146392027240SEQLIST

SEQUENCE LISTING

<110> Genentech, Inc.
GROGAN, Jane L.

<120> METHODS OF TREATING CANCER USING TIGIT
INHIBITORS AND ANTI-CANCER AGENTS

<130> 146392027240

<150> US 62/025,394
<151> 2014-07-16

<160> 26

<170> FastSEQ for Windows Version 4.0

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<400> 1
Lys Ser Ser Gln Ser Leu Tyr Tyr Ser Gly Val Lys Glu Asn Leu Leu
1 5 10 15
Ala

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<220>
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<400> 2
Ala Ser Ile Arg Phe Thr
1 5

<210> 3

<211> 9
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<400> 3
Gln Gln Gly Ile Asn Asn Pro Leu Thr
 1   5

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<400> 4
Gly Phe Thr Phe Ser Ser Phe Thr Met His
 1   5   10

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<400> 5
Phe Ile Arg Ser Gly Ser Gly Ile Val Phe Tyr Ala Asp Ala Val Arg
 1   5   10   15
Gly

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<400> 6
Arg Pro Leu Gly His Asn Thr Phe Asp Ser
 1   5   10

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<400> 7
Arg Ser Ser Gln Ser Leu Val Asn Ser Tyr Gly Asn Thr Phe Leu Ser
 1   5   10   15

<210> 8
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<400> 8
Gly Ile Ser Asn Arg Phe Ser
 1   5

<210> 9
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<400> 9
Leu Gln Gly Thr His Gln Pro Pro Thr
 1   5

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<400> 10
Gly Tyr Ser Phe Thr Gly His Leu Met Asn
 1   5   10

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<400> 11
Leu Ile Ile Pro Tyr Asn Gly Gly Thr Ser Tyr Asn Gln Lys Phe Lys
 1   5   10   15
Gly

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<400> 12
Gly Leu Arg Gly Phe Tyr Ala Met Asp Tyr
 1   5   10

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<400> 13
Asp Ile Val Met Thr Gln Ser Pro Ser Ser Leu Ala Val Ser Pro Gly
 1   5   10   15

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Glu Lys Val Thr Met Thr Cys Lys Ser Ser Gln Ser Leu Tyr Tyr Ser
            20                  25                  30
Gly Val Lys Glu Asn Leu Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln
        35                  40                  45
Ser Pro Lys Leu Leu Ile Tyr Tyr Ala Ser Ile Arg Phe Thr Gly Val
    50                  55                  60
Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Tyr Thr Leu Thr
65                  70                  75                  80
Ile Thr Ser Val Gln Ala Glu Asp Met Gly Gln Tyr Phe Cys Gln Gln
                85                  90                  95
Gly Ile Asn Asn Pro Leu Thr Phe Gly Asp Gly Thr Lys Leu Glu Ile
            100                 105                 110
Lys Arg
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<400> 14

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Asp Val Val Leu Thr Gln Thr Pro Leu Ser Leu Ser Val Ser Phe Gly
 1               5                  10                  15
Asp Gln Val Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Val Asn Ser
            20                  25                  30
Tyr Gly Asn Thr Phe Leu Ser Trp Tyr Leu His Lys Pro Gly Gln Ser
        35                  40                  45
Pro Gln Leu Leu Ile Phe Gly Ile Ser Asn Arg Phe Ser Gly Val Pro
    50                  55                  60
Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile
65                  70                  75                  80
Ser Thr Ile Lys Pro Glu Asp Leu Gly Met Tyr Tyr Cys Leu Gln Gly
                85                  90                  95
Thr His Gln Pro Pro Thr Phe Gly Pro Gly Thr Lys Leu Glu Val Lys
            100                 105                 110
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<210> 15
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<400> 15
Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Thr Gln Pro Gly Lys
 1   5   10   15
Ser Leu Lys Leu Ser Cys Glu Ala Ser Gly Phe Thr Phe Ser Ser Phe
            20   25   30
Thr Met His Trp Val Arg Gln Ser Pro Gly Lys Gly Leu Glu Trp Val
        35   40   45
Ala Phe Ile Arg Ser Gly Ser Gly Ile Val Phe Tyr Ala Asp Ala Val
    50   55   60
Arg Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Leu Leu Phe
65   70   75   80
Leu Gln Met Asn Asp Leu Lys Ser Glu Asp Thr Ala Met Tyr Tyr Cys
                85   90   95
Ala Arg Arg Pro Leu Gly His Asn Thr Phe Asp Ser Trp Gly Gln Gly
            100   105   110
Thr Leu Val Thr Val Ser Ser
        115

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<220>
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<400> 16
Glu Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Lys Pro Gly Thr
 1   5   10   15
Ser Met Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ser Phe Thr Gly His
            20   25   30
Leu Met Asn Trp Val Lys Gln Ser His Gly Lys Asn Leu Glu Trp Ile
        35   40   45
Gly Leu Ile Ile Pro Tyr Asn Gly Gly Thr Ser Tyr Asn Gln Lys Phe
    50   55   60
Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr
65   70   75   80
Met Glu Leu Leu Ser Leu Thr Ser Asp Asp Ser Ala Val Tyr Phe Cys
                85   90   95
Ser Arg Gly Leu Arg Gly Phe Tyr Ala Met Asp Tyr Trp Gly Gln Gly
            100   105   110
Thr Ser Val Thr Val Ser Ser
        115

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<220>
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<400> 17
gatgttgtgt tgactcaaac tccactctcc ctgtctgtca gctttggaga tcaagtttct 60
atctcttgca ggtctagtca gagtcttgta aacagttatg ggaacacctt tttgtcttgg 120
tacctgcaca agcctggcca gtctccacag ctcctcatct ttgggatttc caacagattt 180
tctggggtgc cagacaggtt cagtggcagt ggttcaggga cagatttcac actcaagatc 240
agcacaataa agcctgagga cttgggaatg tattactgct tacaaggtac gcatcagcct 300
cccacgttcg gtcctgggac caagctggag gtgaaa 336

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<400> 18
gaggtccagc tgcaacagtc tggacctgag ctggtgaagc ctggaacttc aatgaagata 60
tcctgcaagg cttctggtta ctcattcact ggccatctta tgaactgggt gaagcagagc 120
catggaaaga accttgagtg gattggactt attattcctt acaatggtgg tacaagctat 180
aaccagaagt tcaagggcaa ggccacattg actgtagaca agtcatccag cacagcctac 240
atggagctcc tcagtctgac ttctgatgac tctgcagtct atttctgttc aagaggcctt 300
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<210> 19
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<400> 19
Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
 1   5   10   15
Ser Leu Arg Leu Ser Cys Ala Ala Ser
            20   25

<210> 20
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<400> 20
Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 1   5   10

<210> 21
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<212> PRT
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<400> 21
Arg Phe Thr Ile Ser Ala Asp Thr Ser Lys Asn Thr Ala Tyr Leu Gln
 1   5   10   15
Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Arg
            20   25   30

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<400> 22
Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ala
 1   5   10

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<400> 23

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
 1   5   10   15
Asp Arg Val Thr Ile Thr Cys
            20

<210> 24
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<400> 24
Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile Tyr
 1   5   10   15

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<400> 25
Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr
 1   5   10   15
Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys
            20   25   30

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<400> 26
Phe Gly Gln Gly Thr Lys Val Glu Ile Lys Arg
 1   5   10