

Histamine (re)uptake by astrocytes: an experimental and computational study

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Abstract Astrocytes participate in the clearance of neurotransmitters by their uptake and subsequent enzymatic degradation. Histamine as a polar and/or protonated molecule must use a carrier to be transported across the cell membrane, although a specific histamine transporter has not been elucidated, yet. In this work we upgraded the kinetic studies of histamine uptake into neonatal rat cultured type 1 astrocytes with quantum chemical calculations of histamine pKa values in conjunction with Langevin dipoles solvation model as the first step toward microscopic simulation of transport. Our results indicate that astrocytes transport histamine by at least two carrier mediated processes, a concentration gradient dependent passive and a sodium-dependent and ATP-driven active transport. We also demonstrated that histamine protonation states depend on the polarity of the environment. In conclusion we suggest that histamine, a polar molecule at physiological pH uses at least two different mechanisms for its uptake into astrocytes – an electrodiffusion and Na⁺-dependent and ouabain sensitive active process. We emphasize relevance of knowledge of histamines protonation states at the rate limiting step of its transport for microscopic simulation that will be possible when structure of histamine transporter is known.

Keywords Astrocytes · Histamine · Rat · Transport · pKa

Introduction

Histamine [2-(imidazol-4-yl) ethylamine] belongs to a family of biogenic amines and is mostly known for its role as an inflammation mediator, neuromodulator and neurotransmitter. In recent years it has gained importance as a signaling molecule in processes of sleep-wake cycle, appetite control, learning, memory and emotion [1, 2], moreover, its signaling pathways seem to be involved in conditions such as depression, schizophrenia, Alzheimer's disease and epilepsy [2].

In the brain histamine is synthesized from histidine by a specific enzyme L-histidine decarboxylase (HD) [3] and is taken up into synaptic vesicles by the vesicular monoamine transporter 2 (VMAT2) [4]. It is released into synaptic cleft upon depolarization, which is followed by its removal from synaptic cleft in a relatively short time. The primary histamine metabolizing enzyme in the brain seems to be an intracellular histamine-N-methyltransferase (HNMT) [5]. Obviously, in order to be enzymatically degraded or recycled, histamine has to be transported either into the presynaptic neuron or into surrounding glial cells. Among glial cells astrocytes have been proposed to be the main histamine inactivation site [6]. Reports from Huszti's laboratory described bidirectional [7–9] histamine uptake into cultured glial cells prepared from embryonic chicks [7] and neonatal rats [8–10] which was Na⁺-but not Cl⁻-dependent [11]. Glial cells were also found to be involved in the continuous removal of neuronal histamine from the synaptic cleft in *in vivo* conditions [12]. In our previous work, we suggested that histamine is taken up into cultured neonatal rat astrocytes by at least two different transport

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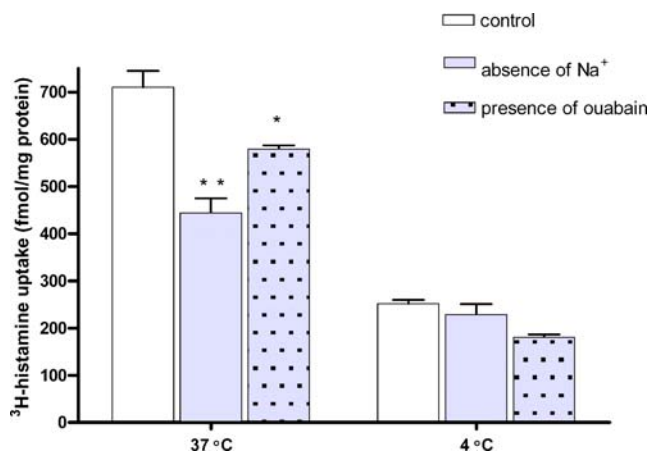


Fig. 1 Histamine uptake into neonatal rat astrocytes at 37 °C and 4 °C. Data for histamine uptake in the absence of Na⁺ (full bar) and in the presence of ouabain (dotted bar) are also presented. N=6, results were considered as statistically significant (*) for P<0.05, df=5 as determined by independent samples t-test. ** designates P<0.0003. Data represent mean +/- SEM

systems [13]. A high affinity, low capacity, energy- and temperature-dependent transport system and a low affinity, high capacity system which shows little energy and temperature dependence and shares some similarities with uptake₂.

When discussing histamine transport through physiological barriers, one should keep in mind that histamine can be

double protonated with pKa values of 5.8 (imidazole ring nitrogen atom) and 9.4 (aliphatic primary amine nitrogen atom) [14]. Therefore, at physiological pH histamine exists as an equilibrium mixture of tautomeric cations, with monocation making approximately 96% and dication 3% and only small amount of nonprotonated species [15].

Structures of histamine tautomers are shown in Fig. 3. Being protonated, it cannot freely cross membranes but needs a carrier protein, which has not been elucidated yet. In order to better understand characteristics of histamine transport it is essential to properly model the protonation states, in particular properties of histamine at physiological conditions, when transported histamine molecule is moving through environments of different polarity. It is expected that the protonation states of histamine will change during the process and it is essential for the transport to evaluate free energy differences associated with protonation and deprotonation.

Protonation states are difficult to study experimentally. When studying the protonation states in different environments it is first necessary to critically examine how different computational methods perform in aqueous solution for which pKa values are already known. Much effort has been spent recently on pKa values calculations [16–25]. Computational practice has proved that in vacuo energetics can be obtained by using high level quantum chemical

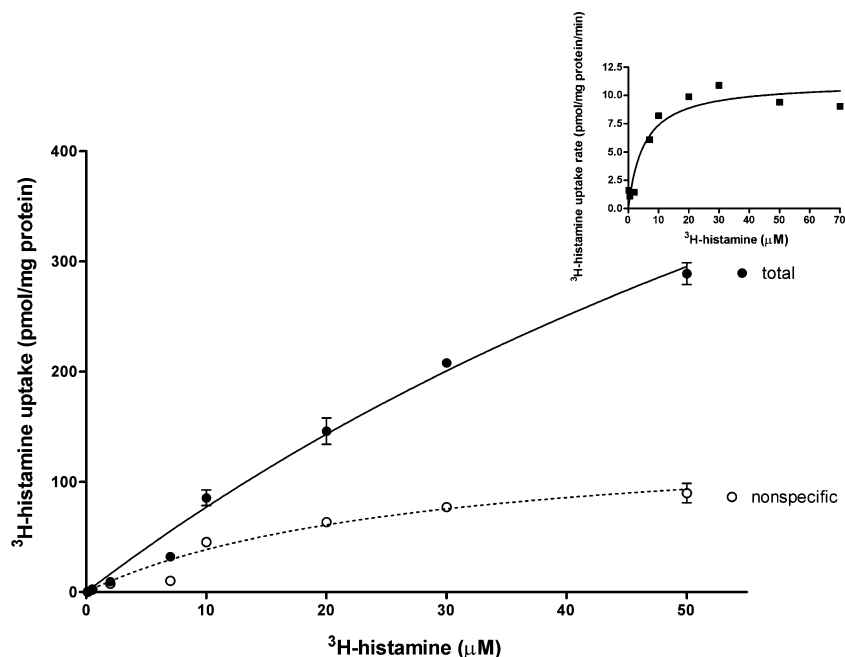


Fig. 2 Concentration dependent [³H]-histamine uptake (main) and specific histamine uptake rate (insert) into cultured rat neonatal astrocytes. Cultured astrocytes were incubated for 20 min at concentrations 0.125–100 μM at 37 °C and 4 °C. Specific uptake of [³H]-histamine was calculated as the difference between the total ([³H]-histamine uptake at 37 °C) and non-specific (uptake of [³H]-histamine at 4 °C). Kinetic

parameter analysis of specific [³H]-histamine uptake revealed K_m of 5.2±2.1 μM of and V_{max} of 11.2±1.0 pmol mg⁻¹ protein/min as determined by the best fit to $V = V_{max} \cdot X / (K_m + X)$, where V is the rate of uptake, V_{max} – maximal rate of uptake, X – concentration of [³H]-histamine and K_m – affinity constant. Each point represents the mean ± SEM of six determinations

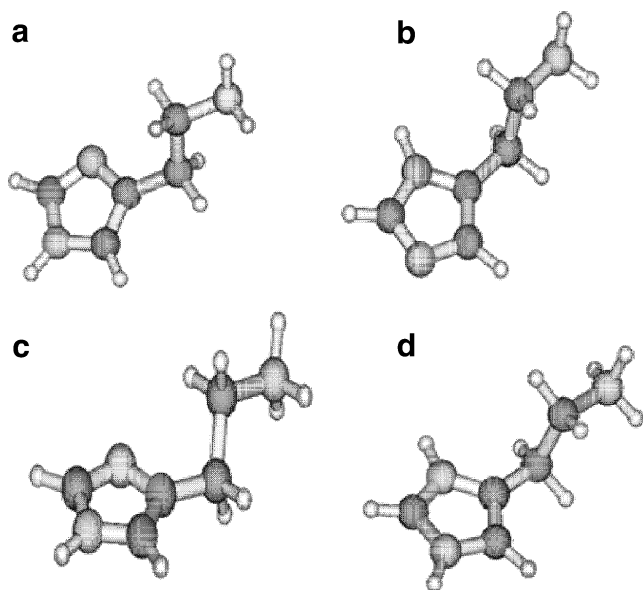


Fig. 3 Structures of various tautomers of histamine calculated on MP2/6-31++G(2d,2p) level. Full geometry optimization was performed for all the species (a) neutral α tautomer, (b) neutral π tautomer, (c) single protonated form, (d) double protonated form

calculations corrected for zero-point vibrational energy which is typically calculated in harmonic approximation. A general rule is that flexible basis should be used in conjunction with correlated wave function. Density functional theory (DFT) methods represent an attractive alternative [26, 27]. As important as in vacuo energetics is evaluation of hydration energies for different species. Critical value is hydration free energy of ionic species. If one is only looking at pKa values relative to other Brønsted acids then the error of hydration free energy for hydronium ion is the same for all the acids and the cation hydration free energy remains a critical component. Since calculation of proton hydration free energy is demanding and subject of significant uncertainty, consideration of its experimental value is a reasonable choice when calculating pKa values. Experimentally free energy for transfer of H^+ from gas phase to aqueous solution is determined more accurately than hydration free energy of hydronium ion. Former value includes formation of hydronium ion. It is interesting that the reported experimental values for proton transfer from gas phase to the aqueous solution span a wide range, e.g., $-252.4 \text{ kcal mol}^{-1}$ [28], $-259.2 \text{ kcal mol}^{-1}$ [29]. The later is an average of values obtained from measurements of standard hydrogen potentials in the range from -253.6 to $-260.8 \text{ kcal mol}^{-1}$. The most recent cluster ion solvation data suggest a hydration free energy of $-264 \text{ kcal mol}^{-1}$ [30], which we used for calculations of histamine pKa values.

Duarte's group [31] carefully examined pKa values of histamine by using solvent reaction field in conjunction with several flexible basis sets on Hartree-Fock (HF),

Møller-Plesset perturbational theory of the second order (MP2) and various DFT levels. The authors emphasized the difficulty to predict a pKa value relative to water because of poor performance of the methods to predict hydration free energy of the hydronium ion. Therefore their calculated pKa values were relative to acetate and a reasonable agreement with the experiment was found. Protonation states and therewith associated pKa values are relevant for enzyme catalysis [32]. Hudaky and Perczel [33] recently performed careful calculations concerning protonation states of histidine involved in catalytic triad of serine proteases. They stressed relevance of conformational changes for pKa values. In the context of calculating the pKa values of flexible systems, it should be noted that the pKa is actually an average of all possible conformations of the protein. Kamerlin *et al.* [25] demonstrated that careful thermal averaging should be performed for both ionized and neutral states of the residue.

To establish whether different protonation states of histamine affect the features of its electrodiffusion and active transport across the cell membrane, we combined uptake kinetic studies with calculation of pKa values of histamine by applying Langevin dipoles (LD) of Florian and Warshel [34–36]. We also critically examined the existing computational methods for calculation of pKa values in aqueous solution. Protonation states were discussed in the context of different environmental polarities histamine faces during the transport.

Materials and methods

Materials

All tissue culture reagents, except fetal bovine serum (Cambrex, Belgium) were obtained from GIBCO, Great Britain. [3H]-histamine ($525.4 \text{ GBq mmol}^{-1}$) was purchased from Perkin Elmer, USA.

Astrocyte culture

Rats were used in accordance with the National Institutes of Health Guidelines for the Care and Use of Laboratory Animals and Permission for Use of Laboratory Animals 323-02-232/2005/2 and 34401-81/2008/5 issued by Veterinary Administration of Republic of Slovenia. Astrocyte cultures were prepared from the cortex of neonatal Wistar rats (3 days old, both sexes) and cultured as previously described [37]. Briefly, cells were grown in high-glucose Dulbecco's modified Eagle's medium (DMEM), containing 10% fetal bovine serum (FBS), 1 mM pyruvate, 2 mM glutamine, and $25 \mu\text{g ml}^{-1}$ gentamycin in 95% air -5% CO_2 . Confluent cultures were shaken at 225 rpm overnight, and

the medium was changed the next morning; this was repeated a total of three times. After the third overnight shaking, the cells were trypsinized and cultured for 24 h in 10 μM cytosine arabinoside. After reaching confluence again the cells were subcultured into 12-well clusters and grown for an additional 3 weeks.

Uptake of [^3H]-histamine into cultured astrocytes

Monolayer cultures in 12-well clusters were preincubated for 30 min in uptake buffer (25 mM HEPES, 125 mM NaCl, 4.8 mM KCl, 1.2 mM KH_2PO_4 , 1.2 mM MgSO_4 , 1.4 mM CaCl_2 , and 5.6 mM glucose, pH 7.4) at 37 $^\circ\text{C}$ (total uptake) and at 4 $^\circ\text{C}$ (non-specific uptake). After preincubation the histamine transport was initialized by addition of radiolabeled histamine, the final radioligand concentration being 0.125 μM –50 μM for the dose response curve and 0.125 μM for other experiments. After 20 min the reaction was stopped by placing the dishes into an ice-water bath. The buffer was quickly removed and the dishes were washed twice with ice-cold uptake buffer. The cells were lysed in 0.6 mL of 0.5 M NaOH. 0.5 mL of each sample was transferred to a scintillation vial and the radioactivity was measured. Protein content was determined in the remaining aliquots (0.1 ml) using Bio-Rad method. To determine the effect of Na^+ on histamine uptake, NaCl in the uptake buffer was replaced with choline chloride. The activity of $\text{Na}^+\text{K}^+\text{ATP}$ -ase was inhibited by adding 1 mM ouabain.

Data analysis

The uptake experiments were routinely carried out in triplicates and each experiment was repeated twice. The results are expressed as means \pm SEM. The kinetic parameters (K_m and V_{max}) were calculated by a non-linear regression method using software Prism4 version 4.00 (GraphPad Software Inc., San Diego, USA). Comparison of data among groups was carried out using independent samples t-test. Differences were considered significant at $P < 0.05$.

Computational methods

All calculations were performed at the National Institute of Chemistry in Ljubljana on the eight processor CHIEFTEC Linux-based personal computer. Series of ab initio and DFT calculations encoded in the Gaussian03 suite of programs had been applied. A full geometry optimization was performed for all the species. We limited our study to two basis sets 6-31+G (d,p) and 6-31++G (2d,2p). Former basis set is double zeta and is augmented with polarization functions of heavy atoms and hydrogen atoms and additionally contains diffuse functions on heavy atoms. To the latter

basis set are added diffuse functions to hydrogen atoms and polarization functions to heavy atoms and hydrogen atoms.

Ab initio calculations were performed on the HF level of theory. In order to include some electron correlation we applied Møller-Plesset perturbation of the second order. Moreover, we applied DFT method B3LYP. The DFT method B3LYP, has the Becke three parameter hybrid gradient-corrected exchange functional [26], combined with the gradient-corrected correlation functional of Lee, Yang and Parr [27]. The same basis sets as for HF level of theory were also used for the B3LYP for the species mentioned above. Full geometry optimization was performed for all the species and vibrational analysis was performed in the harmonic approximation.

The LD by Florian and Warshel [34–36] was used to calculate the solvation free energies for all the species. The Merz-Kollman partial atomic charges also encoded in the Gaussian03 served as an input for the LD model that was later built in the ChemSol 2.1 program [34, 35]. In the LD procedure hydration free energy is calculated as reversible work necessary to build a solute charge distribution in the field of dipoles on a cubic grid. The charge distribution is represented by Merz-Kollman atomic charges and thermal averaging is performed by charging the coordinate origin.

Results

Na^+ -dependence and ouabain sensitivity

Astrocytes were prepared from cortices of neonatal rats (postnatal day 3) as previously described [11]. At that point, the cultures contained 93–100% type 1 astrocytes, determined by immunochemical staining for glial fibrillary acidic protein [37]. Since the time course of histamine uptake has already been reported in our recent paper [13], we proceeded with experiments concerning sodium dependence and ouabain sensitivity. Figure 1 presents the influence of Na^+ replacement with choline and ouabain treatment of cultured astrocytes at two different temperatures. The depletion of Na^+ in cultured medium resulted in a significant decrease of histamine uptake by cultured astrocytes at 37 $^\circ\text{C}$, whereas no Na^+ -dependence of histamine uptake was measured at 4 $^\circ\text{C}$. The addition of ouabain affected only histamine uptake into cultured astrocytes at 37 $^\circ\text{C}$ and had no effect on its transport into astrocytes at 4 $^\circ\text{C}$ (Fig. 1).

Concentration-dependent histamine uptake

Further on, we determined the kinetic properties of histamine uptake (Fig. 2). Cultured astrocytes were incubated for 20 min with ^3H -histamine in a concentration

Table 1 Stabilities of tautomers of neutral histamine. All energies are in kcal mol⁻¹ relative to α tautomer. E is classical energy difference. ZPE stands for zero point energy correction. E_{dry} is sum of E and ZPE. ΔG_{hydr} is hydration contribution to this process and total free energy difference. ΔG stands for free energy of hydration plus E_{dry}

Method	E	ZPE	E_{dry}	ΔG_{hydr}	ΔG
HF/6-31+G(d,p)	0.51	0.08	0.59	-0.54	0.05
HF/6-31++G(2d, 2p)	0.45	0.09	0.54	-0.49	0.05
B3LYP/6-31+G(d,p)	0.52	0.07	0.59	-0.40	0.19
B3LYP/6-31++G(2d,2p)	0.53	0.07	0.60	-0.45	0.15
MP2/6-31+G(d,p)	0.69	0.06	0.75	0.75	0.56
MP2/6-31++G(2d,2p)	0.53	0.09	0.62	0.62	0.33

range from 0.125 – 50 μM at 37 °C (total uptake) and 4 °C (non-specific uptake). Histamine was taken up in a concentration-dependent manner (Fig. 2) at both temperatures used. The specific histamine transport represents the difference between the total (37 °C) and the non-specific one (4 °C). The kinetic analysis of specific histamine transport calculated by non-linear regression analysis revealed Michaelis constant (Km) of $5.2 \pm 2.1 \mu\text{M}$ and maximal rate (Vmax) of $11.2 \pm 1.0 \text{ pmol mg}^{-1} \text{ protein/min}$ (Fig. 2, insert).

Quantum chemical calculations

Quantum chemical calculations in conjunction with free energies of hydration yield free energies of different protonation states. Calculated energies of both neutral tautomers of histamine are collected in Table 1. On all levels of theory the α form is predicted to be more stable in the gas phase for about 0.6 kcal mol⁻¹. Inclusion of solvent decreases the free energy difference but the α form is still more stable. The free energy differences are in the range 0.05 kcal mol⁻¹ (HF/6-31G(d,p) level) to 0.56 kcal mol⁻¹ (MP2/6-31G(d,p)), what gives strong evidence that both neutral forms are present. This fact can be used when

Table 2 The first pKa value for histamine calculated on different levels of theory in conjunction with Langevin dipoles solvation model. The calculated pKa value is for the process where monoprotonated histamine donates a proton to a water molecule. All values are in kcal mol⁻¹. E is classical energy difference for the process. ZPE stands for zero point energy correction. E_{dry}

Method	E	ZPE	E_{dry}	ΔG_{hydr}	ΔG	pKa ⁽¹⁾
HF/6-31+G(d,p)	237.52	-10.22	227.30	-204.67	22.63	16.7
HF/6-31++G(2d, 2p)	237.50	-10.25	227.25	-204.44	22.81	16.8
B3LYP/6-31+G(d,p)	233.48	-9.64	223.84	-205.24	18.60	13.7
B3LYP/6-31++G(2d,2p)	233.02	-9.61	223.41	-204.98	18.43	13.6
MP2/6-31+G(d,p)	234.37	-9.79	224.58	-205.38	19.20	14.1
MP2/6-31++G(2d,2p)	232.34	-9.73	222.62	-205.25	17.37	12.8
Experimental value					12.77	9.4

interpreting experimental data such as spectra that are superpositions of both components.

Table 2 presents the calculated energetics associated with the first pKa value of histamine. The first deprotonation step is proton transfer from the aliphatic protonated amino group of the single charged histamine to a water molecule giving rise to neutral histamine and hydronium ion. In vacuo energetics gives evidence that in the gas phase this is an endothermic process associated with about 220 kcal mol⁻¹ energy difference. Hydration reduces this gap for about 205 kcal mol⁻¹, giving rise to 20 kcal mol⁻¹ free energy difference. The calculated pKa values are in the range between 16.7 and 12.8 and therefore they overestimate the experimental value of 9.4.

Table 3 is a collection of the energetics associated with the second pKa value of histamine. The second deprotonation step is transfer of the proton from the imidazole nitrogen atom of the double charged histamine to a water molecule giving rise to monoprotonated histamine at aliphatic amino group and hydronium ion. In contrast to the first deprotonation step, the gas phase energetics favors this process. The gas phase reaction is exothermic on all levels of theory. The convergence in terms of the level of theory is poorer than for the first deprotonation step. Hydration in this case favors the process for approximately 140 kcal mol⁻¹. The entire process in terms of free energy is uphill for about 9–16 kcal mol⁻¹, giving rise to pKa values of 6.7–12.2. Inspection of Tables 2 and 3 gives confidence that calculation of pKa values on MP2 level in conjunction with flexible basis sets is reliable enough to proceed with replacement of aqueous solution with protein environment.

In order to critically examine the quality of the applied solvation model, we collected the free energies of solvation for different species in Table 4. It is clearly evident that the calculated hydration free energy of hydronium ion is not favorable and is the dominant source of the too high pKa values. Therefore, our strategy to consider the experimental hydration free energy of a proton is fully justified. Zhan and

is the sum of E and ZPE. ΔG_{hydr} is hydration contribution to this process and $\Delta G = E_{\text{dry}} + \Delta G_{\text{hydr}}$ and $\text{pKa}^{(1)} = \Delta G / (\ln(10)k_{\text{B}} T)$. Note that the experimental value for proton hydration free energy of -264 kcal mol⁻¹ was used

Table 3 The second pKa value for histamine calculated on different levels of theory in conjunction with Langevin dipoles solvation model. The calculated pKa value is for the process where diprotonated histamine donates a proton to a water molecule giving rise to monoprotinated form and hydronium ion. All values are in kcal mol⁻¹.

Method	E	ZPE	E _{dry}	ΔG _{hydr}	ΔG	pKa ⁽²⁾
HF/6-31+G(d,p)	165.68	-9.07	156.61	-140.05	16.56	12.2
HF/6-31++G(2d, 2p)	165.57	-9.06	156.51	-139.40	17.11	12.6
B3LYP/6-31+G(d,p)	160.62	-8.61	152.01	-139.67	12.34	9.1
B3LYP/6-31++G(2d,2p)	160.76	-8.62	152.14	-139.11	13.03	9.6
MP2/6-31+G(d,p)	158.58	-8.65	149.93	-139.99	9.94	7.3
MP2/6-31++G(2d,2p)	156.97	-8.59	148.38	-139.32	9.06	6.7
Experimental value					7.88	5.8

E is classical energy difference for the process. ZPE stands for zero point energy correction. E_{dry} is sum of E and ZPE. ΔG_{hydr} is hydration contribution to this process and $\Delta G = E_{\text{dry}} + \Delta G_{\text{hydr}}$. $\text{pKa}^{(2)} = \Delta G / (\ln(10)k_{\text{B}} T)$. Note that the experimental value for proton hydration free energy of -264 kcal mol⁻¹ was used

Dixon emphasized the difficulty to properly measure and calculate hydronium and hydroxide ion hydration free energy [38]. Some authors avoided this difficulty by considering other Brønsted basis as a water molecule giving rise to relative pKa values [31]. The structures of various histamine tautomers are presented in Fig. 3.

Discussion

Histamine acts as a neurotransmitter only in the mature rat brain and as neuromodulator and growth promoting agent in the neonatal rat brain. Regardless the fact that histamine probably adopts its role as a classical signaling molecule not earlier than by postnatal day 20 [39], it is also taken up into neonatal rat astrocytes with a half time of 1.3 min [13] that is similar to its blood inactivation time of 0.5–3.5 min [40, 41]. In addition, astrocytes express mRNA for HNMT, histamine degrading enzyme [13, 42], which supports the notion that uptake and subsequent metabolism by glial cells of released histamine from neurons may play an important role in regulating extracellular histamine concentration.

Astrocytes most probably represent the main histamine inactivation site within CNS. They take up significantly more histamine than synaptosomes [43] and shake off cells (mixture of microglia, oligodendrocytes, O2 progenitor

cells and type 2 astrocytes) [11]. Figure 1 shows that the replacement of sodium chloride in the incubation medium with choline chloride or ouabain treatment significantly, but not completely decrease histamine uptake into cultured neonatal rat astrocytes at 37 °C. This reconfirms our previous finding [13] concerning the involvement of the two different transport pathways, active and passive one by which histamine is carried into astrocytes at 37 °C. The transport of histamine into shake off cells is probably electrodiffusion, as it seems not to be affected either by lack of sodium ions in the incubation medium nor by ouabain treatment [11]. The same phenomenon (electrodiffusion, a process that depends on concentration gradient and membrane potential) might explain the process by which histamine accumulates within astrocytes at 4 °C, which demonstrates concentration dependence [11], but no Na⁺-dependence or ouabain sensitivity (Fig. 1). Saturation profile at 4 °C suggests the presence of a limited number of carrier proteins, that allow just for finite flux and not a simple diffusion process as one might expect. More than one transport pathway probably serves as backup mechanism for situations where high affinity transporter proteins are underrepresented or overloaded.

Carrier-operated histamine uptake into cultured rat type 1 astrocytes revealed the Km value of 5.2±2.1 μM and Bmax of 11.2±1.0 pmol g⁻¹ protein/min (Fig. 2). The

Table 4 Free energies of hydration calculated by Langevin dipoles solvation model. All values are in kcal mol⁻¹. Nα is α tautomer neutral form of histamine. PROT1 is monoprotinated at amino nitrogen atom form and PROT 2 is a double protonated form

Method	Nα	PROT1	PROT2	H ₃ O ⁺	H ₂ O
HF/6-31+G(d,p)	-10.58	-69.91	-193.86	-97.27	-8.44
HF/6-31++G(2d, 2p)	-9.97	-69.53	-194.13	-97.26	-6.94
B3LYP/6-31+G(d,p)	-9.79	-68.55	-192.88	-96.21	-8.04
B3LYP/6-31++G(2d,2p)	-9.11	-68.13	-193.02	-96.80	-6.39
MP2/6-31+G(d,p)	-10.55	-69.17	-193.18	-96.76	-8.37
MP2/6-31++G(2d,2p)	-9.84	-68.59	-193.27	-96.88	-6.58
Experimental value				-105	-6.4

measured K_m value was significantly higher than K_m obtained by Huszti et al. [8] - 0.19 μM in astrocytes and 0.30 μM in rat cerebral endothelial cells [43], but similar to the K_m value of histamine uptake in the P2 fraction of rat synaptosomes [44]. Histamine is a known substrate for organic cation transporters (OCTs) and cultured rat astrocytes express OCT1, OCT2 and OCT3 [45] that transport monoamines in such manner. Nevertheless, the K_m values for histamine transport by OCTs are significantly different (180–940 μM) [44] than the affinity for histamine uptake obtained in our study, so the observed carrier(s) seem(s) to represent a new entity. It remains a major challenge for structural biology to determine the structure of histamine carrier that will pave the way to understanding the transport on a molecular level. Active, ATP-driven component of histamine uptake is highly temperature dependent. As such, it is obviously a process that involves soft modes and is entropy sensitive. To understand the nature of histamine uptake, which includes the temperature dependence, future studies will involve *sine dubio* fine interplay between experimental work and molecular simulation. The latter will require all atom simulation including the transport protein, water and it will be performed in conjunction with chemically reactive species.

While being carried from extracellular to intracellular space histamine molecule is moving through environments of different polarity. We calculated stabilities of two neutral tautomers of histamine and absolute histamine pKa values by using *ab initio* calculations on HF, MP2 and B3LYP level with flexible basis sets in conjunction with LD solvation model. For the neutral form the α tautomer was predicted to be more stable on all applied levels of theory, although the free energy difference is small enough to allow both tautomers to be present and this should be considered when interpreting the experimental data. We analyzed the (free) energy contributions to both histamine pKa values. Moreover, by calculating hydration free energy by LD one is considering all hydration layers around neutral histamine and both protonated forms. Polarity of the interior of the transport protein is different from the polarity of the extracellular as well as the polarity of the cytoplasm. Thus, the protonation states of histamine will presumably change during the transport process. Histamine can be transported in a neutral form or in any of the protonated forms, where pKa is a measure of probability of the protonation states. Our calculations suggest that addition of the second proton in the gas phase is less favorable than addition of the first one. The reduced affinity can be attributed to the repulsion between proton and monoprotonated cation (Fig. 3).

An important message of this study and pKa calculations in general is that proper treatment of hydration is as important as the *in vacuo* energetics. This important issue is

not properly understood by several theoretical chemists who want to explain enzymatic reactions by performing gas phase calculation. Consideration of the corresponding reaction in aqueous solution remains the most successful method of computational enzymology [16].

Summary

We can conclude that histamine; a polar molecule at physiological pH cannot move freely across cell membrane and as such is taken up into neonatal astrocytes by two different processes – an electrodiffusion and an active, Na^+ -dependent and ouabain sensitive process. One must keep in mind that histamine protonation state at the selectivity filter or at the rate limiting step of the histamine transport, such as any transporter involved, is of high relevance for design of the analogs that may act as histamine uptake inhibitors. When structure of histamine transporter is determined, histamine protonation studies will help to clarify the transport mechanisms as well as provide foundation for drug design. We expect that studies of histamine transport will contribute to better understanding of various neurological disorders and their future treatment development.

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Conflict of interests Authors declare no conflict of interests.

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