ProtParam

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User-provided sequence:

```
        10         20         30         40         50         60
MVDQLRERTT MADPLRERTE LLLADYLGYC AREPGTPEPA PSTPEAAVLR SAAARLRQIH

        70         80         90        100        110        120
RSFFSAYLGY PGNRFELVAL MADSVLSDSP GPTWGRVVTL VTFAGTLLER GPLVTARWKK

       130        140        150        160        170        180
WGFQPRLKEQ EGDVARDCQR LVALLSSRLM GQHRAWLQAQ GGWDGFCHFF RTPFPLAFWR

       190        200
KQLVQAFLSC LLTTAFIYLW TRLL
```

References and documentation are available.

Number of amino acids: 204

Molecular weight: 23203.88

Theoretical pI: 9.63

Amino acid composition: CSV format

Ala (A)	21	10.3%
Arg (R)	21	10.3%
Asn (N)	1	0.5%
Asp (D)	8	3.9%
Cys (C)	4	2.0%
Gln (Q)	10	4.9%
Glu (E)	10	4.9%
Gly (G)	14	6.9%
His (H)	3	1.5%
Ile (I)	2	1.0%
Leu (L)	30	14.7%
Lys (K)	4	2.0%
Met (M)	4	2.0%
Phe (F)	12	5.9%
Pro (P)	13	6.4%
Ser (S)	10	4.9%
Thr (T)	14	6.9%
Trp (W)	7	3.4%
Tyr (Y)	5	2.5%
Val (V)	11	5.4%
Pyl (O)	0	0.0%
Sec (U)	0	0.0%
(B)	0	0.0%
(Z)	0	0.0%
(X)	0	0.0%

Total number of negatively charged residues (Asp + Glu): 18
Total number of positively charged residues (Arg + Lys): 25

Atomic composition:

Carbon	C	1055
Hydrogen	H	1635
Nitrogen	N	295
Oxygen	O	281
Sulfur	S	8

Formula: $C_{1055}H_{1635}N_{295}O_{281}S_8$
Total number of atoms: 3274

Extinction coefficients:

Extinction coefficients are in units of M^{-1} cm^{-1}, at 280 nm measured in water.

Ext. coefficient	46200
Abs 0.1% (=1 g/l)	1.991, assuming all pairs of Cys residues form cystines

Ext. coefficient 45950
Abs 0.1% (=1 g/l) 1.980, assuming all Cys residues are reduced

Estimated half-life:

The N-terminal of the sequence considered is M (Met).

The estimated half-life is: 30 hours (mammalian reticulocytes, in vitro).
>20 hours (yeast, in vivo).
>10 hours (Escherichia coli, in vivo).

Instability index:

The instability index (II) is computed to be 42.82
This classifies the protein as unstable.

Aliphatic index: 87.11

Grand average of hydropathicity (GRAVY): -0.098