

SEQ ID NO.: 95

AATTTTATGCTGACTCAGCCCCACTCTGTGTGGGAGTCTCCGGGAAAGACGGTAACCATCTCCTGCACCCGCGAGC
AGTGGCACCATTGCCAGCAACTTTGTGCAGTGGTATCAACAGCGCCCGGGCAGTTGCCCCACCCAGTGATCTTT
GAGAATGACCGAAGACCCTCTGGGGTCCCTGATCGGTTCTCTGGCTCCATCGACAGCTCCTCCAATTCTGCCTCC
CTCACCATTTCGTCACTGAACACTGAGGACAAGGCTGACTACTACTGTCAGTCCTATGATAGCAGCACTCATGGG
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SEQ ID NO.: 96

TCCTATGTGCTGACTCAGCCACCCTCAGCGTCTGGGACCCCCGGGCAGAGGGTCACCATCTCTTGTCTGGAAGC
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AGGAATCATCAGCGCCCTCAGGGGTCCCTGACCGATTCTCTGGCTCCAAGTCTGGCACCTCAGCCTCCCTGGCC
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>CD47-2G4_VL

SEQ ID NO.: 97

GAAACTIACACTCAGCGAGTCTCCATCCTCCCTGTCTGTCATCTGTAGGAGACAGAGTCACCATCACTTGCCGGGCC
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GCATCTAATTTGCAGAGTGGGGTCCCTCAGGATTCAGCGGCGGTGGCTCTGGCTCCGAATTCACCTCTCACAATC
CACAGCCTGGAGTCTGAAGATTTTGC/AACTTACTACTGTCAACAGAGTTACATTACCCCTCCTTGGACGTTCCGGC
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>CD47-2G11_VL

SEQ ID NO.: 98

GAAATTGTGTTGACGAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAGCCACCCTCTCCTGCAGGGCC
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TATGCATCCAGCAGGGCCACTGGCATCCAGACAGGTTCAGCGGCAGTAGGTCTGGGACAGACTTCACTCTCACC
ATCAGCAGACTGGAGCCTGAAGATTTTGCAGTGTATTCTCTGTCAGCAGTATGATACCTCACCTCCACCTTCGGC
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>CD47-6F4_VL

SEQ ID NO.: 99

AATTTTATGCTGACTCAGCCCCACTCTGTGTGGGAGTCTCCGGGGAAGACGGTAACCATCTCCTGCACCCGCGAGC
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GAGGATAACCAAAGACCCTCTGGGGTCCCTGATCGGTTCTCTGGCTCCATCGACAGCTCCTCCAACCTCTGCCTCC
CTCACCATCTCTGGACTGAAGACTGAGGACGAGGCTGACTACTACTGTCAGTCTTATGATAGCAGCAATGTGATA
TTCGGCGGAGGGACCAAGGTCACCGTCCTA

>CD47-5H1_VL

SEQ ID NO.: 100