

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

Nombre	Cadena	Secuencia de ácido nucleico	SEQ ID NO
SC27.104	Light Chain	GACATCCAGATGACACAATCTTCATCCTCC TTTTCTGTATCTCTAGGAGACAGAGTCACCATTA CTGCAAGGCAAGTGAGGACATATATAATCGGTTAGCCTGGTATCAGCAGAAACCAGGAAA TGCTCCCAGGCTCTTAATATCTGGTGCAACAGTTTGGAACTGGGGTTCCTTCAAGATTCAGTGGCAGTGATCTGGAAAGGATTACACTCTCAGCATTACCAGTCTTCAGACTGAAGATGTTG CTACTTACTGTCAACAGTATTGGAGTAATCCTCCGACGTTCTGGTGGAGGACCAAGCTGGAATCAAA	32
SC27.104	Heavy Chain	GAGGTCCAGCTGCAACAGTCTGGACCTGAGCTGGTGAAGCCTGGGGCTTCAGTGAAGATATCCTGCAAGACTTCTGATACACATTCACTGAATACACCTGCACTGGGTGAAGCAGAGCCATGG AAAGAGCCTTGAGTGGATTGGAGGTGTTTATCCTAAGAATGGTGATACTACCTACAAACAGAAAGTTCAAGGCAAGGCCACATTGACTGTAGACAAGTCCTCAACACAGCCTATATGGAACCTC GAGCCTGACATCTGAGGATTCTGCAGTCTATTACTGTACAGGAAGGATGGGTACGACGGGTTTGTACTGGGGCCAAGGGACTCTGGTCACTGTCTCTGCA	34
SC27.105	Light Chain	GATGTTCAAATGACCCAGTCTCCATCCTCCCTGCTGCACTTTGGGAGAGAGAGTC TCCCTGACTTGCCAGGCAAGTCAGAGTGTTAGCAATAA TTTAACTGGTATCAGCAAACACAGGGAA AGCTCCTAGGCTCTTGATCTATGGTGCAAGCAAATGGAAAGATGGGGTCTCTTCAAGGTTCAAGTGGCAGTGGATATGGGACAGATTTCACTTTCAATCAGCAGCCTGGAGGAAGAAGATGTTG CACTTATTTTGTCTACAGATAGGTATCTGTGGACGTTCTGGTGGAGGACCAAGCTGGAATCAAA	36
SC27.105	Heavy Chain	GAGGTCCAGCTGCAGCAGTCTGGACCTGAGTTGGTGAAGCCTGGGGCTTCAGTGAAGATATCCTGCAAGGCTTCTGTTACTCATTCACTGGCTACTACATGAACTGGGTGAAGCAAAGTCTTGA AAAGAGCCTTGAGTGGATTGGAGATTAATCCTAGCACTGGTAGTACTACTACAAACAGAAAGTTCAAGGCAAGGCCACATTGACTGTAGACAATCCTCCAGCACAGCCTACATGCAGCTCA AGAGCCTGACATCTGAGGACTCTGCAGTCTATTACTGTGCAAGAAGGATTATTA CTACGGTAGTGGTTTCTATGCTATGGACTACTGGGGTCAAGGAACCTCAGTCACCGTCTCCTCA	38
SC27.106	Light Chain	GACATCCAGATGACACAATCTTCATCCTCC TTTTCTGTATCTCTAGGAGACAGAGTCACCATTA CTGCAAGGCAAGTGAGGACATATATAATCGGTTAGCCTGGTATCAGCAGAAACCAGGAAA TGCTCCTAGGCTCTTAATATG TGGTGCAACAGTTTGGAACTGGGGTTCCTTCAAGATTCAGTGGCAGTGATCTGGAAAGGATTACACTCTCAGCATTACCAGTCTTCAGACTGAAGATGTTG CTACTTACTGTCAACAGTATTGGAGTACTCCGCTCACGTTCTGGTCTGGGACCAAACTGGAGCTGAAA	40
SC27.106	Heavy Chain	GAGGTCCAGCTGCAACAGTCTGGACCTGAGCTGGTGAAGCCTGGGGCTTCAGTGAAGATATCCTGCAAGACTTCTGATACACATTCACTGAATACACCTGCACTGGGTGAAGCAGAGCCATGG AAAGAGCCTTGAGTGGATTGGAGGTATTAATCCTAACAATGGTGTA CTAACTACAAACAGAAAGTTCAAGGCAAGGCCACATTGACTGTTGACAAGTCCTCCAGCACAGCCTACATGGAGCTCC GAGCCTGACATCTGAGGATTCTGCAGTCTATTACTGTGCAAGAAGGCTTATTACTTACTATGCTATGGACTACTGGGTCAAGGAACCTCAGTCACCGTCTCCTCA	42

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FIG. 3C CONT.