-continued

```
agacggctgt cctgcgctgc ctctgggttt actttctcaa gtttcggtat gcactgggtg    180

cgtcaggccc ccgggaaggg cctggaatgg gttgcttata tatcatctgg cagctccacc    240

atttcttatg ctgattccgt taagggacgc ttcaccattt ccagagacaa cagtaagaaa    300

acccttctgc tgcagatgac ctctctccgc gccgaagaca ccgcaatgta ttattgtgct    360

agagaggcct acggcagtag tatggaatac tgggggcagg ggaccctggt gaccgtgtct    420

tccgcatcta ctaagggccc                                                440
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<210> SEQ ID NO 126
<211> LENGTH: 396
<212> TYPE: DNA
<213> ORGANISM: Artificial sequence
<220> FEATURE:
<223> OTHER INFORMATION: huFR1-65_VL

<400> SEQUENCE: 126

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gaattcgcca ccatgggctg gtcttgcatt attctgttcc tggttgcaac agccactggc     60

gtccattccg aaatcgtgat gacccaatct cccgccacca tgtctacctc tcccggggac    120

cgggtgtctg tgacctgcaa ggcctctcag aatgttggcc caaacgtggc atggtatcaa    180

cagaaaccag ggcagtcacc cagagccctg atttactccg cttcttacag atattcagga    240

gttcccgccc ggttcacagg tagtgggtcc ggcactgact ttaccttgac catttccaac    300

atgcaatccg aggacctggc cgaatacttc tgtcagcagt acaattcata tccctataca    360

ttcggccagg ggaccaagct ggaaataaag cgtacg                              396
```

<210> SEQ ID NO 127
<211> LENGTH: 446
<212> TYPE: DNA
<213> ORGANISM: Artificial sequence
<220> FEATURE:
<223> OTHER INFORMATION: huFR1-65_VH

<400> SEQUENCE: 127

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aagcttgcca ccatgggctg gtcatgcata atcctgttcc tggtcgcaac cgctacaggt     60

gtacactccc aggtgcagtt ggtgcagagc ggggccgaag ttgctaagcc cggtgcaagt    120

gtaaaaatgt cctgcaaagc tagcgggtac acattcacat cctatactat gcattgggta    180

aaacagcgcc caggacaggg gctcgcctgg ataggctata ttaacccaat atcaggatac    240

acaaactaca atcagaaatt tcagggaaag gcaaccctga ccgccgacaa gtcctcttct    300

accgcatata tgcagctcaa ctccctgacc agtgaagata gcgcagtgta ttactgtgcc    360

tccggcggtg cttatggccg gaaacccatg gattactggg gacaaggcac ctccgtcaca    420

gtgagtagcg cctcaaccaa gggccc                                         446
```

<210> SEQ ID NO 128
<211> LENGTH: 17
<212> TYPE: PRT
<213> ORGANISM: Artificial sequence
<220> FEATURE:
<223> OTHER INFORMATION: Kabat Defined Mov19 HC CDR2 Murine

<400> SEQUENCE: 128

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Arg Ile His Pro Tyr Asp Gly Asp Thr Phe Tyr Asn Gln Asn Phe Lys
1               5                   10                  15

Asp
```