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ProtParam

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EAA3_HUMAN (P43005)

Excitatory amino acid transporter 3 (Excitatory amino-acid carrier 1) (Neuronal and epithelial glutamate transporter) (Sodium-dependent glutamate/aspartate transporter 3) (Solute carrier family 1 member 1) Homo sapiens (Human)

The computation has been carried out on the complete sequence (524 amino acids).

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

References and documentation are available.

Number of amino acids: 524 Molecular weight: 57100.38

Theoretical pI: 5.56

```
CSV format
Amino acid composition:
Ala (A) 41
                 7.8%
Arg (R)
        16
                  3.1%
Asn (N)
        20
                  3.8%
Asp (D)
                 4.2%
        22
Cys (C)
         6
                 1.1%
Gln (Q) 14
                  2.7%
Glu (E)
        27
                  5.2%
Gly (G)
        37
                 7.1%
His (H)
         2
                  0.4%
        50
Ile (I)
                 9.5%
Leu (L)
                10.5%
        55
Lys (K)
        29
                  5.5%
Met (M)
                 4.2%
        22
Phe (F)
        26
                 5.0%
Pro (P)
         17
                  3.2%
Ser (S)
        33
                  6.3%
Thr (T)
        38
                 7.3%
Trp (W)
         4
                 0.8%
                 2.1%
Tyr (Y)
        11
        54
                10.3%
Val (V)
Pyl (0)
         0
                 0.0%
Sec (U)
         0
                 0.0%
 (B)
                  0.0%
 (Z)
       0
                  0.0%
       0
                  0.0%
 (X)
```

Total number of negatively charged residues (Asp + Glu): 49 Total number of positively charged residues (Arg + Lys): 45

Atomic composition:

Carbon	C	2593
Hydrogen	Н	4195
Nitrogen	N	643
0xygen	0	739
Sulfur	S	28

Formula: $C_{2593}H_{4195}N_{643}O_{739}S_{28}$ Total number of atoms: 8198 15.11.2023 15:55 Expasy

Extinction coefficients:

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

Ext. coefficient 38765

Abs 0.1% (=1 g/l) 0.679, assuming all pairs of Cys residues form cystines

Ext. coefficient 38390

Abs 0.1% (=1 g/l) 0.672, 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 27.23 This classifies the protein as stable.

Aliphatic index: 115.86

Grand average of hydropathicity (GRAVY): 0.514



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