-continued

```
tgcaaagtgt cgaacaaggc cctgcccgct cctatcgaaa agaccatctc caaggccaag   1080

ggccagccgc gggaacccca ggtctacact ctcccaccga gccgcgacga actgactaag   1140

aatcaagtgt cgctgacttg cctcgtcaag ggcttctacc cgtccgacat cgccgtggaa   1200

tgggagagca acggccagcc ggaaaacaac tacaagacca cccctcccgt gctggattcc   1260

gacgggtcct tcttcctgta ctcaaaactg accgtggata agtccagatg gcagcagggc   1320

aatgtctttt catgctccgt gatgcacgag gctctgcata accactacac ccagaagtcg   1380

ctgtccctgt ccccggggaa gtga                                          1404
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<210> SEQ ID NO 28
<211> LENGTH: 720
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: L2M2 light chain pro sequence

<400> SEQUENCE: 28

atggtgtcat ccgctcaatt tctcggtttg cttctcctgt gtttccaagg cacccgctgc     60

gacatcgtca tgacccagac cccattgagc ctttccgtca cgccgggaca gcccgcctcc    120

atttcctgcc gctcaagcca gtccctggag aactcaaacg ccaagaccta cctgaattgg    180

tatctgcaga agcctggaca gagcccgcag ctgctgatct acggcgtcag caacaggttc    240

tcgggcgtgc cggacagatt ctccggctcc ggaagcggaa ctgacttcac cctgaaaatc    300

tcacgcgtgg aagccgagga cgtgggcgtg tactactgcc tccaagtcac ccacgtgccg    360

tacactttcg gacaagggac taaggtcgag atcaagcgga ccgtggcggc cccctctgtg    420

ttcattttcc ctccctcgga cgaacagctg aagtcgggaa cagcctccgt cgtgtgcctg    480

ctcaacaact tctacccccg ggaagcgaag gtccagtgga aagtggataa cgcactccaa    540

tcggggaact cccaggaatc cgtgactgag caggactcga aggattccac ttactccctg    600

tcgtccaccc tgactctgag caaggccgac tacgagaagc ataaggtcta cgcctgcgaa    660

gtgacccacc agggtctgag ctcccctgtg accaagagct ttaatcgggg cgaatgttga    720
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<210> SEQ ID NO 29
<211> LENGTH: 1404
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: H2 heavy chain pro sequence

<400> SEQUENCE: 29

atgggatgga ctctcgtgtt cctttttctc ctctctgtca ctgccggggt gcattcgcaa     60

gtccagctgg tgcagtcggg accagaggtg aaaaagcccg gagagtcact taagatcagc    120

tgcaaaggct cgggatacac tttcaccaac tactggattg gttgggtcag acaggccccc    180

ggcaaaggac tggagtggat gggcgacatc taccctgggg gcgactatac caactacaac    240

gaaaagttca agggacaagt gacaatttcg gccgataaga gcattagcac tgcatacctt    300

cagtggagct cattgaaggc ctccgatacc gctatctact actgtgcgcg gtggggcctg    360

ggatactact tcgactactg gggaaggggt accttggtca cggtgtcctc cgcgtccacc    420

aagggtccct ccgtgttccc tctcgcgccg tcctcaaagt ctacctccgg tggaactgcc    480

gcgctcggtt gtctcgtgaa ggactacttc ccggagcctg tgactgtctc ctggaactcc    540
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