

Sequence listing 50 870 K_ST25
SEQUENCE LISTING

<110> QIAGEN GmbH
 <120> Method for analysing a complex sample by mass spectrometry
 <130> 50 870 K
 <150> EP08014583.2
 <151> 2008-08-15
 <160> 8
 <170> PatentIn version 3.5
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 <212> PRT
 <213> Homo sapiens

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Ser	Lys	Cys	Gly	Met	Cys	Cys	Lys	Thr
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Phe Pro Ile Cys Ile Phe Cys Cys Gly Cys Cys His Arg Ser Lys Cys
1 5 10 15

Gly Met Cys Cys Lys Thr
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Cys Cys Lys Thr
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<212> PRT
<213> Homo sapiens

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<223> N-terminal sequence tag determined by MS/MS

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Asp Thr His Phe
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<211> 18
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<213> Homo sapiens

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<221> site

<222> (1)..(1)

<223> restriction site for trypsin after arginin (not shown, located before serin in position 1 of this sequence - see Fig. 5b)

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<221> site

<222> (18)..(18)

<223> restriction site for trypsin after arginin (position 18 of this sequence - see Fig. 5b)

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Ser Gly Ser Val Ile Asp Gln Ser Arg Val Leu Asn Leu Gly Pro Ile
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Thr Arg

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<211> 15

<212> PRT

<213> Homo sapiens

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Val Arg Tyr Thr Lys Lys Val Pro Gln Val Ser Thr Pro Thr Leu
1 5 10 15

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<211> 16

<212> PRT

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Leu Val Arg Tyr Thr Lys Lys Val Pro Gln Val Ser Thr Pro Thr Leu
1 5 10 15

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<212> PRT

<213> Homo sapiens

<400> 8

Asp Ala His Lys Ser Glu Val Ala His Arg Phe Lys Asp Leu Gly Glu
1 5 10 15

Glu Asn Phe Lys Ala Leu Val Leu
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