SEQUENCE LISTING

<110> MINA ALPHA LIMITED

<120> C/EBP ALPHA COMPOSITIONS AND METHODS OF USE

<130> 2058.1008PCT

<140> PCT/US2014/067132
<141> 2014-11-24

<150> 61/907,732
<151> 2013-11-22

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cagugagcgc ggcggucauu gucacugguc auagugaagc cacagaugua ugaccaguga 180

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Phe Pro Arg Gly Ala Gly Pro Ala Gln Pro Pro Ala Pro Pro Ala Ala
35 40 45

Pro Glu Pro Leu Gly Gly Ile Cys Glu His Glu Thr Ser Ile Asp Ile
50 55 60

Ser Ala Tyr Ile Asp Pro Ala Ala Phe Asn Asp Glu Phe Leu Ala Asp
65 70 75 80

Leu Phe Gln His Ser Arg Gln Gln Glu Lys Ala Lys Ala Ala Val Gly
85 90 95

Pro Thr Gly Gly Gly Gly Gly Gly Asp Phe Asp Tyr Pro Gly Ala Pro
100 105 110

Ala Gly Pro Gly Gly Ala Val Met Pro Gly Gly Ala His Gly Pro Pro
115 120 125

Pro Gly Tyr Gly Cys Ala Ala Ala Gly Tyr Leu Asp Gly Arg Leu Glu
130 135 140

Pro Leu Tyr Glu Arg Val Gly Ala Pro Ala Leu Arg Pro Leu Val Ile
145 150 155 160

Lys Gln Glu Pro Arg Glu Glu Asp Glu Ala Lys Gln Leu Ala Leu Ala
165 170 175

Gly Leu Phe Pro Tyr Gln Pro Pro Pro Pro Pro Pro Ser His Pro
180 185 190

His Pro His Pro Pro Pro Ala His Leu Ala Ala Pro His Leu Gln Phe
195 200 205

Gln Ile Ala His Cys Gly Gln Thr Thr Met His Leu Gln Pro Gly His
210 215 220

Pro Thr Pro Pro Pro Thr Pro Val Pro Ser Pro His Pro Ala Pro Ala
225 230 235 240

Leu Gly Ala Ala Gly Leu Pro Gly Pro Gly Ser Ala Leu Lys Gly Leu
245 250 255

Gly Ala Ala His Pro Asp Leu Arg Ala Ser Gly Gly Ser Gly Ala Gly
260 265 270

Lys Ala Lys Lys Ser Val Asp Lys Asn Ser Asn Glu Tyr Arg Val Arg
275 280 285

Arg Glu Arg Asn Asn Ile Ala Val Arg Lys Ser Arg Asp Lys Ala Lys
290 295 300

Gln Arg Asn Val Glu Thr Gln Gln Lys Val Leu Glu Leu Thr Ser Asp
305 310 315 320

Asn Asp Arg Leu Arg Lys Arg Val Glu Gln Leu Ser Arg Glu Leu Asp
325 330 335

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340 345 350

Ala Met Gly Asn Cys Ala
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20 25 30

Ala Pro Ala Leu Arg Pro Leu Val Ile Lys Gln Glu Pro Arg Glu Glu
35 40 45

Asp Glu Ala Lys Gln Leu Ala Leu Ala Gly Leu Phe Pro Tyr Gln Pro
50 55 60

Pro Pro Pro Pro Pro Pro Ser His Pro His Pro His Pro Pro Pro Ala
65 70 75 80

His Leu Ala Ala Pro His Leu Gln Phe Gln Ile Ala His Cys Gly Gln
85 90 95

Thr Thr Met His Leu Gln Pro Gly His Pro Thr Pro Pro Pro Thr Pro
100 105 110

Val Pro Ser Pro His Pro Ala Pro Ala Leu Gly Ala Ala Gly Leu Pro
115 120 125

Gly Pro Gly Ser Ala Leu Lys Gly Leu Gly Ala Ala His Pro Asp Leu
130 135 140

Arg Ala Ser Gly Gly Ser Gly Ala Gly Lys Ala Lys Lys Ser Val Asp
145 150 155 160

Lys Asn Ser Asn Glu Tyr Arg Val Arg Arg Glu Arg Asn Asn Ile Ala
165 170 175

Val Arg Lys Ser Arg Asp Lys Ala Lys Gln Arg Asn Val Glu Thr Gln
180 185 190

Gln Lys Val Leu Glu Leu Thr Ser Asp Asn Asp Arg Leu Arg Lys Arg
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cuuucagcug gcgcuggccu u 21

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aaacacttgc tgggattact tcttcaggtt aacccaacag aaggctcgag aaggtatatt 120

gctgttgaca gtgagcgcgg cggtcattgt cactggtcat agtgaagcca cagatgtatg 180

accagtgaca atgaccgcct tgcctactgc ctcggaattc aaggggctac tttaggagca 240

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gaccagugac aaugaccgcc uugccuacug ccucgga 97