Although described some time ago, gamma-butyrobetaine esters and related compounds have not gained much attention from researchers, and their physiological function remains obscure. Formerly we detected GBB-esterase enzymatic activity in rat blood serum using phenylated gamma-butyrobetaine as an artificial substrate of the enzyme and HPLC. The aim of the present work was to develop an assay that would enable spectrophotometric or colorimetric determination of the reaction products of GBB-esterase activity and to reveal individual proteins performing GBB-esterase activity in rat blood serum. For this purpose gamma-butyrobetaine 1-naphthyl ester was synthesised. Hydrolysis of this ester releases 1-naphthol, which increases the optical absorbance at 322 nm. We have shown that the enzymatic hydrolysis of GBB 1-naphthyl ester in rat blood serum is due to GBB-esterase activity. An attempt was done to purify the enzyme from rat blood serum. By combining DEAE Sepharose at pH 4.2 and affinity chromatography with procainamide we achieved a 68-fold enrichment of GBB-esterase activity in our preparations. Separation of fraction proteins in 2D protein electrophoresis with following mass-spectrometry indicated that GBB esterase activity in rat blood serum is performed in part by carboxylesterase.

KEY WORDS: 1-naphtol, 2D electrophoresis, colorimetry, HPLC, spectrophotometry

Although described some time ago, gamma-butyrobetaine esters and related compounds have not gained much attention from researchers. Numerous studies performed by E. Hosein et al. in the 1960s and 1970s suggest that these compounds have multiple functions (1-4). For example, in addition to being a carnitine precursor, gamma-butyrobetaine (GBB) can undergo esterification in mammalian brain tissue (4). The structure of GBB ethyl ester strikingly resembles that of acetylcholine. The distance between positively and negatively charged poles in both molecules is almost identical. The existence of a specific signal transfer system based on GBB esters was suggested recently (5-7), hypothesis was based on increase in GBB levels in stressed animals (8) and cholinergic activity of GBB esters (3). Seeking for arguments that support this hypothesis, in a series of experiments we established GBB-esterase enzymatic activity in rat blood serum using phenylated gamma-butyrobetaine as an artificial substrate of the enzyme (9). The aim of this study was to develop an assay for spectrophotometric or colorimetric determination of GBB-esterase reaction products and to identify individual proteins performing GBB esterase activity in rat blood serum.

MATERIALS AND METHODS

Reagents. DEAE-Sepharose Fast Flow, Heparin Sepharose CL- B and CNBr-activated Sepharose TM
4B were supplied by GE Healthcare Life Sciences (Sweden). Bio Gel P 150 was from BioRad, USA. Polygram SIL G/UV254 plastic sheets (40 mm x 80 mm) were from Macherey-Nagel (Germany) and polyethylene glycol 20000, 6-aminocaproic acid from Roth (Germany). Fast Blue RR salt and carboxylesterase from horse liver was from Fluka (USA), and acetylthiocholine iodide from Acros organics (USA).

**BW 284c51 (1,5 bis {4-allyldimethyl ammonium phenyl} pentane-3-one-dibromide), tetraisopropylpyrophosphoramide (iso-OMPA), butyrylcholine chloride, and procainamide, 1-(3-dimethylamino-propyl)-3-ethylcarbodiimide hydrochloride were from Sigma (Germany).**

**Blood serum**

All manipulations with animals were performed upon Ethics Committee approval and in accordance with Lithuanian and Latvian regulations, which are compliant with EU rules. Blood was obtained from male Wistar rats decapitated under slight ether narcosis. Collected blood was stored at 37 °C for 2 h and then centrifuged at 5000 g for 20 min to obtain serum in the supernatant. Serum was stored at 4 °C until used (up to 3 to 10 days).

**GBB esterase activity measurement**

GBB esterase activity was measured on a Perkin Elmer Lambda-20 spectrophotometer using GBB-O-NF as substrate essentially as described by Morgan et al. (10). Briefly, the hydrolysis of GBB-O-NF to 1-naphthol was observed through increase in absorbance at 322 nm (the extinction coefficient for 1-naphthol is 2.2 L mmol$^{-1}$ cm$^{-1}$). The reactions were conducted in a cuvette (V=1 mL) and initiated by adding substrate (10 µL of 10 mmol L$^{-1}$ stock in water). Initial velocities were measured in 1 mL of the assay mixture composed of 100 mmol L$^{-1}$ of sodium phosphate buffer, pH 7.4; 0.1 mmol L$^{-1}$ GBB-O-NF, and serum or fractionated enzyme preparation at 25°C for 10 min. In the measurements, the spontaneous hydrolysis of GBB-O-NF and the effects of additives on substrate hydrolysis were eliminated by using blank tubes containing appropriate combinations of compounds. The unit of GBB-O-NF esterase was defined as the amount of enzyme which catalyzes the formation of 1 µmol of the product per min under these conditions.

The effects of potential GBB-O-NF esterase inhibitors were measured on a Perkin Elmer Lambda-20 spectrophotometer using GBB-O-NF as substrate. The reactions were performed after preincubation of enzymatically active fractions with additive (0.1 mmol L$^{-1}$) for 30 min. Subsequently, GBB-O-NF was added to a final concentration of 0.1 mmol L$^{-1}$, and absorbance was monitored at 322 nm to determine hydrolysis products. Initial velocities were measured in 1 mL of the assay mixture composed of 100 mmol L$^{-1}$ sodium phosphate buffer, pH 7.4 and tested compounds at 25°C. 50 µL (30 µg of protein) of the fraction fractionated on DEAE Sepharose and 10 µL (10 µg of protein, 0.5 U mg$^{-1}$) of carboxylesterase from horse liver were taken for measurements.

**Dialysis of rat serum**

10 mL of serum was dialysed against 0.02 mol L$^{-1}$ sodium acetate buffer pH 4.5 containing 1 mmol L$^{-1}$ EDTA for 12 h, and against 0.02 mol L$^{-1}$ sodium acetate pH 4.2 containing 1 mmol L$^{-1}$ EDTA for 3 h.

**DEAE Sepharose chromatography at pH 4.2.**

10 mL of dialysed serum (700 mg of total protein) were applied to a DEAE Sepharose column (0.8 cm x 14 cm), eluted with 25 mL of 0.02 mol L$^{-1}$ sodium acetate buffer at pH 4.2, then with 1:1 gradient of (0 to 0.15) mol L$^{-1}$ NaCl in 0.02 mol L$^{-1}$ sodium acetate buffer pH 4.2, containing 1 mmol L$^{-1}$ EDTA. The flow rate was 0.3 mL min$^{-1}$. Fractions were concentrated with polyethylenglycol 20000.

**Procainamide - Sepharose 4B chromatography**

Procainamide gel was synthesised according to the method of Cuatrecasas (11). Briefly, 3 g of water-swollen CNBr-activated Sepharose 4B was washed with water and 1 mmol L$^{-1}$ HCl (V=20 mL) three times each. Washed CNBr-activated Sepharose 4B was then coupled with 6-aminohexanoic acid (0.26 g) in 0.2 mol L$^{-1}$ Na$_2$CO$_3$/NaHCO$_3$ pH 8.3 buffer containing 0.5 mol L$^{-1}$ NaCl. Excess of ligand was removed by washing, and the remaining active groups were blocked with 0.1 mol L$^{-1}$ Tris-HCl buffer at pH 8.0. Procainamide (0.275 g; 100 µmol mL$^{-1}$ of gel) was coupled with 10 mL of 6-aminohexanoic Sepharose 4B by adding 10 mL of 0.2 mol L$^{-1}$ 1-(3-dimethylamino-propyl)-3-ethylcarbodiimide hydrochloride (0.38 g), maintaining the pH at 4.5 with 1 mol L$^{-1}$ HCl for 2 h and then stirring at room temperature (20 to 25) °C for 24 h. The gel was washed three times with 20 mL of
The concentration of procainamide bound to the gel was determined by measuring the absorbance of the washings ($\epsilon = 16150 \, \text{L mol}^{-1} \text{cm}^{-1}$ at 278 nm). The difference between the starting amount and the amount in the washings gave the gel-bound concentration. Affinity columns (0.8 cm x 5 cm) were equilibrated with 20 mmol L$^{-1}$ potassium phosphate and 1 mmol L$^{-1}$ EDTA pH 7, and washed with 20 mL of 20 mmol L$^{-1}$ potassium phosphate and 1 mmol L$^{-1}$ EDTA pH 7.0. The column was then eluted with 38 mL of 0.2 mol L$^{-1}$ NaCl in buffer. Finally, the proteins more tightly bound to procainamide were eluted with 40 mL of 1 mol L$^{-1}$ sodium acetate, pH 6.0. They were then incubated overnight in a buffer containing 65 mmol L$^{-1}$ sodium acetate (pH 6.0), 0.5 mg mL$^{-1}$ acetyliothiocholine, 5 mmol L$^{-1}$ sodium citrate, 3 mmol L$^{-1}$ cupric sulphate, and 0.5 mmol L$^{-1}$ potassium ferricyanide. The reactions were stopped by washing the gels with water.

**Preparation of samples for MALDI-TOF**

The excised gel spots were crushed to 1 mm$^2$ sized slices and dehydrated with 50 % acetonitrile. The gel slices were then dried under vacuum and rehydrated with 30 µL of 25 mmol L$^{-1}$ NH$_4$HCO$_3$ (pH 8.3). Proteins in the gel slices were digested overnight with 25 mg mL$^{-1}$ of modified trypsin (Promega, Madison, WI, USA) at 37 °C. The peptides were then washed twice from the gel with 50 µL 5 % TFA in 50 % acetonitrile. The wash-outs were collected and dried under vacuum. For MALDI-TOF, peptides were diluted in 3 µL of 50 % acetonitrile solution which contained 0.01 % trifluoroacetic acid (TFA). 0.8 µL of each sample with matrix (alpha-cyano-4-hydroxycinnamic acid) were loaded on a MALDI plate. Analysis was performed on a MALDI-TOF mass spectrometer (Perspective Biosystems, Framingham, MA, USA). Data from the EMBL database PeptIdent were used for peptide identification.

**Determination of protein concentrations**

During purification, protein content in chromatography fractions was determined by measuring absorbance at 280 nm. In the pooled samples, protein concentrations were determined with Bradford’s method (15), using BSA as a standard.
RESULTS

Applicability of the assay

GBB-esterase activity in rat blood serum used to be determined with phenylated gamma-butyrobetaine as an artificial substrate. This assay was rather complicated and time-consuming, as the reaction products were detected by HPLC (9). We have developed a new assay which enables spectrophotometric or colorimetric determination of the reaction products. For this purpose, we synthesized gamma-butyrobetaine 1-naphthyl ester. Hydrolysis of this ester should lead to a release of 1-naphthol, which increases optical absorbance at 322 nm (Figure 1). As 1-naphthol forms an insoluble complex with Fast Blue RR dye it has found its application in determining GBB-esterase activity in gels. Hydrolysis of 1-naphtylacetate is successfully used to determine acetylcholinesterase activity in gels. At the initial stage of the study, it was important to test the applicability and specificity of gamma-butyrobetaine 1-naphthyl ester. This was performed using inhibitor analysis. Native or partially fractionated rat blood serum can hydrolyse gamma-butyrobetaine naphtyl ester, and the reaction can be observed by monitoring the absorbance at 322 nm. Acetylcholine (0.1 mmol L\(^{-1}\)) had no effect on the reaction rate and did not compete with the enzyme substrate. In contrast, phenylated gamma butyrobetaine ethyl ester in the same concentration significantly decreased the reaction rate, as did gamma butyrobetaine ethyl and methyl esters. This indicated that the enzyme which hydrolysed gamma-butyrobetaine 1-naphtyl esters was inhibited by GBB-esters, and therefore confirmed GBB-esterase activity. In the remainder of this paper, the GBB-O-NF esterase activity will be called GBB-esterase activity.

Figure 1 Hydrolysis of naphthylated gamma-butyrobetaine ester.

Purification of GBB-esterase: selection of optimal protocol

Several purification protocols were applied during the early stages of the study. Salting out with (NH\(_4\))\(_2\)SO\(_4\) turned out to be ineffective, as enzymatic activity was lost in high salt concentrations. Enzyme activity was also lost during fractionation on DEAE Sepharose at pH 7.4. Fractionation on Biogel P150 was also ineffective (not shown). After fractionation on DEAE Sepharose at pH 4.2, GBB-esterase activity was found in a fraction eluted by 0.05 mol L\(^{-1}\) NaCl. If the active fraction obtained in the previous procedure was applied on the DEAE Sepharose at pH 6.5 and eluted with (0 to 0.5) mol L\(^{-1}\) NaCl gradient, activity was detected in the fraction eluted at 0.315 mol L\(^{-1}\). Protein composition of the fraction was still complex (not shown). When the activity-containing fraction obtained after fractionation on DEAE Sepharose at pH 4.2 was dialysed, concentrated, and/or applied to a heparin Sepharose column at pH 7, proteins were eluted by (0 to 0.5) mol L\(^{-1}\) NaCl gradient. Enzyme activity was detected in a fraction eluted with 0.125 mol L\(^{-1}\) NaCl. In this case only four protein bands were detected by Coomassie staining (not shown). Fractionation on DEAE Sepharose at pH 4.2 with procainamide - Sepharose 4B chromatography turned out to be the most effective protocol (16).

Inhibitory analysis of GBB-esterase activity of the fraction obtained on DEAE Sepharose chromatography at pH 4.2

Inhibitory analysis of the enzyme hydrolysing gamma-butyrobetaine 1-naphtyl esters in the partially purified fraction of rat blood serum on DEAE Sepharose column at pH 4.2 showed that GBB-esterase was not inhibited by acetylthiocholine iodide, which is known as a substrate of cholinesterases, or
by BW 284c51 (1,5 bis {4-allyldimethyl ammonium phenyl} pentane-3-one dibromide), which is a specific inhibitor of acetylcholinesterase. However iso-OMPA, a specific vertebrate butyrylcholinesterase inhibitor, partly inhibited this activity. Inhibition was obtained with butyrylcholine, GBB-methylester, GBB-ethylester, and phenylated GBB ethyl ester. All these data preclude any role of acetylcholinesterase in the hydrolysis of GBB-esters, but point to another specific enzyme. Involvement of butyrylcholinesterase cannot be excluded (Table 1).

**Purification of the protein fraction enriched with GBB-esterase activity for 2D electrophoresis**

Serum was dialysed against acetate buffer at pH 4.2. The dialysate was fractionated by ionic exchange chromatography on DEAE Sepharose at pH 4.2. Fractions manifesting GBB-esterase activity (Figure 2) were pooled and subjected to chromatography on a procainamide column. Procainamide is a ligand

**Table 1** Effect of potential inhibitors on GBB-esterase activity in rat blood serum fraction obtained on DEAE Sepharose and on purified carboxylesterase from horse blood serum. All substances were applied in a 0.1 mmol L<sup>-1</sup> concentration; experiments were repeated in triplicate.

<table>
<thead>
<tr>
<th>Compound</th>
<th>Effect on GBB-esterase activity (remaining activity)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Acetylthiocholine</td>
<td>No effect</td>
</tr>
<tr>
<td>Butyrylcholine</td>
<td>20</td>
</tr>
<tr>
<td>Gamma butyrobetaine methyl ester</td>
<td>58</td>
</tr>
<tr>
<td>Gamma butyrobetaine ethyl ester</td>
<td>47</td>
</tr>
<tr>
<td>Phenyl gamma butyrobetaine ethyl ester</td>
<td>21</td>
</tr>
<tr>
<td>Izo-OMPA</td>
<td>50</td>
</tr>
<tr>
<td>BW 284c51</td>
<td>No effect</td>
</tr>
</tbody>
</table>

**Figure 2** Fractionation on DEAE Sepharose column at pH 4.2. Active fraction of the enzyme was eluted with 0.05 mol L<sup>-1</sup> NaCl in 25 mL of 0.02 mol L<sup>-1</sup> Na acetate buffer at pH 4.2. The bracket indicates pooled active fractions used for further purification.
Procainamide affinity chromatography was found to be efficient for purifying most cholinesterases (16), as procainamide is structurally related to choline and also to gamma-butyrobetaine.

Proteins with GBB-esterase activity were eluted with 0.2 mol L\(^{-1}\) NaCl. Proteins bound more tightly to procainamide were eluted with 1 mol L\(^{-1}\) NaCl (Figure 3).

Table 2: Steps in the partial purification of GBB-esterase activity. The unit of GBB-O-NF esterase is defined as the amount of enzyme which catalyses the formation of 1 µmol of 1-naphthol (2.2 L mmol \(^{-1}\) cm \(^{-1}\) min\(^{-1}\) under optimal conditions (100 mmol L\(^{-1}\) sodium phosphate buffer, pH 7.4; 0.1 mmol L\(^{-1}\) GBB-O-NF).

<table>
<thead>
<tr>
<th>Step</th>
<th>Total protein / mg</th>
<th>Total activity / U</th>
<th>Specific activity / U mg(^{-1})</th>
<th>Yield / %</th>
<th>Increase in activity</th>
</tr>
</thead>
<tbody>
<tr>
<td>Dialysed blood serum, pH 4.2</td>
<td>700</td>
<td>1190</td>
<td>1.7</td>
<td>100</td>
<td>1 - fold</td>
</tr>
<tr>
<td>DEAE Sepharose chromatography</td>
<td>29.6</td>
<td>855</td>
<td>28.9</td>
<td>72</td>
<td>17 - fold</td>
</tr>
<tr>
<td>Procainamide sepharose chromatography</td>
<td>3.96</td>
<td>456</td>
<td>115.2</td>
<td>38</td>
<td>68 - fold</td>
</tr>
</tbody>
</table>

We achieved a 68-fold increase in specific activity. The active fraction in SDS-gels was represented by several polypeptides (Figure 4). Staining native gels of the active fraction for GBB-esterase activity revealed several bands (Figure 5A). One of them contained about 95 % of the total GBB-esterase activity. Four distinct bands of 160 kDa, 70 kDa, 56 kDa, and 27 kDa were observed.

Figure 6 2D electrophoresis of 150 µg of active fraction after DEAE Sepharose fractionation. First dimension - Immobiline Dry Strips; pH 3 to 10; Coomassie stained gel.

Figure 4 10 % SDS-PAGE stained with Coomassie dye. 1 - active fraction after DEAE chromatography; 2 - active fraction eluted with 0.2 mol L\(^{-1}\) NaCl from procainamide gel; 3 - fraction eluted from procainamide gel with 1 mol L\(^{-1}\) NaCl. Protein molecular weight standard positions are indicated on the left.

Figure 5 A - 7.5 % native PAGE of the fraction eluted with 0.2 mol L\(^{-1}\) NaCl from procainamide gel stained for GBB-O-NF esterase activity; B - 10 % SDS PAGE of the fraction eluted with 0.2 mol L\(^{-1}\) NaCl from procainamide gel stained for GBB-O-NF esterase activity.

Specific for the choline binding site. Procainamide affinity chromatography was found to be efficient for purifying most cholinesterases (16), as procainamide is structurally related to choline and also to gamma-butyrobetaine.
Figure 7 2D electrophoresis patterns: A - 180 µg of active fraction after DEAE Sepharose fractionation; B - 150 µg of active fraction after procainamide gel fractionation. First dimension - Immobiline Dry Strips; pH 4 to 7; a-r denotes spots which were subjected to MALDI-TOF

Table 3 Characterised proteins of the partially purified GBB-esterase fraction after separation with 2D electrophoresis.

<table>
<thead>
<tr>
<th>Spot</th>
<th>Protein name</th>
<th>MW / kDa</th>
<th>pI</th>
<th>Protein function</th>
</tr>
</thead>
<tbody>
<tr>
<td>a</td>
<td>Ubiquitin carboxyl-terminal hydrolase</td>
<td>128 5.3</td>
<td></td>
<td>Cleaves ubiquitin fusion protein substrates</td>
</tr>
<tr>
<td>b</td>
<td>Liver carboxylesterase</td>
<td>58 5.5</td>
<td></td>
<td>Involved in the detoxification of xenobiotics and in the activation of ester and amide prodrugs. Catalyses the following reaction: A carboxylic ester + H₂O = an alcohol + a carboxylate. Belongs to the type-B carboxylesterase/lipase family</td>
</tr>
<tr>
<td>c</td>
<td>Calretinin</td>
<td>31 4.9</td>
<td></td>
<td>Calcium-binding protein abundant in auditory neurons</td>
</tr>
<tr>
<td>d</td>
<td>Ras-related protein Rab-3A</td>
<td>25 4.9</td>
<td></td>
<td>Involved in exocytosis by regulating a late step in synaptic vesicle fusion. Could play a role in neurotransmitter release by regulating membrane flow in the nerve terminal I</td>
</tr>
<tr>
<td>e</td>
<td>Cytosol aminopeptidase</td>
<td>56 6.8</td>
<td></td>
<td>Presumably involved in the processing and regular turnover of intracellular proteins. Catalyzes the removal of unsubstituted N-terminal amino acids from various peptides</td>
</tr>
<tr>
<td>f</td>
<td>Acetylcholinesterase</td>
<td>65 5.7</td>
<td></td>
<td>Rapidly hydrolyzes choline released into the synapse</td>
</tr>
<tr>
<td>g</td>
<td>Alpha-2-HS-glycoprotein</td>
<td>36 6.0</td>
<td></td>
<td>Presumably inhibits both insulin-receptor tyrosine kinase activity and insulin-stimulated receptor autophosphorylation</td>
</tr>
<tr>
<td>h</td>
<td>Stress-induced-phosphoprotein 1</td>
<td>63.6 4</td>
<td></td>
<td>Mediates the association of the molecular chaperones HSC70 and HSP90</td>
</tr>
<tr>
<td>k</td>
<td>Alpha-1-antiproteinase</td>
<td>44 5.6</td>
<td></td>
<td>Proteinase inhibitor</td>
</tr>
<tr>
<td>l</td>
<td>Alpha-1-antiproteinase</td>
<td>44 5.6</td>
<td></td>
<td>Proteinase inhibitor</td>
</tr>
<tr>
<td>m</td>
<td>Cytosol aminopeptidase</td>
<td>56 6.8</td>
<td></td>
<td></td>
</tr>
<tr>
<td>n</td>
<td>Haptoglobin beta chain peptidase S1</td>
<td>27 6.3</td>
<td></td>
<td>Transport of released hemoglobin</td>
</tr>
<tr>
<td>o</td>
<td>Haptoglobin beta chain</td>
<td>27 6.3</td>
<td></td>
<td></td>
</tr>
<tr>
<td>p</td>
<td>Haptoglobin beta chain</td>
<td>27 6.3</td>
<td></td>
<td></td>
</tr>
<tr>
<td>r</td>
<td>Haptoglobin</td>
<td>37 6.0</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>
35 kDa were seen on SDS-PAGE gels stained for GBB-esterase activity. The 70 kDa band contained about 95% of the total activity (Figure 5B). Some of the above bands also manifested cholinesterase activity; however, the distribution of both activities was not identical (not shown).

2D electrophoresis and characterisation of proteins

The results of two-dimensional electrophoresis of the fraction obtained with DEAE Sepharose chromatography are shown in Figure 6. Figure 7 shows two-dimensional electrophoretic separation of active fractions after DEAE Sepharose chromatography and chromatography on a procainamide column in a narrow pH gradient (pH 4 to 7). The marked spots were excised and taken for MALDI-TOFF analysis. Table 3 summarises information about the characterised proteins. Carboxylase appeared to be the most suitable for hydrolysing GBB-esters in blood serum. To test this possibility, we measured the activity of commercially available horse liver carboxylase using the standard assay. Carboxylase manifested GBB-esterase activity, but unlike the GBB-esterase-containing fraction of the blood serum, it was sensitive to all inhibitors tested (Table 1). Its high sensitivity to gamma butyrobetaine methyl ester suggests that the enzyme does indeed possess GBB-esterase activity.

DISCUSSION

This study has shown that enzymatic hydrolysis of GBB naphthyl ester to 1-naphthol in rat blood serum is due to GBB-esterase activity. Our previous results (9) indicated that acetylcholine esterase and butyrylcholinesterase were incapable of performing this reaction. We therefore looked further by combining DEAE Sepharose at pH 4.2 and affinity chromatography with procainamide to achieve a 68-fold enrichment in GBB-esterase activity. Further characterization indicated that carboxylase was an important component of this fraction. We observed high GBB-esterase activity in a band of about 70 kDa in one-dimensional native and SDS electrophoresis (Figure 5). The molecular weight of carboxylase released in serum is 71 kDa (13). In 2D electrophoresis the protein was detected in a high-molecular weight complex; the reason for this might be the level of glycosylation. Alternatively, we may have detected some other carboxylase isoenzyme. Carboxylases are homologous to a large extent (17). This conclusion was confirmed by the fact that partially purified enzyme preparations from rat blood serum and purified horse liver carboxylesterase were sensitive to different inhibitors. We believe that there may be one or more other enzymes which are also able to hydrolyse GBB-esters. These could form the high-molecular complex observed on 2D gel together with carboxylesterase.

The hydrolysis of GBB-esters by carboxylesterase indicates the physiological significance of the substance. Along with blood acetylcholinesterase, butyrylcholinesterase, and paraoxonase, this enzyme is involved in detoxifying numerous hazardous substances, including organophosphates (18, 19). It interacts with a broad range of substrates and performs arylesterase activity (20). It is likely to protect against excessive amounts of GBB-esters synthesised in other organs for functional purposes. As GBB-esters trigger a strong cholinergic reaction (21–23), in excess they could be dangerous. An investigation of GBB-esterases in other organs, especially in the brain, might shed more light on the functional significance of GBB-esters.

Acknowledgements

This work was supported by the Taiwanese-Latvian-Lithuanian project “Purification and characterisation of the gamma-butyrobetaine esterase”.

REFERENCES

Sažetak

KARBOKSILESTERAZA DJELOMICE HIDROLIZIRA ESTERE GAMA-BUTIROBETAINA U SERUMU ŠTAKORA


KLJUČNE RIJEĆI: 2D elektroforeza, HPLC, kolorimetrija, 1-naftol, spektrofotometrija

CORRESPONDING AUTHOR:

Nikolajs Sjakste
Latvian Institute of Organic Synthesis,
Aizkraukles Street 21, Riga
LV1006, Latvia
E-mail: Nikolajs.Sjakste@lu.lv