Original article

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Micronuclei and other nuclear anomalies in buccal epithelial cells of children with chronic kidney disease

Banu Aykanat^{1*}, Gonca Cakmak Demircigil^{1*}, Necla Buyan², Esra Baskin⁴, Kaan Gulleroglu⁴, Kibriya Fidan², Umut Selda Bayrakci⁴, Aydin Dalgic³, Hamdi Karakayali⁵, Mehmet Haberal⁵, and Sema Burgaz¹

Department of Toxicology, Faculty of Pharmacy¹, Department of Pediatric Nephrology², Department of General Surgery³, Faculty of Medicine, Gazi University, Department of Pediatric Nephrology⁴, Department of General Surgery⁵, Faculty of Medicine, Baskent University, Ankara, Turkey

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The objective of this study was to reveal the likely genomic instability in children with chronic kidney disease (CKD) using micronucleus (MN) assay on buccal epithelial cells (BEC). We investigated the frequencies of micronuclei and other nuclear anomalies, such as nuclear buds, binucleated cells, condensed chromatin, and karyorrhectic and pyknotic cells in BEC. Children with CKD were grouped as follows: children in the pre-dialysis (PreD) stage (N=17), children on regular haemodialysis (HD) (N=14), and children who have undergone transplantation (Tx) (N=17). As a control group, twenty age- and gender-matched healthy children were selected. The MN frequency in BEC of all groups of children with CKD was significantly elevated (5- to 7-fold) as compared to the control group (p<0.001). In contrast, the frequencies of nuclear buds were not significantly higher in the study groups compared to the control group. The frequencies of binucleated cells and condensed chromatin cells were significantly higher in all subgroups of children with CKD relative to the control group (p<0.001). Our results show that the BEC of pediatric PreD, HD, and Tx patients with CKD display increased cytogenetic, cytokinetic, and cytotoxic effects. They also point to the sensitivity and usefulness of the BEC MN assay in the assessment of genetic susceptibility of patients with CKD.

KEY WORDS: buccal micronucleus assay; genotoxicity; dialysis; renal transplantation; nuclear anomalies

Chronic kidney disease (CKD) is a major public health concern (1). Recent reviews indicate that the adverse health-related outcomes of CKD and/or end-stage renal disease (ESRD) are characterised by high incidences of cardiovascular morbidity and mortality (2), liver damage due to viral infections (3), chronic inflammation (4), oxidative stress (5), and an increased incidence of cancer (6, 7).

Extensive research is still required to clarify the mechanisms underlying these outcomes of CKD and treatments like haemodialysis and transplantation. The causes, development, and treatment of CKD in paediatric patients differ from those in adult patients (8). Long periods of haemodialysis treatment have been linked to DNA damage, mainly due to oxidative stress (9). In adult patients with CKD, increased cancer incidence and cytogenetic effects in peripheral lymphocytes have been observed (7). In addition, the patient's susceptibility to neoplastic disorders increased as a consequence of prolonged immunosuppression after transplantation (10).

Correspondence to: Sema Burgaz, Department of Toxicology, Faculty of Pharmacy, Gazi University, Ankara, Turkey, Tel: +90 312 2023087, Fax: +90 312 222 2326, Email: burgaz@gazi.edu.tr

Latency periods in most types of cancer are usually 15-20 years or longer, and conducting prospective epidemiological studies over long time periods is not always practical. Therefore, it is necessary to use the known biomarkers of DNA damage resulting from genetic instability to predict cancer risks and to identify high-risk individuals (11). Cytogenetic biomarkers are the most frequently used end points in human studies. In addition to peripheral blood lymphocytes, measurements of MN frequencies as well as other nuclear anomalies in buccal epithelial cells are increasingly used in molecular epidemiologic studies to investigate the impact of nutrition, lifestyle factors, genotoxic exposures, and genotype on DNA damage and cell death (12-14).

Studies from our laboratory (15, 16) have recently demonstrated significant increases in MN frequencies and basal DNA damage (determined by comet assay) in peripheral blood lymphocytes of children with CKD.

To the best of our knowledge, no genotoxicity studies of buccal cells of children with CKD have been reported so far. Therefore, we decided to investigate the frequencies of MN and other nuclear anomalies, such as nuclear buds (NBs), binucleated cells (BN), condensed chromatin (CC), karyorrhectic (KR), and pyknotic (PK) cells, in buccal epithelial cells (BEC) in children with CKD. There were

^{*}These authors contributed equally to this work.

three groups: children in the pre-dialysis (PreD) stage, children on regular hemodialysis (HD), and children who have had transplantation (Tx).

This study aims to reveal the possible genomic instability of CKD alone and treatment-related genotoxic and other effects (i.e. cytokinetic defects and cell death) in children with CKD using a non-invasive method: micronucleus assay on buccal tissue.

SUBJECTS AND METHODS

Study population

The study group consisted of 48 children (age range: 2-19 years) diagnosed with CKD. All of them were also involved in our previous studies (15, 16), where the causes of primary kidney disease and medication(s) received were described in more detail. Of these children, 17 (9 males and 8 females) were in the pre-dialysis stage, 14 (6 males and 8 females) were on regular haemodialysis, and 17 (10 males and 7 females) had kidney transplantation.

The study groups were: the children with blood creatinine levels higher than 1 mg dL⁻¹ and glomerular filtration rates (GFR) in the range of 15-90 mL min⁻¹ were selected for PD group. HD group consisted of children who underwent regular haemodialysis treatment for four hours three times per week. Tx group consisted of children who underwent kidney transplantation and were under immunosuppressant treatment, as described in our previous study (16). The duration of being diagnosed with PreD, HD, and Tx, expressed in months (\pm SD) was 41.35 \pm 38.83, 29.75 \pm 27.68, and 22.35 \pm 19.43, respectively.

For the purpose of the study, we selected non-smokers who had never consumed alcohol and who were without any malignancies. The matched control group consisted of 20 healthy children (11 males and 9 females; age range 4-18 years) who were free from medication and infection, as confirmed by face-to-face interviews with their parents.

This study was approved by the local ethical committee of the Baskent University, Faculty of Medicine (03/07/2007-07/39). Furthermore, the parents of all children gave their informed consent prior to involving their children in the study. Detailed questionnaire forms completed by the parents included age, height, weight, and education, cigarette smoking and environmental tobacco smoke (ETS) exposure, recent diagnostic X-ray examination (three months prior to the sampling), recent vaccination, sports activities of the children, and the use of vitamin supplementations. Moreover, for those children who were diagnosed with kidney disease, information regarding the kidney disease type, parameters, grade, treatment type, medication, and viral infection were completed by their physicians. The causes of primary kidney disease and medication received among children have also been described in detail elsewhere (15, 16).

Biological sampling

The sampling of buccal epithelial cells from both the CKD and control groups was done between 16 Jul 2008 and 29 Jul 2009. For each subject, we simultaneously collected BEC for the MN-assay and blood samples which had been used for the purposes of other genotoxicity studies. Thus, data on genotoxicity endpoints (e.g. micronucleus and comet) in peripheral lymphocytes and routine biochemical parameters of the same populations have been published elsewhere (15, 16).

Buccal epithelial cells were obtained by scraping both sides of the inside of the cheeks using pre-moistened wooden tongue depressors after the subjects had rinsed their mouths with water.

The collected cells were first smeared directly onto pre-cleaned and pre-moistened microscope slides and were then left to dry. Within the same day, the slides were transferred to the laboratory where they were further processed.

MN assay in buccal epithelial cells

Preparations of buccal epithelial cells were fixed in 80 % methanol (Merck, Germany) and air-dried. Slides were stained by the Feulgen reagent (pararosaniline, Merck, Germany) and then counterstained with Fast Green (Merck, Germany) (17, 18).

Microscopy analyses were performed using a light microscope (Zeiss Axioscope 2, Goettingen, Germany) at 400x magnification and confirmed at 1000x magnification. The coded slides were evaluated blindly by a single scorer.

The identification criteria of MN and other nuclear anomalies, previously described by Thomas et al. (19), were followed. For each subject, a total of 2000 differentiated and basal cells were analysed for the presence of the MN frequency and micronucleated cell and other nuclear anomaly (nuclear buds, condensed chromatin, binucleated, karyorrhectic and pyknotic cells) frequencies. Karyolytic cells were not scored. Basal and differentiated cell frequencies were not evaluated. The frequency of each parameter has been expressed as per thousand (‰).

Statistical analysis

Data were analysed using SPSS for Windows, version 11.5 (SPSS Inc., Chicago, IL, United States). The Shapiro-Wilk test was used to determine whether the distributions of continuous variables were normal. The Levene test was used to evaluate the homogeneity of variances. The data for continuous variables were expressed as the means ± standard deviations (SD) or medians (min.-max.), and for nominal variables as the number of cases and/or percentage (%). Regarding the number of independent groups, the mean differences were compared by Student's t test or One-Way ANOVA. Where applicable, the Mann-Whitney U or Kruskal-Wallis tests were applied to compare the median values. When the *p*-values (from One-Way ANOVA or

Kruskal-Wallis test statistics) were statistically significant, a *post-hoc* Tukey or Conover's multiple comparison test was applied to determine which groups differed. Where applicable, nominal data were evaluated by the Pearson Chi-square or Fisher's exact test.

Degrees of association between continuous variables were evaluated using the Spearman's rank correlation test. Correlation analyses were performed to explore the association (if any) between buccal MN frequencies in the study subjects and corresponding lymphocyte MN frequencies and routine biochemical parameters data from recent publications (15, 16). Stepwise linear regression analyses were used to define predictive factors and dependent variables (e.g. MN frequency in BEC, micronucleated cell frequency in BEC), nuclear buds, binucleated cells, condensed chromatin cells, karyorrhectic and pyknotic cells). For those dependent variables that were not normally distributed, log-transformation was applied via multiple linear regression analyses. Any variable determined by univariate tests to be statistically significant and clinically important was accepted as a candidate for the multivariable model. After adjusting for factors, such as age, gender, body mass index, smoking, X-ray exposure, drug usage, and sports activities, multiple linear regression analyses were performed to determine whether there were any statistically significant differences among the study groups. Coefficients of regression and 95 % confidence intervals were calculated for all independent variables. A p-value of less than 0.05 was considered statistically significant.

RESULTS

The demographic characteristics of the children with CKD and control group have already been reported in our previously published studies (15, 16). Considering that the present study and both of the previously mentioned ones were conducted simultaneously, here we briefly report the most important results which could help the reader to better understand the results of the MN assay on BEC.

As reported previously (15, 16), age, gender, and duration of sports activities among the study groups did not significantly vary. X-ray exposure three months prior to the sampling was significantly higher in CKD subgroups. None of them had a dental radiography examination. Vaccinations within a year were significantly more frequent in CKD subgroups, except HD vs. control group. Exposure to environmental tobacco smoke was significantly lower in CKD subjects, and differences in CKD subgroups vs. control group were statistically significant (p<0.05). The Tx group exhibited the highest BMI values, and these values were significantly different from the control and HD groups (p<0.05). In each CKD subgroup, either the MN frequency or the micronucleated cell frequency in BEC was significantly (5- to 7- fold) elevated in comparison with the control group (p < 0.001; Table 1).

Cell anomalies also revealed differences between the groups of children studied. The frequency of nuclear buds in the study group was not significantly different from the control group (Table 1) even though the children in the HD subgroup demonstrated a 23 % increase in this parameter. Binucleated cell frequencies were significantly higher in all CKD subgroups relative to the control group (p<0.001; Table 1). The % increases for binucleated cells were 79 %, 56 %, and 86 % in the Pre-D, HD, and Tx subgroups, respectively.

Table 1 MN and other nuclear anomalies (%) (Mean±SD) in buccal epithelial cells of children with CKD and control group

		Control group			
N	TS	PreD	HD	Tx	Control group
	48	17	14	17	20
MN frequency in BEC (‰)	8.34±7.33	8.32±8.54	9.57±7.61	7.35±5.95	1.45±1.30*
Micronucleated cell frequency in BEC (%)	5.36±3.99	5.09±4.22	5.79±3.93	5.29±4.03	1.03±0.95*
Nuclear buds (‰)	0.33±0.43	0.26±0.31	0.43±0.58	0.30 ± 0.40	0.33±0.29
Binucleated cells (%)	30.66±9.20	31.32±7.29	27.43±7.99	32.65±11.39	17.55±5.25*
Condensed chromatin cells (%)	20.93±8.49	24.18±12.12	20.36±6.17	18.15±3.77	8.75±3.70*
Karyorrhectic cells (%)	8.23±5.70	9.00±5.34	11.21±6.94	5.00±2.91***	3.25±1.97**
Pyknotic cells (‰)	24.41±14.83	29.18±16.82	29.04±11.28	15.82±11.76***	14.55±6.17**

Abbreviations: TS=total study group of children with disease and under treatment; PreD=children in predialysis; HD=children in hemodialysis; Tx=children who underwent renal transplantation. *p<0.001; control group vs. TS, PreD, HD, and Tx. **p<0.05; control group vs. TS, PreD and HD. ***p<0.05; Tx vs. PreD and HD

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	PreD	HD	Tx				
MN frequency (R ² =0.617)	1.418	1.573	1.474				
p	0.000	0.002	0.001				
B (95 % CIs)	(0.683 to 2.152)	(0.612 to 2.534)	(0.665 to 2.283)				
Micronucleated cell frequency (R ² =0.634)	1.271	1.489	1.421				
p	0.000	0.001	0.000				
R (05 % CIs)	(0.625 to 1.917)	(0.644 to 2.333)	(0.710 to 2.132)				

Table 2 Results on the multiple linear regression analysis of MN frequency and micronucleated cell frequency in BEC of study subjects (patients+controls, N=68)

Independent variables: Tx group, HD group, PreD group, age, gender, BMI, environmental tobacco smoke exposure, sports activity (hours), use of vitamin (yes/no), X-ray exposure, antihypertensive drugs, erythropoietin, osteoporosis drugs, antibiotics, iron drugs, phosex. **Abbreviations:** CIs=confidence intervals, B=regression coefficient (slope)

Significantly increased frequencies of condensed chromatin were found in each subgroup of children with CKD (p<0.001; Table 1).

Frequencies of karyorrhectic and pyknotic cells were significantly higher in the PreD and HD subgroups compared to the control group (p<0.05; Table 1). However, the frequencies of karyorrhectic and pyknotic cells in the Tx group were significantly lower than those in the PreD and HD groups (p<0.05; Table 1), and exhibited no significant differences relative to the control group.

Regression analysis indicates that every patient subgroup exhibited significant increases in buccal MN frequencies as well as micronucleated cell frequencies compared to control values (p<0.01; Table 2). This increase in the MN frequency was the highest for the HD subgroup (B=1.573, p=0.002), followed by the Tx (B=1.474, p=0.001) and PreD (B=1.418, p=0.000) subgroups, respectively. The same trend was observed with regard to BEC micronucleated cell frequencies (Table 2).

A PreD or HD status significantly influenced condensed chromatin and pyknotic cell frequencies in BEC (p<0.01; Table 3), whereas a Tx status significantly influenced condensed chromatin cell frequencies and binucleated cells in BEC (p<0.01, Table 3). The effect on the frequencies of condensed chromatin cells was the highest in the PreD group (B=18.471, p<0.00001), followed by the HD (B=15.045, p<0.00001) and Tx (B=12.154, p=0.001) groups, respectively (Table 3), while the effect on the frequencies of pyknotic cells was the highest in the HD group (B=20.353, p=0.003), followed by the PreD group (B=16.993, p=0.001) (Table 3). A significant effect on the frequencies of binucleated cells was only observed for the Tx group (B=12.915, p=0.007) (Table 3).

Diagnostic X-ray examination had a significant negative effect on the frequencies of condensed chromatin (B=-7.735 p<0.00001) and a positive significant effect on the frequencies of nuclear buds (B=0.272, p<0.05). Other independent variables, such as age, gender, BMI, ETS exposure, sports activity (hours), vitamin usage (yes/no), routine biochemical parameters and medication usage, had no significant effect on MN frequencies or nuclear anomalies (Tables 2 and 3). When patients and control subjects were taken as a whole, correlation analysis showed

significant differences regarding the MN frequencies in BEC and routine biochemical parameters, such as creatinine, BUN, parathyroid hormone, ferritin, albumin, uric acid (r=0.653, r=0.616, r=0.600, r=0.534, r=-0.450, r=0.436, respectively; p<0.00001), homocysteine, triglycerides, CRP, total protein, haematocrit, calcium and total cholesterol levels (r=0.408, r=0.365, r=0.336, r=-0.306, r=-0.302, r=-0.292, r=0.254, respectively; p<0.05). Similar correlations were obtained with regard to micronucleated cell frequencies in BEC.

Correlation analyses showed significant differences between the frequencies of nuclear anomalies in BEC and routine biochemical parameters. There was a significant correlation between the nuclear bud frequencies and ALT (r=0.387, p<0.05). The frequency of binucleated cells was significantly correlated to triglycerides (r=0.300, p<0.00001), creatinine, BUN, total cholesterol, CRP, ferritin, homocysteine, parathyroid hormone, and uric acid levels (r=0.445, r=0.260, r=0.361, r=0.265, r=0.342, r=0.253, r=0.426, r=0.268, respectively; p<0.05). Significant correlations were also obtained between the condensed chromatin cell frequency and creatinine, BUN, triglycerides, and parathyroid hormone levels (r=0.523, r=0.542, r=0.595, r=0.609, respectively; p<0.00001) as well as total cholesterol, albumin, ferritin, homocysteine, alkaline phosphatase, and uric acid levels (r=0.292, r=-0.247, r=0.410, r=0.403, r=0.340, r=0.348, respectively; p<0.05). Significant correlations between karyorrhectic cell frequencies and creatinine, BUN, and triglyceride levels (r=0.555, r=0.549, r=0.595, respectively; p<0.00001), as well as albumin, ferritin, homocysteine, parathyroid hormone, and potassium levels (r=-0.313, r=0.263, r=0.361, r=0.369, r=0.299, respectively; p<0.05) were also found. Pyknotic cell frequencies also correlated with creatinine, BUN, iron, parathyroid hormone, and sodium levels (r=0.374, r=0.385, r=0.318, r=0.260, r=-0.293, respectively; p < 0.05).

There was a significant correlation between the MN frequency in BEC and the MN frequency in peripheral lymphocytes (in the whole study population; r=0.609; p=0.000). A similar significant correlation was found with regard to the micronucleated cell frequency in BEC (r=0.613; p<0.00001).

DISCUSSION

Recent data from the Turkish Society of Nephrology (20) reported 1643 haemodialyses, 2325 peritoneal dialyses, and 546 transplantations in children in our country. The increasing numbers of CKD patients and concurrently developing secondary diseases (especially cancer) indicate the importance of understanding this chronic disease, both in Turkey and all over the world. Children with CKD have an increased incidence of cancer (21), which is about ten times more frequent compared to healthy children of the same age (22). Thus, long-term survival and preventive strategies are crucial for children who have CKD and are undergoing treatment (23). Immunosuppressive drugs used to prevent graft rejection in recipients of kidney transplants and/or prolonged haemodialysis treatments have been linked to DNA damage and malignancies (8, 24).

Previous studies on chromosomal and/or DNA damage resulting from CKD and its treatment modes were based on adults. The biomarkers used in these studies were micronuclei in peripheral blood lymphocytes (PBL) (7, 25-27) and BEC (9), chromosome aberrations and sister chromatid exchanges (28, 29), and DNA strand breaks (comet assay) (26-28, 30, 31). The only reports regarding children with CKD were our recent studies using the MN and comet assays in PBL of children (15, 16).

Although PBL are generally accepted as appropriate surrogate tissue for biomarker studies, exfoliated epithelial cells, such as buccal, nasal, or urothelial cells constitute an alternative option, especially because the latter can be obtained with non-invasive methods. Over 90 % of cancers occur in epithelial tissues, and, in many cases, epithelial

tissues are the actual targets of carcinogens (32). The time required for BEC cells to migrate from the oral cavity to the basal layer is approximately 1-3 weeks (33). There are no definitive data regarding the cell turnover rates of buccal cells that influence a patient's disease status and/or the effects of several treatment chemicals. Nonetheless, we used this tissue to determine the short-term effects of exposures and/or disease-treatment status, in part because it is the most easily accessible tissue (particularly for children).

Recent meta-analysis results (34) suggest that the MN frequency in BEC can be used in the prescreening and follow-up of precancerous oral lesions and may reflect the chromosomal instability of other organs. Increased buccal MN frequencies have also been observed in small-sized studies on patients with chronic diseases, with Alzheimer's disease and Down syndrome (34). Scoring MN and other nuclear anomalies such as nuclear buds (for DNA damage), binucleated cells (for cytokinetic defects), condensed chromatin, and karyorrhectic and pyknotic (for cell death) cells in BEC, known as the cytome approach, has also been found to be associated with Down syndrome and Alzheimer's disease (19). Though there have been a limited number of studies on nuclear anomalies other than the MN frequency, it has been suggested that the cytome approach may improve the predictive value of the MN assay in BEC (34).

Therefore, using the cytome approach in BEC, the monitoring of children with CKD alone and treatment-related effects can be of particular significance.

In the present study, cytogenetic, cytokinetic, and cytotoxic (cell death) effects in the BEC of children

Table 3 Results on the multiple linear regression analysis of other nuclear anomaly frequencies in BEC of study subjects (patients+controls, N=68)

	PreD	HD	Tx	X-ray exposure
Nuclear buds (R ² =0.279)	-0.203	-0.298	-0.322	0.272
p	0.162	0.143	0.072	0.021
B (95 % CIs)	(-0.490 to 0.084)	(-0.699 to 0.104)	(-0.674 to 0.030)	(0.042 to 0.502)
Binucleated cells (R ² =0.531)	7.748	7.052	12.915	-0.513
p	0.110	0.199	0.007	0.857
B (95 % CIs)	(-1.831 to 17.327)	(-3.867 to 17.972)	(3.750 to 22.080)	(-6.216 to 5.189)
Condensed chromatin cells (R ² =0.582)	18.471	15.045	12.154	-7.735
p	0.000	0.000	0.001	0.000
B (95 % CIs)	(12.932 to 24.010)	(8.320 to 21.770)	(5.352 to 18.955)	(-11.837 to-3.634)
Karyorrhectic cells (R ² =0.488)	3.010	3.425	-1.589	-0.448
p	0.180	0.265	0.480	0.746
B (95 % CIs)	(-1.432 to 7.452)	(-2.670 to 9.520)	(-6.068 to 2.890)	(-3.208 to 2.311)
Pyknotic cells (R ² =0.384)	16.993	20.353	4.994	-6.988
p	0.001	0.003	0.373	0.073
B (95 % CIs)	(7.485 to 26.500)	(7.250 to 33.456)	(-6.167 to 15.156)	(-14.652 to 0.675)

Independent variables: Tx group, HD group, PreD group, age, gender, BMI, environmental tobacco smoke exposure, sports activity (hours), use of vitamin (yes/no), X-ray exposure, antihypertensive drugs, osteoporosis drugs, antibiotics, shohl solution, other drugs, triglycerides, parathyroid hormone. **Abbreviations:** CIs=confidence intervals, B=regression coefficient (slope)

diagnosed with CKD were clearly observed in the PreD stage and during treatment (HD and Tx), in agreement with our recent studies on peripheral lymphocytes (15, 16). In the scientific literature, we could not find any study in which the BEC MN assay has been used for children with CKD. In fact, only Roth et al. (9) have used the BEC MN assay in adults with CKD. This study reported higher MN frequencies for HD patients, which supports the findings of our study. These authors also indicated that the duration of treatment was a significant factor (9).

In our study, a cell death effect was shown in the PreD and HD groups, while in the Tx group only condensed chromatin was observed (Table 1). Karyorrhectic and pyknotic cell frequencies were not different for the Tx group compared to control group children. This might be because. compared to the PreD and HD groups, most of the CKD biochemical parameters improved in the Tx-group children (15). Uremic toxins, reactive carbonyl compounds, longterm haemodialysis treatment and the accumulation of advanced glycation end-products (AGEs) may be responsible for the effects observed in the PreD and HD groups, due to their suppression of DNA repair (35). Because the cellular anomalies reflecting the cytokinetic and cytotoxic effects in BEC were not comprehensively determined via the mechanistic approach, our interpretation of the data could be rather speculative.

Nuclear buds are considered to be a biomarker for the elimination of amplified DNA and/or DNA repair complexes and thus thought to play an important role in tumour progression (36). In our study, nuclear buds were the only parameter that showed no significant differences between patients and controls. A similar result was also found for nuclear buds in the PBL MN assay performed in the same subjects (16). Therefore, it is possible to link CKD and treatment statuses in children to genome damage. In the same subjects, the MN frequencies in lymphocytes correlated with those in BEC suggesting that the assessments of BEC and lymphocytes may complement each other. Considering the The HUman MicroNucleus project on eXfoLiated buccal cells (HUMN_{XI}) project directions (37), more studies could be useful to prove the relationship between lymphocytes and BEC and to determine whether BEC may be used to predict the risk of cancer.

Considering that involvement in the PreD, HD, or Tx subgroups was the most significant factor linked to increased MN frequencies, as well as nuclear anomalies in the BEC of CKD children, an observation that supports the link between genetic instability and the development and treatment of this chronic disease is evident. Additionally, the significant associations observed among the biochemical parameters of CKD and BEC assay parameters further support this association. Also, a higher frequency of binucleated BEC observed in the present study and lower nuclear division index (NDI) in PBL (16) in the Tx group could be a signal of genome instability and cytokinetic defects that need further clarification.

It is known that several factors, such as age, gender, smoking, alcohol intake, and dietary habits, may contribute to spontaneous MN levels in exfoliated buccal cells (14). However, due to the limitations of the associated studies, the most important confounding factors have not yet been determined. These issues have recently been addressed in two review articles (14, 38).

The consistencies in age and gender between the study and control groups and the participation of non-smokers in the present study eliminated the most crucial of these confounding factors.

From a clinical point of view, diagnostic X-ray exposure in children and adults with CKD unavoidably occurred during their medical surveillance. Of course, this issue was also acknowledged by the clinician authors of our study. In the available literature, there are no cytogenetic effect studies on CKD which presented the X-ray exposure data of their populations. In the present study, 41.6 % of children with CKD had an X-ray exposure history (mostly chest radiography) in the last three months prior to the biological sampling. Therefore, as explained in the statistical analysis section, multiple regression analysis was performed after adjusting for factors such as X-ray exposure, age, gender, body mass index, smoking, drug usage, and sports activities (Tables 2, 3). Accordingly, X-ray exposure issue seemed to have no apparent confounding effect on buccal MN frequencies. Similar results have also been demonstrated in peripheral lymphocytes of the same children with CKD (15, 16). On the other hand, the X-ray exposure effect on nuclear anomalies other than MN, a positive X-ray effect, was only found for the formation of nuclear buds in BEC, while negative X-ray effects were found for condensed chromatin cells in BEC. The previously published studies that have investigated directly the effects of several radiographic and cephalographic techniques in dental practices (e.g. panoramic dental radiography, lateral cephalography, post anterior cephalography, and full-mouth X-ray) on the target tissue BEC collected ten days after X-ray exposure, which is the likely maximal rate of MN occurrence, have linked X-ray exposure to cytotoxicity but not genotoxicity determined as MN frequencies (39-42). In all these studies, an X-ray dose was between 0.003 mSv and 0.02 mSv (1-4). Choi (2011) compared the panoramic radiography for oral examination (0.014-0.024) to the effective doses of chest radiography (0.06-0.25 mSv in two views) and looked at variability (43). Therefore, the type, dose, target tissue, target tissue turnover of X-ray exposure, and time after exposure are crucial parameters. From this point of view, the interpretation of the X-ray exposure effect as a confounding factor in biomonitoring studies is quite difficult and providing detailed information and statistical analysis could be the appropriate approach, as we have done in this study. Obviously, the association between the ionising radiation effects and outcomes of buccal micronucleus assay has to be more extensively evaluated in future biomonitoring studies, in contrast with micronucleus assay on lymphocytes, for which this association has nowadays been fairly well established (44).

From several previously published studies (45-49), including the one conducted in our laboratory, the 46 buccal MN frequencies reported for healthy children ranged from 0.10 to 4.03 \%. The mean MN frequency for our control subjects (1.45 %) fell in that range. However, in our previous study, the background level variability observed in the buccal cells of healthy subjects (49) confirmed the need for method standardisation, which is an initiative of the HUMN_{x1} project (37). When frequencies of nuclear anomalies of condensed chromatin and karyorrhectic and pyknotic cells were compared in different healthy children populations, large variabilities were observed. While some of them (40, 50), including our study, had much higher values, others (51) had lower values than the levels obtained in the study by Thomas et al. (52). The causes of this variability may, in part, reflect genetic and non-specific exposure differences, or in part, technical factors and differences in the sample preparation and interpretation of the scoring criteria between laboratories. Therefore, it was suggested that an automated scoring system should be developed to overcome this issue (53).

CONCLUSION

Increased cytogenetic, cytokinetic, and cytotoxic effects have been found in the buccal epithelial cells of children with CKD who underwent different treatment modes. Both the disease and treatment of CKD contribute to increased cytogenetic, cytokinetic, and cytotoxic effects on buccal epithelial cells of children. We believe that our data will add to the validation studies in the scope of the HUMN_{XL} Project (37). Finally, they point to the sensitivity and usefulness of the BEC MN assay in the assessment of genetic susceptibility of patients with CKD.

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Declaration of interest

The authors declare that they have no competing interests.

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Mikronukleusi i ostala citogenetička oštećenja u epitelnim stanicama bukalne sluznice djece koja boluju od kroničnih bubrežnih bolesti

Cilj ovoga istraživanja bio je utvrditi moguću genomsku nestabilnost u djece koja boluju od kroničnih bubrežnih bolesti (KBB). Korišten je mikronukleus (MN) test na epitelnim stanicama bukalne sluznice. Ispitivali smo učestalost mikronukleusa i ostalih citogenetičkih oštećenja poput jezgrinih pupova, binuklearnih stanica, kondenziranoga kromatina, kariorektičnih i piknotičnih stanica u stanicama bukalne sluznice. Djeca s KBB-om bila su podijeljena u sljedeće skupine: djeca u fazi prije dijalize (PreD) (N=17), djeca koja su redovito na hemodijalizi (HD) (N=14) i djeca s transplantiranim bubregom (Tx) (N=17). Kontrolnu skupinu činila su zdrava djeca iste dobi i spola. Učestalost MN-a u stanicama bukalne sluznice u svim skupinama djece s KBB-om bila je značajno povećana (petero- do sedmerostruko) u usporedbi s kontrolnom skupinom (p<0,001). Suprotno tome, učestalost binuklearnih stanica i kondenziranoga kromatina bila je značajnije veća u svim podskupinama djece s KBB-om u odnosu na kontrolnu skupinu (p<0,001). Naši rezultati upućuju ne samo na povećanu razinu citogenetičkih, citokinetičkih i citotoksičnih promjena u stanicama bukalne sluznice u djece PreD, HD i Tx skupina nego i na osjetljivost i korisnost MN-testa na stanicama bukalne sluznice u procjeni genetičke osjetljivosti pacijenata koji boluju od KBB-a.

KLJUČNE RIJEČI: citogenetička oštećenja, dijaliza; genotoksičnost; mikronukleus-test na stanicama bukalne sluznice; transplantacija bubrega