

Figura 36K

Cadena 3 - CD20-B_L1 (SEQ ID NO: 429)

EIVLTQSPATLSLSPGERATLSCRASLSVSSMHWYQQKPGQAPRLLIYATSNLASGIPARFSGSGSGTDFTLTIS
SLEPEDFAVYYCQQWIFNPPTFGGGTKVEIK/RTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKV
DNALQSGNSQESVTEQDSKDSSTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC

**>XENP42164 CD20-B_H1L1 Fab-CD20-B_H1L1 Fab-IgG1_L234A/L235A/D265S/T366S/L368A/Y407V-
IgG1_L234A/L235A/D265S/T366W/H435R/Y436F/K447del_(G4S)3_1A7[CD28]_H1_L1.71_scFv(GKPGS)4
Cadena 1 - CD20-B_H1_IgG1_L234A/L235A/D265S/T366S/L368A/Y407V (SEQ ID NO: 516)**

QVQLVQSGAEVKKPGSSVKVSCKASGYTFSSYNMHWVRQAPGQGLEWMGAIYEGAGDTSYAQKFQGRVTITADES
TSTAYMELSSLRSEDTAVYYCARSNYGSSGWYFDVWGKGTTTVTVSS/ASTKGPSVFPLAPSSKSTSGGTAALGC
LVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLYSLSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKS
CDKTHHTCPPEAPEAAGGPSVFLFPPKPKDTLMISRTPEVTCVVVSVSHEDPEVKFNWYVDGVEVHNAKTKPREE
QYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLSCAV
KGFYPSDIAVEWESNGQPENNYKTPPVLDSDGSFFLVSKLTVDKSRWQQGNVFCSCVMHEALHNHYTQKSLSLS
PGK

Cadena 2 - CD20-

**B_IgG1_L234A/L235A/D265S/T366W/H435R/Y436F/K447del_(G4S)3_1A7[CD28]_H1_L1.71_scFv(GKPGS)4
(SEQ ID NO: 14)**

QVQLVQSGAEVKKPGSSVKVSCKASGYTFSSYNMHWVRQAPGQGLEWMGAIYEGAGDTSYAQKFQGRVTITADES
TSTAYMELSSLRSEDTAVYYCARSNYGSSGWYFDVWGKGTTTVTVSS/ASTKGPSVFPLAPSSKSTSGGTAALGC
LVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLYSLSVVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKS
CDKTHHTCPPEAPEAAGGPSVFLFPPKPKDTLMISRTPEVTCVVVSVSHEDPEVKFNWYVDGVEVHNAKTKPREE
QYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLWCLV
KGFYPSDIAVEWESNGQPENNYKTPPVLDSDGSFFLYSKLTVDKSRWQQGNVFCSCVMHEALHNRTQKSLSLS
PGGGGSGGGGSGGGGS/EVQLLESGGGLVQPGGSLRLSCAASGFTSSYAMSWVRQAPGKLEWVSTISGSGDS
TYYADSVKGRFTISRDN SKNTLYLQMNSLRAEDTAVYYCAKSGEGLRQVGFQYWGQGLTVTVSS/GKPGSGKPGS
GKPGSGKPGS/DIQMTQSPSSLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLIYAASSLQSGVPSRFSG
SGSGTDFTLTISLQPEDFATYYCQQVYSTPFTFGQGTKLEIK

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EIVLTQSPATLSLSPGERATLSCRASLSVSSMHWYQQKPGQAPRLLIYATSNLASGIPARFSGSGSGTDFTLTIS
SLEPEDFAVYYCQQWIFNPPTFGGGTKVEIK/RTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKV
DNALQSGNSQESVTEQDSKDSSTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC