

ProtParam

User-provided sequence:

102030405060

FKTNYEKRSF KVPIQANETL VGAVINNVSE AMETLTRITE ELVPVPGSVN GVNALGLVVF

708090100110120

SMCFGFVIGN MKEQGQALRE FFDSLNEAIM RLVAVIMCSA TLPITFKCLE ENNGVDKRV

130140150160

RFVLPVGATI NMDGTALYEA LAAIFIAQVN NFELNFGQII TI

[References](#) and [documentation](#) are available.

Number of amino acids: 162

Molecular weight: 17757.68

Theoretical pI: 4.87

Amino acid composition: CSV format

| | | |
|---------|----|-------|
| Ala (A) | 14 | 8.6% |
| Arg (R) | 6 | 3.7% |
| Asn (N) | 14 | 8.6% |
| Asp (D) | 3 | 1.9% |
| Cys (C) | 3 | 1.9% |
| Gln (Q) | 5 | 3.1% |
| Glu (E) | 13 | 8.0% |
| Gly (G) | 11 | 6.8% |
| His (H) | 0 | 0.0% |
| Ile (I) | 13 | 8.0% |
| Leu (L) | 14 | 8.6% |
| Lys (K) | 6 | 3.7% |
| Met (M) | 6 | 3.7% |
| Phe (F) | 12 | 7.4% |
| Pro (P) | 5 | 3.1% |
| Ser (S) | 6 | 3.7% |
| Thr (T) | 11 | 6.8% |
| Trp (W) | 0 | 0.0% |
| Tyr (Y) | 2 | 1.2% |
| Val (V) | 18 | 11.1% |
| Py1 (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): 16
Total number of positively charged residues (Arg + Lys): 12

Atomic composition:

| | | |
|----------|---|------|
| Carbon | C | 798 |
| Hydrogen | H | 1275 |
| Nitrogen | N | 205 |
| Oxygen | O | 233 |
| Sulfur | S | 9 |

Formula: C₇₉₈H₁₂₇₅N₂₀₅O₂₃₃S₉
Total number of atoms: 2520

Extinction coefficients:

This protein does not contain any Trp residues. Experience shows that this could result in more than 10% error in the computed extinction coefficient.

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

Ext. coefficient 3105
Abs 0.1% (=1 g/l) 0.175, assuming all pairs of Cys residues form cystines

Ext. coefficient 2980
Abs 0.1% (=1 g/l) 0.168, assuming all Cys residues are reduced

Estimated half-life:

The N-terminal of the sequence considered is F (Phe).

The estimated half-life is: 1.1 hours (mammalian reticulocytes, in vitro).
 3 min (yeast, in vivo).
 2 min (Escherichia coli, in vivo).

Instability index:

The instability index (II) is computed to be 22.75
This classifies the protein as stable.

Aliphatic index: 105.86

Grand average of hydropathicity (GRAVY): 0.399



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