Expasy - ProtParam 2.05.2025, 11:59



ProtParam (https://web.expasy.org/protparam)

★ Home (/protparam/)

Documentation (/protparam/protparam-doc.html)

Reference (/protparam/protpar-ref.html)

ProtParam - Results

A0A7P0T911 HUMAN (https://www.uniprot.org/uniprotkb/A0A7P0T911) (A0A7P0T911)

Description:

Solute carrier family 1 member 3

Organism:

Homo sapiens (Human)

The computation has been carried out on the complete sequence (70 amino acids).

Warning: All computation results shown below do not take into account any annotated post-translational modification.

[Documentation (/protparam/protparam-doc.html) / Reference (/protparam/protpar-ref.html)]

Number of amino acids: 70

Molecular weight: 7921.41 Theoretical pI: 10.17

Amino acid composition: CSV format Ala (A) 5 7.1% Arg (R) 5 7.1% Asn (N) 3 4.3% Asp (D) 1 1.4% Cys (C) 1 1.4% Gln (Q) 3 4.3% Glu (E) 5 7.1% Gly (G) 4 5.7% His (H) 0 0.0% 2 Ile (I) 2.9% Leu (L) 5 7.1% 9 Lys (K) 12.9% 5 Met (M) 7.1% 3 Phe (F) 4.3% 2 Pro (P) 2.9% Ser (S) 4 5.7% Thr (T) 5 7.1% Trp (W) 0 0.0% Tyr (Y) 1 1.4% Val (V) 7 10.0% Pyl (0) 0 0.0% Sec (U) 0.0% (B) 0.0% 0 (Z) 0 0.0%

Total number of negatively charged residues (Asp + Glu): 6 Total number of positively charged residues (Arg + Lys): 14

Atomic composition:

0

(X)

Carbon	C	346
Hydrogen	Н	584
Nitrogen	N	100
0xygen	0	99
Sulfur	S	6

0.0%

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Formula: $C_{346}H_{584}N_{100}O_{99}S_6$ Total number of atoms: 1135

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 $\rm M^{-1}~cm^{-1}$, at 280 nm measured in water.

Ext. coefficient 1490

Abs 0.1% (=1 g/l) 0.188, assuming all pairs of Cys residues form cystines

Ext. coefficient 1490

Abs 0.1% (=1 g/l) 0.188, 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 55.78 This classifies the protein as unstable.

Aliphatic index: 75.14

Grand average of hydropathicity (GRAVY):-0.366



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