

Anti-A594 mAbs

1A HC	ATGTA CT TGGGACTGAACTGTGTATTCATAGTTTTCTCTTAAAAGGTGCCAGAGTGAAGTGAAGCTTGAGGAGTCTGGAGGAGGCTTGGTGCAACCTGGAGGATCCATGAAACTCTCTTGTGCTGCCTCTGGATTCACTTTTGTAGTGACGCCTGGATGGACTGGGTCCGCCAGTCTCCAGAGAAGGGGCTTGAGTGGGTTGCTGAAATTAGAAACAAAGCTAATAATCATGCAACATACTATGCTGAGTCTGTGAAAGGGAGGTTCAACCATCTCAAGAGATGATTCCAAAAGTAGTGTCTACCTGCAAATGAATAGCTTAAGAGCTGAAGACACTGGCATTATTACTGTACGTCGGTCTGGCTTACTGGGGCCAAGGGACTCTGGTCACTGTCTCTGCA
1A LC	ATGAAGTTGCCTGTTAGGCTGTTGGTGCTGATGTTCTGGATTCCCTGCTTCCAGCAGTGATGTTGTGATGACCCAAACTCCACTCTCCCTGCCTGTCAGTCTTGGAGATCAAGCCTCCATCTCTTGCAGATCTAGTCAGAGCCTTGACACAGTAATGGAAACACCTATTTACATTGGTACCTGCAGAAGCCAGGCCAGTCTCCAAAGCTCCTGATCTACAAAGTTTCCAACCGATTTTCTGGGGTCCCAGACAGGTTTCAGTGGCAGTGGATCAGGGACAGATTTCACTCAAGATCAGCAGAGTGGAGGCTGAGGATCTGGGAGTTTATTTCTGCTCTCAAAGTACACATGTTCCGCTCACGTTCCGGTGCTGGACCAAGCTGGAGCTGAAA
32B HC	ATGGGATGGAGCTGGATCTTTCTCTCTTCTGTCAGGAACTGCAGGTGTCTCTCTGAGGTCCAGCTGCAACAGTCTGGACCTGAGCTGGTGAAGCCTGGGGCTTCAGTGAAGATACCCTGCAAGGCTTCTGGATACACATTCAGTACTACAACATGGACTGGGTAAAGCAGAGCCATGGAAAGAGCCTTGAGTGGATTGGAGATTAATCCTAACAAATGGTGGTCTTACTACAACAGAAAGTTCAAGGGCAAGGCCACATTGACTATAGACAAGTCTCCAGCACAGCCTACATGGAGCTCCGACGCTGACATCTGAGGACACTGCAGTCTATTACTGTGCAGGCCGGTGGTTTCTTACTGGGGCCAAGGGACTCTGGTCACTGTCTCTGCA
32B LC	ATGAAGTTGCCTGTTAGGCTGTTGGTGCTGATGTTCTGGATTCCCTGCTTCCAGCAGTGATGTTGTGATGACCCAAACTCCACTCTCCCTGCCTGTCAGTCTTGGAGATCAAGCCTCCATCTCTTGCAGATCTAGTCAGAGCCTTGACACAGTAATGGAAACACCTATTTACATTGGTACCTGCAGAAGCCAGGCCAGTCTCCAAAGCTCCTGATCTACAAAGTTTCCAACCGATTTTCTGGGGTCCCAGACAGGTTTCAGTGGCAGTGGATCAGGGACAGATTTCACTCAAGATCAGCAGAGTGGAGGCTGAGGATCTGGGAGTTTATTTCTGCTCTCAAAGTACACATGTTCCGTACACGTTCCGAGGGGGGACCAAGCTGGAAATAAAA
49C HC	ATGGGATGGAGCTCTATCATCTCTTCTTGGTAGCAACAGCCTCAGGTGTCCACTCCCAGGTCCAAGTGCAGCAACCTGGGTCTGAGCTGGTGAGGCCTGGAGCTTCAGTGAAGCTGCCTGCAAGGCTTCTGGCTACACATTCACCAGTACTGGATGGACTGGGTGAAGCAGAGGCCTGGACAAGGCCTTGAGTGGATTGGAAATATTTATCTGGTAGTGGTAGTACTAACTACGATGAGAAGTTCAAAGCAAGGCCACACTGACTGTAGACACATCTCCAGCACAGCCTACATGCAGCTCAGCAGCCTGACATCTGAGGACTCTGCGGTCTATTACTGTACCTGTACCTGGTTTCTTACTGGGGCCAAGGGACTCTGGTCACTGTCTCTGCA
49C LC	ATGAAGTTGCCTGTTAGGCTGTTGGTGCTGATGTTCTGGATTCCCTGCTTCCAGCAGTGATGTTGTGATGACCCAAACTCCACTCTCCCTGCCTGTCAGTCTTGGAGATCAAGCCTCCATCTCTTGCAGATCTAGTCAGAGCCTTGACACAGTAATGGAAACACCTATTTACATTGGTACCTGCAGAAGCCAGGCCAGTCTCCAAAGCTCCTGATCTACAAAGTTTCCAACCGATTTTCTGGGGTCCCAGACAGGTTTCAGTGGCAGTGGATCAGGGACAGATTTCACTCAAGATCAGCAGAGTGGAGGCTGAGGATCTGGGAGTTTATTTCTGCTCTCAAAGTACACATGTTCCGTACACGTTCCGAGGGGGGACCAAGCTGGAAATAAAA