First *IKBKG* Gene Mutation Study in Serbian Incontinentia Pigmenti Patients

Snežana Minić1,2, Dušan Trpinac1, Heinz Gabriel4, Martin Gencik4, Miljana Obradović3

1Clinics of Dermatovenerology, Clinical Center of Serbia, Belgrade, Serbia; 2School of Medicine, University of Belgrade, Belgrade, Serbia; 3Institute of Histology and Embryology, School of Medicine, University of Belgrade, Belgrade, Serbia; 4Diagenos, Center for Medical Genetics, Osnabruceck, Germany

**SUMMARY**

**Introduction** Incontinentia pigmenti (IP) is a rare X-linked dominant genodermatosis. Mutations of the *IKBKG* gene are the only known cause of IP. The presence of other than skin changes is important in the diagnosis of atypical IP cases when skin changes are discrete.

**Objective** The study was designed to analyze clinical manifestation, family histories and the frequency of *IKBKG* gene mutation in IP patients in Serbia for the first time and to compare them with other reported findings.

**Methods** Two Serbian unrelated families with eight female subjects were investigated. Blood samples were used for *IKBKG* exon 4-10 deletion testing using modified PCR protocol. For probands pathohistological and ultrastructural analyses of skin biopsies were done.

**Results** Positive clinical diagnosis according to IP criteria was present in seven cases. In six of them, including probands, positive molecular gene testing for *IKBKG* exon 4-10 deletion was present.

**Conclusion** This is the first report of genetically confirmed IP in two Serbian families. The IP patients investigated patients in Serbia were similar to results of other studies. Various clinical features of investigated patients have allowed us to demonstrate that molecular genetic testing which specifically detects the common *IKBKG* mutations, the only known cause of IP, is useful in diagnosing IP especially in mild or atypical cases. The molecular genetic testing of the *IKBKG* mutations may be helpful for rapid confirmation of IP diagnosis, prenatal diagnosis and carrier detection.

**Keywords:** Incontinentia pigmenti; *IKBKG* gene; *IKBKG* exon 4-10 deletion; X-chromosome; X-chromosome inactivation; phenotype

**INTRODUCTION**

The X-chromosome holds a unique place in medical genetics since a disproportionately large number of disease conditions have been associated with the X-chromosome because the phenotypic consequence of a recessive mutation is revealed directly in males for any gene that has no active counterpart on the Y-chromosome [1]. Gene expression on one (maternal or paternal) of the female X-chromosomes is random. Skewed X-chromosome inactivation is a marked deviation from a 50:50 ratio [4]. It may occur under genetic influence or under selection that will favor cells in which the normal X is the active X-chromosome [4].

Incontinentia pigmenti (IP; Bloch-Sulzberger syndrome; MIM 308300) is a rare X-linked dominant genodermatosis [5]. The estimated prevalence for IP is 0.2/100,000 [6]. It appears almost exclusively in females and is usually lethal *in utero* for males [5]. It is a multi-system disorder predominantly affecting ectodermal tissues: skin, hair, nails, teeth, eyes and central nervous system [5]. Criteria for IP proposed by Landy and Donnai [7] are in routine practice since 1993. The presence of other than skin changes is important in atypical IP cases when skin changes are discrete. They can be of great prognostic and diagnostic value because they will be present throughout the patient’s whole life, while skin changes usually fade [8]. The prognosis of IP is generally good and depends on extracutaneous manifestations that may also affect patients’ quality of life. According to Landy and Donnai’s criteria [7], skin lesions as well as multiple male miscarriages were classified as IP major criteria, while dental, hair, nails and retinal anomalies were classified as IP minor criteria. Dental abnormalities were registered in 54.38% of IP patients and comprise dental shape anomalies, hypodontia, and delayed dentition [9]. A typical hair change is scarring alopecia. Ophthalmologic findings occurred in 36.5% IP patients and include retinal anomalies, strabismus, vitreous and lens anomalies, optical nerve atrophy, and microphthalmus [10].
IKBKG (Inhibitor of Kappa light polypeptide gene enhancer in B-cells, Kinase Gamma, previously NEMO) is the only gene known to be associated with IP [11]. Mutations of the IKBKG gene are responsible for IP [5]. The IKBKG gene is composed of 10 exons. Located at Xq28, IKBKG has a unique genomic organization, as it is part of a segmental duplication or low-copy repeat 1 and 2 (LCR1 and LCR2) containing the gene and its pseudogene copy (IKBKGP1). The two LCRs in the IKBKG locus are able to recombine producing a pathological recurrent IKBKG exon 4-10 deletion [12].

Within cells, the IKBKG protein interacts with two enzymes, IKK-alpha and IKK-beta, to activate NF-κB (Nuclear Factor-kappa-B). The activated factor then moves into the nucleus and binds to DNA. NF-κB regulates the activity of multiple genes, including genes that control the body’s immune responses and inflammatory reactions. It also protects the cell from certain signals that would otherwise undergo apoptosis [13]. Its misregulation is involved in many diseases [14]. However, failure to identify IKBKG mutations does not rule out the diagnosis of IP [11]. The phenotypic expression of IKBKG mutations does not rule out the diagnosis of IP [11]. The phenotypic expression of IKBKG mutation is highly variable, even among related patients with the same mutation [5]. Because females with IP have skewed X-chromosome inactivation in which the X-chromosome with the mutant IKBKG allele is preferentially inactivated [15] it was thought that this variability was likely to be result of skewed X-chromosome inactivation [16]. It was also suggested that the phenotype of IP might be due to the pleiotropic role of the NF-κB [12].

**OBJECTIVE**

This study was designed to analyze clinical manifestation, family histories and identify the existence and frequency of IKBKG gene mutation in IP patients in two Serbian families and to compare them with other reported findings.

**METHODS**

We investigated two Serbian unrelated families with eight subjects, including two probands, all of them females. Pedigrees for both families were made (Figure 1). Probands were first examined by a dermatologist, since they had obvious skin changes, and then sent for additional examinations to stomatologists, ophthalmologists etc. Routine laboratory analyses were done for all patients.

**Pathohistological and ultrastructural analysis**

Skin biopsies were taken and pathohistological and ultrastructural analysis were done to confirm the diagnosis of IP in both probands. For pathohistological analyses, skin biopsies from affected areas were fixed in formaldehyde and stained with hematoxyline and eosin in a routine manner. For transmission electron microscopy investigation, skin biopsies were fixed in glutaraldehyde, postfixed in osmium tetroxide and embedded in araldite resin in a routine manner [17]. The ultra-thin sections stained with uranyl acetate and lead citrate [17] were analyzed with transmission electron microscopes Philips EM 300 (Philips, Eindhoven, The Netherlands) and LEO 912AB (Carl Zeiss SMT, Oberkochen, Germany).

**DNA sampling and genotyping**

Blood samples were collected and used to extract DNA using standard protocols. Molecular genetic testing was done at Diagenos, Center for Medical Genetics, Osnabrueck (Germany). For deletion testing a modified PCR protocol according to Smahi et al. 2000 [5] was performed for all patients except III 2 Family 2. For III 2 Family 2 who had no clinical signs of IP all coding exons of IKBKG gene were amplified and subsequently sequenced. Obtained sequences were analyzed by comparing to data base entries. Additional deletion testing was done.

<table>
<thead>
<tr>
<th>Family</th>
<th>Subject</th>
<th>Age at onset</th>
<th>Age of patients</th>
<th>IP stage at first exam</th>
<th>IKBKG exon 4-10 deletion positive</th>
<th>Skin pathohistology</th>
<th>Skin ultrastructure</th>
<th>Clinical findings</th>
<th>Miscarriages</th>
</tr>
</thead>
<tbody>
<tr>
<td>Family 1</td>
<td>III 1</td>
<td>5 days</td>
<td>1980</td>
<td>1, 2, 3*, 4*</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td></td>
<td>II 2</td>
<td>/</td>
<td>1968</td>
<td>3, 4</td>
<td>+</td>
<td>/</td>
<td>/</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td></td>
<td>II 4</td>
<td>/</td>
<td>1964</td>
<td>3, 4</td>
<td>+</td>
<td>/</td>
<td>/</td>
<td>-</td>
<td>+</td>
</tr>
<tr>
<td></td>
<td>12</td>
<td>/</td>
<td>1940</td>
<td>-</td>
<td>+</td>
<td>/</td>
<td>/</td>
<td>-</td>
<td>+</td>
</tr>
<tr>
<td>Family 2</td>
<td>III 1</td>
<td>39 days</td>
<td>1996</td>
<td>1, 2, 3*, 4*</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>4*</td>
</tr>
<tr>
<td></td>
<td>III 2</td>
<td>/</td>
<td>1998</td>
<td>-</td>
<td>-</td>
<td>/</td>
<td>/</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td></td>
<td>II 3</td>
<td>/</td>
<td>1977</td>
<td>-</td>
<td>+</td>
<td>/</td>
<td>/</td>
<td>-</td>
<td>+</td>
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<tr>
<td></td>
<td>12</td>
<td>/</td>
<td>1952</td>
<td>-</td>
<td>-</td>
<td>/</td>
<td>/</td>
<td>-</td>
<td>+</td>
</tr>
</tbody>
</table>

* At the age of 14 years
The investigation protocol followed the guidelines of the Declaration of Helsinki and was approved by the Clinical Center of Serbia Ethics Committee. Written informed consent was obtained from all participants or their parental guides.

RESULTS

Basic subjects’ data are presented in Table 1. Out of eight subjects, positive clinical diagnosis of IP according to Landy and Donnai’s criteria [7] were present in seven cases. Positive molecular gene testing for IKBKG mutation was present in six subjects - in both probands and four relatives from two families (Figure 2A). All of them were with positive clinical diagnosis of IP according to Landy and Donnai’s criteria [7]. Both families were presented in our previously published articles in context of ocular [18], oral and dental anomalies in IP [19].

Probands had skin (Figure 2B, 2C), dental and oral changes typical for IP while eosinophilia was not registered. Also, they had pathohistological (Figure 2D) and ultrastructural skin findings characteristic for appropriate stage of IP (Figure 2E). Common IKBKG exon 4-10 deletion was detected in six of seven clinically diagnosed IP patients and no IKBKG exon 4-10 deletion was found in the remaining case. One of the examinees, the sister of proband in Family 2 (III 2), had none of the phenotypical signs of IP. Both probands inherited the IKBKG exon 4-10 deletion from their mothers.

In Family 1 three out of four examined patients had typical skin changes. All four tested patients were positive for IKBKG exon 4-10 deletion. In Family 2 only proband had typical skin changes and two out of four family members were positive for IKBKG exon 4-10 deletion.

DISCUSSION

Although numerous molecular genetic confirmations of IP have been described in the world literature [20], up to date there were not genetically confirmed patients from Serbia. In this study, a genetic analysis of two unrelated Serbian families clinically diagnosed according to Landy and Donnai’s criteria [7] with IP was performed.

Skin abnormalities are consistent IP features and usually occur in four stages that evolve sequentially [11]. The pattern of skin changes follows lines of embryonic and fetal skin development known as Blaschko’s lines that correspond with cell migration and growth pathways that are established during embryogenesis and represent functional X-chromosome mosaicism [21]. Blaschko’s lines are linear on the limbs and circumferential on the trunk [11]. Stage 1, the bullous stage, is characterized by erythema and blistering within the first few weeks of life. Stage 2, is characterized by a hypertrophic rash within the first few months of life. Stage 3, is characterized by hyperpigmentation that occurs along Blaschko’s lines. It starts usually as stage 2, begins to resolve and persists into adulthood. The hyperpigmentation usually begins to fade in the teens and early twenties. It is the reason why IP patients in their thirties may have no skin changes associated with IP. Stage 4, the hypopigmented stage, opposite to stage 3 is characterized by linear hypopigmentation and alopecia. Stage 4 does not occur in all patients. The onset, duration and overlapping of IP stages vary among patients, and not all patients experience all four stages [11]. Besides variable appearance of skin phenotype in IP patients, significant clinical heterogeneity exists with regard to eye, dental, and neurological abnormalities, even within families [22]. These factors are the cause of diagnostic dilemma in atypical or mild IP cases.

In the process of confirming IP diagnosis reliance on the skin biopsy is important. However, one must consider that IP skin changes have clinical course with four different stages with different histological features in each stage [23]. As in IP is not always possible to detect IKBKG mutation, in such situations of hidden IKBKG mutations, histology can be very helpful for confirmation of IP [24].

Clinical features of IP in seven diagnosed patients were diverse. Both probands inherited common IKBKG exon 4-10 deletion from their mothers. They had skin changes typical for IP and previously confirmed IP diagnosis using pathohistology and ultrastructural analyses of skin biopsies because, at that time, IKBKG was not discovered as a

Figure 2. A. Result of deletion testing in the investigated Family 1. A set of deletion spanning PCR primer were used for the detection of the common exon 4-10 deletion. The deletion is characterized by additional PCR product of 1.8 kb and 1045 bp not seen in control samples. B. Hyperpigmented stage (stage 3) maculas along Blaschko’s lines on the trunk. C. Hypopigmented, atrophic, hairless arias on the shin (stage 4). D. Histopathological findings in proband 1 (stages 3 and 4). Pigment localized in dermal cells (arrows) and epidermal hyperkeratosis (asterisk) (hematoxyline and eosin, magnification 150×). E. Ultrastructural appearance of dermis of proband 1 (stages 3 and 4). phagocytosed melanosome clusters (arrows) localized in cytoplasm of dermal cells (magnification 4800×)

Table 1. Basic subjects’ data.

<table>
<thead>
<tr>
<th>Subject</th>
<th>Age</th>
<th>Gender</th>
<th>Diagnosis</th>
<th>IKBKG Mutation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Proband 1</td>
<td>12</td>
<td>F</td>
<td>Yes</td>
<td>Yes</td>
</tr>
<tr>
<td>Proband 2</td>
<td>14</td>
<td>M</td>
<td>Yes</td>
<td>No</td>
</tr>
<tr>
<td>Family 1</td>
<td>11</td>
<td>M</td>
<td>Yes</td>
<td>Yes</td>
</tr>
<tr>
<td>Family 2</td>
<td>13</td>
<td>F</td>
<td>Yes</td>
<td>Yes</td>
</tr>
</tbody>
</table>

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causative gene for IP and molecular genetic testing was not available. Six of seven IP affected individuals showed an identical genomic alteration, common IKBKG exon 4-10 deletion. Four IP patients with IKBKG exon 4-10 deletion showed the classical cutaneous signs of IP. Two patients with IKBKG exon 4-10 deletion had no IP skin changes, but had noncutaneous anomalies typical for IP and fulfilled Landy and Donnai’s IP criteria [7]. Only one examinee, III 2 Family 2, had no phenotypic signs of IP and was negative for IKBKG mutation. Though she did not show phenotypic signs of IP, she was analyzed as at-risk female relative since the IKBKG gene mutation had been identified in her family and to discover if she was a carrier of the disease. In this study, no IKBKG exon 4-10 deletion was found in the remaining IP case, I 2 Family 2. As failure to identify IKBKG mutation does not rule out the diagnosis of IP [11], in this case IP may be attributed to the somatic mosaicism which has been reported in some studies [11, 12, 25].

Most of the studies on IP with molecular genetic confirmation were reported from Europe and North America [12, 20, 25]. A few reports of IP were available from Asia [26, 27, 28] and Australia [29]. A large deletion of IKBKG exons 4 to 10 is found in approximately 80% of IP patients [5, 20]. In general, the frequency of IKBKG mutation in IP patients was similar in all studies irrespective of the ethnic background [26]. The frequency of IKBKG exon 4-10 deletion in Serbian population (6/7) was similar to the results in other ethnic groups [5, 19, 26]. Besides the most frequently found IKBKG exon 4-10 deletion [20], a total of 79 different small IKBKG mutations (missense, frameshift, nonsense, and splice-site mutations) have been reported [12, 25, 30, 31].

As a consequence of skewed X-chromosome inactivation [16] and consequent mosaicism, and the pleiotropic role of the NF-κB [12] phenotypical features of IP are variable and it is difficult to diagnose cases with mild manifestations [8]. It is likely that IP in mildly affected patients is often undiagnosed and underrepresented because older patients have poor recall, and their mothers may no longer be alive [29]. Genetic testing is especially helpful in diagnosing IP in such patients. Molecular genetic diagnosis may help to confirm the clinical suspicion of IP and is essential for providing definite genetic counseling and prenatal diagnosis [26]. Carrier testing of at-risk female relatives is possible if the IKBKG mutation has been identified in the family. Preimplantation and prenatal genetic diagnosis may be available for families in which the IKBKG mutation has been identified [11].

Our study is the first to report genetically confirmed IKBKG exon 4-10 deletion in six IP patients from two families in Serbian population. We investigated IP patients for more than 20 years and registered 9 families with 22 subjects, two of them males [18, 19]. In the present study we analyzed two families willing to cooperate for further investigations, molecular genetic testing, that became available long after they have been diagnosed for IP. The most frequent molecular cause, the IKBKG exon 4–10 deletion, is the same as in other populations. Six out of seven IP patients had a common genomic rearrangement involving the deletion of exons 4 to 10 in IKBKG and of mutation incidence similar to previously reported [20].

A relatively small number of IP patients were investigated but the variety of their clinical features have allowed us to demonstrate that molecular genetic testing which specifically detects the common IKBKG exon 4-10 deletion is useful to confirm the diagnosis in IP patients with typical, and especially with mild or atypical phenotype.

**CONCLUSION**

This is the first report of genetically confirmed IP patients in Serbian population. Six of seven investigated IP patients presented a common IKBKG exon 4-10 deletion. The frequency and type of IKBKG mutation found in IP patients in Serbia were similar to the results of other studies irrespective of the ethnic background. Although a relatively small number of patients were investigated, this study indicates that molecular genetic analysis of IKBKG is helpful for rapid confirmation of IP diagnosis particularly in mild or atypical cases and in prenatal diagnosis and carrier detection.

**ACKNOWLEDGEMENTS**

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**NOTE**


**REFERENCES**

Minić S. et al. First IKBKG Gene Mutation Study in Serbian Incontinentia Pigmenti Patients


Прва студија мутације гена IKBKG код болесника са инконтинемцијом пигменти у Србији

Снежана Минић1,2, Душан Трпинац3, Хајнц Габијел4, Мартин Генцик4, Миљана Обрадовић3

1Клиника за дерматовенерологију, Клинички центар Србије, Београд, Србија;
2Медицински факултет, Универзитет у Београду, Београд, Србија;
3Клиника за дерматовенерологију, Клинички центар Србије, Београд, Србија;
4Медицински факултет, Универзитет у Београду, Београд, Србија;

КРАТАК САДРАЖ
Увод Инконтинемција пигменти (ИП) је ретка генодермата која се наслеђује доминантно везано за X-хромозом. За појаљу ИП одговарају мутација гена IKBKG. У дијагностикованим нетичним случајевима ИП, када су промене које дискретне, важно је постојање промена на другим органима. Циљ рада Циљ истраживања је била анализа клиничких манифестација ИП, породичне историје болести, молекуларногенетичких тестирања ради утврђивања постојања и унучеталости мутације гена IKBKG код болесника са ИП из Србије (првих пута) и упоређивање са резултатима других студија. Метода рада Испитиване су две несрдне породице из Србије са осам испитивана. Узорци крви су коришћени за молекуларногенетичко тестирање болесница. Резултати На основу критеријума за утврђивање ИП, код седам испитанца је била позитивна клиничка дијагноза ИП. Позитивно генетично тестирање делеције гена 4–10 IKBKG је показано код шест болесница. Закључак Ово је први извештај о ИП код болесника из две породице у Србији којима је дијагноза потврђена и молекуларногенетичким тестирањем. Болесници су имали типичну делецију гена 4–10 IKBKG. Учесталост и тип мутације овог гена код испитиваних болесника из Србије слични су резултатима других студија. Различите клинички симптоми код испитиваних болесника се омогућиле да пажљиво доживелим о томе да је молекуларногенетичко тестирање које специфично открива мутације гена IKBKG, јединог гена што код болесника ИП, врло корисно для брзо потврђивање дијагнозе, препрека транзиторним и откривањем преносацама гена. Кључне речи: инконтинемција пигменти; ген IKBKG; делеција гена 4–10 IKBKG; Х-хромозом; инактивација Х-хромозома; фенотип