

(1) GENERAL INFORMATION

<160> NUMBER OF SEQ ID NOS: 4

(2) INFORMATION OF SEQ ID NO:1:

<210> SEQ ID NO 1

<211> LENGTH: 11 amino acids

<212> TYPE: amino acid

<223> OTHER INFORMATION:

(i) STRANDEDNESS: single

(ii) TOPOLOGY: linear

(iii) MOLECULE TYPE: peptide

<400> SEQUENCE: 1

Trp	Leu	Leu	Ala	Leu	Gly	Ser	Leu	Leu	Ala	Leu
1		5			10					

(2) INFORMATION OF SEQ ID NO:2:

<210> SEQ ID NO 2

<211> LENGTH: 9 amino acids

<212> TYPE: amino acid

<223> OTHER INFORMATION:

(iv) STRANDEDNESS: single

(v) TOPOLOGY: linear

(vi) MOLECULE TYPE: peptide

<400> SEQUENCE: 2

Arg	Leu	Ser	Leu	Leu	Leu	Val	Ile	Leu
1				5				

(2) INFORMATION OF SEQ ID NO:3:

<210> SEQ ID NO 3

<211> LENGTH: 9 amino acids

<212> TYPE: amino acid

<223> OTHER INFORMATION:

(i) STRANDEDNESS: single

(ii) TOPOLOGY: linear

(iii) MOLECULE TYPE: peptide

<400> SEQUENCE: 3

Gly	Ile	Leu	Gly	Phe	Val	Phe	Thr	Leu
1		5						

(2) INFORMATION OF SEQ ID NO:4:

<210> SEQ ID NO 4

<211> LENGTH: 9 amino acids

<212> TYPE: amino acid

<223> OTHER INFORMATION:

(i) STRANDEDNESS: single

(ii) TOPOLOGY: linear

(iii) MOLECULE TYPE: peptide

<400> SEQUENCE: 4

Ser	Leu	Tyr	Asn	Thr	Tyr	Ala	Thr	Leu
1		5						

(1) GENERAL INFORMATION

(2) INFORMATION OF PRIMERS USED IN QUANTITATIVE PCR (qPCR)

FJX1: sense 5' CCCGCAAAGGTGTCTAAAAACT 3'
 antisense 5' GTGCTGGCACAGTAAAGAATCCT 3'

MAGED4b: sense 5' CCAGAATCAGAACCGAGA 3'
 antisense 5' CCAAATCTCCGTCCTCA 3'

GAPDH: sense 5' GAAGGTGAAGGTCGGAGTC 3'
 antisense 5' GAAGATGGTGATGGGATT TC 3'