-continued

```
Thr Ile Gly Ile Arg Pro Ser Ser Ser Gln Ile Pro Ser Ser Ser Pro
    370                 375                 380

Thr Thr Ala Pro Ser Pro Glu Ala Gln Thr Pro Thr Thr His Thr Ser
385                 390                 395                 400

Gly Pro Ser Val Met Ala Thr Glu Glu Pro Thr Thr Pro Pro Gly Ser
                405                 410                 415

Ser Pro Gly Pro Thr Thr Glu Ala Pro Thr Leu Thr Thr Pro Glu Asn
            420                 425                 430

Ile Thr Thr Ala Val Lys Thr Val Leu Pro Gln Glu Ser Thr Ser Asn
        435                 440                 445

Gly Leu Ile Thr Ser Thr Val Thr Gly Ile Leu Gly Ser Leu Gly Leu
    450                 455                 460

Arg Lys Arg Ser Arg Arg Gln Thr Asn Thr Lys Ala Thr Gly Lys Cys
465                 470                 475                 480

Asn Pro Asn Leu His Tyr Trp Thr Ala Gln Glu Gln His Asn Ala Ala
                485                 490                 495

Gly Ile Ala Trp Ile Pro Tyr Phe Gly Pro Gly Ala Glu Gly Ile Tyr
            500                 505                 510

Thr Glu Gly Leu Met His Asn Gln Asn Ala Leu Val Cys Gly Leu Arg
        515                 520                 525

Gln Leu Ala Asn Glu
    530


<210> SEQ ID NO 20
<211> LENGTH: 7
<212> TYPE: PRT
<213> ORGANISM: artificial
<220> FEATURE:
<223> OTHER INFORMATION: Synthetic heavy chain of Mab 17F6 amino acid

<400> SEQUENCE: 20

Ala Gly Asn Tyr Arg Trp Ser
1               5


<210> SEQ ID NO 21
<211> LENGTH: 16
<212> TYPE: PRT
<213> ORGANISM: artificial
<220> FEATURE:
<223> OTHER INFORMATION: synthetic ehavy chain construct of the Mab 17F6

<400> SEQUENCE: 21

Asn Ile Tyr Tyr Ser Gly Thr Ile Ala Tyr Asn Pro Ser Leu Thr Ser
1               5                   10                  15


<210> SEQ ID NO 22
<211> LENGTH: 9
<212> TYPE: PRT
<213> ORGANISM: artificial
<220> FEATURE:
<223> OTHER INFORMATION: synthetic construct of Mab 17F6 Heavy variable
      chain

<400> SEQUENCE: 22

Asp Arg Gly Trp Leu Leu Leu Asp Tyr
1               5


<210> SEQ ID NO 23
<211> LENGTH: 10
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