

Secuencias de ácido nucleico de región variable de anticuerpo anti-CLDN humanizado

Nombre	Cadena	Secuencia de ácido nucleico	SEQ ID NO
hSC27.1	Light Chain	GACATCCAGATGACCCAGTCTCCATCTCCGTGCTGCACTCTGAGGAGACAGAGTCCACCTCACTTGTAAAGGAGTGAAGATATTACAAACCAGTTAGCCTGGTATCAGCAGAAACCA GGGAAAGCCCCTAAGCTCCTGATCTATGGTGAACAGTTGGAACTGGGGTCCATCAAGGTTCAAGGGCAGTGGATCTGGGACAGATTACACTCTCACCATCAGCAGCCTGCAGCCT GAAGATTGTCAACTTACTATTGTCAACAGTATTGGAGTACTCCACTACGTTCCGTCAAGGACCAAGCTGGAGATTAAA	60
hSC27.1	Heavy Chain	CAGGTCAGCTTGTGCAAGTCTGGGCTGAGGTGAAGAAGCCTGGGGCTCAGTGAAGGTTTCCTGCAAGGCTCTGGATACACCTTCAGTACTGCAATGGGTGCGCCAGGCC CCCGGACAAAGCTTGAGTGGATGGGAGGGATCAACCTTAAATGGTGACACAATATATAACCAAGAACTTCAAGGGCAGAGTCAACATTACCAAGGACACATCGCGAGCACAGCCTAC ATGGAGCTGAGCAGCCTGAGATCTGAAGACACGGCTGTGATTACTGTGCSAGAAGSGCATACGGTCTATGCTATGGACTACTGGGTCAAGGTACCCTAGTCAACGCTCTCAGC	62
hSC27.22	Light Chain	GACATGTGATGACCCAGTCCCTGACAGTGTGGCGTTAGCTTGGGGAGCGTGCCACCATCAACTGTAGGGCTAGTCAAACTGTTCTACATCCTCTACTCTTACATGCAATGGTAT CAGCAGAAACCTGGTCAAGCTCCAAACTGCTGATTTATTTGCACTCAACGTCGAGTCCGGAGTTCTGACCGGTTCAGCGGATCAGGAAGCGGTACAGATTATACATTACATCTCA TCTCTGCAAGCAGAAGATGTGGCGGTATATTGTGTCAGCATCTCTGGAGATCCCCTGAGACTTCGGGCAAGGAAACCAAGCTCGAGATTAAA	64
hSC27.22	Heavy Chain	CAGGTGCAAGTGGTGACAGCGGCGCCGAAGTCAAGAAACAGGAGCTTCTGTCAAGTCTCTGTAAAGCTCCGGATATACCTTCACAGCTACTGGATGAATGGGTAAAGACAGGCC CCCGGACAGAGCTTGAGTGGATGGGAATGATCATCCCTCTGACAGCGAGATTCGGCTCAACCAAGAACTTAAAGACCGAGTGAATATCACACGGATACCAAGTGCTAGCACAGCCTAC ATGGAGTTGAGTTCTCTCGTAGCGAGGACACTGCCGTGATTATTGCGCCCGCATCGACTATATATGGTTATCTGTTCTACTTCGACTATTGGGCGAGGGGACCAACCGTGACTGTG TCTTCC	66
hSC27.108	Light Chain	GAAATCGTGCTTACACAATCCCTGCCACTCTGAGCCTTCTCCAGGCGAGCAGCAACCTTCTCTGAGTGTTCCTCTCAATCAGTCCAGCAATTTGCACTGGTACCAAGCAGAAAG CCTGGTCAGGCAACCCGATTGTTGATCTATGGCACATCTAACCTGGCCAGCGGCATCTCTGCTCGGTTCAAGTGGATCTGGCTCCGGAACAGATTTCACCTCTACATCAGCTCCCTTGAG CCTGAAGATTTGCGGTGTACTACTGTCAAGTGGAGTCTCTACCCCCACACCTTTGGGCGGGACAAAGTCAAGATAAAA	68
hSC27.108	Heavy Chain	CAGGTACAGCTGGTCCAGTCCGCGCTGAGGTTAGAGGCCGGTGCTCCGTGAAGGTATCTGTAAAGCTCAGGTTACACCTTTACAATTATCTGATCGAATGGGTGAGACAGGCC CCAGGTCAAGGTCTGGAATGGATGGGACTCATCAACCTGGAGTGGCGGACCAACTACAAAGAAAGTTAAAGGGAGAGTGAATGACCAAGATACAGTACCTCCACCGCATAT ATGGAGCTGCGAAGCTTGAGGTCCGATGACACTGCTGTGACTATTGCGCCCGTAGAAGCCACTCGGCTCTGGATCTATTACGCATACGATGGTGTGGCCTATTGGGGCCAGGGCACC CTGGTGACAGTCAGCTCC	70
hSC27.204	Light Chain	GACATCCAGATGACCCAGTCCCTCCAGCTGTCTGCTCCGTGGGCGACAGAGTACCATCACATGCAAGGCGGCGCAGAACGTGGGACCTCTGTGGCCTGGTTCCAGCAGAAACCT GGCAAGGCCCCCAAGTCCCTGATCTACTCCGCCTCTACAGATACTCCGCGTGCCCCTCAGATTCTCCGGCTCTGGCTCTGGCAACGACTTTACCTTGACCATCAGTCCCTGCAGCCCC GAGGACTTCGCCACCTACTACTGCCAGCAGTACATCACCACCTTACACCTTCGGCGGAGGACCAAGGTGGAAATCAAG	72
hSC27.204	Heavy Chain	GAAAGTCAAGCTGCTGGAATCTGGCGCGGACTGGTGAGCCTGGCGGATCTCTGAGACTGTCTGTGCTCCGCTCCCGCTTCACTTCTCCCGGTACTGGATGTCCTGGGTGCGACAGGCT CTGGCAAGGGCCTGGAATGGGTGCTCCGAGATCAACCCGACTCCTCCACCATCAACTACACCCAGCTGAAAGGCCGGTTCAACATCTCTCGGCAACTCAAGAACACCTGTAC CTGCAGATGAACCTCCGCGGGCGAGGACACCGCGTGACTACTGTACCGGCCCTGCTATTGGGGCCAGGGACCTCTGTGACCGTGTCTCT	74

FIG. 3C CONT.