Supplementary Table 4
Alignment of IC1 and TM2 sequences in S1P₁₋₅
Residues in bold represent residues identical to S1P₁ sequence but different from S1P₂ sequence in motifs *a*, *b*, *c*, or *d*.

Receptor subtype	IC1 and TM2 sequence			
S1P ₁	WKTK a	FH <u>RPMYY</u> I b	F <u>IGNLAL</u> SI <i>C</i>	DLLAGVA <u>YTANL</u> LLSG d
S1P ₂	<u>ARNS</u> K	FH <u>SAMYL</u> 1	F <u>LGNLAA</u> SI	DLLAGVA <u>FVANT</u> LLSG
S1P ₃	<u>wk</u> nnk	FH <u>NRMYF</u> 1	F <u>IGNLA</u> LCI	DLLAGIA <u>Y</u> KVNI <mark>LMSG</mark>
S1P ₄	TSHMR:	SR <u>RWVYY</u> (C <u>LVNITL</u> SI	DLLTGAA <u>Y</u> LANVLLSG
S1P ₅	<u>GRHP</u> R:	FH <u>APMFL</u> I	L <u>LGSLTL</u> SI	DLLAGAA <u>Y</u> AANILLSG