

311SG445-seq1-000001
SEQUENCE LISTING

<110> Agency for Science, Technology and Research

<120> Anti-PD-L1 Antibodies

<130> RIC/FP7082969

<160> 36

<170> PatentIn version 3.3

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Asp Val Phe Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Ile Val
35 40 45

Met Tyr Glu Thr Asn Lys Arg Pro Trp Gly Ile Pro Asp Arg Phe Ser
50 55 60

Gly Ser Lys Ser Gly Thr Ser Ala Thr Leu Asp Ile Ala Gly Leu Gln
65 70 75 80

Thr Gly Asp Glu Ala Asp Tyr Tyr Cys Gly Ala Trp Asp Ser Gly Leu
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His Trp Tyr Gln Gln Arg Pro Gly Gln Ala Pro Leu Leu Leu Val Tyr
35 40 45

Asp Asp Gly Asp Arg Pro Ser Gly Ile Pro Asp Arg Phe Ser Gly Ser
50 55 60

311SG445-seql-000001

Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser Gly Thr Gln Ala Met
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35 40 45

Ile Tyr Asp Asn Asn Glu Arg Leu Ser Gly Ile Pro Asp Arg Phe Ser
50 55 60

Gly Ser Lys Ser Gly Thr Ser Ala Thr Leu Gly Ile Ser Gly Leu Gln
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Thr Gly Asp Glu Ala Asp Tyr Tyr Cys Gly Thr Trp Asp Ser Ser Leu
85 90 95

Ser Val Val Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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Tyr Asp Val His Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu
35 40 45

311SG445-seq1-000001

Leu 50 Ile Tyr Gly Asn Ser 55 Asn Arg Pro Ser Gly 60 Val Pro Asp Arg Phe

Ser 65 Gly Ser Lys Ser 70 Gly Thr Ser Ala Ser 75 Leu Ala Ile Thr Gly 80 Leu

Gln Ala Glu Asp 85 Glu Ala Asp Tyr Tyr 90 Cys Gln Ser Tyr Asp 95 Ser Ser

Leu Ser Gly 100 Ser Tyr Val Val Phe 105 Gly Gly Gly Thr Lys 110 Leu Thr Val

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Gly 50 Arg Ile Ile Pro Ile 55 Leu Gly Ile Ala Asn 60 Tyr Ala Gln Lys Phe

Gln Gly Arg Val Thr 70 Ile Thr Ala Asp Lys 75 Ser Thr Ser Thr Ala Tyr 80

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Gln Val Gln Leu 5 Gln Gln Ser Gly Ala 10 Glu Val Lys Lys Pro Gly Ser 15

311SG445-seq1-000001

Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Ser Tyr
20 25 30

Ala Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
35 40 45

Gly Arg Ile Ile Pro Ile Leu Gly Ile Ala Asn Tyr Ala Gln Lys Phe
50 55 60

Gln Gly Arg Val Thr Ile Thr Ala Asp Lys Ser Thr Ser Thr Ala Tyr
65 70 75 80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

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Ala Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
35 40 45

Gly Arg Ile Ile Pro Ile Leu Gly Ile Ala Asn Tyr Ala Gln Lys Phe
50 55 60

Gln Gly Arg Val Thr Ile Thr Ala Asp Lys Ser Thr Ser Thr Ala Tyr
65 70 75 80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Arg Gly Gly Tyr Gly Gly Asn Ser Leu Tyr Ala Phe Asp Ile Trp
100 105 110

Gly Gln Gly Thr Met Val Thr Val Ser Ser
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311SG445-seq1-000001

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Ala Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
 35 40 45

Gly Arg Ile Ile Pro Ile Leu Gly Ile Ala Asn Tyr Ala Gln Lys Phe
 50 55 60

Gln Gly Arg Val Thr Ile Thr Ala Asp Lys Ser Thr Ser Thr Ala Tyr
 65 70 75 80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
 85 90 95

Ala Arg Ser Gly His Gly Tyr Ser Tyr Gly Ala Phe Asp Tyr Trp Gly
 100 105 110

Gln Gly Thr Leu
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311SG445-seq1-000001

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311SG445-seq1-000001

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ccaggaacag	ccccaaaat	cgtcatgtat	gaaactaata	aacgtccctg	ggggattcct	180
gaccgattct	ccggctccaa	gtctggcacg	tccgccaccc	tggacatcgc	cggactccag	240
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ccaggaacag	ccccaaaact	cctcatttat	gacaataatg	agcgactctc	agggattcct	180

311SG445-seq1-000001

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ttcggcggag ggaccaagct gaccgtccta 330

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Ala Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
 35 40 45

Gly Arg Ile Ile Pro Ile Leu Gly Ile Ala Asn Tyr Ala Gln Lys Phe
 50 55 60

Gln Gly Arg Val Thr Ile Thr Ala Asp Lys Ser Thr Ser Thr Ala Tyr
 65 70 75 80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
 Page 11

Ala Arg Ser Gly His Gly Tyr Ser Tyr Gly Ala Phe Asp Tyr Trp Gly
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Gln Gly Thr Leu Val Thr Val Ser Ser
 115 120

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