ProtParam

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User-provided sequence:

```
10 20 30 40 50 60

MVPHAILARG RDVCRRNGLL ILSVLSVIVG CLLGFFLRTR RLSPQEISYF QFPGELLMRM

70 80 90 100 110 120

LKMMILPLVV SSLMSGLASL DAKTSSRLGV LTVAYYLWTT FMAVIVGIFM VSIIHPGSAA

130 140 150

QKETTEQSGK PIMSSADALL DLIRNMFPAN LVEATFKQ
```

References and documentation are available.

```
Number of amino acids: 158
Molecular weight: 17422.90
```

Theoretical pI: 9.82

```
Amino acid composition:
                        CSV format
Ala (A) 12
                7.6%
Arg (R) 10
                 6.3%
Asn (N)
         3
                 1.9%
Asp (D)
         4
                 2.5%
Cys (C)
         2
                 1.3%
Gln (Q)
                 3.2%
        5
Glu (E)
        5
                3.2%
Gly (G)
        10
                6.3%
His (H)
        2
                 1.3%
Ile (I)
        11
                 7.0%
Leu (L)
        24
                15.2%
                3.2%
Lys (K)
        5
Met (M)
        10
                 6.3%
                5.1%
Phe (F)
        8
Pro (P)
         7
                 4.4%
Ser (S)
        15
                 9.5%
Thr (T)
                 5.1%
         8
Trp (W)
                 0.6%
         1
Tyr (Y)
                 1.9%
                 8.2%
Val (V) 13
        0
                 0.0%
Pyl (0)
Sec (U)
                 0.0%
                 0.0%
 (B)
      0
 (Z)
      0
                 0.0%
 (X)
                 0.0%
```

```
Total number of negatively charged residues (Asp + Glu): 9
Total number of positively charged residues (Arg + Lys): 15
```

Atomic composition:

Carbon	C	789
Hydrogen	Н	1290
Nitrogen	N	206
0xygen	0	211
Sulfur	S	12

Formula: $C_{789}H_{1290}N_{206}O_{211}S_{12}$ Total number of atoms: 2508

Extinction coefficients:

```
Extinction coefficients are in units of \,\mathrm{M}^{-1} cm^{-1}, at 280 nm measured in water.
                    10095
Ext. coefficient
Abs 0.1% (=1 g/l) 0.579, assuming all pairs of Cys residues form cystines
                     9970
Ext. coefficient
Abs 0.1\% (=1 g/l) 0.572, 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 52.04
This classifies the protein as unstable.
Aliphatic index: 117.85
Grand average of hydropathicity (GRAVY): 0.603
```



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