

Table S19: Positive PCR Control Synthetic Constructs

Synthetic Construct (5' to 3')	
IgH	<p>GATCGATATGGA CTGGACCTGGAGGATGATCGATACAGGTGCCCACTCCCAGGTGCAG GATCGATCAATTAGCGTACCGTTATGTGGCGAAGTACCGGCGGTACCAGCGCGGCGG AAGGTCTAGTTAGTGGCGTAGGCCTTAGTGCGTCGCGTGTAGTTAGCGTTACAATACT TACAACGATACTACAGGTGAAGAAGCCTGGGGCCTCAGTGAAGGTCTCCTGCAAGGCT TCTGGATACACCTTCACCGGCTACTATATGCACTGGGTGCGACAGGCCCCCTGGACAAG GGCTTGAGTGGATGGGATGGATCAACCCTAACAGTGGTGGCACAACTATGCACAGAA GTTTCAGGGCAGGGTCACCATGACCAGGGACACGTCCATCAGCACAGCCTACATGGAG CTGAGCAGGCTGAGATCTGACGACACGGCCGTGTATTACTGTGCGAGAGGCGTGCGCT GCGGCGTGATTCCGGGCGATCTGATCTGGGGCCAAGGGACAATGGTCACCGTCTCTCA GGCATCATAGCAACATTTCATAACATTGCGATTGATGTGCGCTGCGTGATTCCGGATGC GGTGATTGATCTGGAAGGCGGCGCGACCATTTGGCGGCCATGAAGCGGTGTATTGCCAT GCGATTAACGATCGATGTCAAGGACTACTTCCCCGAACGATCGATGACCAGCGGCGGTG CACACCTTCCGATCGAT</p>
IgK	<p>GATCGATATGAGGGTCCCTGCTCAGCTGCTGGGATCGATTGCTGCTCTGGTTCCCAGA TCGATGCCTTAGTGCGTCGCGTGTAGTTAGCGTTACAATACTTACAACGATACTACCA ATTAGCGTACCGTGCGGCCGCCGCGAAACCAGCGCGGCGGAAGGTCTAGTTAGTGGCG TAGGATGCGGTGAAAAAGCGTGGAACCCATGCGCGCATGGCGTGCCATATGGCGTGCC ATACCCATATTAGCATTAGCGCGAGCCCGGCGTGCGAACGCGGCGAACTGAGCCATAT TTTTACCGAAGTTTTTATGACCCAGTCTCCAGTCACCCTGTCTGTGTCTCCAGGGGAA AGAGCCACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGCAGCAACTTAGCCTGGTACC AGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCACCAGGGCCAC TGGTATCCAGCCAGGTTCA GTGGCAGTGGGTCTGGGACAGAGTTCACTCTCACCATC AGCAGCCTGCAGTCTGAAGATTTTGCAGTTTATTACTGTCAGCAGAAAGTGCCTGCA AAGTGATTCCGAAAGATCTGGCGTTTCGGCCAAGGGACCAAGGTGGAATCAAACGATG CGGTGATTGATCTGGAAGGCGGCGCGACCAAAGCGCCGCGGCGTGCCATGCGATTAA CCATCATAGCAACATTTCATAACATTGCGATTGATGTGCGCTGCGTGATTCCGGATCGA TAGCAGGACAGCAAGGACAGCACGATCGATTGAGCAAAGCAGACTACGAGAAACGATC GAT</p>
Igλ	<p>GATTAATCCTCCTCACTCACTGTGCAGGATCGATGGTCCTGGGCCCAGTCTGTGCTGG ATCGATCAATTAGCGTACCGTGCGTAGGTAGCGGTCCCAGCGCGGCGGAAGGTCTAGT TAGTGCGGTAGGCCTTAGTGCGTCGCGTGTAGTTAGCGTTACAATACTTACAACGATA CTACAGCGGCAGCCCGGGCCAGAGCGTGACCATTAGCTGCACCGGCACCAGCAGCGAT GTGGGCGGCTATAACTATGTGGCGTGGTATCAGCAGCATCCGGGCAAAGCGCCGAAAC TGATGATTTCATGAAGTGAGCAAACGCCCGAGCGGCGTGCCGGATCGCTTTAGCGGCAG CAAAAGCGGCAACACCGCGAGCCTGACCGTGAGCGGCCTGCAGGCGGAAGATGAAGCG GATTATTATTGCAGCAGCCTGGTGCCTGCTGCTGGTGATTCCGCTGGATCTGGTGTTTG GCGGCGGCACCAAACCTGACCGTGCTGGGCCAGCCGAAAGGCTGCCCCGTTTCATCATAG CAACATTTCATAACATTGCGATTGATGTGCGCTGCGTGATTCCGGATGCGGTGATTGAT CTGGAAGGCGGCGCGACCCCTGGCGATGGATGCGTGCCATGCGATTAAACGATCGATCAC TCTGTTCCCACCCTCGAGTGAGGAGGATCGATCAAGCCAACAAGGCCACACTTGGTGG ATCGAT</p>