

FIGURA 9A

Alineaciones de Secuencia de Región Variable de Cadena Ligera

SEQ ID NO.	NOMBRE DEL CLON	CDR1	CDR2
3	1N23VL5-5	FIVMTQTPLSLPVS LGDQASISCR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKVS NRF	60
4	1N23VL5-8_0816	SIVMTQTPLSLPVS LGDQASISCR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKVS NRF	60
5	1N23VL1-8	LVLMTQTPLSLPVS LGDQASISCR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKVS NRF	60
6	1N23VL1-2_0820	SVLMTQTPLSLPVS LGDQASISCR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKVS NRF	60
7	1N23VL1-4_0820	YVLMTQTPLSLPVS LGDQASISCR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKVS NRF	60
8	1K23VL2-5	QVLMTQAPLSLPVS LGDQASISCR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKVS NRF	60
9	1N23VL1-2	RFQVSQTPLSLPVS LGDQASISCR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKVS NRF	60
10	1K23VL2-6	VFVMTQAPLSLPVS LGDQASISCR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKVS NRF	60
11	1K23VL2-8_0822	LIVMTQAPLSLPVS LGDQASTSCR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKVS NRF	60
12	1K23VL2-3_0829	SIVMTQAPLSLPVS LGDQASISCR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKVS NRF	60
13	2K4VL1-3_0820	SVLMTQTPLSLPVS LGDQASISSR-SSQSLVHSNGNTYLHWYLQKPGQSPKLLIYKIS NRF	60
14	2K4VL1-4	LVLMTQTPLSLPVS LGDQASISCR-SSQNIVHSNGNTYLEWYLQKPGQSPKLLIYKVS NRF	60
15	2K4VL1-1	LVLMTQTPLSLPVS LGDQASISCR-SSQNIVHSNGNTYLEWYLQKPGQSPKLLIYKVS NRF	60
16	2K4VL1-6_0820	LVLMTQTPLSLPVS LGDQASISCR-SSQNIVHSNGNTYLEWYLQKPGQSPKLLIYKVS NRF	60
17	1C9VL2-4	LIVMTQAAPSV PVT PGESVSISCR-STKSLLSHNGNTYLSWFLQRPQSPQLLIYRMS NLA	60
18	1C9VL2-6	YIVMTQAAPSV PVT PGESVSISCR-STKSLLSHNGNTYLSWFLQRPQSPQLLIYRMS NLA	60
19	1C9VL2-3_0816	SGLMTQAAPSV PVT PGESVSISCR-STKSLLSHNGNTYLSWFLQRPQSPQLLIYRMS NLA	60
20	2K4VL2-5_0816	PGLMTQAAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
21	2K4VL2-4	SLVMTQAAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
22	2K4VL2-6_0816	SIVMTQAAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
23	1J20VL5-2_0907	DIVMTQSAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
24	1J20VL5-6_0907	DIVMTQSAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
25	1J20VL4-3_0907	DIVLTQSAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
26	1L20VL5-0912_0917	DGVMTQSAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
27	1K3VL2-5	LIVMTQAAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
28	1K3VL2-3	LIVMTQAAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
29	1K3VL2-4	SIVMTQAAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
30	2K4VL2-5	VFVMTQAAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
31	1L5VL2-4	DIVMTQAAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
32	1L5VL3-1	LIVITQAAPSV PVT PGESVSISCR-SSKSLLSHNGNTYLYWFLQRPQSPQLLIYRMS NLA	60
33	CTC6_VL	DIVMTQAAPSVSVTPGESVSISCR-STKSLLSHNGNTYLYWLLQRPQSPQRLIYHMS NLA	60
34	CTD6_VL	DIVMTQAAPSVSVTPGESVSISCR-STKSLLSHNGNTYLYWLLQRPQSPQRLIYHMS NLA	60
35	CTA4_VL	DIKMTQSPSSVFASLGERVTITCK-ASQ-----NINSFLTWFHQKPGKSPTTLIYRTNRLL	55
36	CTB4_VL	DIKMTQSPSSVFASLGERVTITCK-ASQ-----NINSFLTWFHQKPGKSPTTLIYRTNRLL	55
37	1E9_VL	SYVLTQ-PPSVSAAPGQKVTISCSGSSSNI-----GNNYVSWYQQLPGTAPKLLIYDNNKRP	56
38	1A12_VL	SYVLTQ-PPSVSVSPGQTASVTCSGD-KL-----GNKYASWYQKPGQSPVLVIYQDKKRP	55