ProtParam

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User-provided sequence:

```
10 20 30 40 50 60

MATPASAPDT RALVADFVGY KLRQKGYVCG AGPGEGPAAD PLHQAMRAAG DEFETRERT

70 80 90 100 110 120

FSDLAAQLHV TPGSAQQRFT QVSDELFQGG PNWGRLVAFF VFGAALCAES VNKEMEPLVG

130 140 150 160 170 180

QVQEWMVAYL ETRLADWIHS SGGWAEFTAL YGDGALEEAR RLREGNWASV RTVLTGAVAL

190

GALVTVGAFF ASK
```

References and documentation are available.

Number of amino acids: 193
Molecular weight: 20774.45

Theoretical pI: 5.40

Amino acid composition: CSV format

| Amir | no ac | id (| composition: | CSV format |
|------|-------|------|--------------|------------|
| Ala | (A) | 30 | 15.5% | |
| Arg | (R) | 13 | 6.7% | |
| Asn | (N) | 3 | 1.6% | |
| Asp | (D) | 8 | 4.1% | |
| Cys | (C) | 2 | 1.0% | |
| Gln | (Q) | 9 | 4.7% | |
| | (E) | | | |
| Gly | (G) | 21 | 10.9% | |
| His | (H) | 3 | 1.6% | |
| | (I) | | 0.5% | |
| | (L) | 17 | 8.8% | |
| Lys | (K) | 4 | 2.1% | |
| | (M) | | | |
| Phe | | 12 | | |
| Pro | | 8 | 4.1% | |
| | (S) | | 4.7% | |
| Thr | | 11 | 5.7% | |
| Trp | | 5 | 2.6% | |
| | (Y) | | 2.1% | |
| | (V) | | 8.3% | |
| Pyl | | 0 | 0.0% | |
| Sec | (U) | 0 | 0.0% | |
| | | | | |
| (B) | | | 0.0% | |
| (Z) | | | 0.0% | |
| (X) |) (|) | 0.0% | |
| | | | | |

Total number of negatively charged residues (Asp + Glu): 21 Total number of positively charged residues (Arg + Lys): 17

Atomic composition:

| Carbon | С | 930 |
|----------|---|------|
| Hydrogen | Н | 1421 |
| Nitrogen | N | 259 |
| 0xygen | 0 | 272 |
| Sulfur | S | 6 |

 $\label{eq:condition} \begin{array}{lll} \textbf{Formula:} & \mathsf{C}_{930}\mathsf{H}_{1421}\mathsf{N}_{259}\mathsf{O}_{272}\mathsf{S}_6 \\ \\ \textbf{Total number of atoms:} & 2888 \\ \end{array}$

Extinction coefficients:

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

```
Ext. coefficient 33585 Abs 0.1\% (=1 g/l) 1.617, assuming all pairs of Cys residues form cystines
```

```
Ext. coefficient 33460
Abs 0.1% (=1 g/l) 1.611, 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 18.14
This classifies the protein as stable.

Aliphatic index: 75.96

Grand average of hydropathicity (GRAVY): -0.044
```



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