

DNA Damage in Perinatal Asphyxia using Micronucleus Assay

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Abstract

This study was conducted to determine the level of DNA damage in perinatal asphyxia using micronucleus (MN) assay and correlate with severity of asphyxia. Eighty term asphyxiated and sixty term non-asphyxiated neonates were enrolled in this case/control study. Blood samples were collected within 24 hours of birth for micronucleus assay and estimation of serum malondialdehyde (MDA). Micronucleus score was correlated with severity of asphyxia and serum MDA level. There was a significant difference of micronucleus (MN) score and serum MDA level in cases and controls (p value <0.0001). Among babies in various hypoxic ischemic encephalopathy (HIE) stages the micronucleus score significantly increased with severity of asphyxia (p value <0.0001). There was a significant correlation between Apgar score, HIE stages, seizures, micronucleus score and serum MDA level (p value <0.0001). Micronucleus score and serum MDA level correlated well with severity of asphyxia

Key words: DNA damage, Micronucleus, MDA, Perinatal asphyxia

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Introduction

Perinatal asphyxia is a global problem resulting in neonatal morbidity and mortality [1]. In developing countries the incidence is much higher. Hypoxia leads to specific cellular changes affecting enzymatic activities, mitochondrial function, cytoskeletal structures, membrane transport and antioxidant defences. The highly reactive oxygen species (ROS) and reactive nitrogen species (RNS) react with several living cell contents eg phospholipids, sugars, amino acids and nucleic acids leading to lipid peroxidation DNA strand breaks, base methylation etc. The consequence could be immediate and long term complications [2]. The DNA damage can be quantified by cytokinesis block micronucleus assay (CBMN). Micronucleus can be originated by fragments or lagging of chromosome during cell division. It is morphologically similar to main nuclei but smaller in size hence the name. It can be made visible by freezing the cytoplasmic division of dividing cell by addition of Cytochalasin (Cyto-B) [3]. Oxidative stress can be assessed by serum malondialdehyde (MDA) level [4]. We tried to correlate the severity of asphyxia to oxidative stress and DNA damage.

Patients and Methods

The study was conducted in the Cytogenetic unit of Department of Anatomy in collaboration with Departments of Pediatrics and Biochemistry from February 2008 to July 2010. The study was approved by the institute research council and human ethical committee. Term asphyxiated appropriate for gestational age babies were taken as cases. Gestational age and weight matched babies without asphyxia were taken as controls. Perinatal asphyxia was diagnosed when more than three of the following criteria were present viz: (1) Apgar score less than 6 at 5 mts. (2) meconium stained liquor (3) changes in the fetal heart rate (4) clinical evidence of HIE (5) evidence of multiorgan dysfunction. Preterm or post term babies, large (LGA) or small (SGA) for gestational age babies, those with congenital malformations and delivered of mothers with significant illness were excluded. The CBMN assay was

carried out in conventional RPMI supplemented with phytohaemagglutinin (Sigma), on blood lymphocytes using cytochalasin-B (Sigma) [5]. Heparinized whole blood 0.2 ml of was inoculated in the culture media and incubated at 37°C for 44 hours. The Cytochalasin-B (Sigma) was prepared at a final concentration of 3µl/ml. Binucleated lymphocytes were harvested for 72 hours. The buffy coat was dispersed with hypotonic solution (0.075 M KCl). The cell pellets were fixed with 3:1 methanol acetic acid. The slides were stained with Giemsa stain for three minutes. The micronucleus index for each sample was analysed by counting 1000 binucleated cells based on the scoring criteria outlined by HUMN project [6]. The binucleated cells were scored blindly using 400X magnification of the Epifluorescent microscope BX 51 (Olympus). Thiobarbituric acid reactive substances (TBRAS) which measure MDA present in the serum was estimated for assessment of oxidative stress.

Statistical Analysis

The statistical analysis has been performed using unpaired student t-test for comparison of parametric test, Mann-Whitney for nonparametric test and One way ANOVA (Kruskal Wallis test) for multiple comparison. Pearson correlation coefficient was used for association between the groups. All the data was analysed by Graph Pad (In-Stat, San Diego, USA)

Results

There were 80 cases and 60 control babies. Among the cases there were 50 male babies 30 females while there were 32 male and 28 female babies among controls. Birth weight and gestational age of the asphyxiated babies were not significantly different from the controls. The age, parity and mode of delivery were comparable among the groups. The mean and SD of Apgar score [7] in cases were significantly lower than the controls (4.9±1.624 vs 8.633±0.604). Based on Sarnat and Sarnat score [8] 21 babies were in HIE stage -1, 40 in HIE stage-2, 19 in HIE stage-3. Among the cases 59 (73.75%) developed seizure, 50 were discharged, 2 left against medical advice (AMA) and 28 expired. All the control babies were discharged.

The serum MDA level in asphyxiated babies was found to be significantly increased (6.709±1.695 vs 3.683 ± 0.536). The serum MDA level (P value <0.0001) positively correlated with severity of asphyxia.

Table 1. Micronucleus score and serum MDA in Asphyxiated and control babies.

Groups	Micronucleus Score	Serum MDA
Control (n=60)	1.212±0.775	3.683±0.536
Hypoxia (n=80)	3.959±2.816	6.709±1.69
HIE-1 (n=21)	1.579±0.901	4.667±0.950
HIE-2 (n=40)	2.944±1.013	7.117±0.529
HIE-3 (n=19)	8.263±1.327	8.99±0.5395
Seizure present(n=59)	4.782±2.787	1.780±0.86
Seizure absent(n=21)	1.579±0.901	4.667±0.950

*P value <0.0001 **P value <0.0001

It has been observed that there was a significant difference between micronucleus score in cases and controls (3.959±2.816 vs 1.212±0.775). MN score was significantly increased with severity of asphyxia (P value <0.0001). In babies with seizure there was a significant increase of MN score compared to non-seizure babies

Table 2. Correlation coefficient[®] among various parameters

Parameter	Correlated Variables			
	HIE	Seizure	MNi	Serum MDA
Apgar Score (n=80)	-0.6314**	-0.7580**	-0.7570**	-0.8325**
HIE Staging (n=80)		0.9599*	0.8123*	0.9004*
Seizure (n=59)			0.8232*	0.7702*
MNi (n=80)				0.9201*

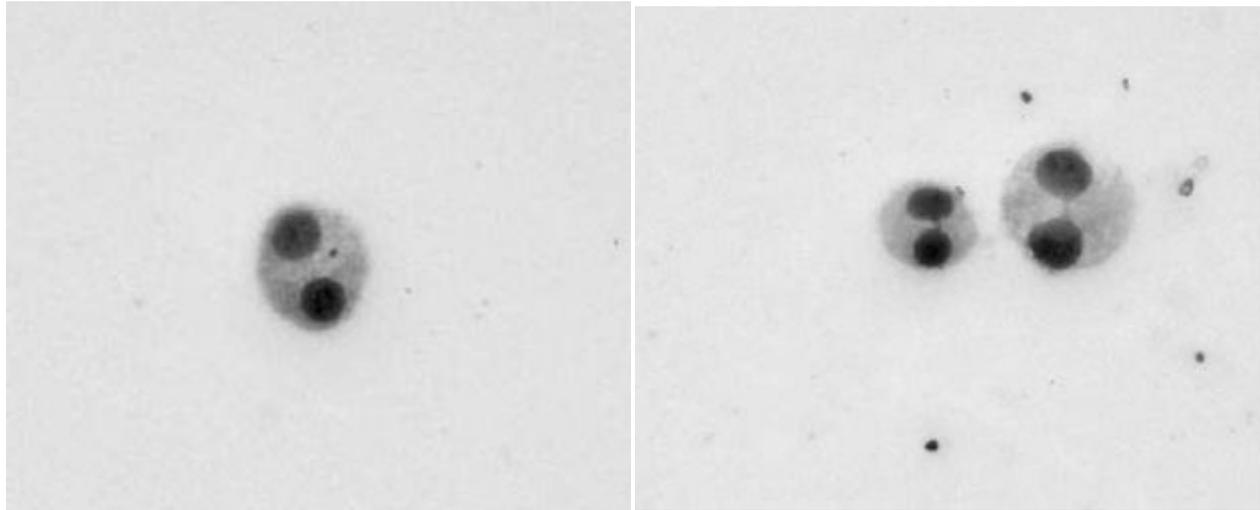


Figure 1. Images of 40 X Geimsa stained A – Binucleated lymphocytes with micronucleus (←) and B- Nucleoplasmic bridge (→) in Perinatal asphyxia. $y = 0.3818x + 5.0162R^2 = 0.52030123456789100246810$ Micronucleus Score Serum MDA level in micromol/lit.

Figure 2. Correlation Coefficient between micronucleus and serum MDA level r value 0.9201 and p value <0.0001

Discussion

Perinatal asphyxia is one of the major causes of neonatal mortality in developing countries like India. Perinatal asphyxia is characterised by impaired gas exchange which leads to hypoxemia, hypercarbia and metabolic acidosis.

Impaired gas exchange and insufficient perfusion to vital organs leads to severe neurological insult resulting in death and long term disability. The molecular mechanism behind the neurological deficit is not fully evaluated.

Two mechanisms of the DNA damage have been suggested. It is envisaged that H₂O₂ which crosses the biological membrane easily can penetrate the nucleus and react with ions of iron or copper to form hydroxyl radical (OH \cdot) [9]. Other explanation is the ability of oxidative stress to cause DNA damage by triggering a series of metabolic events within the cell that lead to the elevation of nuclease enzymes which cleave the DNA backbone. The intracellular free Ca²⁺ interact with Ca²⁺ dependent endonuclease leading to fragmentation of DNA, resembling the mechanism that of apoptosis (programmed cell death) [10].

The micronucleus score of cases and controls were significantly different. Normal and HIE-1 babies have only single strand breaks which could be repaired and the chance of formation of micronuclei is less. There was significant increase in MN score in higher HIE stages. The DNA damage in severe hypoxic ischemic encephalopathy is unable to repair. There were deletions, interchromatid and intrachromatid exchanges. These drastic instability in the genome was visible as micronucleus, nucleoplasmic bridges and nuclear buds in the next generations of cells. This will arrest further growth of cell and the normal functioning leading to apoptosis and necrosis of the cell. This also leads to alteration in the normal synthesis of DNA and gene encoding reflected in the neurological behaviour of the child [11].

Increased micronucleus score has been observed in women having ovarian cyst and after chemical exposure [12,13]. There was a significant increase in the micronuclei in cerebral palsy children compared with normal children and in double hemiplegia compared to hemiparesis patients. The endometagen cause cerebral palsy in birth asphyxia with excess generation of glutamate. These excess glutamate production leads to the formation of clastogenesis and aneuploidogenesis [14].

Peroxidation of lipids have been reported to damage DNA. It produces wide range of reactive oxygen species including OH \cdot , H₂O₂, singlet oxygen, peroxy radicals and alkoxy radicals. Lipid peroxidation also decomposes to give a huge range of products including carbonyl compounds such as malondialdehyde and the unsaturated aldehyde 4-hydroxy-2 transnonenal. If the aldehydes are generated in the vicinity of DNA, they may be able to combine with it to form distinctive products. The association between oxidative stress and DNA damage have been observed by many authors. MDA react with adenine, cytosine and guanine, the guanine MDA adduct has been identified in the urine. The above findings is a strong evidence of peroxidation of lipids causing DNA damage [15]. In Perinatal asphyxia we found that the serum MDA level was significantly increased compared to controls. The oxidants and antioxidant level in hypoxic ischemic encephalopathy were found to be significantly higher [16,17]. The MDA level in cord blood of babies with meconium stained liquor with or without birth asphyxia was significantly higher than the controls [18]. The urinary creatinine and MDA values were significantly high in asphyxiated babies and their levels correlated with outcome [19]. Markers like serum MDA and protein carbonyl will be beneficial in predicting the outcome of perinatal asphyxia [20].

There was a negative correlation between micronucleus score and Apgar score and also MDA level and Apgar score. It has been observed that Apgar score inversely related to DNA damage and oxidative stress. The correlation between micronucleus index and HIE stages was positive and also there was a positive correlation between HIE stages and serum MDA values signifying that oxidative stress induces DNA damage in perinatal asphyxia. This results in conditions like epilepsy, motor dysfunction, loss of memory, abnormalities in sympathetic and parasympathetic system in later life. Immediate newer therapies like cerebroprotective agents being tried following asphyxial insult to prevent childhood sequelae.

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