TLR5m

Number of amino acids: 619

Molecular weight: 70139.62

Theoretical pI: 6.73

Amino acid composition Ala (A) 28 4.5% Arg (R) 27 4.4% Asn (N) 56 Asp (D) 35 9.0% 5.7% Cys (C) 11 1.8% Gln (Q) 25 4.0% Glu (E) 18 2.9% 5.8% Gly (G) 36 His (H) 15 2.4% Ile (I) 38 6.1% Leu (L) 105 17.0% Lys (K) 24 3.9% Met (M) 8 1.3% Phe (F) 43 6.9% Pro (P) 24 3.9% Ser (S) 49 Thr (T) 27 7.9% 4.4% Trp (W) 6 1.0% Tyr (Y) 20 3.2% Val (V) 24 3.9% Pyl (0) 0 0.0% Sec (U) 0 0.0% (B) 0 0.0% 0 (Z) 0.0% (X) 0 0.0%

Total number of negatively charged residues (Asp + Glu): 53 Total number of positively charged residues (Arg + Lys): 51

Atomic composition:

Carbon	С	3190
Hydrogen	Н	4949
Nitrogen	N	841
Oxygen	0	903
Sulfur	S	19

Formula: $C_{3190}H_{4949}N_{841}O_{903}S_{19}$ Total number of atoms: 9902

Extinction coefficients:

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

Ext. coefficient 63425

Abs 0.1% (=1 g/l) 0.904, assuming all pairs of Cys residues form cystines

Ext. coefficient 62800 Abs 0.1% (=1 g/l) 0.895, assuming all Cys residues are reduced

Estimated half-life:

The N-terminal of the sequence considered is ${\tt M}$ (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 37.29 This classifies the protein as stable.

Aliphatic index: 105.86

Grand average of hydropathicity (GRAVY): 0.016

TLR5s

Number of amino acids: 626

Molecular weight: 70264.83

Theoretical pI: 6.60

Amino acid composition 2.9% Ala (A) 18 Arg (R) 24 3.8% Asn (N) 50 Asp (D) 35 8.0% 5.6% Cys (C) 11 1.8% Gln (Q) 26 4.2% Glu (E) 23 3.7% Gly (G) 38 6.1% His (H) 21 3.4% Ile (I) 31 5.0% Leu (L) 110 17.6% Lys (K) 30 4.8% Met (M) 11 1.8% Phe (F) 34 5.4% Pro (P) 24 Ser (S) 61 Thr (T) 27 3.8% 9.7% 4.3% Trp (W) 6 1.0% Tyr (Y) 11 1.8% Val (V) 35 5.6% Pyl (0) 0 0.0% Sec (U) 0 0.0% (B) 0 0.0% 0 (Z) 0.0% (X) 0 0.0%

Total number of negatively charged residues (Asp + Glu): 58 Total number of positively charged residues (Arg + Lys): 54

Atomic composition:

Carbon	С	3156
Hydrogen	Н	4992
Nitrogen	N	852
Oxygen	0	918
Sulfur	S	22

Formula: $C_{3156}H_{4992}N_{852}O_{918}S_{22}$ **Total number of atoms:** 9940

Extinction coefficients:

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

Ext. coefficient 50015

Abs 0.1% (=1 g/l) 0.712, assuming all pairs of Cys residues form cystines

Ext. coefficient 49390

Abs 0.1% (=1 g/l) 0.703, 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 39.75 This classifies the protein as stable.

Aliphatic index: 106.93

Grand average of hydropathicity (GRAVY): -0.035