

eolf-seql (4)
SEQUENCE LISTING

<110> Syncerus S.à r.l.

<120> Compositions and methods for assessing the risk of cancer occurrence

<130> B371682D35884

<150> EP 16305133.7
<151> 2016-02-05

<150> EP 15307190.7
<151> 2015-12-31

<160> 39

<170> PatentIn version 3.5

<210> 1
<211> 80
<212> PRT
<213> Homo sapiens

<220>
<221> SITE
<222> (1)..(80)
<223> Human progastrin

<400> 1

Ser Trp Lys Pro Arg Ser Gln Gln Pro Asp Ala Pro Leu Gly Thr Gly
1 5 10 15

Ala Asn Arg Asp Leu Glu Leu Pro Trp Leu Glu Gln Gln Gly Pro Ala
20 25 30

Ser His His Arg Arg Gln Leu Gly Pro Gln Gly Pro Pro His Leu Val
35 40 45

Ala Asp Pro Ser Lys Lys Gln Gly Pro Trp Leu Glu Glu Glu Glu
50 55 60

Ala Tyr Gly Trp Met Asp Phe Gly Arg Arg Ser Ala Glu Asp Glu Asn

65	70	eolf-seql (4)	80
		75	

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 <223> Amino acids 1-14, N-terminal extremity of human progastrin

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Ser	Trp	Lys	Pro	Arg	Ser	Gln	Gln	Pro	Asp	Ala	Pro	Leu	Gly
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 <213> Artificial

<220>
 <223> Amino acids 55-80, C-terminal extremity of human progastrin

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Gln	Gly	Pro	Trp	Leu	Glu	Glu	Glu	Glu	Glu	Ala	Tyr	Gly	Trp	Met	Asp
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Phe Gly Arg Arg Ser Ala Glu Asp Glu Asn
 20 25

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<220>
 <223> 6B5B11C10 CDR-H1

<400> 4

Gly	Tyr	Ile	Phe	Thr	Ser	Tyr	Trp
1				5			

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<210> 5
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Phe Tyr Pro Gly Asn Ser Asp Ser
1 5

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<400> 6

Thr Arg Arg Asp Ser Pro Gln Tyr
1 5

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<223> 6B5B11C10 CDR-L1

<400> 7

Gln Ser Ile Val His Ser Asn Gly Asn Thr Tyr
1 5 10

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<220>

<223> 6B5B11C10 CDR-L2

<400> 8

Lys Val Ser

1

<210> 9

<211> 9

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<223> 6B5B11C10 CDR-L3

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Phe Gln Gly Ser His Val Pro Phe Thr

1 5

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<211> 7

<212> PRT

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<223> 20D2C3G2 CDR-H1

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Gly Tyr Thr Phe Ser Ser Trp

1 5

<210> 11

<211> 8

<212> PRT

<213> Artificial

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<223> 20D2C3G2 CDR-H2

<400> 11

Phe Leu Pro Gly Ser Gly Ser Thr

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1 5

<210> 12
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<223> 20D2C3G2 CDR-H3

<400> 12

Ala	Thr	Asp	Gly	Asn	Tyr	Asp	Trp	Phe	Ala	Tyr
1				5						10

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<223> 20D2C3G2 CDR-L1

<400> 13

Gln	Ser	Leu	Val	His	Ser	Ser	Gly	Val	Thr	Tyr
1				5						10

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Lys	Val	Ser
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Ser Gln Ser Thr His Val Pro Pro Thr
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<223> 1E9D9B6 CDR-H1

<400> 16

Gly Tyr Thr Phe Thr Ser Tyr Tyr
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Ile Asn Pro Ser Asn Gly Gly Thr
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<223> 1E9D9B6 CDR-H3

<400> 18

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Thr Arg Gly Gly Tyr Tyr Pro Phe Asp Tyr
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<223> 1E9D9B6 CDR-L1

<400> 19

Gln Ser Leu Leu Asp Ser Asp Gly Lys Thr Tyr
1 5 10

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Leu Val Ser
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Trp Gln Gly Thr His Ser Pro Tyr Thr
1 5

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Gly Tyr Ser Ile Thr Ser Asp Tyr Ala
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<400> 23

Ile Ser Phe Ser Gly Tyr Thr
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<400> 24

Ala Arg Glu Val Asn Tyr Gly Asp Ser Tyr His Phe Asp Tyr
 1 5 10

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<400> 25

Ser Gln His Arg Thr Tyr Thr
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<210> 26

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<223> 1B3B4F11 CDR-L2

<400> 26

Val Lys Lys Asp Gly Ser His
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<223> 1B3B4F11 CDR-L3

<400> 27

Gly Val Gly Asp Ala Ile Lys Gly Gln Ser Val Phe Val
1 5 10

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<223> 1C10D3B9 CDR-H1

<400> 28

Gly Phe Thr Phe Thr Thr Tyr Ala
1 5

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<210> 29
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<223> 1C10D3B9 CDR-H2

<400> 29

Ile Ser Ser Gly Gly Thr Tyr Thr
1 5

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<223> 1C10D3B9 CDR-H3

<400> 30

Ala Thr Gln Gly Asn Tyr Ser Leu Asp Phe
1 5 10

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<223> 1C10D3B9 CDR-L1

<400> 31

Lys Ser Leu Arg His Thr Lys Gly Ile Thr Phe
1 5 10

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<220>

<223> 1C10D3B9 CDR-L2

<400> 32

Gln Met Ser

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<223> 1C10D3B9 CDR-L3

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Ala Gln Asn Leu Glu Leu Pro Leu Thr

1

5

<210> 34

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<213> Artificial

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<223> 2C6C3C7 CDR-H1

<400> 34

Gly Phe Ile Phe Ser Ser Tyr Gly

1

5

<210> 35

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<212> PRT

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<223> 2C6C3C7 CDR-H2

<400> 35

Ile Asn Thr Phe Gly Asp Arg Thr

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<210> 36
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<400> 36

Ala Arg Gly Thr Gly Thr Tyr
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<400> 37

Gln Ser Leu Leu Asp Ser Asp Gly Lys Thr Tyr
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Leu Val Ser
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eolf-seql (4)

<213> Artificial

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<223> 2C6C3C7 CDR-L3

<400> 39

Trp	Gln	Gly	Thr	His	Phe	Pro	Gln	Thr
1				5				