

Evaluation of energetic metabolism in the rat brain after meningitis induction by *Klebsiella pneumoniae*

Barichello T, Simões LR, Generoso JS, Carradore MM, Moreira AP, Panatto AP, Costa CS, Filho AS, Jeremias IC, Bez GD, Streck E. Evaluation of energetic metabolism in the rat brain after meningitis induction by *Klebsiella pneumoniae*.

Background: Bacterial meningitis is an infection of the central nervous system characterised by strong inflammatory response. The brain is highly dependent on ATP, and the cell energy is obtained through oxidative phosphorylation, a process which requires the action of various respiratory enzyme complexes and creatine kinase (CK) as an effective buffering system of cellular ATP levels in tissues that consume high energy.

Objectives: Evaluate the activities of mitochondrial respiratory chain complexes I, II, III, IV and CK activity in hippocampus and cortex of the Wistar rat submitted to meningitis by *Klebsiella pneumoniae*.

Methods: Adult Wistar rats received either 10 µl of sterile saline as a placebo or an equivalent volume of *K. pneumoniae* suspension. The animals were killed in different times at 6, 12, 24 and 48 h after meningitis induction. Another group was treated with antibiotic, starting at 16 h and continuing daily until their decapitation at 24 and 48 h after induction.

Results: In the hippocampus, the meningitis group without antibiotic treatment, the complex I was increased at 24 and 48 h, complex II was increased at 48 h, complex III was inhibited at 6, 12, 24 and 48 h and in complex IV all groups with or without antibiotic treatment were inhibited after meningitis induction, in the cortex there was no alteration.

Discussion: Although descriptive, our results show that antibiotic prevented in part the changes of the mitochondrial respiratory chain. The meningitis model could be a good research tool to study the biological mechanisms involved in the pathophysiology of the *K. pneumoniae* meningitis.

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Keywords: creatine kinase; *Klebsiella pneumoniae*; meningitis; mitochondrial respiratory chain

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Accepted for publication April 26, 2012
First published online 21 February, 2013

Significant outcomes

- The influence of meningitis by *Klebsiella pneumoniae* in the creatine kinase (CK) and mitochondrial respiratory chain activity.
- The meningitis by *K. pneumoniae* is associated with permanent neurological damage; understand the illness pathophysiology is the main factor that contributes to the therapeutic success.
- Our study helps to better understand the pathophysiology of meningitis by *K. pneumoniae*.

Limitations

- Animal model.
- The results in animal models should be interpreted with caution before correlate with the clinic.
- This study evaluates the animal model of adult rats.

Introduction

Bacterial meningitis remains a major cause of death and long-term neurologic sequelae worldwide. However, mortality and morbidity vary by causative organism, age and geographical location of the patient (1,2). *Klebsiella pneumoniae* is a capsulated gram-negative pathogen that is known to cause infection both in community and mainly as a hospital-acquired infection presenting as pneumonia, septicemia and meningitis in patients with some predisposing factors (3,4). *Klebsiella pneumoniae* has recently become an increasingly common cause of the meningitis acute (5) affecting 14–61% of nosocomial meningitis (6), being particularly devastating among immunocompromised patients (7) with mortality rates ranging between 30 and 40% (8), however in Singapore and parts of northern Taiwan the mortality rate is higher. Furthermore, this pathogen was the most frequent causative of the meningitis, bacteremia and septic shock in patients with liver cirrhosis (9). Furthermore, bacterial meningitis in young adults in south Taiwan, and (10) in some Asian areas, there has been an increased incidence in adults (11). Bacterial invasion in the cerebral spinal fluid (CSF) promotes the release of bacterial components like polysaccharide capsule, peptidoglycan, bacterial DNA and lipopolysaccharide (12,13), leading to the activation of the brain innate immune defence, releasing a cascade of inflammatory mediators and leukocytes recruitment (14,15). An excessive release of pro-inflammatory mediators and reactive oxygen species could contribute to interrupt the bioenergetic activity or the metabolic activity in injured neurons (16,17). The CK is vital for normal energy homeostasis by exerting some integrated functions, such as temporary energy buffering, metabolic capacity, energy transfer and metabolic control. The brain, like other tissues with high and variable rates of ATP metabolism, presents high phosphocreatine concentration and CK activity (18–20). Furthermore, another generating source of ATP is the oxidative phosphorylation, that is the predominant mitochondria physiological function but additional functions include the production and detoxification of reactive oxygen species, which is involved in various forms of apoptosis, cytoplasmic regulation and mitochondrial matrix calcium, synthesis and metabolites catabolism, so, abnormality any of these processes

can lead to mitochondrial dysfunction (21). Therefore, the main factor that contributes to the therapy success is to understand the pathogenesis and pathophysiology of the bacteria in the central nervous system (CNS) (16). Thus, to clarify a little more the pathophysiology of this illness, the aim of our study was to investigate the energetic metabolism in the rat brain after meningitis induced by *K. pneumoniae*.

Materials and methods

Infecting organism

Klebsiella pneumoniae was cultured overnight in 10 ml of Todd Hewitt broth, diluted in fresh medium and grown to logarithmic phase. The culture was centrifuged for 10 min at (5000 × g) and resuspended in sterile saline to the concentration of 1×10^6 cfu/ml. The size of the inoculum was confirmed by quantitative cultures (22,23).

Animal model of meningitis

Adult male Wistar rats (250–300 g body weight), from our breeding colony were used for the experiments. All procedures were approved by the Animal Care and Experimentation Committee of UNESC, Brazil, and followed in accordance with the National Institute of Health Guide for the Care and Use of Laboratory Animals (NIH Publications No. 80–23) revised in 1996. All surgical procedures and bacterial inoculations were performed under anaesthesia, consisting of an intraperitoneal administration of ketamine (6.6 mg/kg), xylazine (0.3 mg/kg) and acepromazine (0.16 mg/kg) (24,25). Rats underwent a cisterna magna tap with a 23-gauge needle. The animals received either 10 µl of sterile saline as a placebo or an equivalent volume of *K. pneumoniae* suspension. At the time of inoculation, animals received fluid replacement (2 ml of saline subcutaneously) and were subsequently returned to their cages. The animals were killed in different times at 6, 12, 24 and 48 h after meningitis induction by *K. pneumoniae* ($n = 6$ in each group). Another group was treated with antibiotic (ceftriaxone 100 mg/Kg twice a day, i.p.) starting at 16 h and continuing daily until their decapitation at 24 and 48 h after meningitis induction, $n = 6$ in each group (22,23). The control group did not receive antibiotic treatment (26).

Tissue assessment and homogenate preparation

Hippocampus and cortex were homogenised (1:10, w/v) in SETH buffer, pH 7.4 (250 mM sucrose, 2 mM EDTA, 10 mM Trizma base, 50 IU/ml heparin). The homogenates were centrifuged at $800 \times g$ for 10 min and the supernatants kept at -70°C until used for enzymes activity determination. The maximal period between homogenate preparation and enzyme analysis was always less than 5 days. Protein content was determined by the method described by Lowry et al. (27) using bovine serum albumin as standard.

Activities of mitochondrial respiratory chain enzymes

NADH dehydrogenase (complex I) was evaluated according to the method described by Cassina and Radi (28) by the rate of NADH-dependent ferricyanide reduction at 420 nm. The activities of succinate: DCIP oxidoreductase (complex II) and succinate: cytochrome *c* oxidoreductase (complex III) was determined according to the method of Fischer et al. (29). Complex II activity was measured by following the decrease in absorbance due to the reduction of 2,6-DCIP at 600 nm. Complex III activity was measured by cytochrome *c* reduction from succinate. The activity of cytochrome *c* oxidase (complex IV) was assayed according to the method described by Rustin et al. (30), measured by following the decrease in absorbance due to the oxidation of previously reduced cytochrome *c* at 550 nm. The activities of the mitochondrial respiratory chain complexes were expressed as nmol/min \times mg protein.

CK activity assay

CK was measured in brain homogenates pre-treated with 0.625 mM lauryl maltoside. The reaction mixture consisted of 60 mM Tris-HCl, pH 7.5, containing 7 mM phosphocreatine, 9 mM MgSO₄ and approximately 0.4–1.2 μg protein in a final volume of 100 μl . After 15 min of pre-incubation at 37°C , the reaction was started by the addition of 0.3 μmol of ADP plus 0.08 μmol of reduced glutathione. The reaction was stopped after 10 min by the addition of 1 μmol of *p*-hydroxymercuribenzoic acid. The creatine formed was estimated according to the colorimetric method of Hughes (31). The colour was developed by the addition of 100 μl 2% μ -naphthol and 100 μl 0.05% diacetyl in a final volume of 1 ml and read spectrophotometrically after 20 min at 540 nm. Results were expressed as units/min \times mg protein.

Statistics. Data about CK and mitochondrial respiratory chain complexes were analysed by Student's

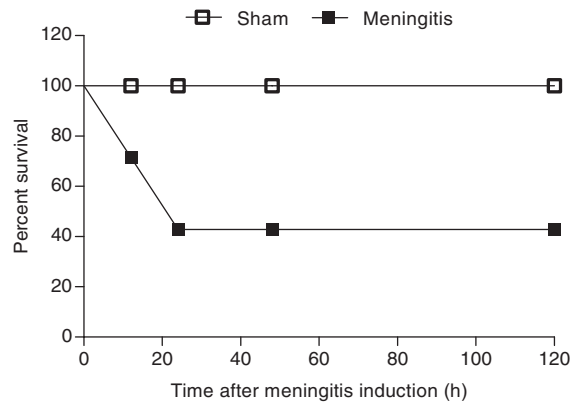


Fig. 1. Kaplan–Meier survival curves of adults Wistar rats infected by intracisternal inoculation with *Klebsiella pneumoniae*. Results are expressed as percentage of survival over time with spontaneous death. Rats had a mortality rate of 57.143%.

t-test and are expressed as mean \pm SME of five to six animals in each group. All analyses were performed using the Statistical Package for the Social Science (SPSS) software version 16.0.

Results

In this study, we evaluated the CK activities and mitochondrial respiratory chain complexes I, II, III and IV in hippocampus and cortex of rats submitted to meningitis by *K. pneumoniae*.

Survival was analysed by Kaplan–Meier curves (Fig. 1), including all infected animals from the time of infection up to 120 h ($n = 38$). The animals started dying at 12 h (28.571%) and 24 h (57.143%), after that time surviving animals were killed until 120 h.

We verified that complex I was increased at 24 and 48 h in hippocampus in meningitis group without antibiotic treatment (Fig. 2a; $p < 0.05$); complex II was increased in hippocampus at 48 h in meningitis group without antibiotic treatment (Fig. 2b; $p < 0.05$); complex III was inhibited at 6, 12, 24 and 48 h in hippocampus in meningitis group without antibiotic treatment (Fig. 2c; $p < 0.05$) and in complex IV all groups were inhibited in hippocampus after meningitis induced by *K. pneumoniae* (Fig. 2d; $p < 0.05$). We also verified that there was no change in CK activity in both structures (Fig. 3).

Discussion

There has been an increased incidence of meningitis by *Klebsiella* sp in adults (11), especially in Asian countries. Among the gram-negative pathogens implicated in bacterial meningitis, in Taiwan, *K. pneumoniae* most common in adults (32), in great part this increase in the number of cases is related to the frequency of neurosurgical procedures,

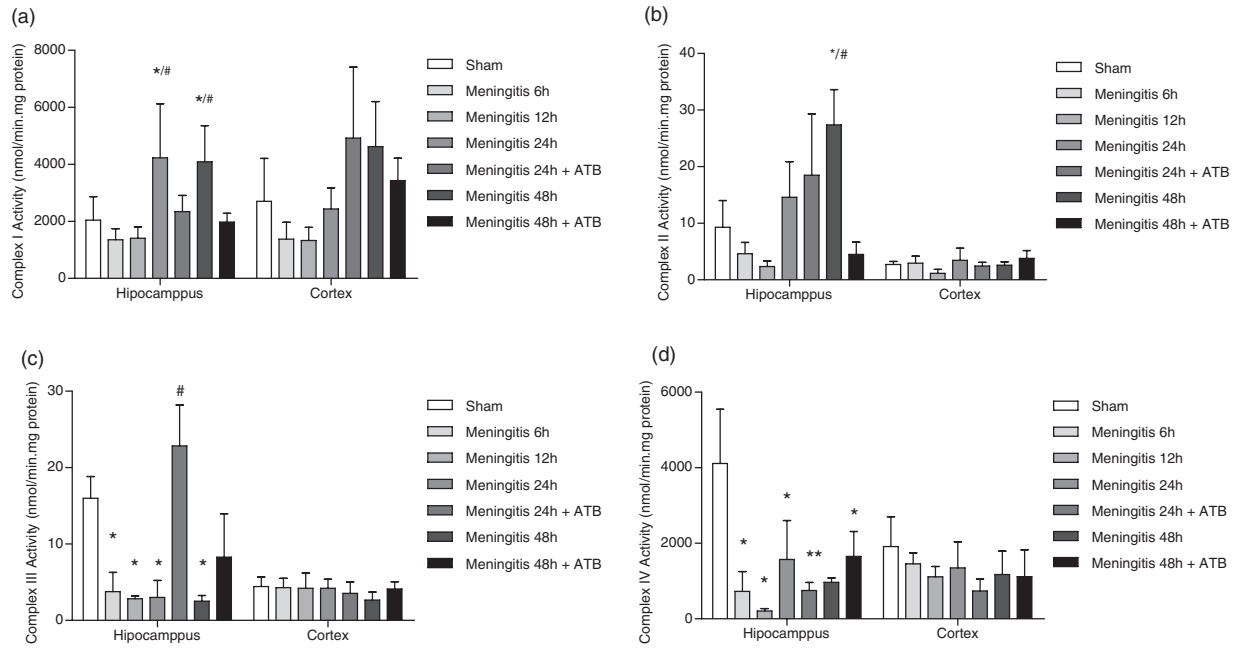


Fig. 2. Activity of the mitochondrial respiratory chain complexes I (a), II (b), III (c) and IV (d) in hippocampus and cortex of rats after meningitis by *Klebsiella pneumoniae*. Results are expressed as mean \pm SD ($n = 6$) (nmol/min \times mg protein). *Statistically significant when compared with sham group, $p < 0.05$. #Statistically significant when compared between meningitis groups with and without antibiotic in the same period.

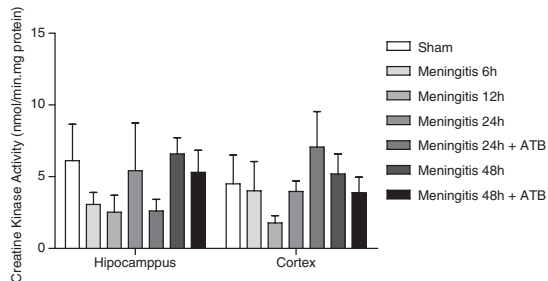


Fig. 3. CK activity in hippocampus and cortex of rats after meningitis by *Klebsiella pneumoniae*. Results are expressed as mean \pm SD ($n = 6$) (nmol/min \times mg protein). *Statistically significant when compared with sham group, $p < 0.05$. #Statistically significant when compared between meningitis groups with and without antibiotic in the same period, $p < 0.05$.

and the large number of patients with head injury from motorcycle accidents (33); however, chronic diseases like diabetes and liver cirrhosis predispose to meningitis (5). Bacterial invasion of the meninges induces a complex immune response (34), being that glial cells are an important early source of pro-inflammatory cytokines during the CNS infection by *K. pneumoniae* (35). The complex host inflammatory response from the white blood cells leads to mitochondrial damage initiating the release of the cytochrome *c* into the cytosol. There are many evidences that mitochondria participates in the caspase-dependent pathway resulting caspase

activation and neuronal damage development in the bacterial meningitis (16,36). Furthermore, the brain is highly dependent on ATP; most cell energy is obtained through oxidative phosphorylation, a process that requires the action of various respiratory enzyme complexes situated in a special structure of the inner mitochondrial membrane, the mitochondrial respiratory chain (37). Another way to get energy is through the creatine/phosphocreatine/CK system that is essential for normal energy homeostasis by exerting some integrated functions, such as, temporary energy buffering, metabolic capacity, energy transfer and metabolic control (18,38). Meningitis caused by *K. pneumoniae* increased complex I at 24 and 48 h and complex II at 48 h in hippocampus. In a previous study, we also showed increased complex II at 24 and 48 h in hippocampus among surviving rats by pneumococcal meningitis (6). The increase of the complexes I and II could be compensation mechanisms, because of the decreased activity of complex III and IV. There was an activity decrease in complex III at 6, 12, 24 and 48 h; however at 24 and 48 h with antibiotic treatment the levels did not change in the hippocampus. In complex IV, there was also an activity decrease in all the times in the hippocampus. Furthermore, complex III deficiencies are among the least common respiratory chain abnormalities identified to date in humans (39), mutations in the cytochrome *b* gene constitute a major cause of complex III deficiency, and underlie a variety of

disorders, such as encephalopathy, optic neuropathy (39), encephalomyopathy (40) although, there are not clinical findings which are specific for complex III deficiency (41), likewise, meningitis caused by *K. pneumoniae* inhibited the activity of complex III and IV in the hippocampus. The mitochondrial dysfunction can be responsible of oxidative stress due to the lack of reactive oxygen species detoxification and neurological clinical symptoms (21). In autopsy studies on patients who died from bacterial meningitis, injury in the CNS was characterised by tissue necrosis in the cortical hemispheres and by apoptotic cell death in the dentate gyrus (42). Hippocampal apoptosis is associated with learning and memory deficits observed in survivors of bacterial meningitis (43).

The meningitis by *K. pneumoniae* also is associated with permanent neurological damage (44), moreover, understand the illness pathophysiology is the main factor that contributes to the therapeutic success (16). Although descriptive and with high rate mortality our results show that antibiotic prevented in part the changes of the mitochondrial respiratory chain. White blood cells and oxidative stress are responsible to apoptosis activation; furthermore, treatment with antibiotics decreases immunogenic components in the CSF. The complete sterilisation of *Neisseria meningitidis* from CSF occurs within 2 h of given a parenteral third-generation cephalosporin and the beginning of sterilisation of *Streptococcus pneumoniae* from CSF by 4 h into treatment (1). In previous studies, we verified that early antibiotic administration prevented cognitive impairment induced by meningitis in rats (45) and prevented in part oxidative stress (22).

We believe that the damage by *K. pneumoniae* meningitis is related to mitochondrial respiratory chain dysfunction. The statistic shows differences between groups; however, the work do not have statistical power to generalise the findings. Although descriptive, our findings suggest that the meningitis model could be a good research tool to study the biological mechanisms involved in the pathophysiology of this illness and the secondary alterations of the *K. pneumoniae* meningitis.

Acknowledgements

This research was supported by grants from work supported by NENASC project (PRONEX program CNPq/FAPESC), CNPq, FAPESC, UNESC, INCT-TM and L'Oréal-UNESCO Brazil Fellowship for Women in Science 2011. The authors declare that they have no conflict of interest.

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