY75_HUMAN	HDCILDEDHSGPWCATTILNYEYDRKWGICLKWPENGCEDNWEKNEQFGSCYQFNTQTALSW	240
MS-Peptides_Elutela	-----	
MS-Peptides_Elutela1b	-----WGICLKWPENGCEDNWEK-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	KEAYVSCQNQGADLLSINSAAELTYLKEKEGIATKIFWIGLNLQLYSARGWEWSDHKPLNFL	300
MS-Peptides_Elutela	-----I F W I G L N L Q L Y S A R-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----I F W I G L N L Q L Y S A R-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	NWDPRPSAPTIGGSSCARMDAESGLWQSFSCEAQLPYVCKKPLNNTVELTDVWVTSYDTR	360
MS-Peptides_Elutela	-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	CDAGWLPNNGFCYLLVNESNSWDKAHAKCKAFSSDLISIHSLADVEVVVTKLHNEDIKEE	420
MS-Peptides_Elutela	-----LHNEDIKEE	
MS-Peptides_Elutela1b	-----LHNEDIKEE	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	VWIGLKNINIPTLFQWSDGTEVTILTYWDENEPNVPYNKTPNCVSYLGEIGQWVQSCEEK	480
MS-Peptides_Elutela	VWIGLK-----TPNCVSYLGEIGQWK-----	
MS-Peptides_Elutela1b	VWIGLK-----TPNCVSYLGEIGQWK-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	

FIGURA 7 (continuación)

sp O60449 LY75_HUMAN	LKYVCKRKGEKLNDASSDKMCPPEDEGWKRHGETICYKIYEDEVFPFGTNCNLTITSRFBQEY	540
MS-Peptides_Elutela	-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	LNDLMKKYDKSLRKYFWTGLRDVDSCGEYNWATVGGRRRAVTFSNWNFLPASPGGCVAM	600
MS-Peptides_Elutela	-----YFWTGLRDVDSCGEYNWATVGGR-----	
MS-Peptides_Elutela1b	-----YFWTGLRDVDSCGEYNWATVGGR-----	
MS-Peptides_Elute2a	-----YFWTGLRDVDSCGEYNWATVGGR-----	
MS-Peptides_Elute2b	-----YFWTGLRDVDSCGEYNWATVGGR-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	STGKSVGKWEVKDCRSFKALSICKKMSGPLGPEEASPKPDDPCPEGWQSFPAASLCYKVF	660
MS-Peptides_Elutela	-----SVGKWEVKDCR-----ALSICK-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----WEVKDCRSFK-----PASLSYKVF	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	HAERIVRKRNWEEAERFCQALGAHLSSFHVDEIKEFHLFLTQDQSGQHWLWIGLNKRSP	720
MS-Peptides_Elutela	-----RNWEEAER-----RSP	
MS-Peptides_Elutela1b	-----RNWEEAER-----RSP	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	HA-----	
sp O60449 LY75_HUMAN	DLQGSWQWSDRTPVSTIIMPNEFQQDYDIRDCAAVKFVHRPWRRGWHFYDDREFIYLRPF	780
MS-Peptides_Elutela	-----TPVSTIIMPNEFQQDYDIR-----	
MS-Peptides_Elutela1b	DLQGSWQWSDRTPVSTIIMPNEFQQDYDIR-----	
MS-Peptides_Elute2a	-----GWHFYDDR-----	
MS-Peptides_Elute2b	-----PWRRGWHFYDDREFIYLRPF	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	ACDTKLEWVCQIPKGRTPKTPDWYNPDRAGIHGPPLIIEGSEYWFVADLHLYNEEAVLYC	840
MS-Peptides_Elutela	-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	ASNHSFLATITSFVGLKAINKNIANSIGDGQKWIRISEWPI	900
MS-Peptides_Elutela	-----ISEWPI	
MS-Peptides_Elutela1b	-----ISEWPI	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	GEECLYMSAKTWLIDLGKPTDCSTKLPTICEKYNSVLSLEKYSPDSAAKVCSEQWIPFQN	960
MS-Peptides_Elutela	GEECLYMSAKTWLIDLGKPTDCSTK-----YNVSSLEK-----VQCSEQWIPFQN	
MS-Peptides_Elutela1b	-----TWLIDLGKPTDCSTK-----YNVSSLEK-----VQCSEQWIPFQN	
MS-Peptides_Elute2a	-----TWLIDLGKPTDCSTK-----	
MS-Peptides_Elute2b	-----TWLIDLGKPTDCSTK-----	
LC-Peptides	G-----	
sp O60449 LY75_HUMAN	KCFIKIKPVSLTFSQASDTCHSYGGTLPVLSLQIEQDFITSLLPDMEATLWIGLRWTAYE	1020
MS-Peptides_Elutela	-----	

FIGURA 7 (continuación)

MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	KINKWTDNRELTYSNFHPLVSGRLRIPENFFEEESRYHCALILNLQKSPPTGTWN	1080
MS-Peptides_Elutela	K-----ELTYSNFHPLVSGRLRIPENFFEEESRYHCALILNLQK-----	
MS-Peptides_Elutela1b	-----YHCALILNLQK-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----RELTYSNFHPLL-----	
sp O60449 LY75_HUMAN	SERHFFVSLCQKYSSEVKSQTLQNASPTVKYLNNNLYKIIPKTLTWHSAKRECLKSNMQLVS	1140
MS-Peptides_Elutela	--HFVSLCQK-----QTLQNASPTVK-----TLTWHSAK-----	
MS-Peptides_Elutela1b	--HFVSLCQK-----QTLQNASPTVK-----TLTWHSAK-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	SERHFFVSLCQKYS-----TVKYLNNNLYKI-----	
LC-Peptides	***** **	
sp O60449 LY75_HUMAN	ITDPYQQAFLSVQALLHNSSLWIGLFSQDDELNFGWSDGKRLHFSRWAETNGQLEDCVVL	1200
MS-Peptides_Elutela	-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	DTDGFWKTVDNCNDNQPGAICYYSGNETEKEVKFPVDSVKCPSPLVNTWPWIPFQNCNCYFI	1260
MS-Peptides_Elutela	-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	TKNRHMATTQDEVHTKCQKLNPKSHILSIRDEKENNFVLEQLLYNFYMASWVMLGITYRN	1320
MS-Peptides_Elutela	-----SHILSIR-----	
MS-Peptides_Elutela1b	--NRHMATTQDEVHTK-----SHILSIR-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	KSLMWFDKTPLSYTHWRAGRPTIKNEKFLAGLSTDGDWIDQTFKVIIEAVYFHQHSILAC	1380
MS-Peptides_Elutela	-SLMWFDKTPLSYTHWR-----	
MS-Peptides_Elutela1b	-SLMWFDK-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----EAVYFHQHSIL-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	KIEMVDYKKEEYNTTLQPFMPYEDGISVIQKKVTWYEALNMCSSQSGGHLASVHNQNGQLF	1440
MS-Peptides_Elutela	-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	LEDIVKRDGPPLWGLSSHDSGSESSFESWDGSTFDYIPWKQQTSPGNCVLLDPKGTWKHE	1500
MS-Peptides_Elutela	-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	

FIGURA 7 (continuación)

sp O60449 LY75_HUMAN	KCNVSKDGAICYKPTKSKKLSRLTYSSRCPAKENGSRWIQYKGHCYKSDQALHSFSEAK	1560
MS-Peptides_Elutela	-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----KKLSRLTYSS-C-----NGSRWIQYKGHCYKSDQALH-----	
sp O60449 LY75_HUMAN	KLCSKHDHSATIVSIKDEDENKFVSRILMRENNNIIMRVWLGLSQHSVDQSWLWLDGSEVT	1620
MS-Peptides_Elutela	-----HDHSATIVSIKDEDENKFVSR-----	
MS-Peptides_Elutela1b	-----HDHSATIVSIKDEDENKFVSR-----	
MS-Peptides_Elute2a	-----HDHSATIVSIKDEDENKFVSR-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	FVKWENKSKSGVGRMCLIASNETWKKVECEHGFGRVCKVPLGPDYTAIAIIVATLSIL	1680
MS-Peptides_Elutela	-----VECEHGFGR-----	
MS-Peptides_Elutela1b	-----VECEHGFGR-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	
sp O60449 LY75_HUMAN	VLMGGLIWFLQRRHRLHLAGFSSVRYAQVNEDEIMLPSFHD	1722
MS-Peptides_Elutela	-----	
MS-Peptides_Elutela1b	-----	
MS-Peptides_Elute2a	-----	
MS-Peptides_Elute2b	-----	
LC-Peptides	-----	