

Isoform 1	000241	1	MPVPASWPHLPSPFLLMTLLGRLTGVAGEDELQVIQPEKSVSVAAGESATLRCAMTSLI	60
Isoform 3	Q5TFQ8	1	MPVPASWPHLPSPFLLMTLLGRLTGVAGEEELQVIQPKSISVAAGESATLHCTVTSLI	60
Isoform 1	000241	61	PVGPIWFRGAGAGRELIIYNQKEGHEFPRVTTVSELTKRNNLDFSISISNIITPADAGTYYC	120
Isoform 3	Q5TFQ8	61	PVGPIQWFRGAGPGRELIIYNQKEGHEFPRVTTVSDLTNRNMDFSIRISNIITPADAGTYYC	120
Isoform 1	000241	121	VKFRKGGSPDDVEFKSGAGTELSVRAKPSAPVWVGPAVRATPEHTVTSFTCESHGFSPRDIT	180
Isoform 3	Q5TFQ8	121	VKFRKGGSPDHVEFKSGAGTELSVRAKPSAPVWVGPAARATPQHTVTSFTCESHGFSPRDIT	180
Isoform 1	000241	181	LKWFKNGNELSDFQTNVDPAGDSVSYSIHSTARVVLTRGDVHSQVCEIAHITLQGDPLR	240
Isoform 3	Q5TFQ8	181	LKWFKNGNELSDFQTNVDPAGDSVSYSIHSTAKVVLTREDVHSQVCEVAHVTLQGDPLR	240
Isoform 1	000241	241	GTANLSEAIRVPPITLEVTQQPMRAENQANVTCQVSNFYPRGLQLTWLENGNVSRTEYAST	300
Isoform 3	Q5TFQ8	241	GTANLSEIRVPPITLEVTQQPVRAENQVNVTCQVRKFPYQRLQLTWLENGNVSRTEYAST	300
Isoform 1	000241	301	LIEKDGTYNMMSWLLVNTCAHRDDVWLTCQVEHDGQQAQVSKSYALEISAHQKEHGSDIT	360
Isoform 3	Q5TFQ8	301	LTEKDGTYNMMSWLLVNVSAHRDDVKLTCQVEHDGQPAVSKSHDLKVSAHPKEQGSNTA	360
Isoform 1	000241	361	HEAALAPTAPLLVALLGPKLLLVWGVSAIYICWKQKA	398
Isoform 3	Q5TFQ8	361	PGPALASAAPLLIAFLGPKVLLVWGVSVIYVWVKQKA	398

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HUMAN	000241	1	MPVPASWPHLPSPFLLMTLLGLRTGVADEDELQVIOPEKSVSVAAGESATLRCAMTSLI	60
MOUSE	Q8BFX8	1	MLLLDANTHIPCVCVLLILLGLKG--AAVRELKVIQPVKSFVAGGSATLNCVTYLL	58
			* ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ;	
HUMAN	000241	61	PVGPIWFRGAGAGRELIYNQKEGHFPRVTTVSELTKRNWDFSSISNIITPADAGTYYC	120
MOUSE	Q8BFX8	59	PVGPIKWYRGVQSRLIYPFTGEYFPRIITSVDVKRKNLDFSIKISNVTADSGTYYC	118
			***** * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ;	
HUMAN	000241	121	VKFRKGSP-DDVEFKSGAGTELSVRAKPSAPVWSGPAVRAITPEHTVSTCESHGFSPRDI	179
MOUSE	Q8BFX8	119	VKFQGSSEPDIEIQSGGGTELSVFAKPSPPMVSGPAARAVPQQTVFTCRSHGFFPQNL	178
			***** * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ;	
HUMAN	000241	180	TLKWFKNELSDFTNVDPAGDSVSYSIHSTARVWLTRGDVHSQVCEIAHITLQGDPL	239
MOUSE	Q8BFX8	179	TLKWFKNELSHLETSVEPEETSVYRVSSIVQVLEPRDVRSQIICEVDHVTLDRAPL	238
			***** * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ;	
HUMAN	000241	240	RGIANLSEAIRVPPTLEVTQQPMRAENQANVTQVSNFYPRGLQLTWLENGNVSRTETAS	299
MOUSE	Q8BFX8	239	RGIAHISEIIQVPPTLEISQQTVMVNVINVTQIQKFYPRFQVTWLENGNISRREVPF	298
			***** * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ;	
HUMAN	000241	300	TLIENKDGTYNNMSWLLVNTCAHRDDVVLTCQVEHDGQQAQVSKSYALEISAHQKEHGSDI	359
MOUSE	Q8BFX8	299	THIVNKDGTYNNISWLLVNISALEENMVVTCQVEHDGQAEVIEHTVVVTEHQRVKGTST	358
			***** * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ;	
HUMAN	000241	360	THEAALAPTAPLLVALLGPKLLLVWGVSAIYICWKQKA	398
MOUSE	Q8BFX8	359	MSELKTAGIAKIPVAVLLGSKILLIAATVIYMRKKQNA	397
			* ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ; * ;	

FIGURE 2

BWZ-Human SIRPB1

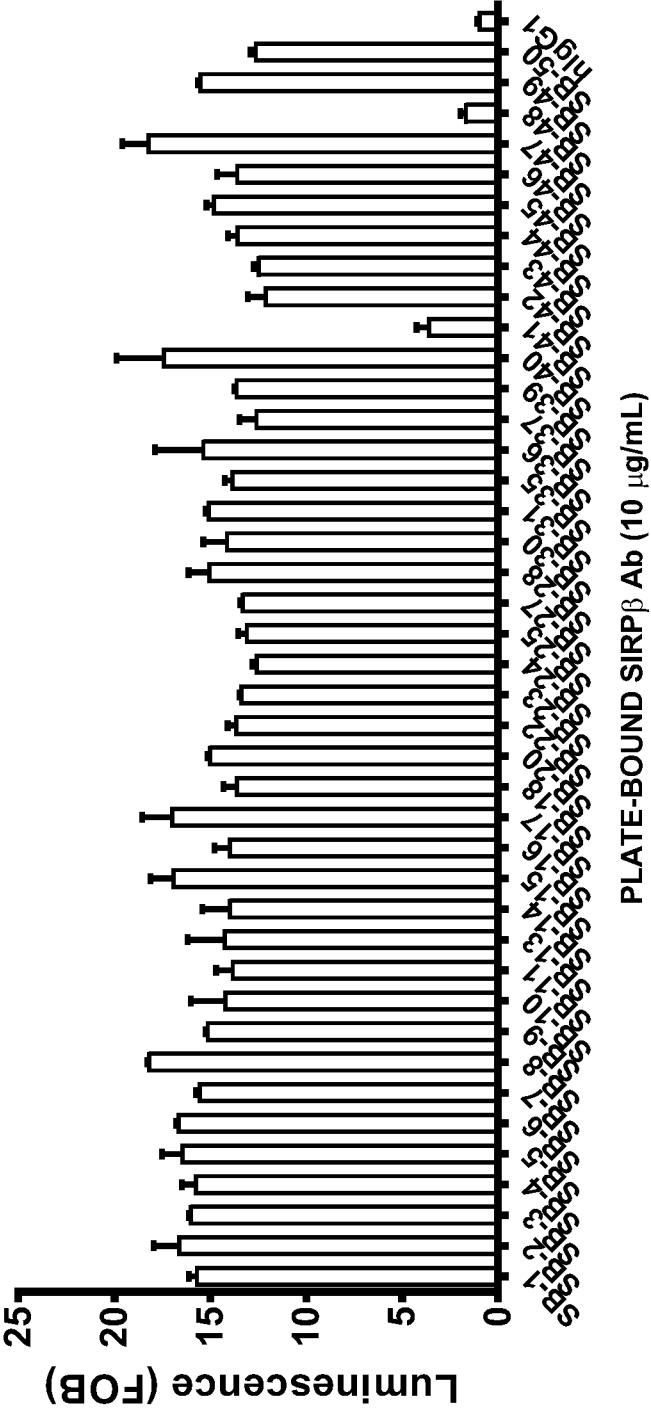


FIGURE 3A

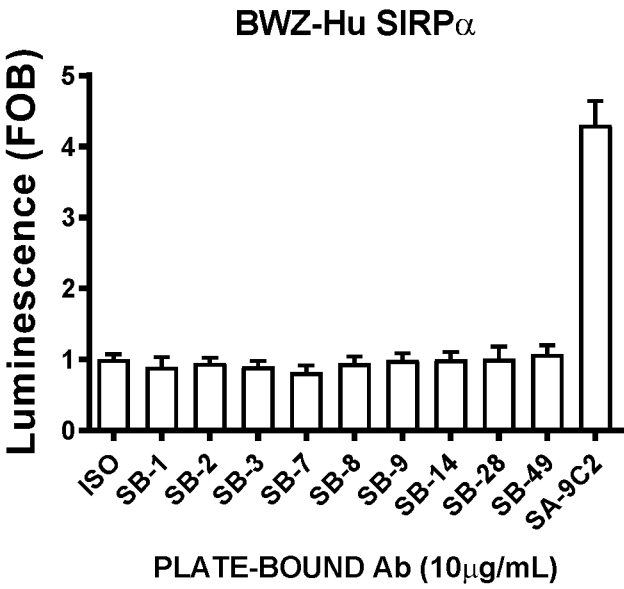


FIGURE 3B

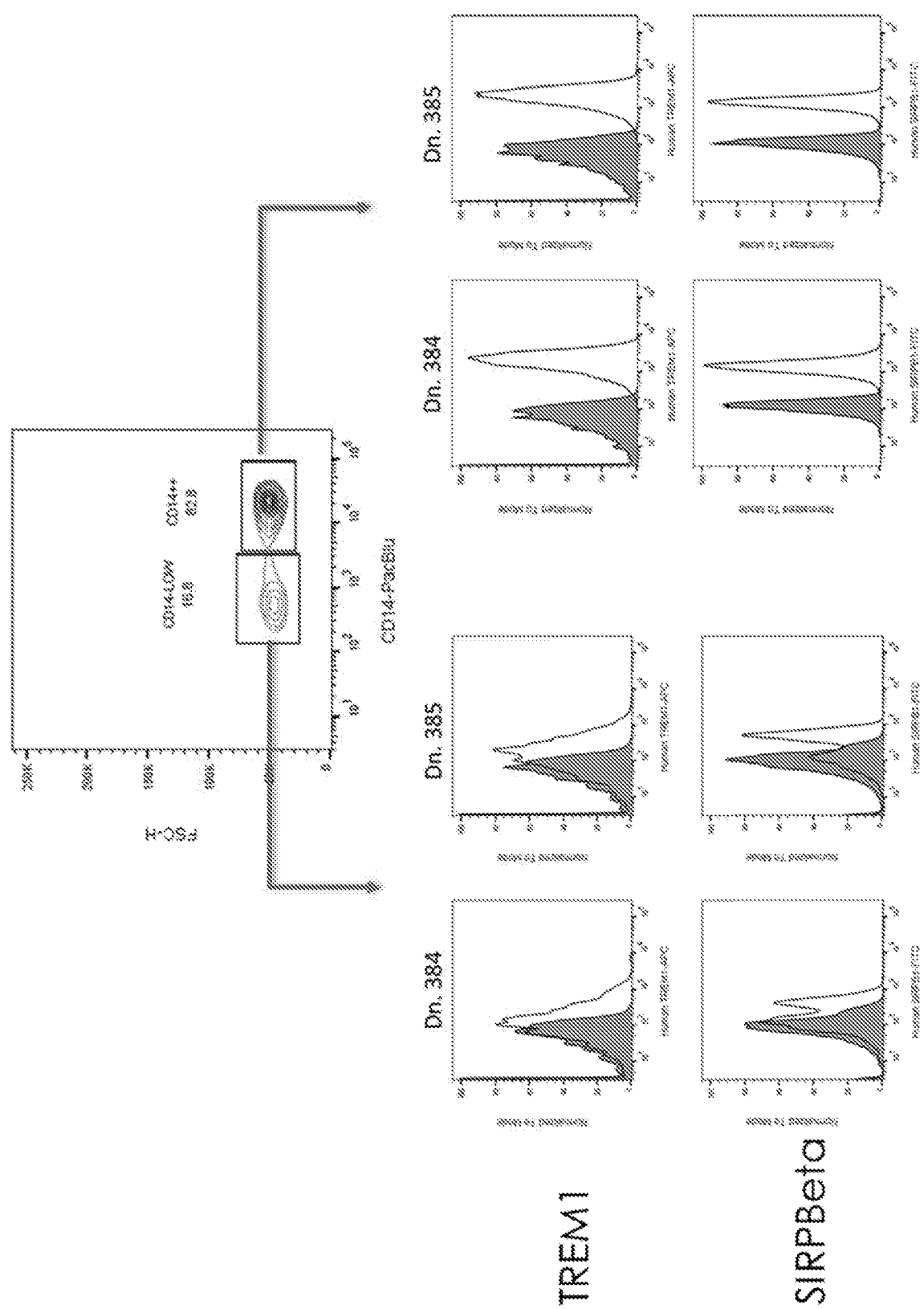


FIGURE 4A

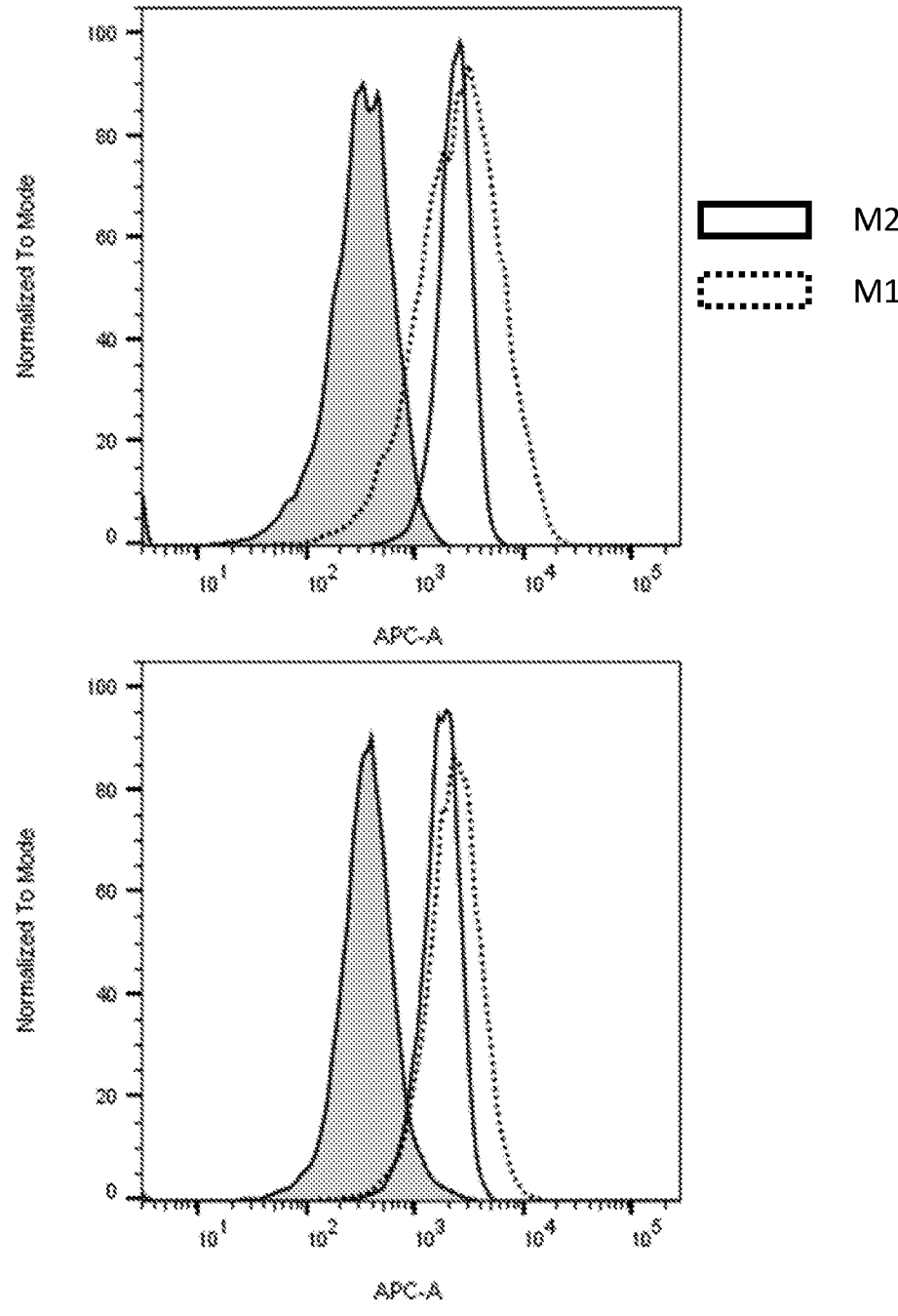


FIGURE 4B

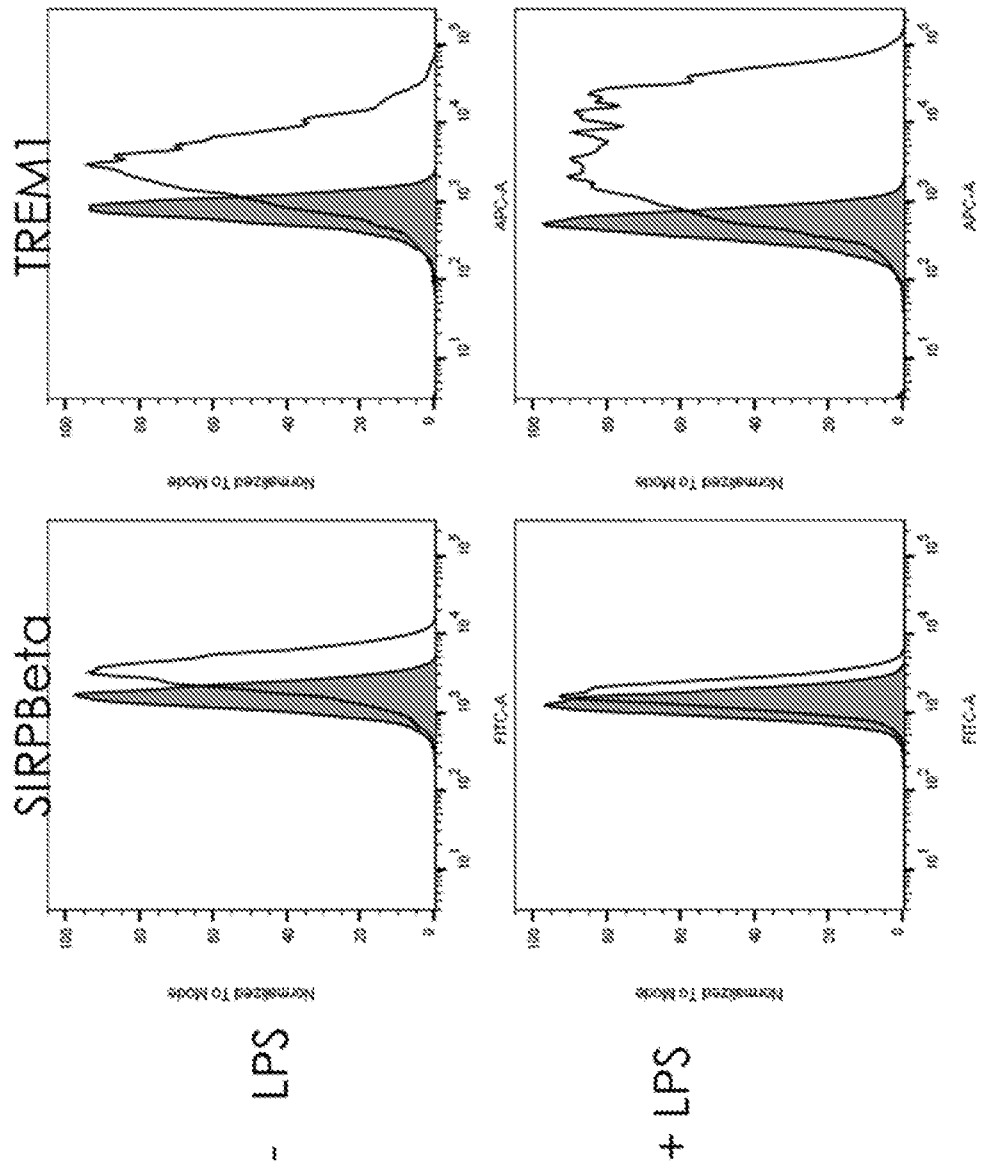
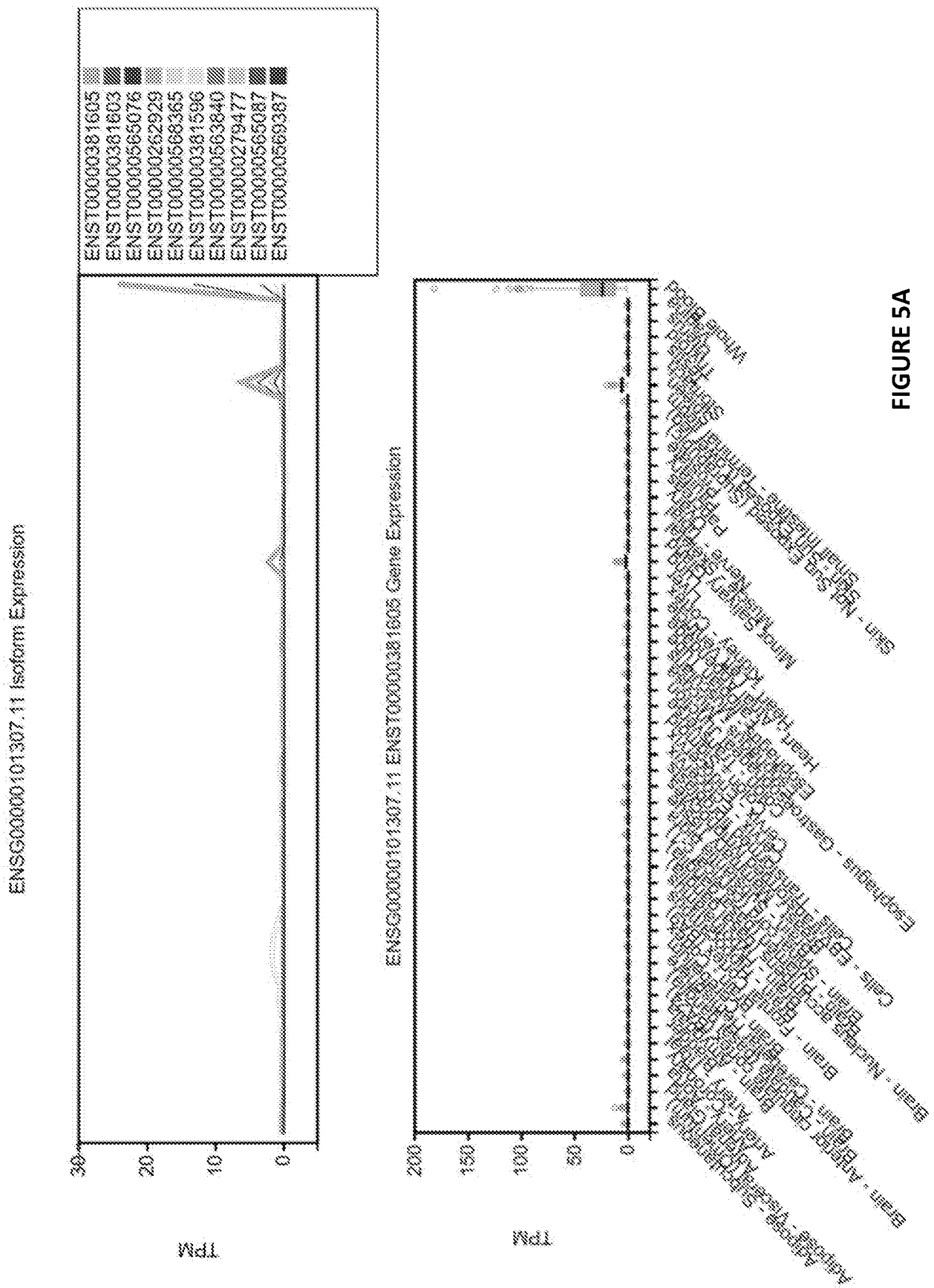
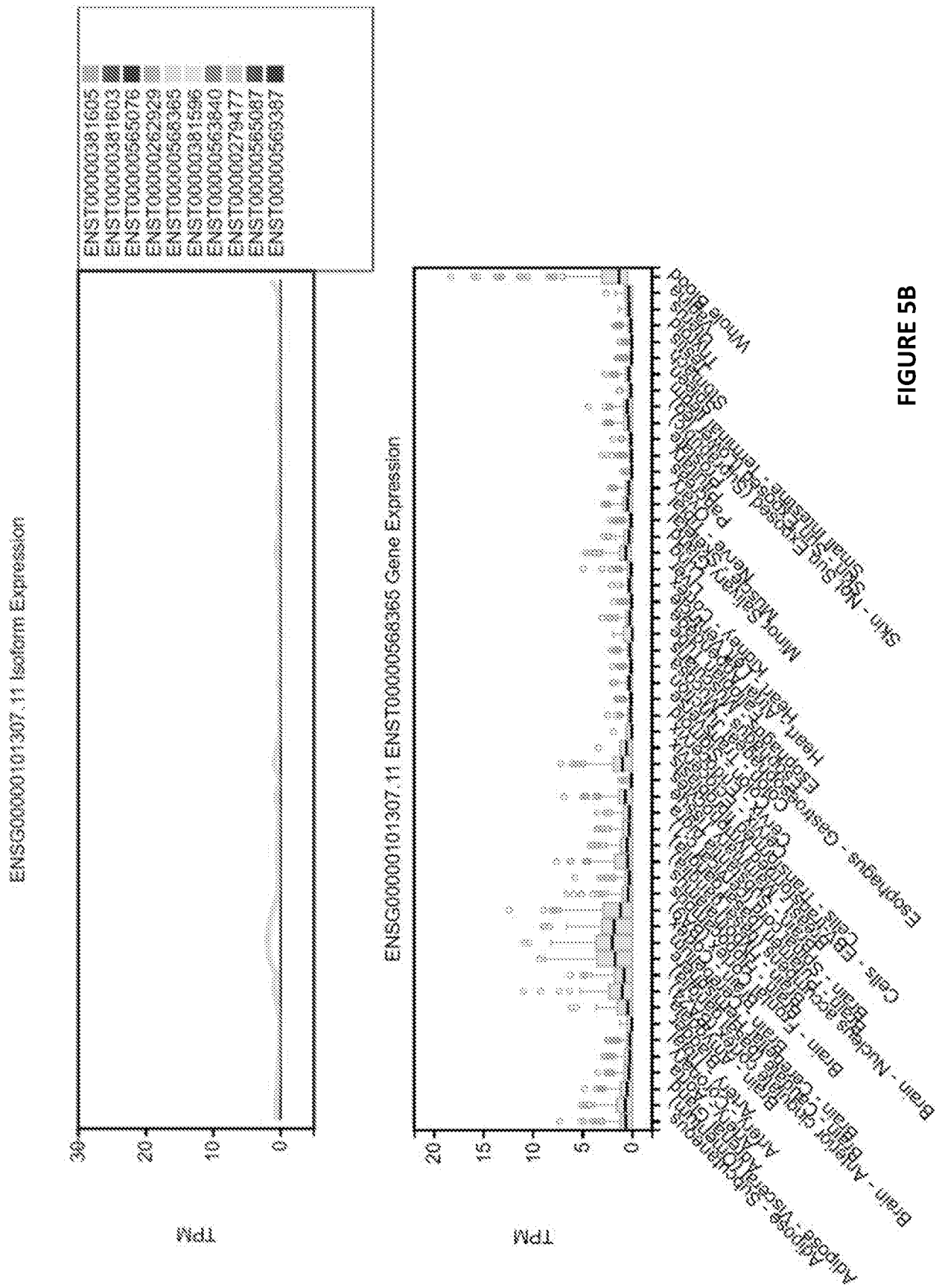


FIGURE 4C





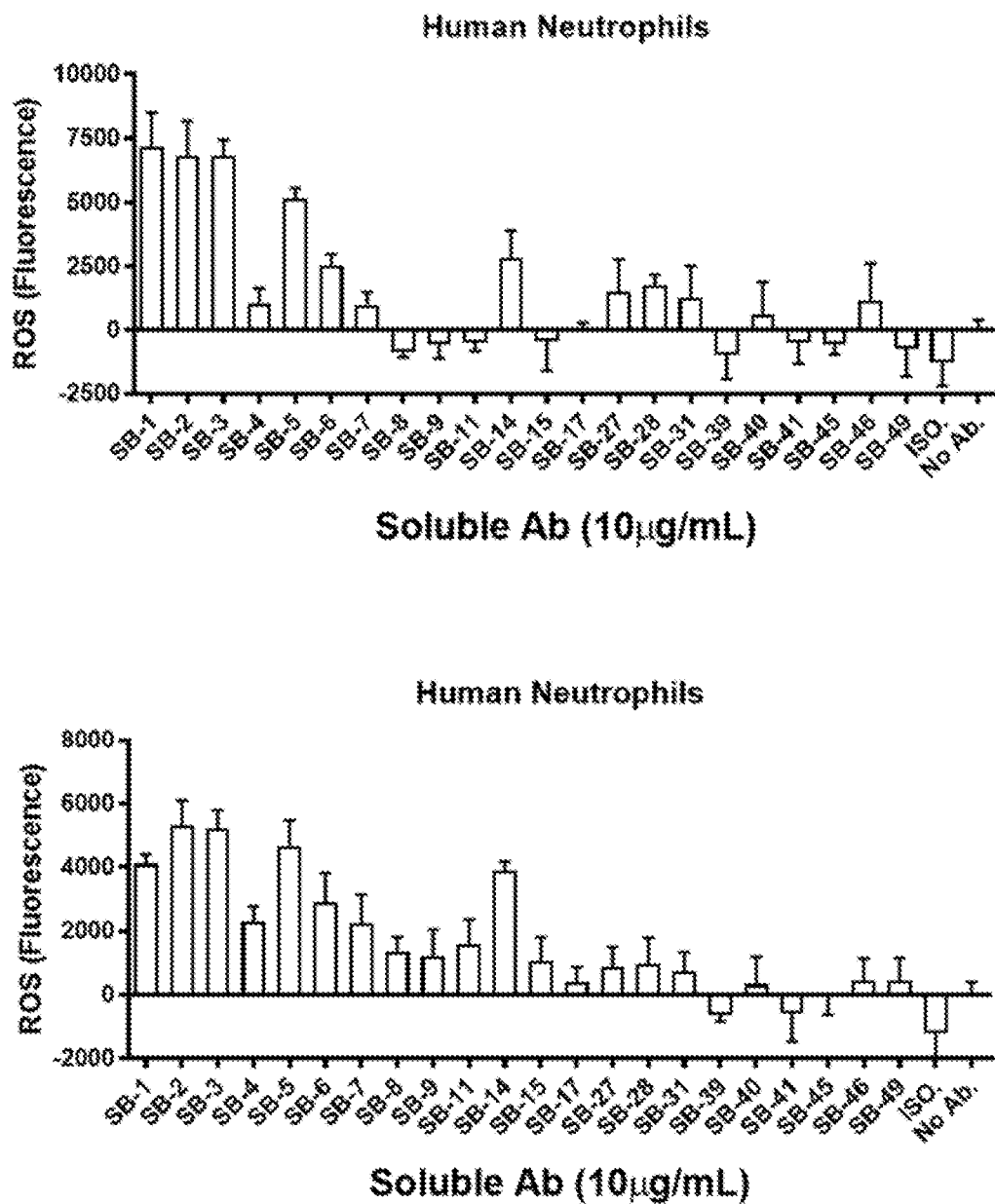


FIGURE 6A

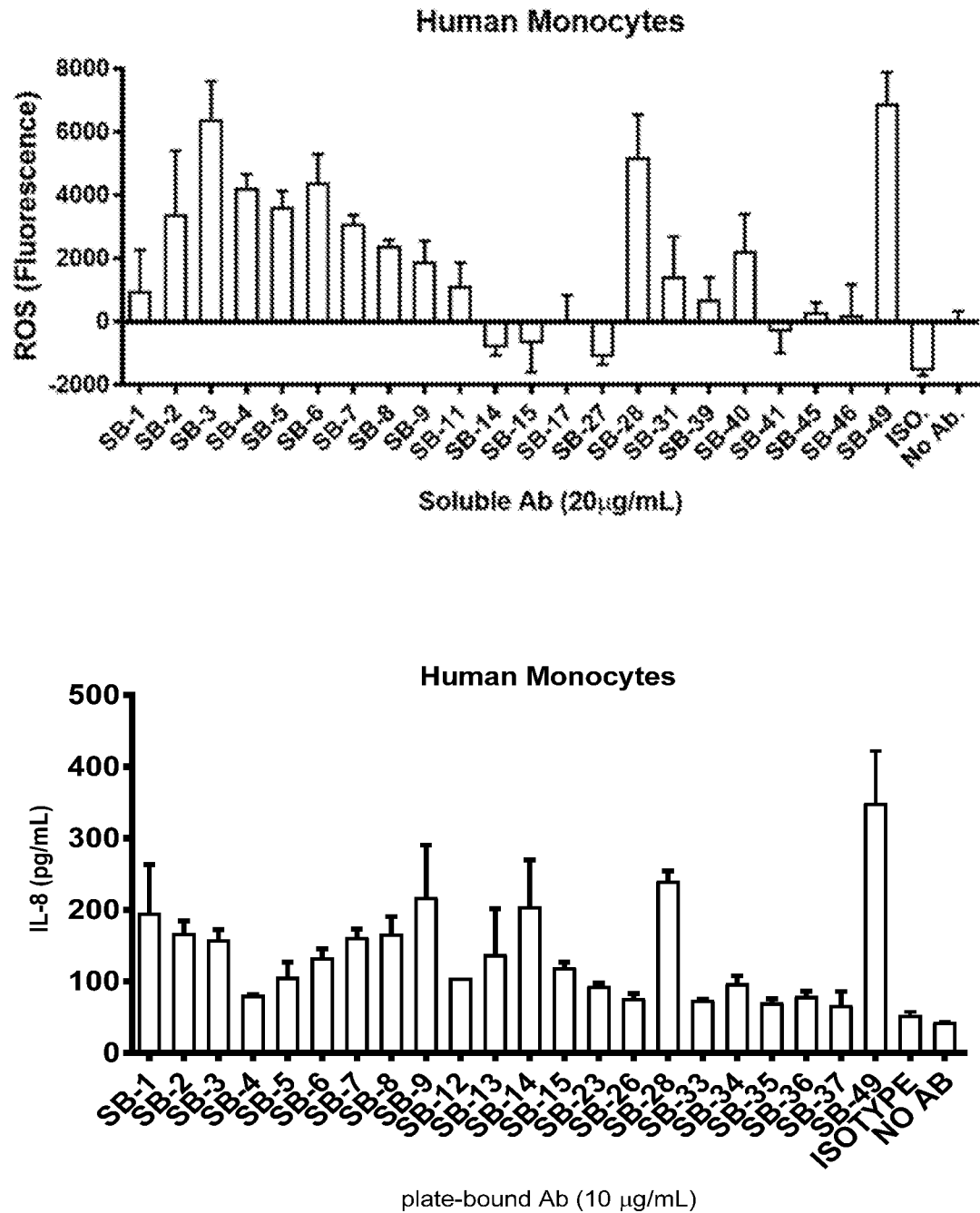


FIGURE 6B

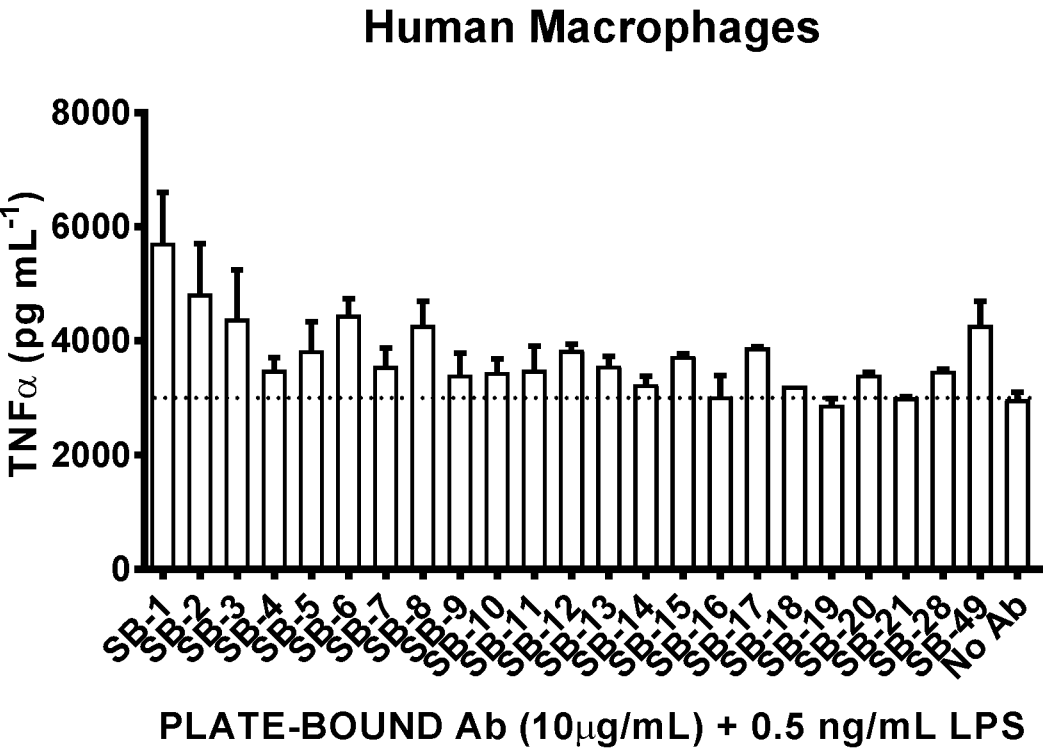
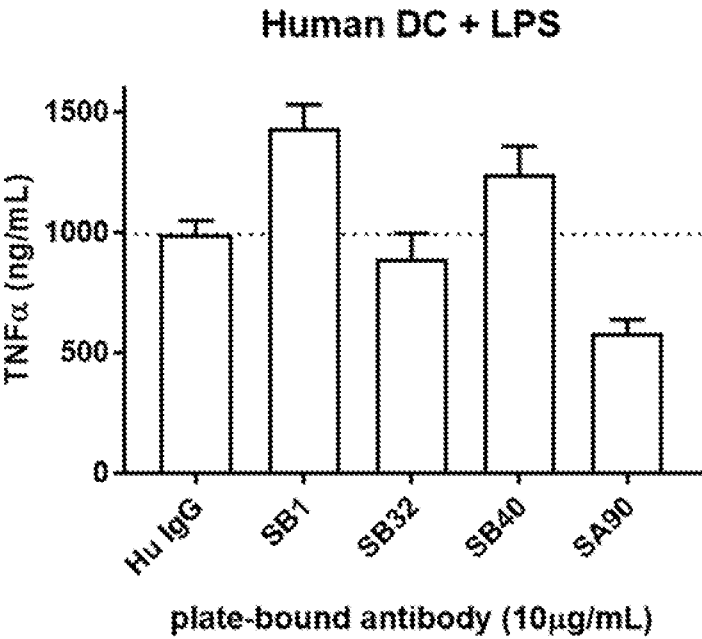


FIGURE 6C

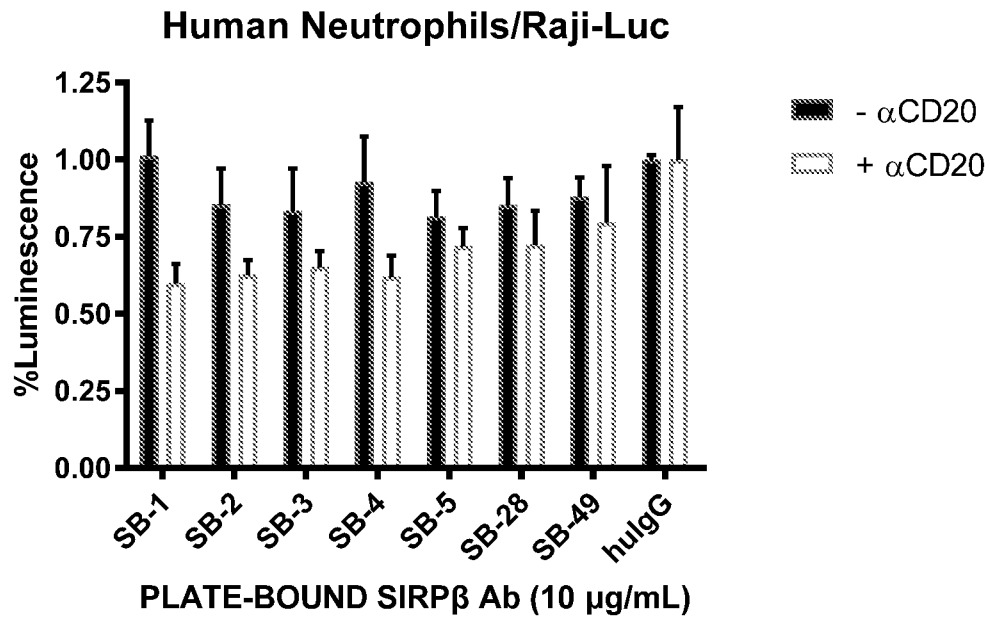


FIGURE 7

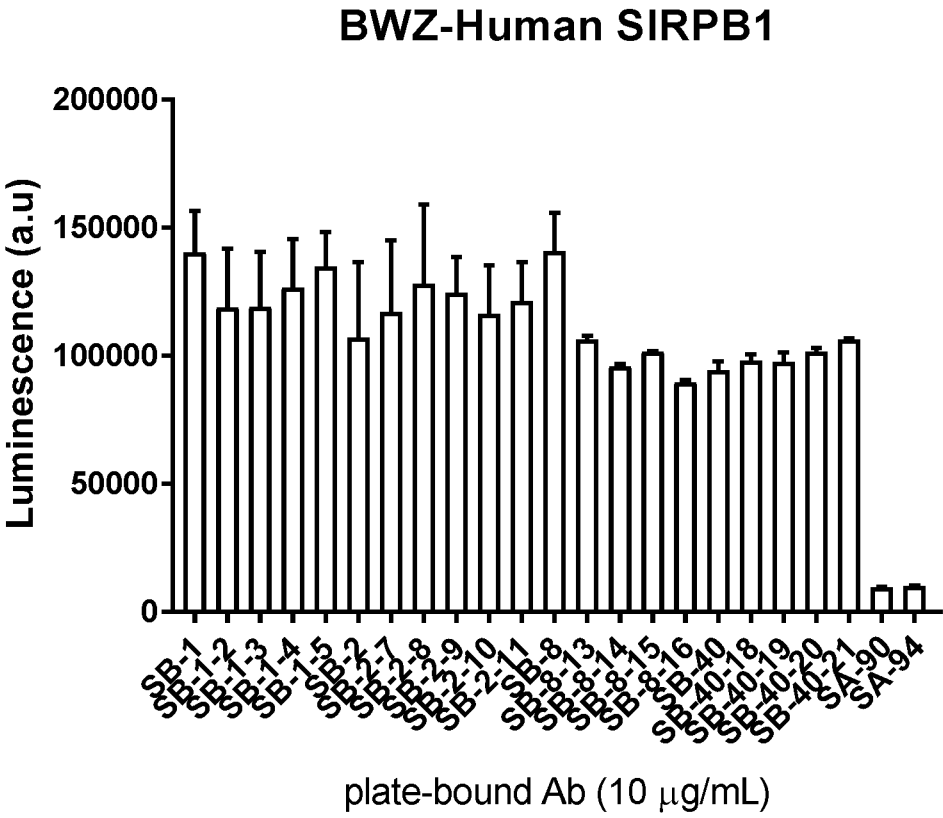
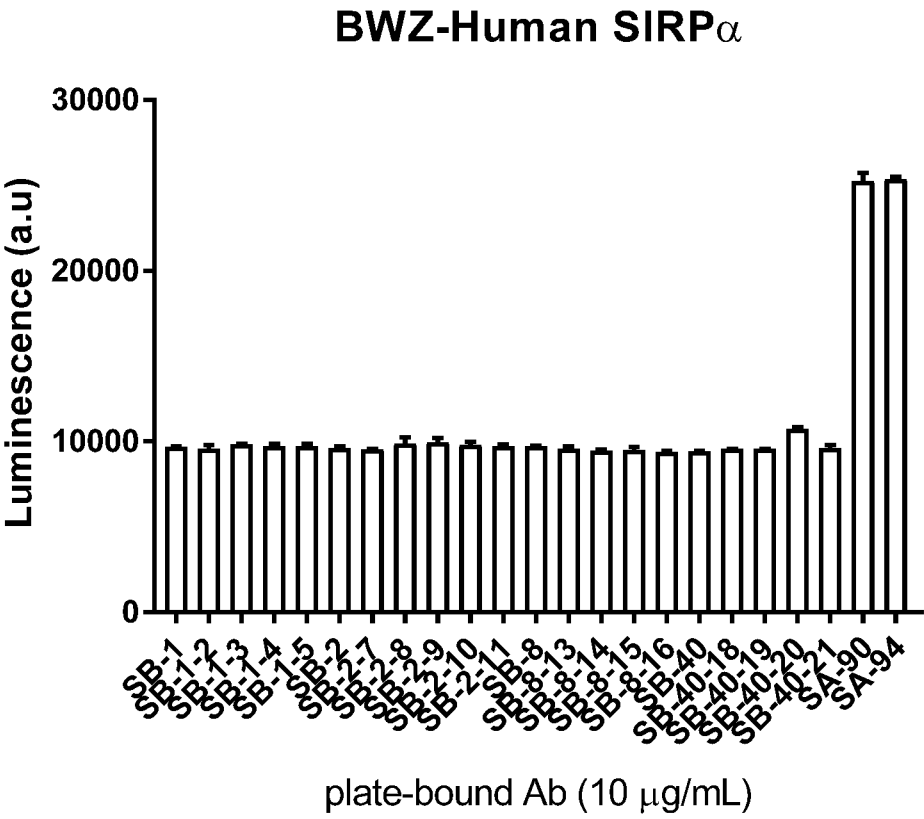


FIGURE 8

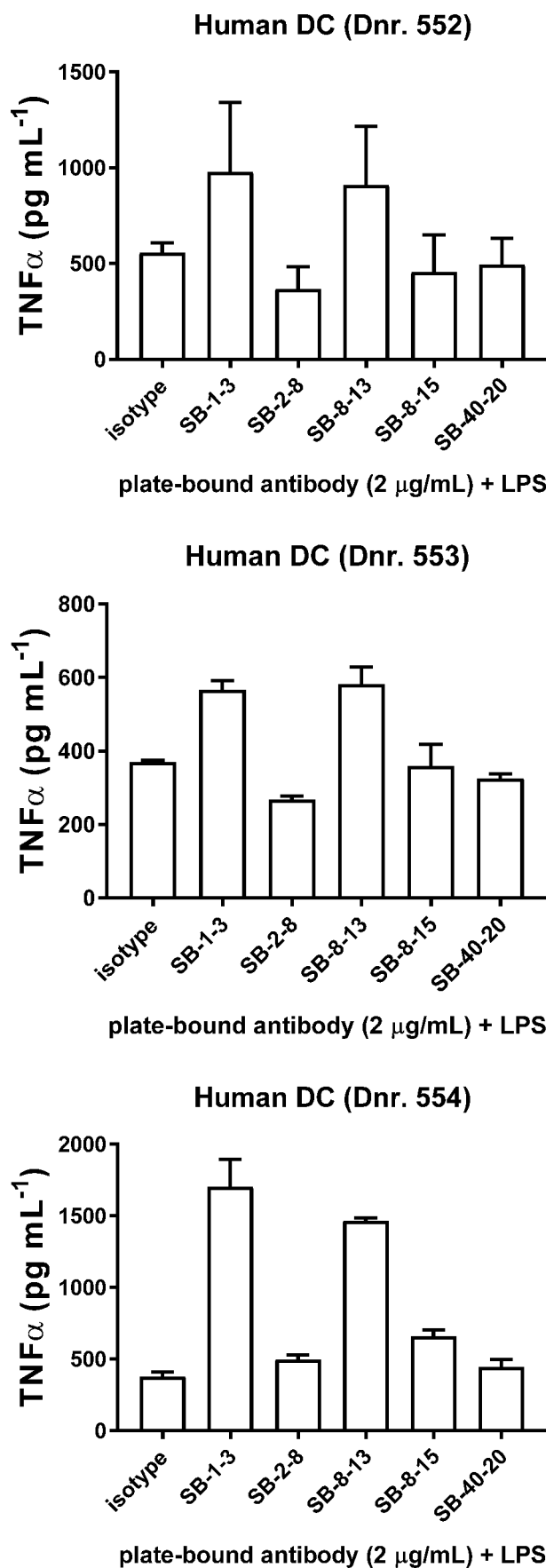


FIGURE 9

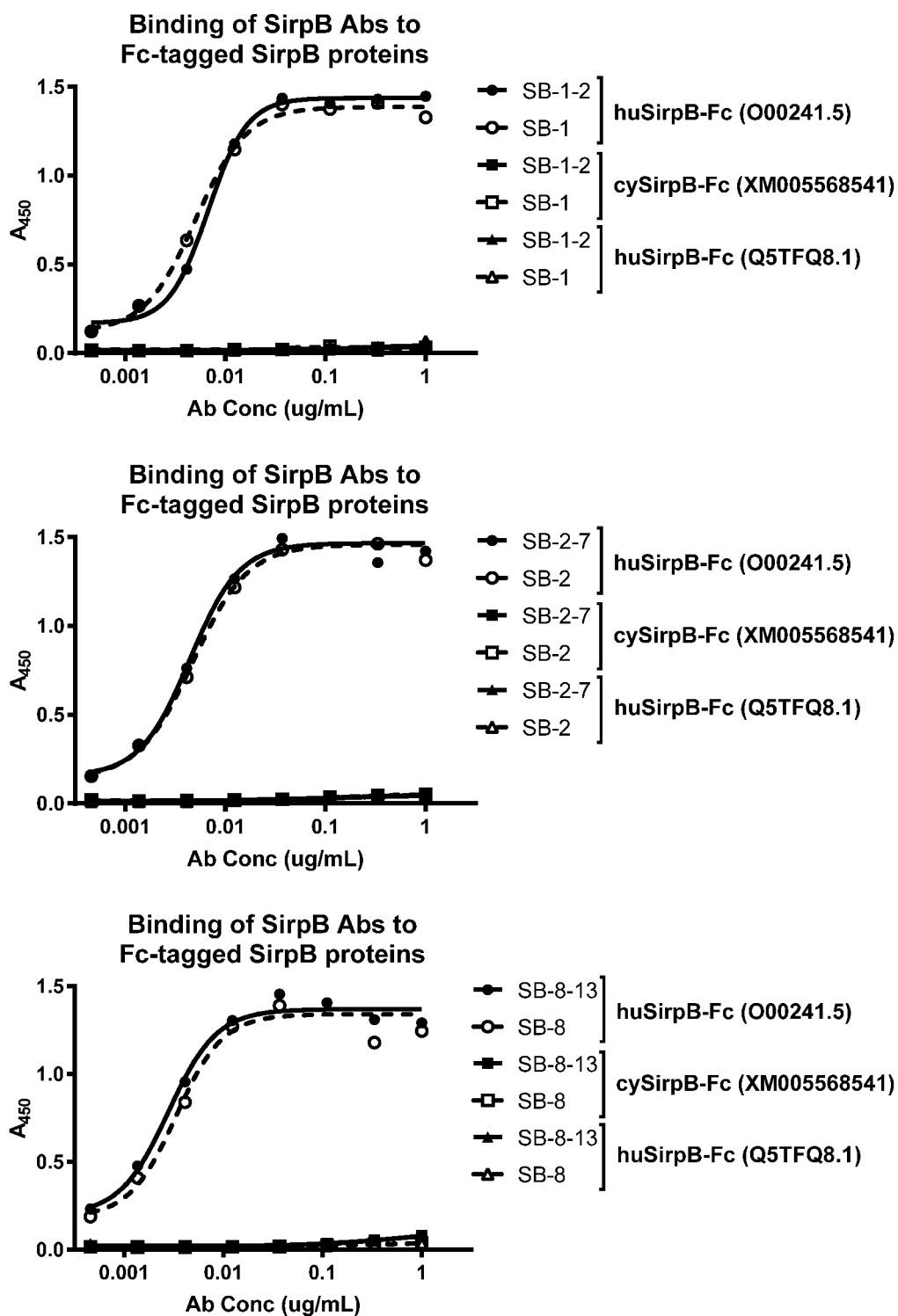


FIGURE 10A

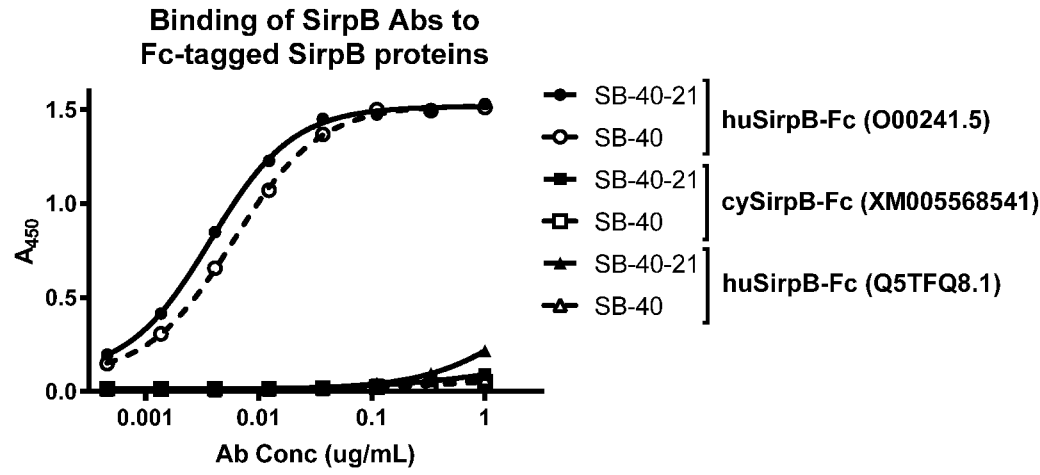


FIGURE 10B

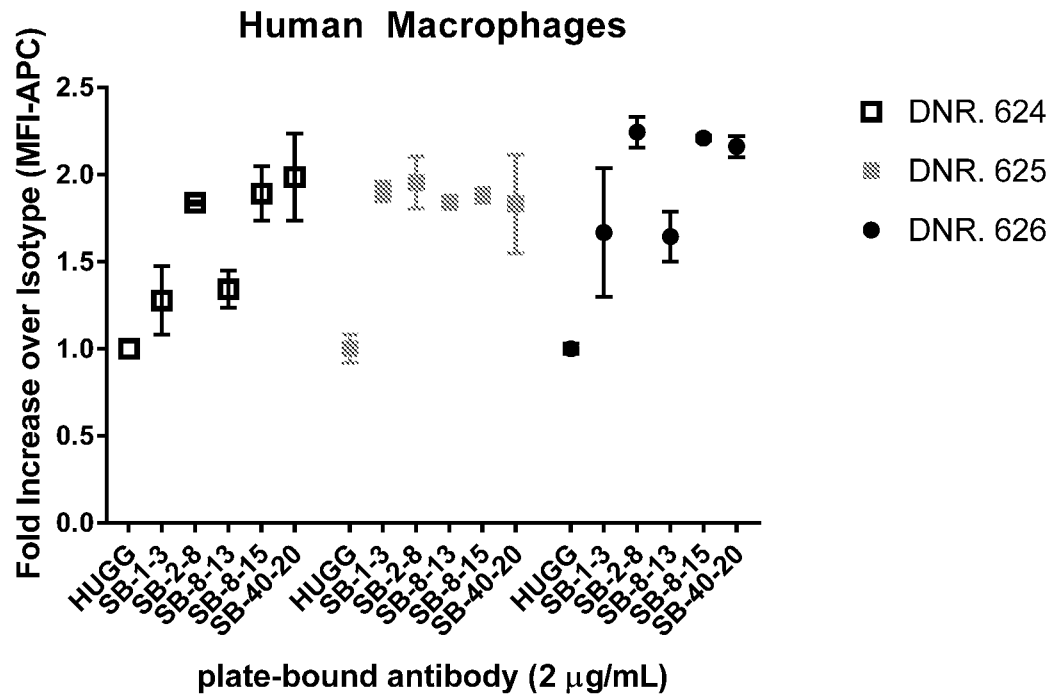


FIGURE 11A

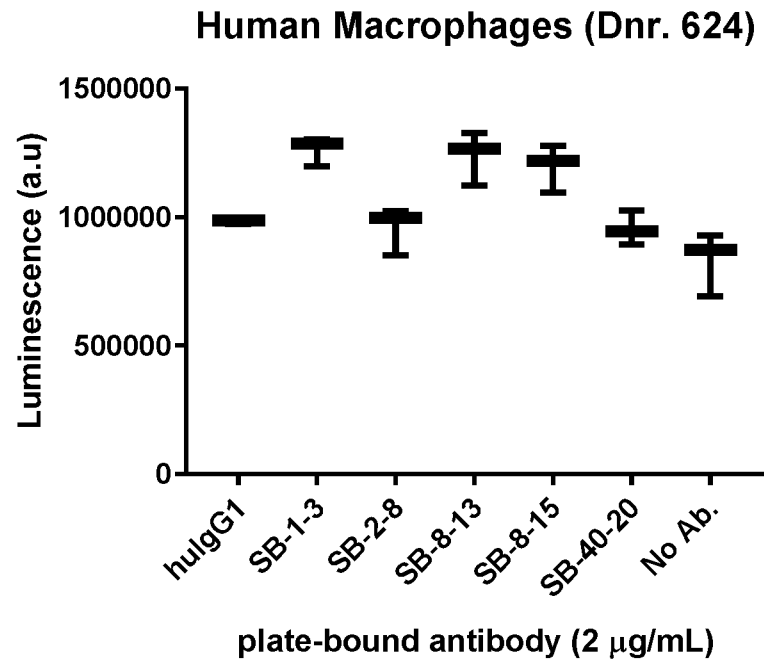


FIGURE 11B

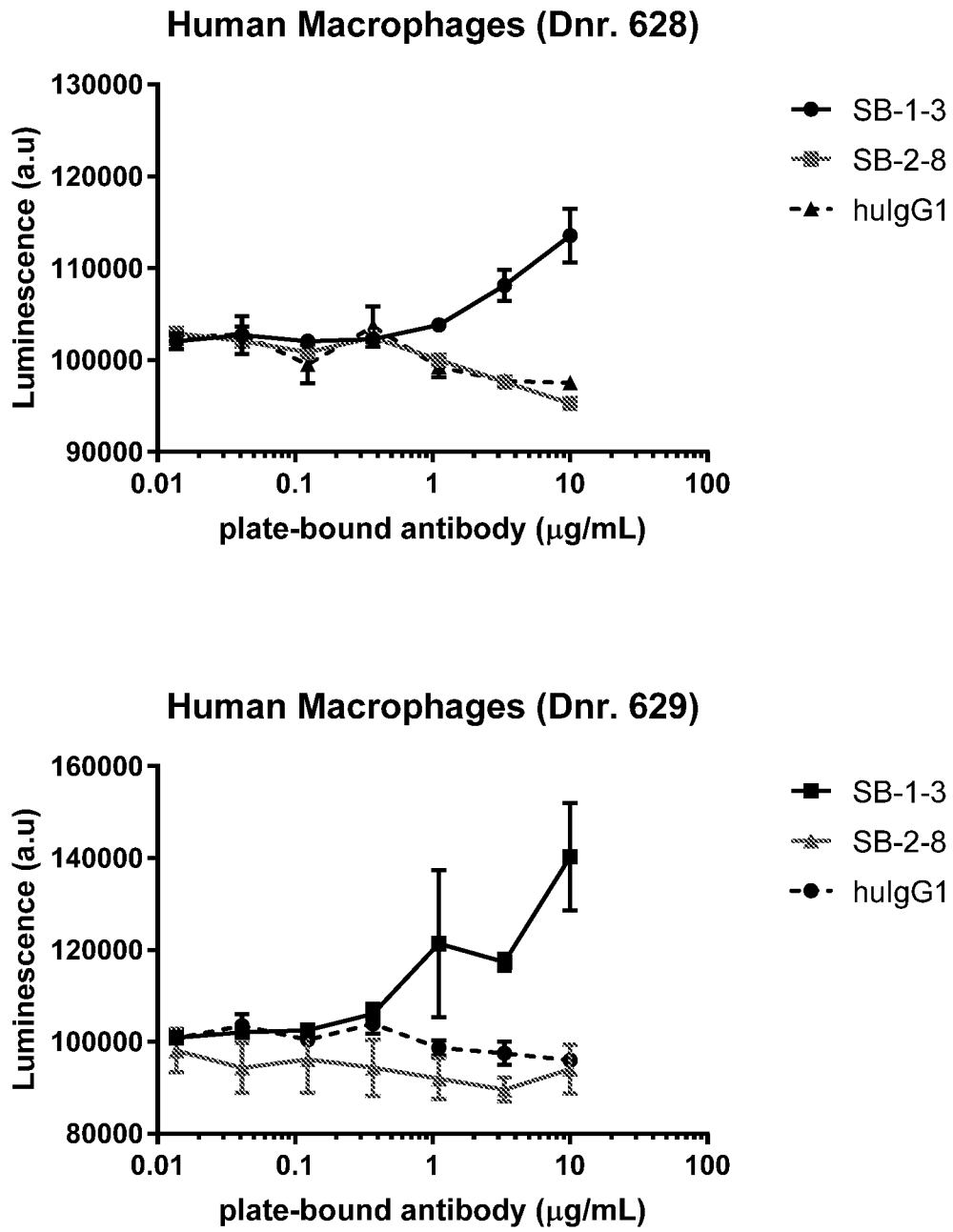


FIGURE 12

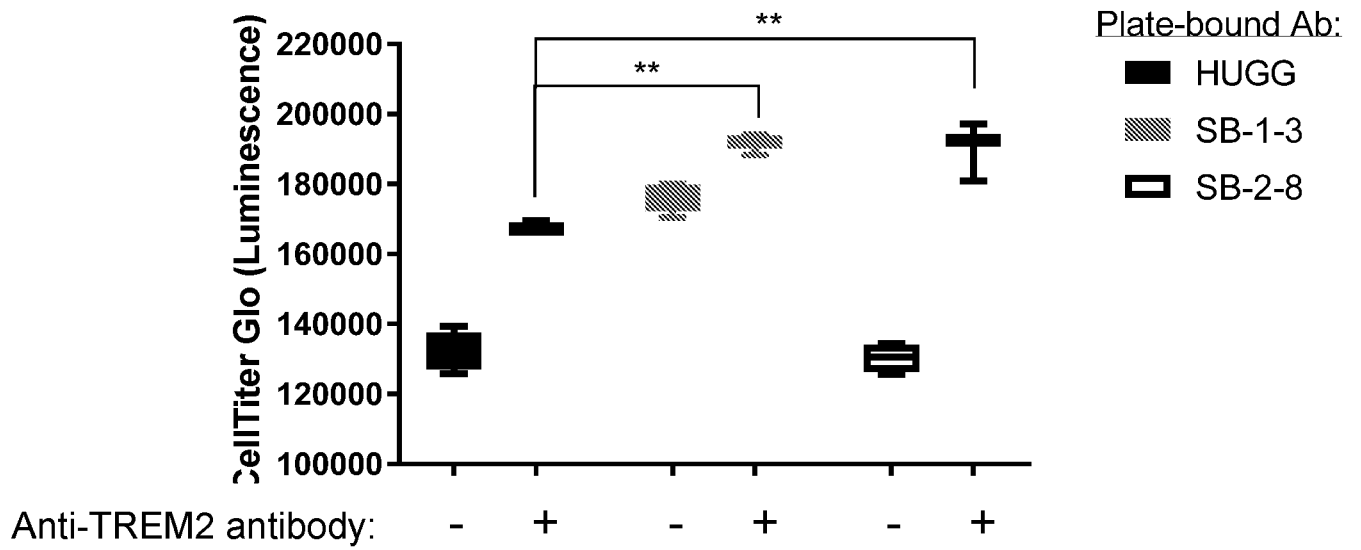
Human Macrophages (Dnr. 632)

FIGURE 13

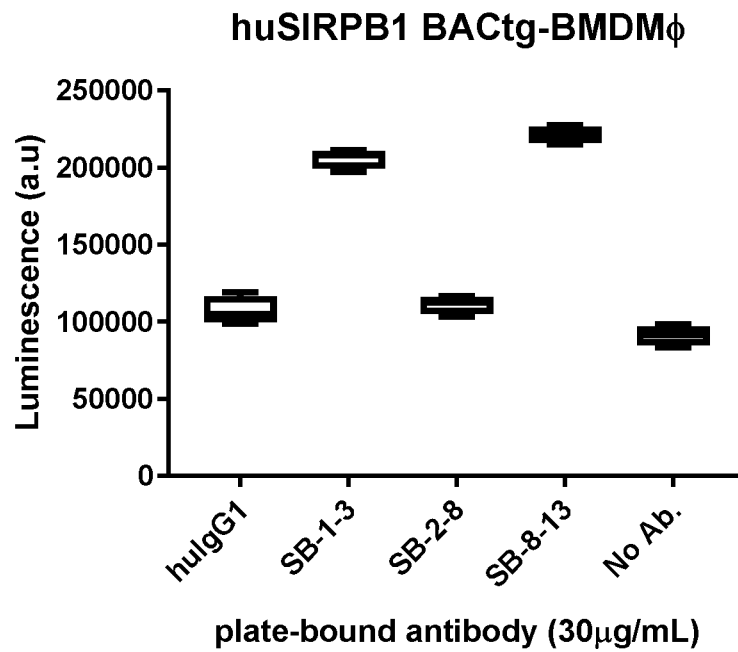


FIGURE 14A

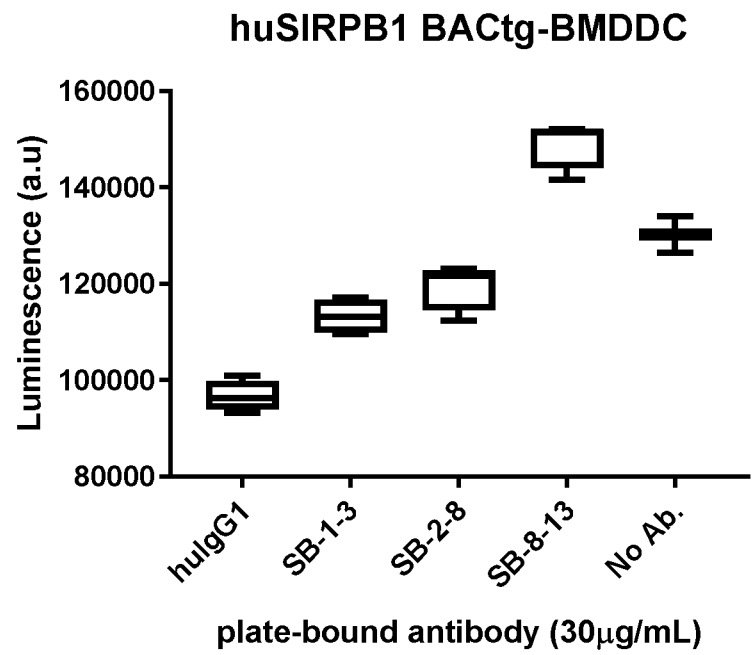


FIGURE 14B

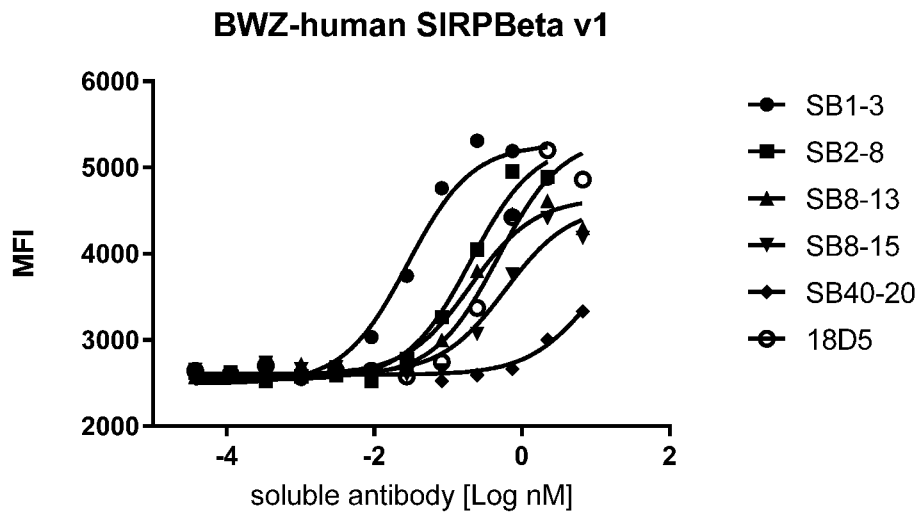


FIGURE 15A

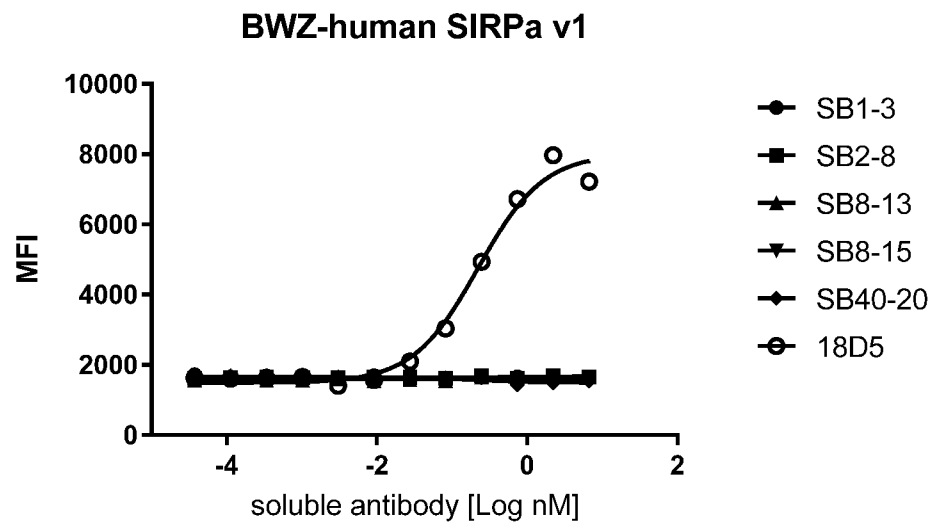


FIGURE 15B

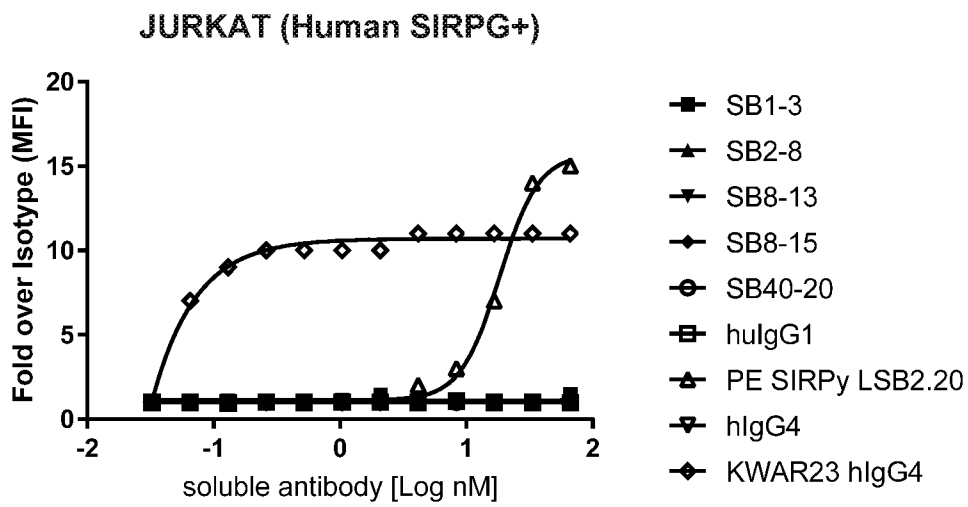


FIGURE 15C