

SIRPBeta:	Isoform 1		1	MPVPASWPHLPSPFLMTLLGLRTGVAGEDELQVIOPEKSVSVAAGESATLRCAMTSLI	60
	Isoform 3		1	MPVPASWPHLPSPFLMTLLGLRTGVAGEEELQVIOPEKSVSVAAGESATLHCTVTSLI	60
Isoform 1	Isoform 3		61	PVGPIMWFRGAGAGRELIYNQKEGHFPRVTTVSELTKRNLDFSISISNITPADAGTYYC	120
	Isoform 3		61	PVGPIMWFRGAGAGRELIYNQKEGHFPRVTTVSDLTKNWMDFSIRISNITPADAGTYYC	120
Isoform 1	Isoform 3		121	VKFRKGSPODVEFKSGAGTELSVRAKPSAPVWSGPAVRATPEHTVSVFTCESHGFSPRDIT	180
	Isoform 3		121	VKFRKGSPODVEFKSGAGTELSVRAKPSAPVWSGPAARATPQHTVSVFTCESHGFSPRDIT	180
Isoform 1	Isoform 3		181	LKWFKNIGNELSDFQTNVDPAGDSVSYSIHSTARVWLTRGDVHSQVCEIAHITLQGDPLR	240
	Isoform 3		181	LKWFKNIGNELSDFQTNVDPAGDSVSYSIHSTAKVWLTRGDVHSQVCEIAHITLQGDPLR	240
Isoform 1	Isoform 3		241	GTANLSEAIRVPPTLEVTQOPMRAENQANVTCQVSNFYPRGLQLTWLENGNVSRTEIAT	300
	Isoform 3		241	GTANLSEAIRVPPTLEVTQOPMRAENQANVTCQVRKLYPQRLQLTWLENGNVSRTEIAT	300
Isoform 1	Isoform 3		301	LIENKDGTYNMMSWLLVNTCAHRDDWLTCCQVEHDGQQAQVSKSYALEISAHQKEHGS	360
	Isoform 3		301	LIENKDGTYNMMSWLLVNVSAHRDDVCLTCQVEHDGQQAQVSKSHDLKVSAPKPEQGS	360
Isoform 1	Isoform 3		361	HEAALAPTAPLLVALLGPKLLLVGVSAIYICWKQKA	398
	Isoform 3		361	PGPALASAAPLLIAFLGPKLLLVGVSVIYVYWKQKA	398

FIGURE 1A

[illegible]

FIGURE 1B

HUMAN	000241	1	MPVPASWPHLPSPFLMTLLGLRLTGVAGEDELQVIQPEKSVSVAAGESATLRCAMTSLI	60
MOUSE	Q88FX8	1	MLLLDAWTHIPHCVLILLGLKG--AAVRELKVIQPVKSFVVGAGGSATLNCVTYLL	58
			* ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ;	
HUMAN	000241	61	PVGPIWFRGAGAGRELIYNQKEGHFPRVTTVSELTKRNNLDFSISNITPADAGTYYC	120
MOUSE	Q88FX8	59	PVGPIKWYRGVQSRLLIYPFTGEYFPRIITSVDVKRRNLDIFSIRISNVTPADSGTYYC	118
			***** * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ;	
HUMAN	000241	121	VKFRKGGSP-DDVEFKSGAGTELSVRAKPSAPVWSGPAVRATPEHTVSFTCESHGFSPRDI	179
MOUSE	Q88FX8	119	VKFQRGSEPDIEIQSGGGTELSVFAKPSPPMVSGPAARAVPQQTVTFTCRSHGFFPQNL	178
			***** * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ;	
HUMAN	000241	180	TLKWFKNGNELSDFTNVDPAGDSVSYSIHSTARVVLTRGDVHSQVICEIAHITLQGDPL	239
MOUSE	Q88FX8	179	TLKWFKNGNELSHLETSVEPEETSVSYRVSSIVQVWLEPRDVRSQIICEVDHVTLDRAPL	238
			***** * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ;	
HUMAN	000241	240	RGIANLSEAIRVPPTLEVTQQPMRAENQANVTQVSNFYPRGLQTLWLENGNVSRTETAS	299
MOUSE	Q88FX8	239	RGIAHISEIIQVPPPTLEISQQPTMVMNVINVTQIQKFYPRRFQVTLWLENGNISRRVPE	298
			***** * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ;	
HUMAN	000241	300	TLIENKDGTYNWMWSLLVNTCAHRDDVVLTCQVEHDGQQAQVSKSYALEISAHQKEHGSDI	359
MOUSE	Q88FX8	299	THIVNKDGTYNWMISWLLVNISALEENMVVTCQVEHDGQAEVIEHTVWVTEHQRVKGTST	358
			***** * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ;	
HUMAN	000241	360	THEAALAPTAPLLVALLGPKLLLVVGVSAIYICWKQKA	398
MOUSE	Q88FX8	359	MSELKTAGIAKIPVAVLLGSKILLIAATVIYMRKKQNA	397
			* * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ; * * ;	

FIGURE 2

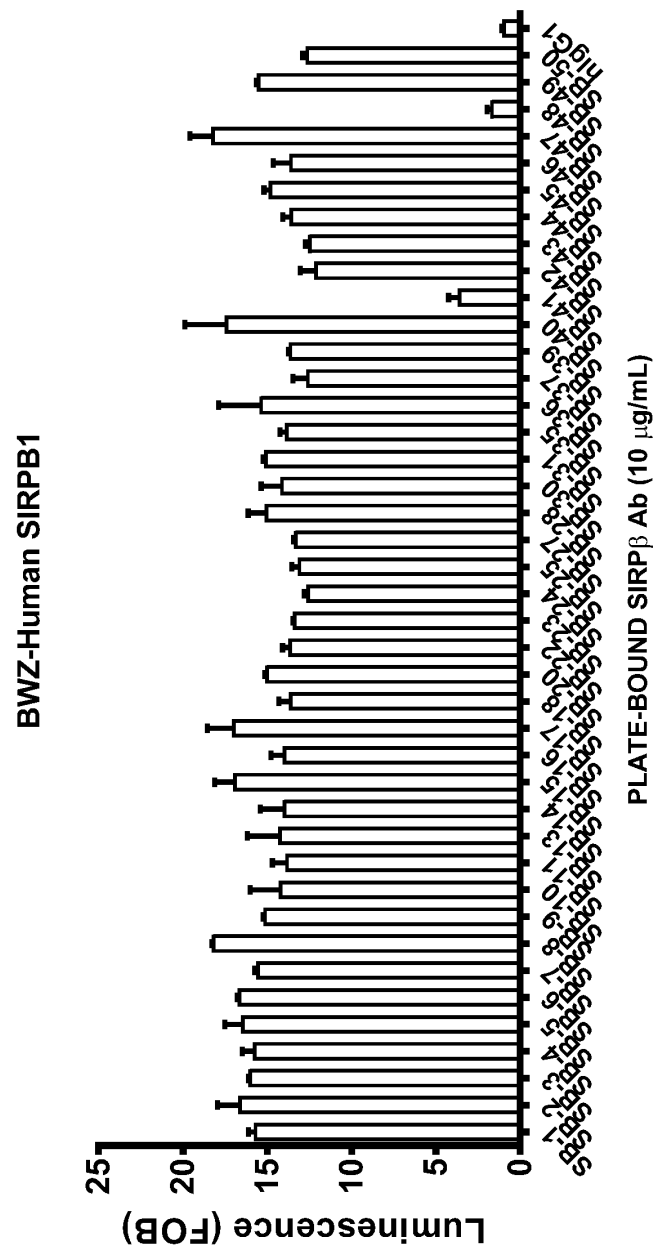
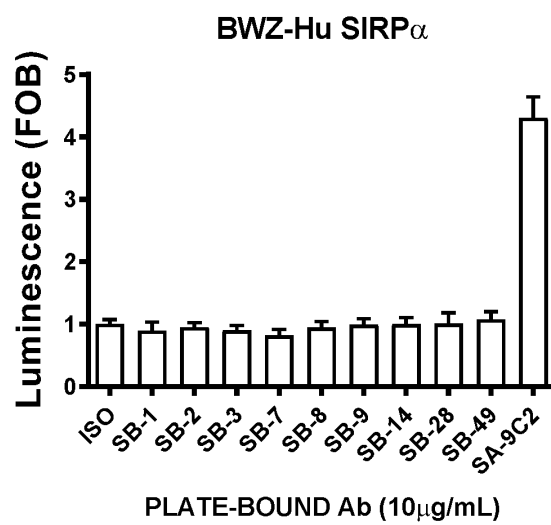


FIGURE 3A

**FIGURE 3B**

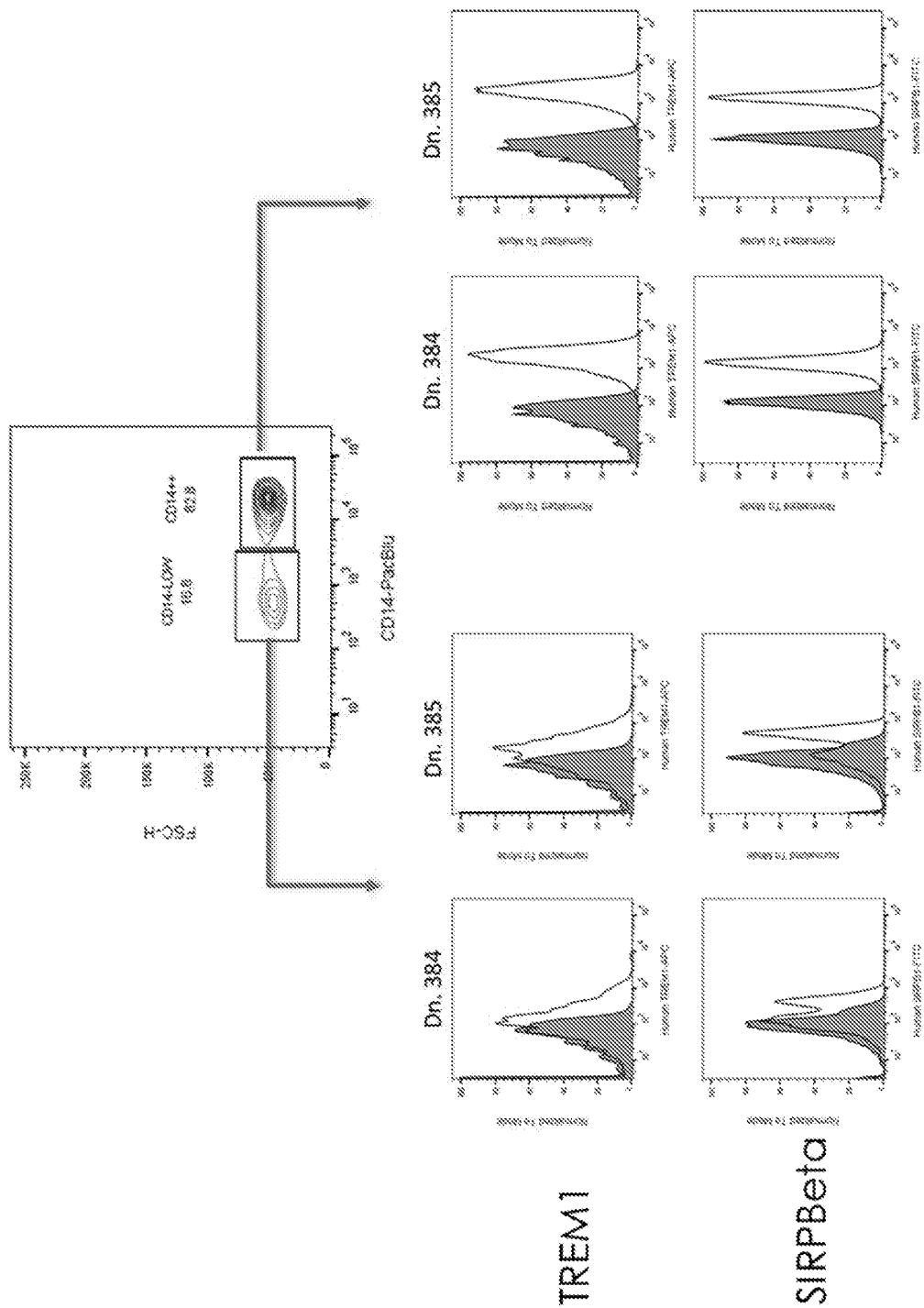


FIGURE 4A

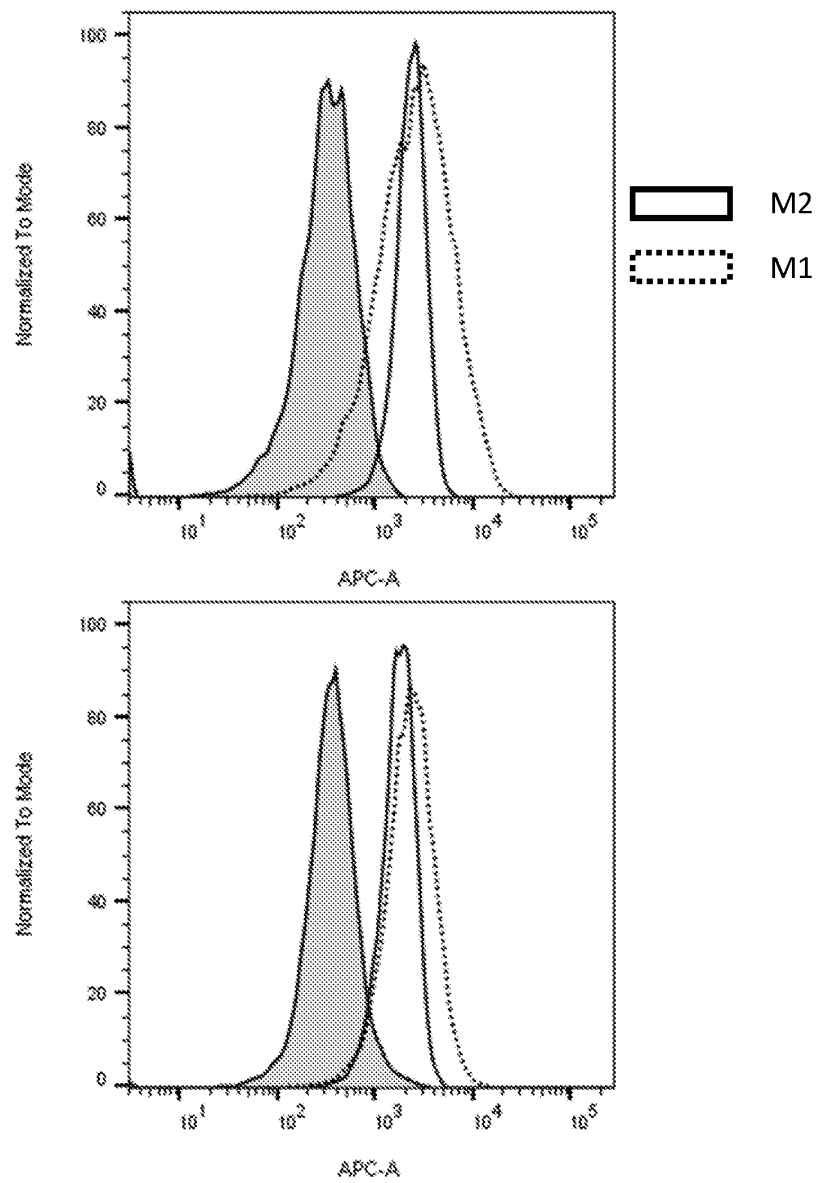


FIGURE 4B

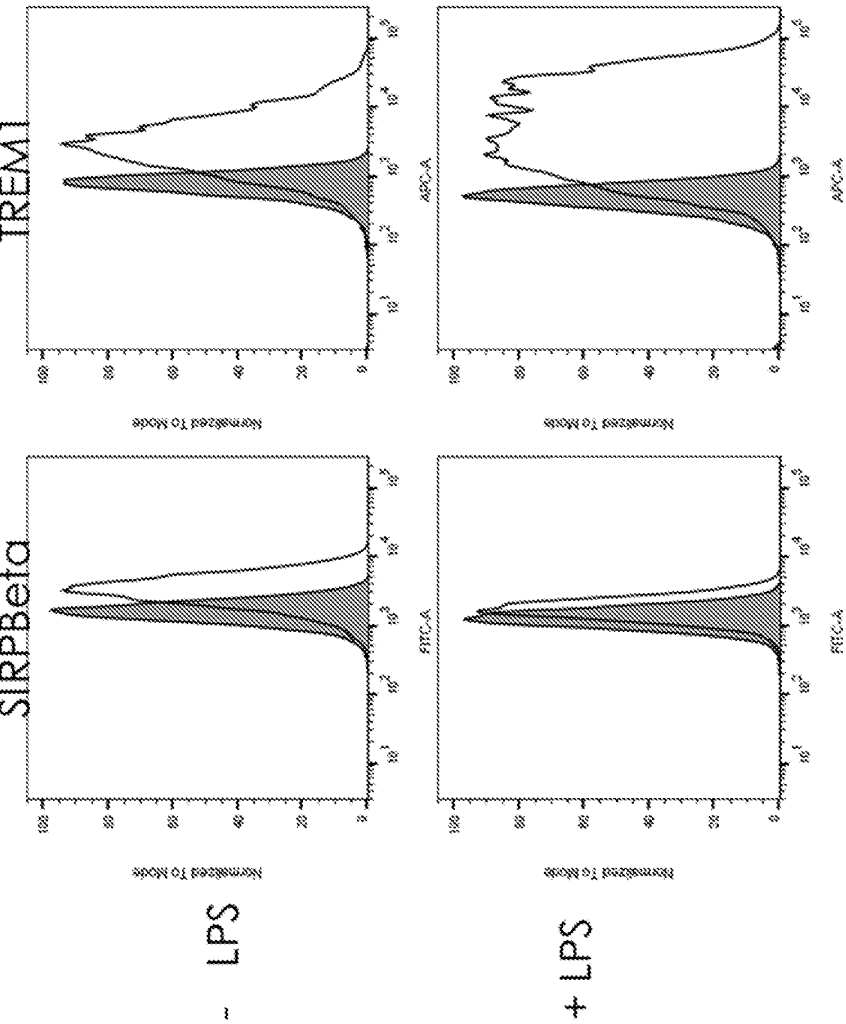
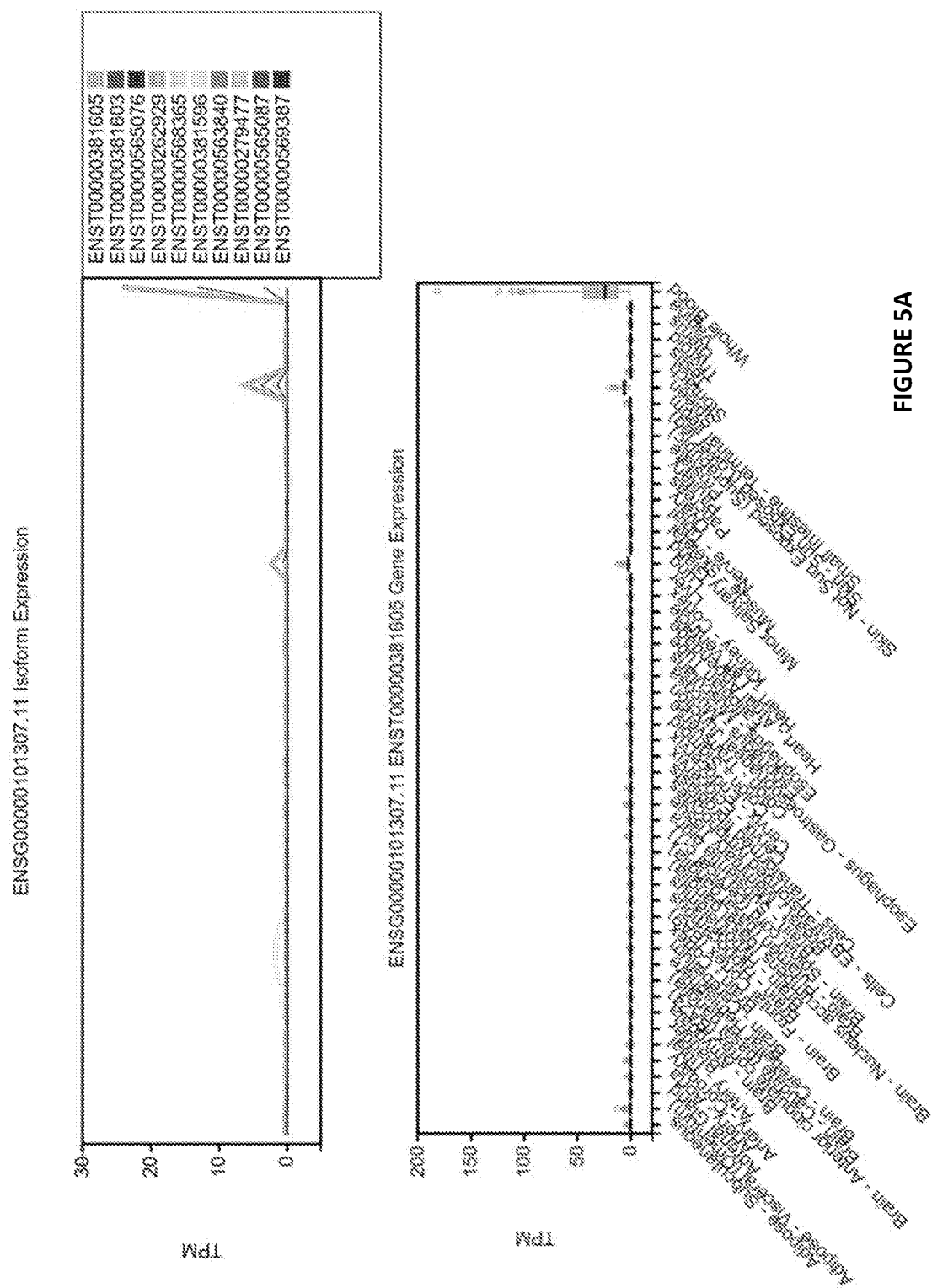
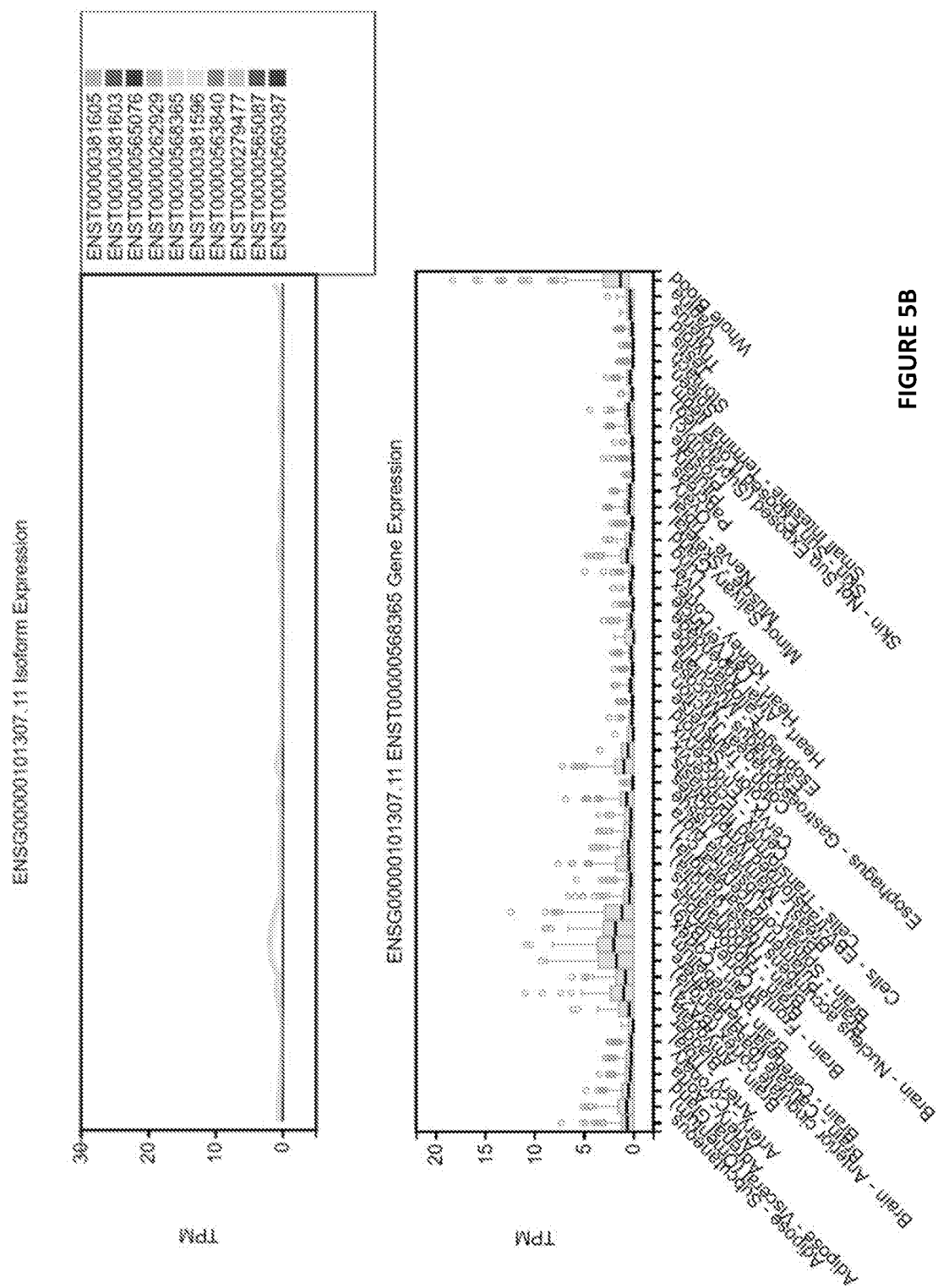


FIGURE 4C





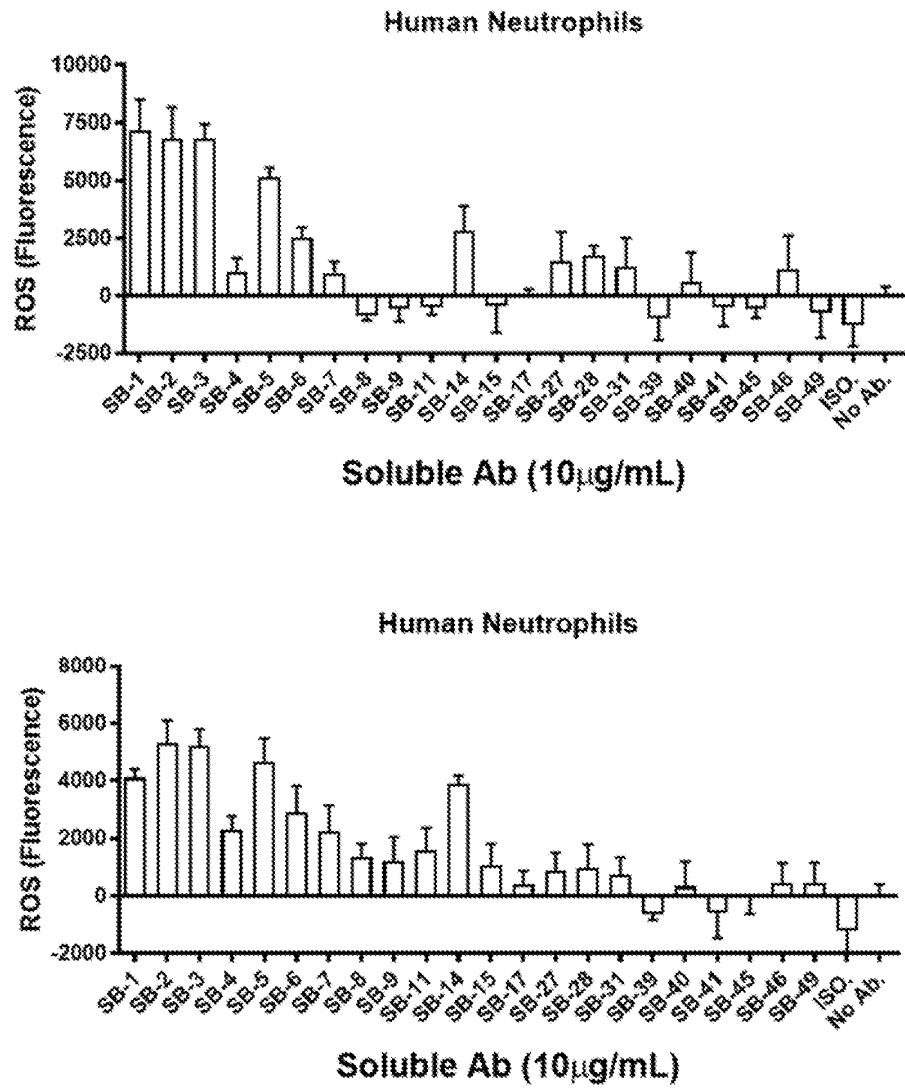


FIGURE 6A

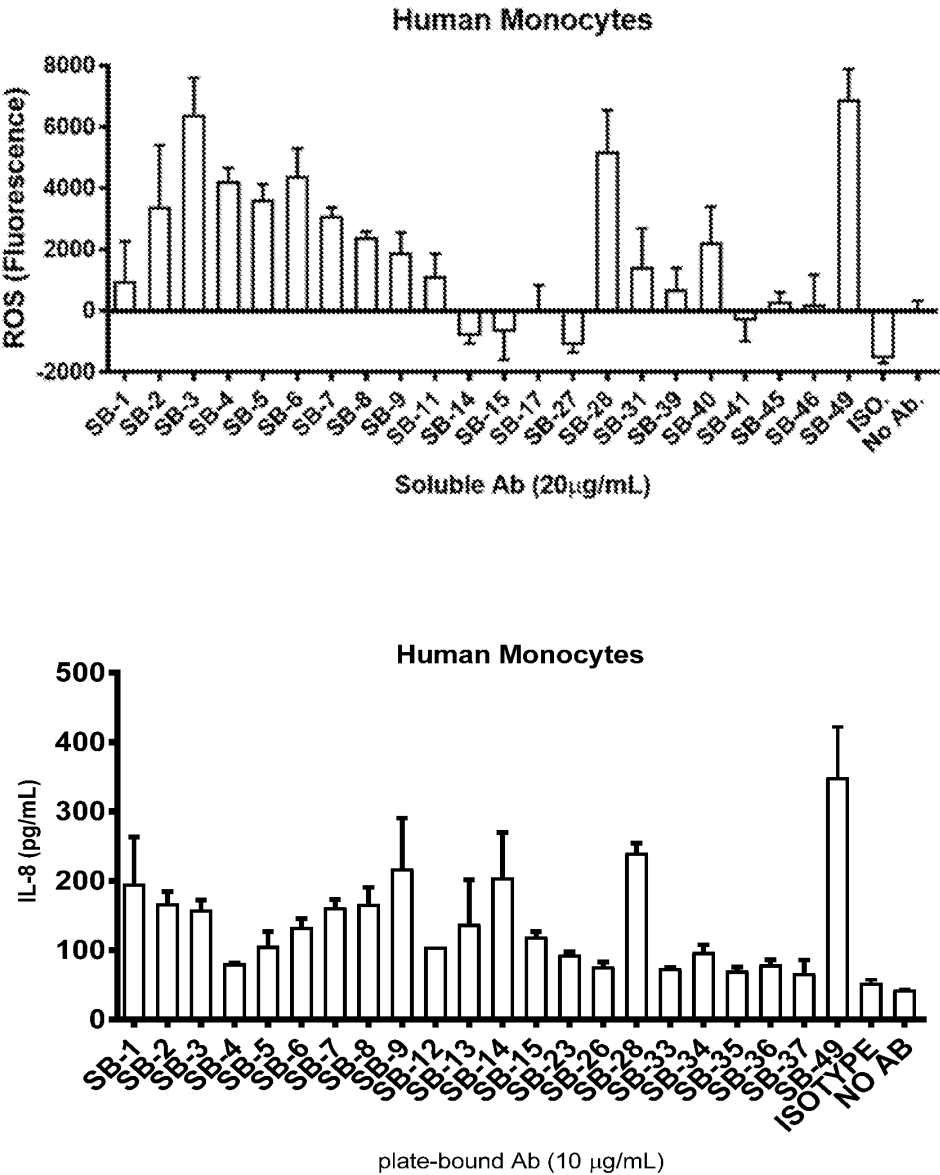


FIGURE 6B

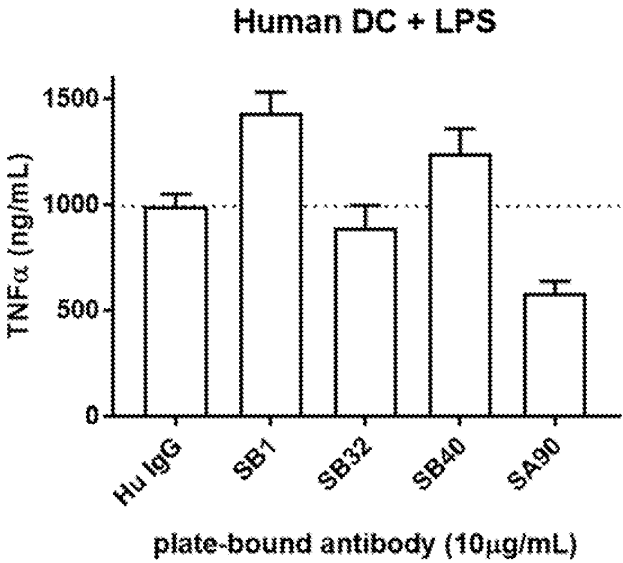
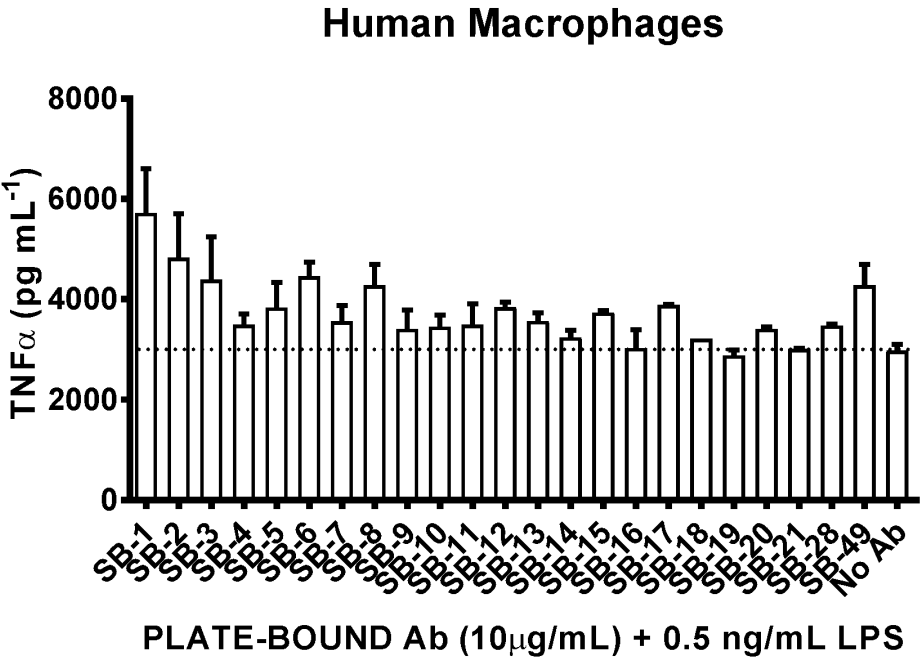


FIGURE 6C

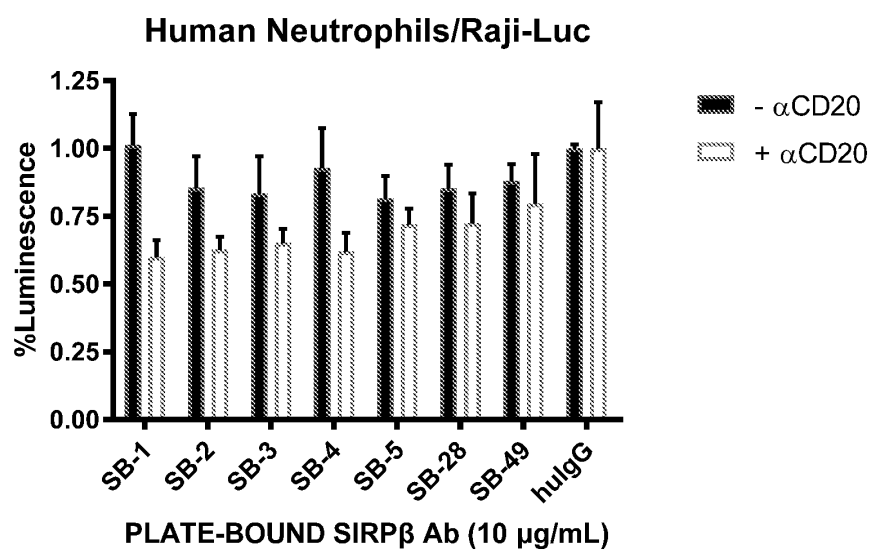


FIGURE 7

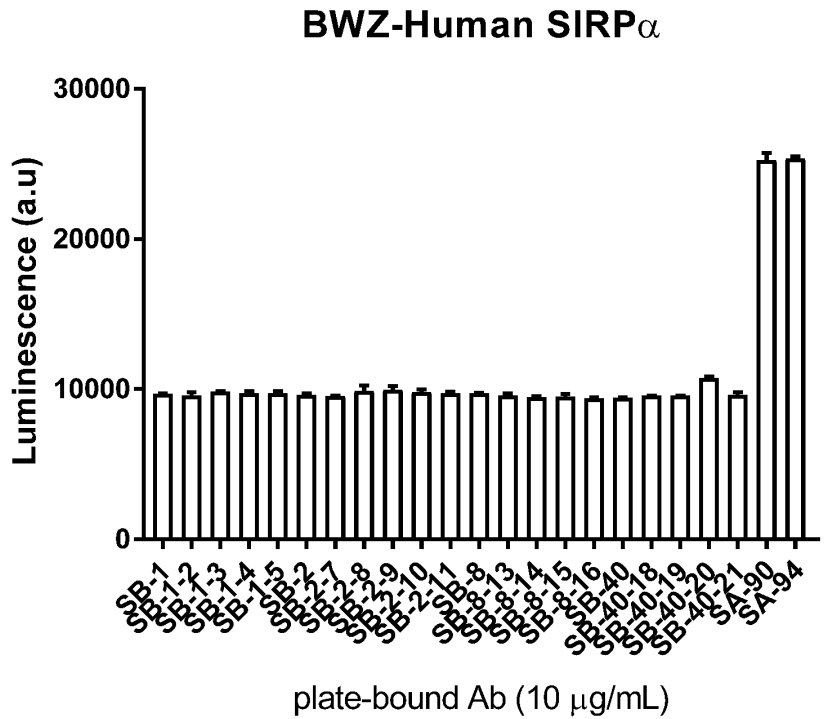
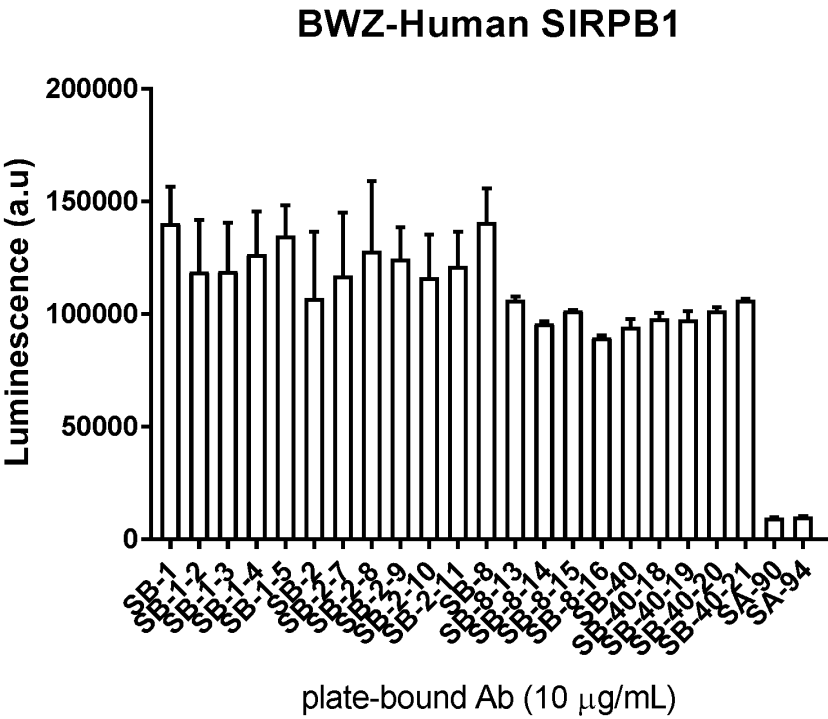


FIGURE 8

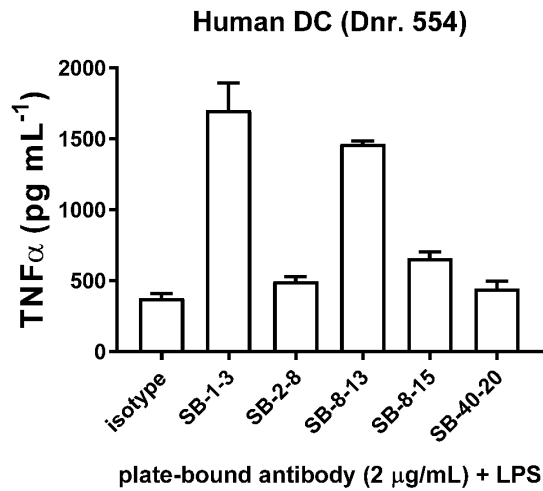
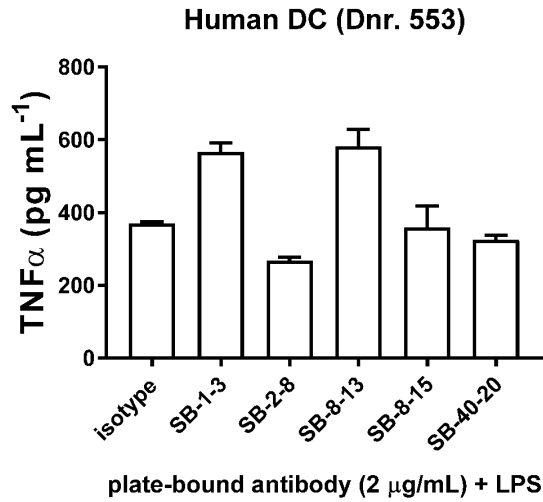
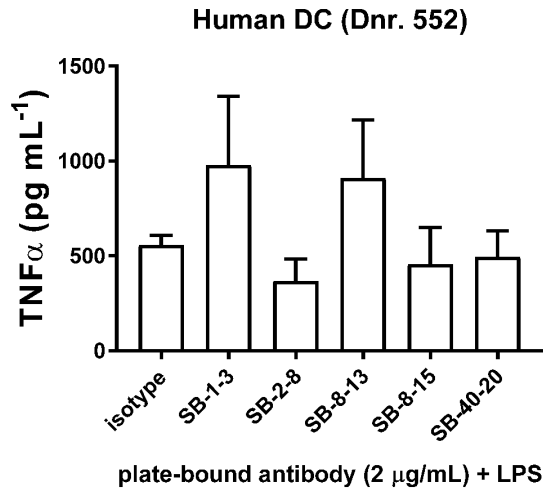


FIGURE 9

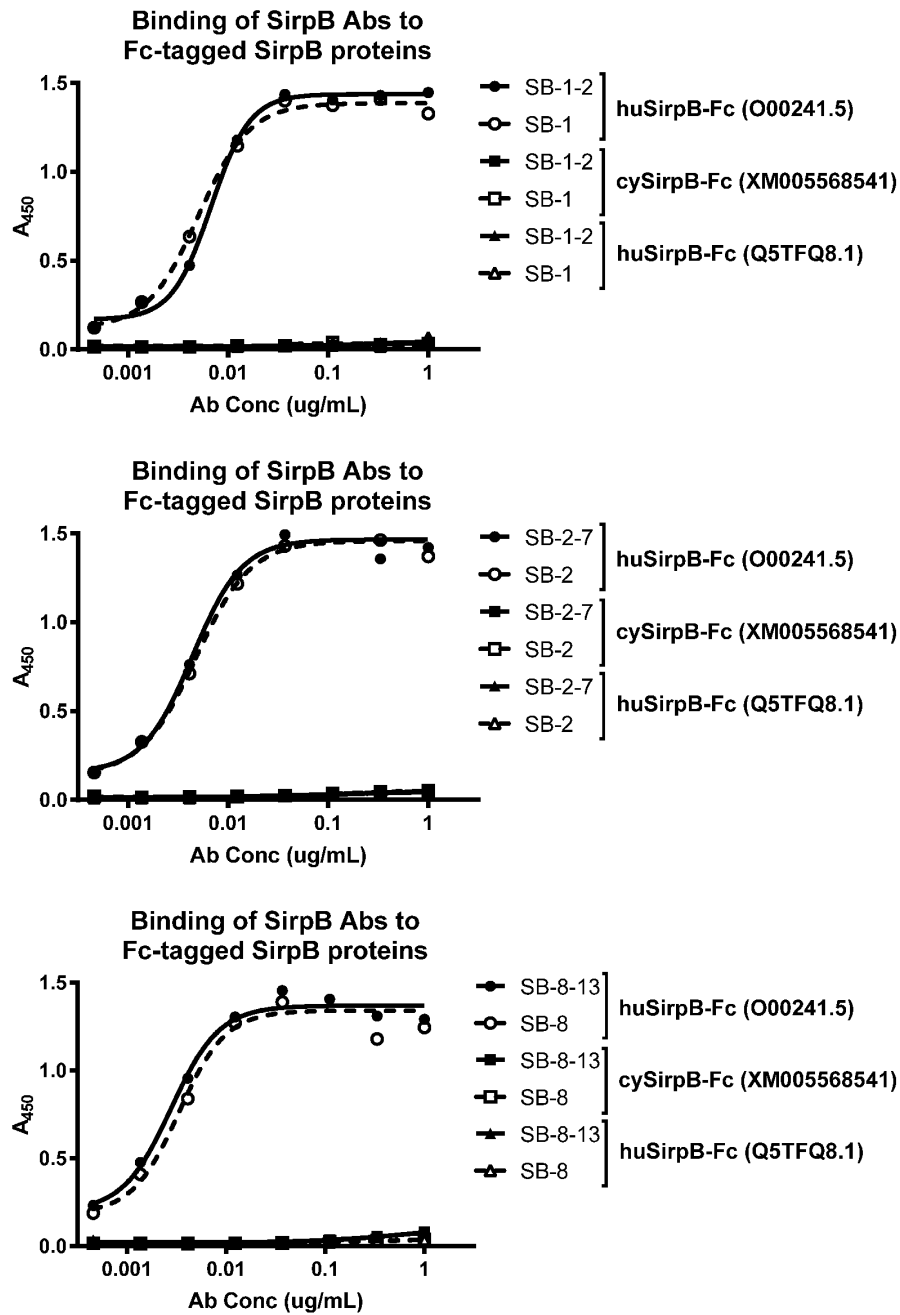


FIGURE 10A

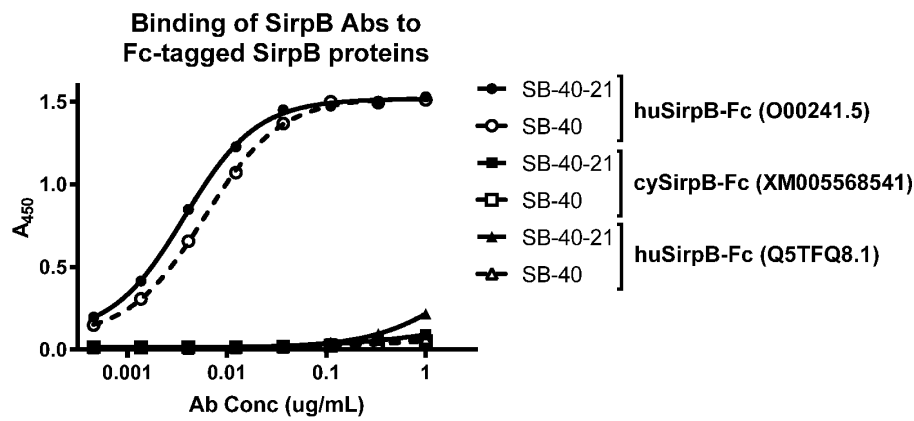


FIGURE 10B

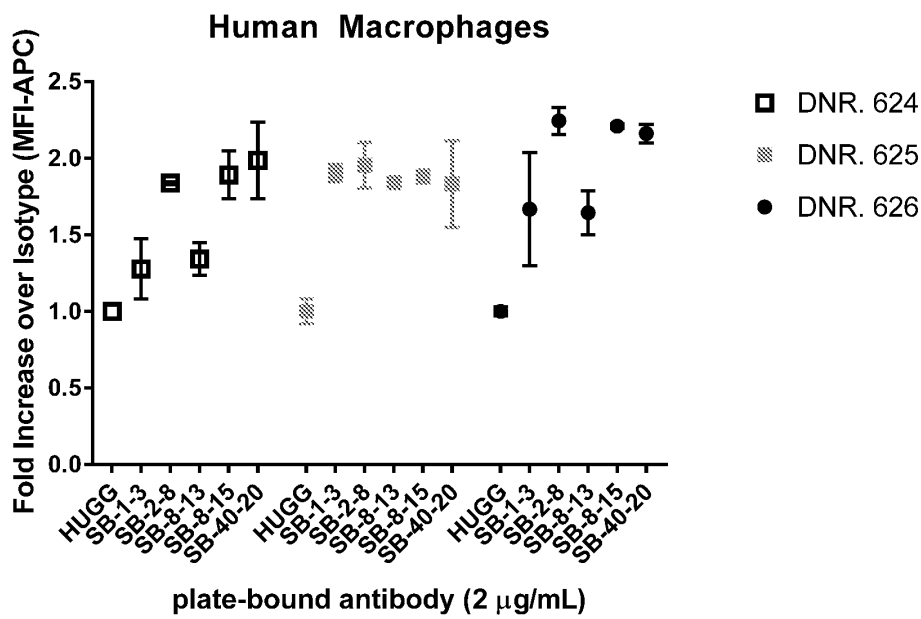


FIGURE 11A

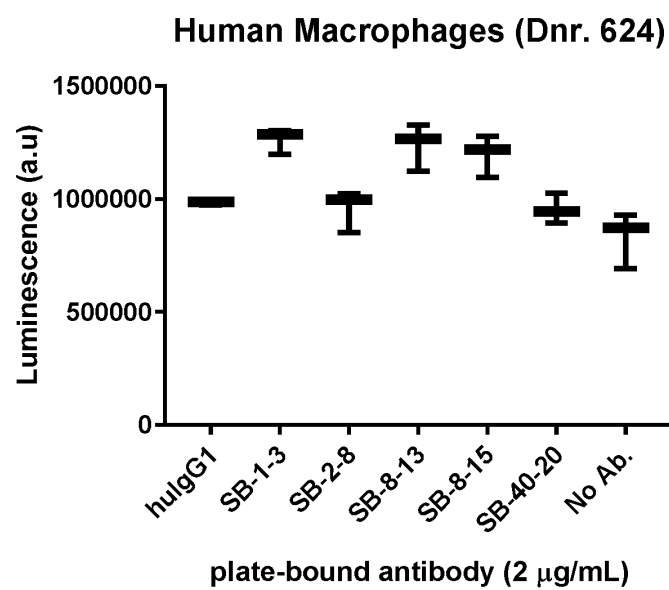


FIGURE 11B

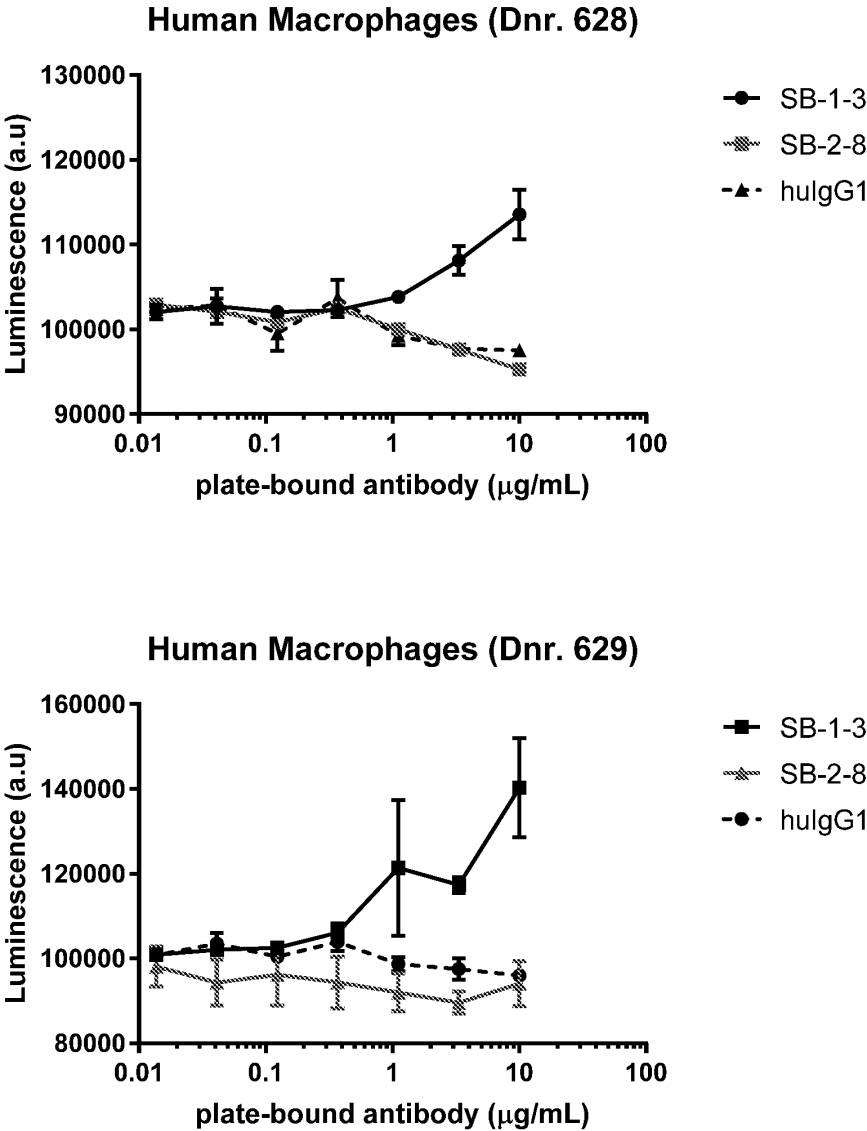


FIGURE 12

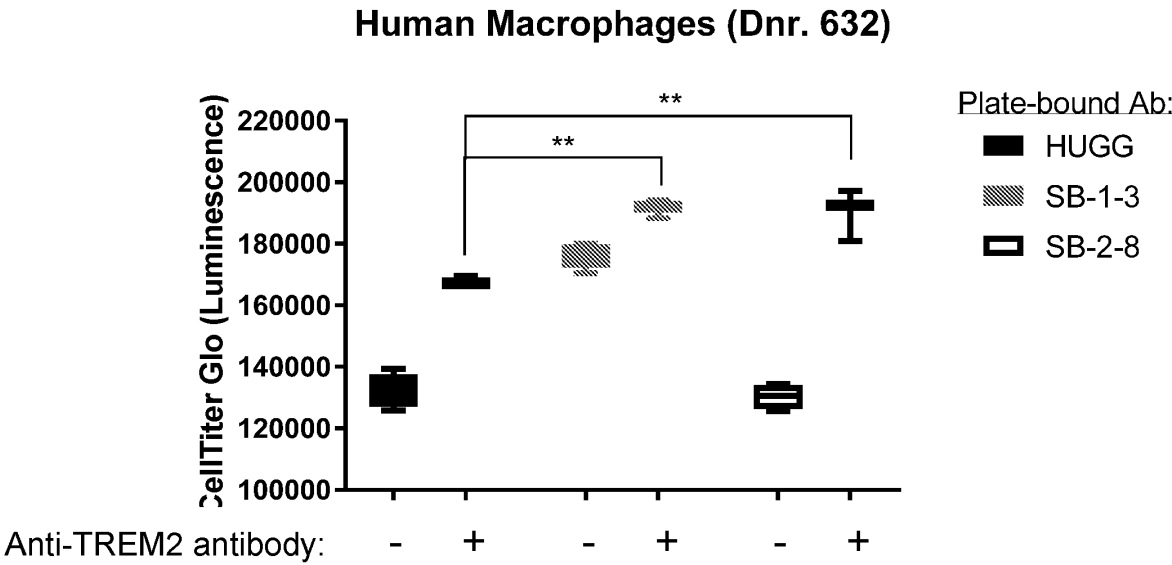


FIGURE 13

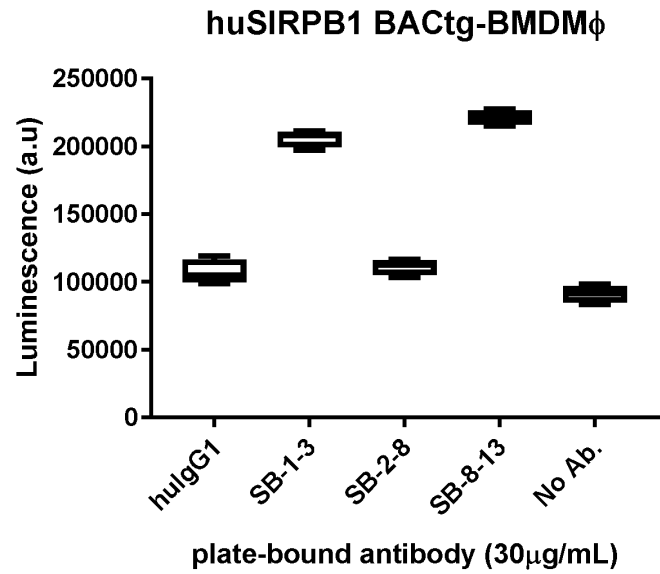


FIGURE 14A

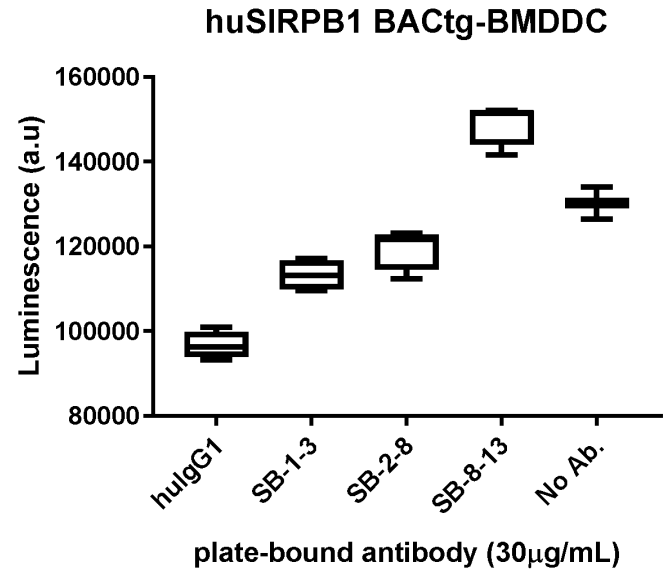


FIGURE 14B

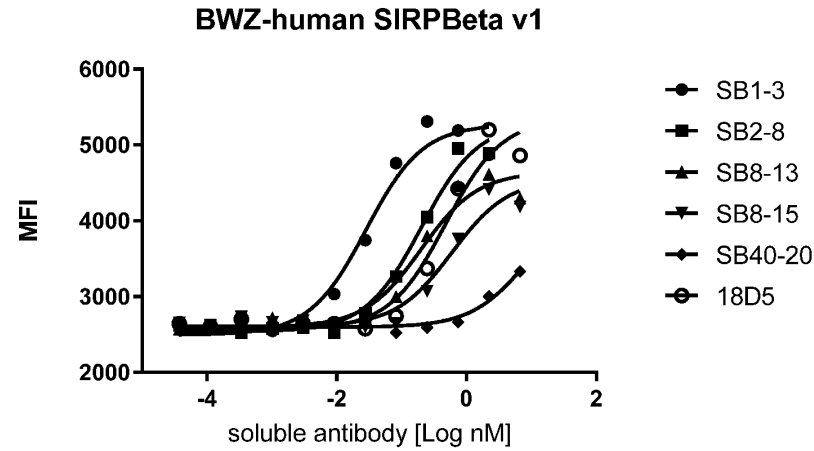


FIGURE 15A

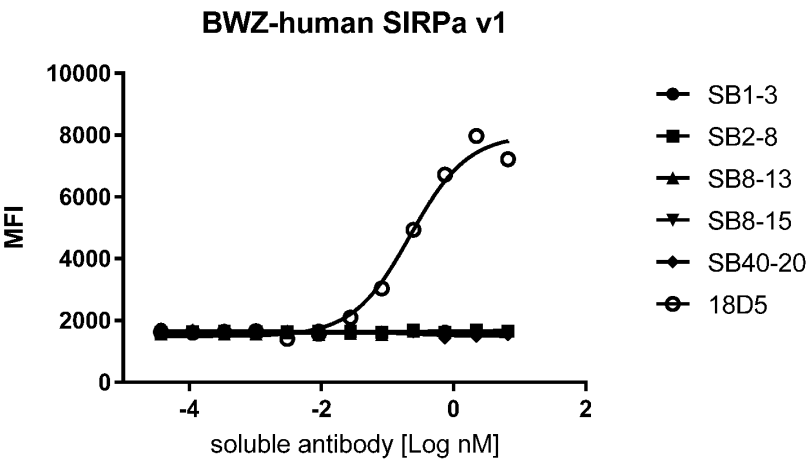


FIGURE 15B

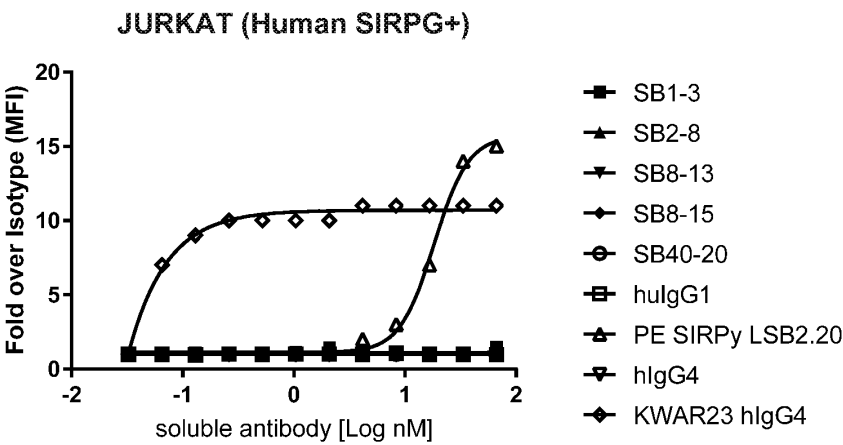


FIGURE 15C