

Description	Sequence
Bis2 PD-1/CTLA-4 HC	DIQMTQSPSSLSASVGDRVITTCRASQSINSYLDWYQQKPGKAPKLLIY AASSLQSGVPSRFSGSGSGTDFTLTISLQPEDFATYYCQQYYSTPFTFG CGTKVEIKGGGGSGGGGSGGGGSGGGGSQVQLVESGGGVVQPGRSLR LSCAASGFTFSSYGMHWVRQAPGKCLEWVAVIWDGGSNKYYADSVK GRFTISRDNKNTLYLQMNSLRAEDTAVYYCARDPRGATLYYYYYYGM DVWGQGTITVTVSSGGGGSGGGGSEVQLVESGGGLVQPGGSLRLSCAA SGFTFSDYGMHWVRQAPGKGLEWVAYISSGSYTIYSADSVKGRFTISR DNAKNSLYLQMNSLRAEDTAVYYCARRAPNSFYEYYFDYWGQGTITV TVSSASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGA LTSGVHTFPAVLQSSGLYSLSSVTVPSSSLGKTYTCNVDHKPSNTKV DKRVESKYGPPCPPCPAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVV DVSQEDPEVQFNWYVDGVEVHNAKTKPREEQFNSTYRVVSVLTVLHQ DWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEEMT KNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPVLDSDGSFFLY SRLTVDKSRWQEGNVFSCSVMHREALHNHYTQKSLSLGLGK (SEQ ID NO:1)
Bis2 PD-1/CTLA-4 LC	QIVLTQSPATLSLSPGERATLSCSASSKHTNLYWSRHYWYQQKPGQA PRLLIYLTSNRATGIPARFSGSGSGTDFTLTISLQPEDFAVYYCQQWSS NPFTFGQGTKLEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPRE AKVQWKVDNALQSGNSQESVTEQDSKDYSLSSLTLSKADYEKHK VYACEVTHQGLSPVTKSFNRGEC (SEQ ID NO:2)
Bis3 PD-1/CTLA-4 HC	EVQLVESGGGLVQPGGSLRLSCAASGFTFSDYGMHWVRQAPGKGLE WVAYISSGSYTIYSADSVKGRFTISRDNKNSLYLQMNSLRAEDTAVY YCARRAPNSFYEYYFDYWGQGTITVTVSSASTKGPSVFPLAPCSRSTSES TAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLYSLSSV VTVPSSSLGKTYTCNVDHKPSNTKVDKRVESKYGPPCPPCPAPEFLGGP SVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHN AKTKPREEQFNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEK TISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEWES NGQPENNYKTTTPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMH ALHNHYTQKSLSLGLGKGGGGSGGGGSDIQMTQSPSSLSASVGDRVIT TCRASQSINSYLDWYQQKPGKAPKLLIYAASSLQSGVPSRFSGSGSGTD FTLTISLQPEDFATYYCQQYYSTPFTFGCGTKVEIKGGGGSGGGGSGG GGGGGGGSQVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQ APGKCLEWVAVIWDGGSNKYYADSVKGRFTISRDNKNTLYLQMNSL RAEDTAVYYCARDPRGATLYYYYYYGM DVWGQGTITVTVSS (SEQ ID NO:3)
Bis3 PD-1/CTLA-4 LC	QIVLTQSPATLSLSPGERATLSCSASSKHTNLYWSRHYWYQQKPGQA PRLLIYLTSNRATGIPARFSGSGSGTDFTLTISLQPEDFAVYYCQQWSS NPFTFGQGTKLEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPRE AKVQWKVDNALQSGNSQESVTEQDSKDYSLSSLTLSKADYEKHK VYACEVTHQGLSPVTKSFNRGEC (SEQ ID NO:6)