

IL21R_HUMAN/190-223	FRKDSSEYELQVRAGPMGPSSSYQGT.....	WSEWSDPVI
IL21R_MOUSE/190-223	FHKDSSEYQLQVRAAPQPGTSFRGT.....	WSEWSDPVI
IL2RB_HUMAN/198-229	LTPDQTQYEFQVRVKPLQGEFTT.....	WSPWSQPLAF
IL2RB_MACFA/198-229	LTPDQTQYEFQVRVKPLQGEFTT.....	WSPWSQPLAF
IL2RB_MOUSE/199-230	LIPSTSYEVQVRVKAQRNNTGT.....	WSPWSQPLTF
IL2RB_PANTR/198-229	LTPDQTQYEFQVRVKPLQGEFTT.....	WSPWSQPLAF
IL2RB_RAT/199-230	LTPDTSYELQVVRVIAQRGKTRT.....	WSPWSQPMAF
IL2RG_BOVIN/220-253	VDAQKLYTFRVRSRYNPLCGSAQH.....	WSDWSYPIHW
IL2RG_CANFA/213-246	VDGQKFYTFRVRSSRYNPLCGSAQR.....	WSEWSHPPIHW
IL2RG_HUMAN/213-246	VDGQKRYTFRVRSRYNPLCGSAQH.....	WSEWSHPPIHW
IL2RG_MOUSE/214-247	VDELKRYTFRVRSRYNPLCGSSQQ.....	WSKWSQPVHW
IL3B2_MOUSE/405-436	LEPDTSYTCARVRVKPISDYGDI.....	WSEWSNEYTW
IL3RB_HUMAN/402-434	LEPSTRYWARVRVTRTSRTGYNGI.....	WSEWSEARSW
IL3RB_MOUSE/406-437	LEPDTSYTCARVRVKPISNYDGI.....	WSKWSEETTW
IL4RA_HORSE/190-220	LKSRATYSARVKARAQNYNST.....	WSEWSPSTTW
IL4RA_HUMAN/191-221	LKSGISYRARVRAWAQCYNNT.....	WSEWSPSTKW
IL4RA_MOUSE/192-222	LMSGVYYTARVRVRSQILTGT.....	WSEWSPSITW
IL4RA_PIG/199-229	LKSGAAYSARVKAWAQRYNST.....	WSEWSPSVKW
IL4RA_RAT/191-221	LTSGVRYRARVRVLSQSFPGI.....	WSEWSPSITW
IL7RA_HUMAN/195-226	LQPAAMYEIKVRSIPDHYFKGF.....	WSEWSPSYFF
IL7RA_MOUSE/195-227	LRPKAMYEIKVRSIPHDYFKGF.....	WSEWSPSSTF
IL9R_HUMAN/214-254	LDPGFIHEARLRVQMATLEDDVVEEERYTGQ.....	WSEWSQPVCF
IL9R_MOUSE/212-253	LNPGSYIEARLRVQMTLESYEDKTEGEYYKS.....	HWSEWSQPVSF
LEPR_HUMAN/298-328	ILPGSSYEVQVRGKRLDGPGI.....	WSDWSTPRVF
LEPR_MACMU/296-326	ILPGSSYEVQVRGKRLDGPGI.....	WSDWSTPHVF
LEPR_MOUSE/296-326	VLPGSSYEVQVRGKRLDGSGV.....	WSDWSSPQVF
LEPR_PIG/298-328	VLPGSSYEVQVRGKRLDGPGI.....	WSDWSTPFTF
LEPR_RAT/296-326	VLPGSSYEVQVRGKRLDGSGV.....	WSDWSLPQLF

5 10 15 20 25 30 35 40

	Asp, Glu
	Arg, Lys, His
	Phe, Tyr, Trp
	Ala, Gly
	Cys, Met
	Ser, Thr
	Asn, Gln
	Leu, Val, Ile
	Pro