

Figura 28 - Secuencia de CD45 humano SEQ ID NO: 127 (SEQ ID NO: 41 de WO 2022/079199)

Met	Tyr	Leu	Trp	Leu	Lys	Leu	Leu	Ala	Phe	Gly	Phe	Ala	Phe	Leu	Asp
1				5					10					15	
Thr	Glu	Val	Phe	Val	Thr	Gly	Gln	Ser	Pro	Thr	Pro	Ser	Pro	Thr	Gly
			20					25					30		
Leu	Thr	Thr	Ala	Lys	Met	Pro	Ser	Val	Pro	Leu	Ser	Ser	Asp	Pro	Leu
			35				40					45			
Pro	Thr	His	Thr	Thr	Ala	Phe	Ser	Pro	Ala	Ser	Thr	Phe	Glu	Arg	Glu
						55					60				
Asn	Asp	Phe	Ser	Glu	Thr	Thr	Ser	Leu	Ser	Pro	Asp	Asn	Thr	Ser	
65					70				75					80	
Thr	Gln	Val	Ser	Pro	Asp	Ser	Leu	Asp	Asn	Ala	Ser	Ala	Phe	Asn	Thr
				85					90					95	
Thr	Gly	Val	Ser	Ser	Val	Gln	Thr	Pro	His	Leu	Pro	Thr	His	Ala	Asp
				100				105					110		
Ser	Gln	Thr	Pro	Ser	Ala	Gly	Thr	Asp	Thr	Gln	Thr	Phe	Ser	Gly	Ser
				115				120				125			
Ala	Ala	Asn	Ala	Lys	Leu	Asn	Pro	Thr	Pro	Gly	Ser	Asn	Ala	Ile	Ser
						135					140				
Asp	Val	Pro	Gly	Glu	Arg	Ser	Thr	Ala	Ser	Thr	Phe	Pro	Thr	Asp	Pro
145					150					155					160
Val	Ser	Pro	Leu	Thr	Thr	Leu	Ser	Leu	Ala	His	His	Ser	Ser	Ala	
				165				170						175	
Ala	Leu	Pro	Ala	Arg	Thr	Ser	Asn	Thr	Thr	Ile	Thr	Ala	Asn	Thr	Ser
				180				185					190		
Asp	Ala	Tyr	Leu	Asn	Ala	Ser	Glu	Thr	Thr	Thr	Leu	Ser	Pro	Ser	Gly
				195			200					205			
Ser	Ala	Val	Ile	Ser	Thr	Thr	Thr	Ile	Ala	Thr	Thr	Pro	Ser	Lys	Pro
					215						220				
Thr	Cys	Asp	Glu	Lys	Tyr	Ala	Asn	Ile	Thr	Val	Asp	Tyr	Leu	Tyr	Asn
225					230					235					240
Lys	Glu	Thr	Lys	Leu	Phe	Thr	Ala	Lys	Leu	Asn	Val	Asn	Glu	Asn	Val
				245					250					255	
Glu	Cys	Gly	Asn	Asn	Thr	Cys	Thr	Asn	Asn	Glu	Val	His	Asn	Leu	Thr
				260				265					270		
Glu	Cys	Lys	Asn	Ala	Ser	Val	Ser	Ile	Ser	His	Asn	Ser	Cys	Thr	Ala
				275			280					285			
Pro	Asp	Lys	Thr	Leu	Ile	Leu	Asp	Val	Pro	Pro	Gly	Val	Glu	Lys	Phe
					295						300				
Gln	Leu	His	Asp	Cys	Thr	Gln	Val	Glu	Lys	Ala	Asp	Thr	Thr	Ile	Cys
305					310					315					320
Leu	Lys	Trp	Lys	Asn	Ile	Glu	Thr	Phe	Thr	Cys	Asp	Thr	Gln	Asn	Ile
				325					330					335	
Thr	Tyr	Arg	Phe	Gln	Cys	Gly	Asn	Met	Ile	Phe	Asp	Asn	Lys	Glu	Ile
				340				345					350		
Lys	Leu	Glu	Asn	Leu	Glu	Pro	Glu	His	Glu	Tyr	Lys	Cys	Asp	Ser	Glu
				355			360					365			
Ile	Leu	Tyr	Asn	Asn	His	Lys	Phe	Thr	Asn	Ala	Ser	Lys	Ile	Ile	Lys
				375							380				
Thr	Asp	Phe	Gly	Ser	Pro	Gly	Glu	Pro	Gln	Ile	Phe	Cys	Arg	Ser	
385					390				395					400	
Glu	Ala	Ala	His	Gln	Gly	Val	Ile	Thr	Trp	Asn	Pro	Pro	Gln	Arg	Ser
				405					410					415	
Phe	His	Asn	Phe	Thr	Leu	Cys	Tyr	Ile	Lys	Glu	Thr	Glu	Lys	Asp	Cys
				420				425					430		
Leu	Asn	Leu	Asp	Lys	Asn	Leu	Ile	Lys	Tyr	Asp	Leu	Gln	Asn	Leu	Lys
				435			440					445			
Pro	Tyr	Thr	Lys	Tyr	Val	Leu	Ser	Leu	His	Ala	Tyr	Ile	Ile	Ala	Lys
					455						460				
Val	Gln	Arg	Asn	Gly	Ser	Ala	Ala	Met	Cys	His	Phe	Thr	Thr	Lys	Ser
465					470					475					480
Ala	Pro	Pro	Ser	Gln	Val	Trp	Asn	Met	Thr	Val	Ser	Met	Thr	Ser	Asp
				485					490					495	
Asn	Ser	Met	His	Val	Lys	Cys	Arg	Pro	Pro	Arg	Asp	Arg	Asn	Gly	Pro
				500				505					510		
His	Glu	Arg	Tyr	His	Leu	Glu	Val	Glu	Ala	Gly	Asn	Thr	Leu	Val	Arg
				515			520					525			
Asn	Glu	Ser	His	Lys	Asn	Cys	Asp	Phe	Arg	Val	Lys	Asp	Leu	Gln	Tyr
				530			535				540				
Ser	Thr	Asp	Tyr	Thr	Phe	Lys	Ala	Tyr	Phe	His	Asn	Gly	Asp	Tyr	Pro
545					550					555					560