

## Short Communication

# G-Protein Linked Effect of Pasteurella Multocida Toxin on GABA Outflow from Neocortical Synaptosomes

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## Abstract

*Pasteurella Multocida* Toxin (PMT) stimulates diverse cellular signal transduction pathways by activating heterotrimeric G proteins through deamidation of a  $\alpha$ -subunit glutamine residue of the G-protein. Since mammalian cells are able to take up PMT probably by receptor-mediated endocytosis, the question was raised whether PMT is capable of modulating  $\gamma$ -aminobutyric acid (GABA) outflow from neocortical synaptosomes of the rat *in vitro*. In our experiments PMT did not modify basal GABA outflow. However, GABA release induced by potassium ions was significantly reduced. Here the toxin effect was not modulated by the GABA<sub>B</sub>-receptor agonist baclofen. These results let us to suggest that PMT activates G-proteins of the inhibitory metabotropic GABA<sub>B</sub> autoreceptor on GABAergic nerve terminals.

**Keywords:** G-protein; Pasteurella multocida toxin;  $\gamma$ -aminobutyric acid; Synaptosomes; Rat

## Abbreviations

PMT: Pasteurella Multocida Toxin; GABA:  $\gamma$ -Aminobutyric Acid; GTP: Guanosine Triphosphate; AUC: Area Under the Curve; GABA-R: GABA Receptor; Bcl: Baclofen

## Introduction

The Gram negative opportunistic bacterium *Pasteurella multocida* produces a 146-kDa protein toxin (*Pasteurella multocida* toxin, PMT) which stimulates diverse cellular signal transduction pathways by activating heterotrimeric G proteins, thereby arresting the G protein in the active state [1]. The toxin substrates primarily are G $\alpha_q$ , G $\alpha_{12/13}$  and the G $\alpha_i$ -family proteins. The toxin deamidates a glutamine of the  $\alpha$ -subunits of heterotrimeric G-proteins [2]. This glutamine is essential for hydrolyzing GTP. The question was raised whether PMT is intracellularly capable of modulating neurotransmitter release from neocortical synaptosomes of the rat *in vitro*, since cellular (or synaptosomal) up-take of PMT by mammalian cells was described through receptor-mediated endocytosis [3].

## Materials and Methods

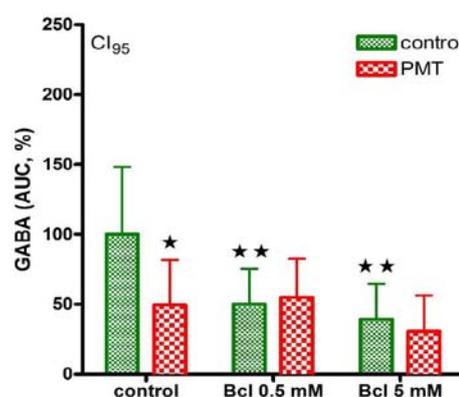
Animal experiments were performed according to the National Institute of Health Guide for the Care and Use of Laboratory Animals and all efforts were made to minimize both suffering and number of animals. Wistar rats (weighing 200-300 g, N=6, University of Freiburg, Germany) were decapitated under CO<sub>2</sub> anesthesia, and the brains were carefully removed. The neocortex was dissected and immediately placed in ice-cold saline.

The buffer used for the preparation, suspension, incubation, and superfusion of synaptosomes contained (mM): NaCl 121, KCl 1.8, CaCl<sub>2</sub> 1.3, KH<sub>2</sub>PO<sub>4</sub> 1.2, MgSO<sub>4</sub> 1.2, NaHCO<sub>3</sub> 25, glucose 10, and was saturated before use with 95% O<sub>2</sub>/5% CO<sub>2</sub>; pH 7.4. Synaptosomes were preincubated during 30 min, 37°C, 95% O<sub>2</sub>/5% CO<sub>2</sub> with Pasteurella multocida toxin (PMT, 100 nM). To study the release of endogenous

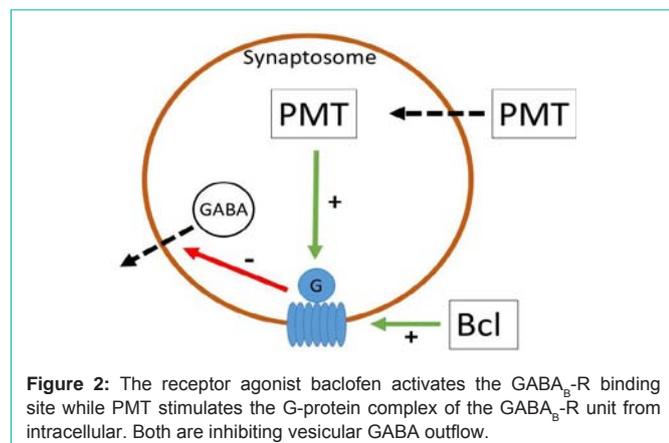
GABA, synaptosomes were then transferred to glass microfiber GF/B filters (Whatman, Dassel, Germany) into 24 superfusion chambers within a water bath (37°C). Synaptosomes were covered with GF/B filters within the superfusion chambers to prevent their wash out.

Then, collection of four fractional samples followed (5 min intervals, 0.2 ml/min flow rate; 37°C, presence of baclofen as indicated). A 10 min interval of superfusion before stimulation corresponded to resting conditions. Synaptosomes were stimulated for 2 min using KCl 15 mM.

The GABA contents of fractional samples and of synaptosomes were determined by High Performance Liquid Chromatography (HPLC). After pre-column derivatization with o-phthalaldehyde and sodium sulfite for 30 min, GABA values were measured using HPLC with electrochemical detection. The HPLC system consisted of a C18 column (Eurospher 100, 5  $\mu$ m, column size 250x4 mm)



**Figure 1:** GABA outflow in the presence of potassium ions (15 mM) was given as area under the curve (%). In the verum group pasteurella multocida toxin (PMT, 100 nM) was preincubated for 30 min. Baclofen (Bcl) 0.5 or 5 mM was added as indicated. \*significant, when compared to control; \*\*significant, when compared to -Bcl; p < 0.05.



and a pre-column (30x4 mm). The isocratic mobile phase (0.1 M sodium phosphate buffer, pH 4.5, containing 0.5 mM EDTA and 25% methanol) was previously degassed using helium and pumped at a flow rate of 1.0 ml/min. The compounds were detected electrochemically using a glassy carbon electrode set at a potential of 900 mV, relative to an Ag/AgCl reference electrode (Waters 460 electrochemical detector, Millipore Corporation, Eschborn/Ts., Germany).

GABA outflow was expressed in nM, as mean  $\pm$  Standard Error of the Mean (SEM). The average of baseline samples was defined as 100%. Differences between the means of treatments and their corresponding controls were tested with one-way Analysis of Variance (ANOVA). Subsequent multiple comparisons were made by the Kruskal-Wallis test and t-test for pair wise comparisons as indicated. Areas Under the Curve (AUC) were calculated by GraphPad Prism software. Statistical significance was set at  $p < 0.05$ .

## Results and Discussion

Synaptosomes are isolated nerve terminals and serve as an important model system for studying the molecular mechanisms of synaptic function in the brain [4]. In our preparation of rat neocortical synaptosomes, GABA outflow was  $15.0 \pm 3$  nM under basal conditions. Preincubation of synaptosomes with *Pasteurella multocida* toxin (PMT, 100 nM) did not modify basal GABA outflow ( $14.2 \pm 2$  nM). The GABA amounts per fraction were comparable to those found by Rassner and co-workers [5] in synaptosomes of rat and also human neocortex. Here fluorescence-activated particle sorting experiments corroborated GABAergic synaptosomes to be highly present in such preparations [5,6].

Potassium ions (15 mM) significantly increased outflow of vesicular GABA to  $28.8 \pm 3.7$  nM, as described earlier [7,8]. This potassium-induced enhancement was completely abolished under PMT conditions (Figure 1), most probably through G-protein activation. GABA<sub>B</sub> Receptors (GABA<sub>B</sub>-R) are metabotropic, composed of 7 transmembrane domain subunits linked to G-proteins [9], and also presynaptically located on GABAergic synaptic terminals as autoreceptors [10,11]. As all GABA<sub>B</sub>-R subtypes these autoreceptors are inhibitory receptors functionally [8,10]. During Baclofen (Bcl) incubation, i.e. by activation of GABA<sub>B</sub>-R, GABA outflow significantly decreased and no further potassium-induced stimulation could be observed (Figure 1). In the case of PMT preincubation, the inhibitory baclofen effect (0.5 or 5 mM, respectively) remained unchanged,

without any further decrease of GABA outflow (Figure 1). The finding that the effects of PMT and baclofen are not additive implicates activation of the same effector cascade. The receptor agonist baclofen activates GABA<sub>B</sub>-R from the extracellular site while PMT stimulates the G-protein complex of the GABA<sub>B</sub>-R unit intracellularly as shown in (Figure 2). For this, it is necessary for PMT to pass through cell or as in our experiments the synaptosomal membrane by receptor-mediated endocytosis as proposed by Pettit and co-workers [3]. Since PMT uptake could be competed with mixed gangliosides, the receptor was suggested to belong to this class [3]. Although the precise mechanism of translocation of PMT into the cytosol remains to be elucidated [2], the effects of PMT can, additionally, be used as a tool for biochemical studies like other bacterial toxins as pertussis or cholera toxin that also modulate G-proteins [12,13].

## Conclusion

The results described above give evidence that *Pasteurella multocida* toxin may pass through cellular membranes into synaptosomes where it activates G-proteins associated to the GABA<sub>B</sub> autoreceptor inhibiting release of vesicular GABA from neocortical synaptosomes of the rat.

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