

DEGENERATIVE DISC DISEASE IN YOUNG ADULTS: CYTOKINE PROFILE AND ANGIOGENIC FACTORS

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Back pain (BP), associated with the degenerative disc disease (DDD), poses a heavy social and economic burden due to early disability and indications to surgery, emerging in young adults. Pathophysiological basis of premature intervertebral disc (IVD) degeneration is being actively studied. The study was aimed to define the profiles of inflammatory cytokines in DDD, as well as their relationship to the structural spine diseases. The molecular genetic analysis of the mRNA gene abundance in patients with BP and herniated IVD after discectomy and healthy individuals was performed by the quantitative polymerase chain reaction method. High expression of TNF α , IL17 was revealed in the IVD tissues of the affected patients ($p < 0.01$); the levels of TNF α and IL1 β correlated with the DDD severity ($r = 0.301$ and 0.37 ; $p < 0.05$). Elevated expression of IL1 β , IL6 was found in peripheral white blood cells ($p < 0.01$); the levels of IL6 negatively correlated with Modic type 1 and 2 changes ($r = -0.31$; $p < 0.05$), and the levels of IL17 positively correlated with the IVD herniation in combination with erosions of the adjacent vertebral body endplates and Modic changes ($r = 0.401$; $p < 0.05$). The expression of VEGF-A in the IVD tissues and white blood cells negatively correlated with the DDD grades ($r = -0.85$; $p < 0.001$), indicating reduced vascularization in the terminal phase of the disease. The findings on DDD demonstrate the contribution of the local low-immune inflammation, coupled with the intense disc vascularization at the earlier stages, and associated with the reactive inflammation in vertebral bodies. The results are prerequisites for developing the anti-inflammatory and reparative therapy based on the DDD grade and the presence of Modic changes in young adults with BP.

Keywords: back pain, degenerative disc disease, young age, discectomy, cytokine expression, mRNA TNF α , IL1 β , IL6, IL17, VEGF-A, Modic-changes

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ДЕГЕНЕРАТИВНАЯ БОЛЕЗНЬ ДИСКА У МОЛОДЫХ: ЦИТОКИНОВЫЙ ПРОФИЛЬ И ФАКТОРЫ АНГИОГЕНЕЗА

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Боль в спине (БС), ассоциированная с дегенеративной болезнью диска (ДБД), — тяжелое социальное и экономическое бремя вследствие ранней инвалидизации и возникновения показаний к оперативному вмешательству уже в молодом возрасте. Патолофизиологические основы преждевременной дегенерации межпозвоночного диска (МПД) находятся на стадии активного изучения. Целью исследования было определить профиль воспалительных цитокинов при ДБД и их связь со структурными нарушениями в позвоночнике. У пациентов молодого возраста с БС и грыжей МПД, подвергшихся дискэктомии, и у здоровых лиц проводили молекулярно-генетический анализ представленности генов мРНК методом количественной полимеразной цепной реакции. У больных в ткани МПД выявлен высокий уровень экспрессии TNF α , IL17 ($p < 0,01$); уровни TNF α и IL1 β коррелировали с тяжестью ДБД ($r = 0,301$ и $0,37$; $p < 0,05$). В лейкоцитах периферической крови обнаружена повышенная экспрессия IL1 β , IL6 ($p < 0,01$); уровень IL6 отрицательно коррелировал с I и II стадиями Modic-изменений ($r = -0,31$; $p < 0,05$), IL17 прямо коррелировал с грыжей МПД в сочетании с эрозией замыкательных пластин и Modic ($r = 0,401$; $p < 0,05$). Экспрессия VEGF-A в ткани МПД и в лейкоцитах крови отрицательно коррелировала со стадией ДБД ($r = -0,85$; $p < 0,001$), указывая на снижение активности васкуляризации в терминальной стадии. Данные, выявленные при ДБД, говорят о вкладе локального низкоиммунного воспаления, сопряженного с активной васкуляризацией диска на более ранних стадиях и ассоциированного с реактивным воспалением тел позвонков. Полученные результаты служат предпосылкой к разработке противовоспалительной и репаративной терапии в зависимости от стадии ДБД и наличия Modic-изменений у лиц молодого возраста с БС.

Ключевые слова: боль в спине, дегенеративная болезнь диска, молодая возраст, дискэктомия, экспрессия цитокинов, мРНК TNF α , IL1 β , IL6, IL17, VEGF-A, Modic-изменения

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Back pain (BP) is one of the main causes of the patients' disability in the developed world, resulting in permanent disability. BP is found in all age groups. Thus, according to a Polish study, BP recurrence within 34 years was observed in 85% of males and 86% of females with the BP onset at the age of 14 [1]. Degenerative disc disease (DDD), resulting from degradation and inflammation of the intervertebral disc (IVD) tissues, is one of the BP variants, associated with unfavourable outcomes [2]. DDD is a chronic condition with a trend towards progression. Despite the fact that there are still no explicit criteria for distinguishing between the "natural" physiological disc ageing and the pathological degeneration, also found in young adults, the term "DDD" is used by both clinicians and pathologists to define the disc extracellular matrix disruption with impaired homeostasis and the inflammatory process induction in the IVD space [3]. The DDD clinical manifestations are well documented: these are BP of mechanical origin associated with axial load (getting worse with physical activity, moving heavy objects, flexion, and improving at rest) and/or with spinal stenosis, radiculopathy, and less often with myelopathy. DDD is one of the causes of chronic segmental instability and early disability in the working age patients. Studies have shown that the IVD degeneration is a multifaceted process, involving apoptosis, inflammation, ageing, and biomechanical dysfunction [4]. Recently, considerable attention has been paid to studying the effects of inflammatory cytokines on the DDD development, as well as to the imbalance between the pro-inflammatory and anti-inflammatory cytokines [5].

It is shown, that inflammatory pattern triggers the catabolic processes in the cartilage matrix of the compromised functional spinal unit, together with further degeneration of extracellular matrix and dehydration of nucleus pulposus (NP) and annulus fibrosus (AF) [5, 6]. The loss of the disc structural integrity and the AF microcracks cause the prolapse of the NP content into the AF tissue and outwards with the formation of protrusion, extrusion, and sequestration. The role of TNF α , IL1 β , overexpression of IL6, and CD16 monocytes in the development and progression of degenerative changes is being discussed in literature. It is important to mention that the cytokine expression may be correlated with the IVD degeneration severity. The role of IL6 in the disc degeneration and herniation was studied by many researchers [7]. The hypothesis about the IL6 involvement in the human IVD degeneration was confirmed by the fact that the abundance of the catabolic gene transcript increased, and the expression of genes encoding proteoglycans was suppressed with the increase in the IL6 expression [8]. Studying the role of IL17 in the DDD-associated inflammatory cascade showed that the exposure of human NP cells to IL17 and TNF α contributed to the increased release of IL6 in vitro and the increased expression of intercellular adhesion molecules (ICAM-1) on the surface of cells in NP and AF [9]. The association between pain intensity, IVD herniation, and inflammatory response was defined. It was suggested to treat the listed above biomarkers as the potential markers of the disease onset, severity, and progression.

The possibility of using the analysis of serum inflammatory biomarkers for identification of degeneration and inflammation of the IVD tissues in patients with DDD was demonstrated [7, 10]. Molecular patterns of degeneration and inflammation in the disc tissues were assessed in order to develop the approaches to the reparative therapy of DDD [3]. IL1 β , TNF α , and VEGF were verified by immunohistochemistry in individuals of various ages having no symptoms of BP [11]; the expression of these markers in the disc tissues showed almost no differences between the groups of young and elderly individuals having no

symptoms of the disease. Accordingly, measuring the cytokine levels in the damaged disc is an important challenge in terms of searching for the pathogenetically substantiated anti-inflammatory and reparative therapy. Recently, the role of the vascular endothelial growth factor (VEGF, VFA) in the disc tissue vascularization (the disc is normally avascular) at all stages of the IVD degeneration was shown. The VEGF activity is realized via regulation of the soluble vascular endothelial growth factor receptor 1 expression [12].

When studying the pattern of the hernia resorption in patients with DDD (confirmed by MRI), Japanese researchers assessed the interaction sequence TNF α -VEGF-MMP (matrix metalloproteinases MMP-3 and MMP-7 involved in degradation of the extracellular matrix proteins, aggrecan and collagen) and found that the expression of mRNA and VEGF protein increased in the situation of the contact between macrophages and human disc tissues in vitro, and positively correlated with the TNF α expression levels [13]. Thus, it was shown that neovascularization promoted the reverse development of intervertebral hernias. The process of the capillary ingrowth into the IVD tissue was confirmed by magnetic resonance imaging (MRI) with the use of gadolinium-based contrast agents, and could serve as an additional determinant of the extrusion resorption [14].

The relationship between the markers of aseptic inflammation and IVD reparation in the disk tissues, and the abundance of those in peripheral blood of young patients with advanced grade DDD have not been fully defined. Various data are available on the preponderant role in altering the IVD immune homeostasis, played by one or another cytokine. Studying the cytokine profile (together with the features of the functional spinal unit lesion) would make it possible to define the immune phenotypes of patients in order to develop the biological targets for therapy and prognosis of the disease. The study was aimed to define the profiles of the key biomarkers of inflammatory damage (TNF α , IL6, IL17, IL1 β) and angiogenesis (isoforms of the vascular endothelial growth factor A, VEGF121, VEGF165, and VEGF189) in the cartilage tissue of IVDs and white blood cells of young patients with DDD, who underwent discectomy, compared to controls.

METHODS

The study was carried out at the Department of Neurosurgery, Pirogov City Clinical Hospital № 1, and A.I. Nesterov Department of Faculty Therapy, Pirogov Russian National Research Medical University, in 2019–2021. A total of 87 young (aged 18–44 in accordance with the WHO classification, 2012) adults were enrolled (40 males and 48 females). Index group inclusion criteria: young patients (median age 37.01 years [35.54–38.49]) having BP associated with DDD, confirmed by instrumental evaluation (MRI) (Table). Exclusion criteria: history of spinal injury or spinal injury at the time of the study, tumors and infections affecting the spine and other organs, inflammatory spondyloarthropathy, surgical interventions in the previous 30 days. All individuals in the index group underwent surgery (microdiscectomy) due to the spinal disc herniation at the corresponding level in the lumbar spine. Pain intensity was measured in millimeters using the Visual Analogue Scale (VAS). The functional limitations in the lumbar spine were assessed based on the Backache Index (BAI) [15]. The control group was represented by healthy volunteers with no BP (20 individuals), comparable in gender and age. All patients and controls underwent MRI of the lumbar spine prior to surgery. IVD degeneration was assessed based on the reduced IVD

Table. Demographic, clinical and instrumental characteristics of the studied groups

Characteristics of patients	Index group		Control group		P I-II
Number of patients	67		20		
females	30		10		
males	37		10		
Average age, years	Me	LQ-UQ	Me	LQ-UQ	
	37,01	[35,54–38,49]	34,5	[29,26–39,74]	< 0,01
Pfirrmann degeneration grade at the level of L1–L5 (mean value)	2,62	[2,4–3,0]	1,2	[1,1–1,8]	< 0,01
Pfirrmann degeneration grade at the level of operated IVD (mean value)	M	σ	M	σ	
	4,26	$\pm 0,59$	2,15	$\pm 1,18$	< 0,01
Number of patients with herniated IVDs at the level of L4–L5 at the level of L5–S1	Abs. (%)		Abs. (%)		
	24 (43) 36 (64)		1 (5) 0		< 0,01
Number of patients with Modic changes (total)	47 (70)		1 (5)		< 0,01
Modic 1	21 (45)		0		< 0,01
Modic 2	26 (55)		1 (5)		< 0,01
Modic 3	0		0		< 0,01
Index group					
Characteristics of patients	Me	LQ–UQ			
Disease duration, years	5	[2,0–10,0]			
Duration of the last painful episode, weeks	6	[3,0–12,0]			
Pain intensity (VAS, mm)	68	[48,0–88,0]			
Variant of pain syndrome acute chronic	Abs.		%		
	12		18		
	55		82		
Functional impairment severity (Backache Index (BAI))					
	mild	11	18		
	moderate	16	24		
	severe	40	58		

signal intensity, disc space narrowing, structural changes in the disk, and blurring between the nucleus pulposus (NP) and the annulus fibrosus (AF). IVD degeneration was graded in accordance with the C.W. Pfirrmann grading system (2001) (grades 1–5). Grades 3, 4, 5 were treated as irreversible damage to the disc, and grades 4, 5 were interpreted as severe DDD.

The fragments of degenerative IVDs were obtained during discectomy. Microparticles of IVD smaller than 1 mm³ were immediately immersed in the IntactRNA stabilization solution (non-toxic aqueous fixative solution for rapid stabilization of cellular RNA in tissues and cell cultures, preserving the cell integrity), and the disc tissue samples were assigned a barcode, identical to the barcode of the patient's blood samples. The same molecular markers were identified in blood and cartilage tissue samples, selected in accordance with the recent data based on the review articles and technical accessibility of the laboratory diagnosis (DNA-Technology; Russia). The abundance of mRNAs of the studied genes in blood cells (white blood cells) and IVD tissue was defined by the reverse transcription-quantitative polymerase chain reaction method with the use of the reagent kits (DNA-Technology; Russia). Amplification was performed in the DTprime 4 PCR system (DNA-Technology; Russia).

Obtaining IVD tissue samples (1 mm³): RNA was extracted after cutting the cartilage with surgical blade (sterile disposable surgical scalpel, manufactured by Huaiyin Medikal Instruments

Co. Ltd., China). After cutting, the fragment sized 1 mm was obtained, which was further grinded by the rubbing the conus of the microcentrifuge tube against the Petri dish bottom in order to obtain fine particles and molds. Then 320 μ L of the lysing solution from the Proba-NK kit (DNA-Technology; Russia) were added to the dish. The contents of the dish together with the crushed sample were transferred to the Axygen microcentrifuge tubes (Axygen, Inc.; USA), and then mixed using the Micro-Spin FV-2400 centrifuge/vortex (Biosan; Latvia) for 5 s, and sedimented. Subsequently, this was left to lyse for 1 h. After that RNA was isolated using the Proba-NK kit (DNA-Technology; Russia) in accordance with the manufacturer's guidelines. The extracted RNA in the amount of 16.5 μ L was immediately used for reverse transcription, which was performed at a temperature of 40 °C for 30 min, with subsequent inactivation of reverse transcriptase at 95 °C for 5 min. The resulting cDNA solution was either immediately used for quantitative PCR, or stored at –20 °C. The volume of 35 μ L was used for amplification with the real-time registration of the results in accordance with the following program: 50 cycles 94 °C — 10 s, 64 °C — 20 s, 72 °C — 10 s. Fluorescence was measured during each cycle at a temperature of 64 °C.

The 4 ml blood samples were collected into the disposable Vacutainer EDTA tubes (Becton Dickinson; USA) 24 hours before surgery with subsequent blood processing in order to extract white blood cells. To obtain the buffy coat, the Proba-Ficoll kit for extraction of lymphocytes from the whole blood was used.



Fig. 1. Patient K, aged 37 with BP and Pfirrmann grade 4 DDD at the level of L5–S1, with IVD herniation (arrow) and Modic type 1 changes in vertebral bodies (yellow contour). T2-weighted and STIR images

RNA was isolated using the Proba-NK kit (DNA-Technology; Russia) in accordance with the manufacturer's guidelines. RNA extracted in the amount of 16.5 μ L was immediately used for reverse transcription, which was performed at a temperature of 40 $^{\circ}$ C for 30 min, with subsequent inactivation of reverse transcriptase at 95 $^{\circ}$ C for 5 min. The resulting cDNA solution was either immediately used for quantitative PCR, or stored at -20° C. The volume of 35 μ L was used for amplification with the real-time registration of the results in accordance with the following program: 50 cycles 94 $^{\circ}$ C — 10 s, 64 $^{\circ}$ C — 20 s, 72 $^{\circ}$ C — 10 s. Fluorescence was measured during each cycle at a temperature of 64 $^{\circ}$ C.

The studied genes included TNF α , IL6, IL17, IL1 β , and isoforms of vascular endothelial growth factor A (VEGF121, VEGF165, VEGF189). Normalization genes were represented by β 2-microglobulin (B2m), and β -glucuronidase (GUSB). Normalization values for each gene mRNA were calculated by the $\Delta\Delta$ Ct method [16]. The expression levels of the gene mRNAs were expressed in arbitrary units in relation to normalization genes (B2m, GUSB), which had the relatively stable expression levels.

Statistical analysis

The nonparametric method (Mann–Whitney U test) was used to assess the statistical significance of the differences obtained. Quantitative indicators were tested for normality using the Shapiro–Wilk test. Qualitative indicators were compared using the chi-squared test and the two-tailed Fisher's test for small samples. The differences between groups were considered significant when $p < 0.01$ and $p < 0.05$. Data analysis was performed using the Statistica v 8.0, SPSS v.10, and Graph Pad Prism software.

RESULTS

Clinical and instrumental characteristics of patients

Clinical characteristics of the young patients with BP are presented in Table. The average pain intensity value corresponded to 68 mm [48.0–88.0]. Acute pain (lasting for a maximum of 12 weeks) was found in 18% of patients, and 82% of patients had chronic pain. The median disease

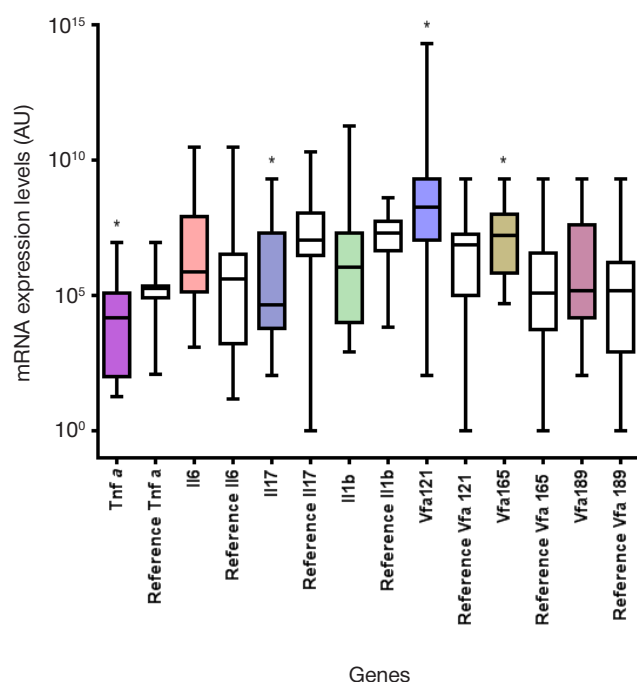


Fig. 2. Expression levels of mRNAs of the studied genes in relation to normalization genes (B2m, GUSB) in the IVD tissues of the index group patients and controls (* — $p < 0.01$)

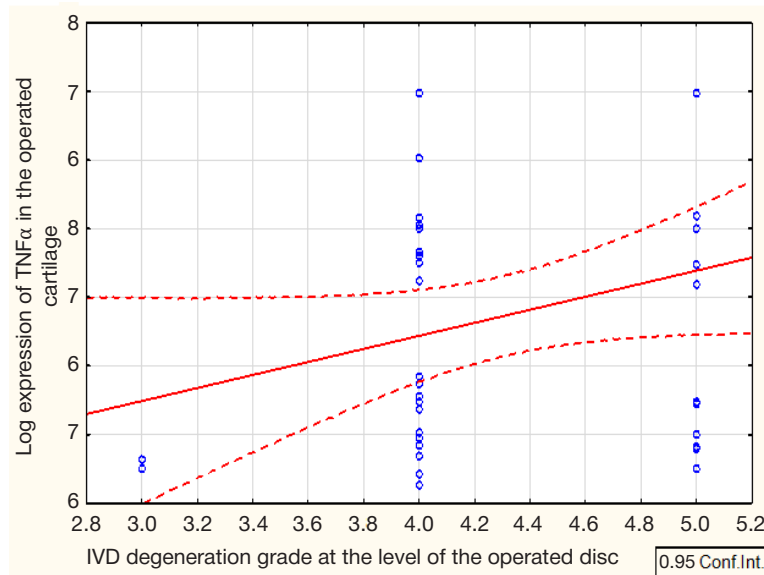


Fig. 3. Correlation between the TNF α mRNA expression levels in the IVD tissues and the Pfirrmann degeneration grades in the index group patients

duration in the latter was 5 years [2.0–10.0]. Based on BAI values, 82% of patients had moderate and severe functional impairment. The lumbar spine MRI showed the significant disc space narrowing, blurring between NP and AF, and reduced NP signal intensity. At the level of the operated disc, the IVD degeneration grade corresponded to grades 4 and 5 of the Pfirrmann grading system in 66% and 33% of the index group patients, respectively. Herniation was localized at the levels of L4–L5, L5–S1 at a ratio of 36 and 64%, respectively. In 70% of patients, MRI at the level of the operated disc revealed the altered intensity of the bone marrow MR signal on the T2-weighted and STIR (short tau inversion recovery) images, which indicated Modic type 1 changes, the bone marrow edema (45% of patients) (Fig. 1), and Modic type 2 changes, the fatty degeneration (55%), in almost equal proportions. In the control group, Modic changes were found in only one patient (5%) out of 20, and these changes were also associated with severe DDD and asymptomatic IVD herniation. MRI revealed the combination of IVD herniation with erosions of the adjacent vertebral body endplates and Modic changes in 37 index group patients (53.6%). Such a "triple combination" was associated with longer BP duration (years), morning pain, and persistent chronic pain with no "lucid intervals" ($p < 0.05$) compared to

patients with no erosions of the vertebral body endplates and/or Modic changes. The index group patients and the controls showed significant differences in the average Pfirrmann DDD grades and the prevalence of Modic changes (4.4 and 2.8 for DDD grades, 70 and 5% for Modic changes; $p < 0.01$). The findings confirm the correlation of Modic changes with IVD herniation and severe DDD.

Abundance of mRNAs of cytokines and vascular endothelial growth factor isoforms in IVD tissues

Of all mRNAs of the studied genes, encoding cytokines and vascular endothelial growth factor isoforms, the TNF α , IL17, VEGF121, and VEGF165 expression in the cartilage tissue was significantly elevated ($p < 0.01$) compared to the control group (Fig. 2). Juxtaposing the mRNA expression levels of all genes, studied in the index group patients, with the IVD degeneration Pfirrmann grades revealed a positive correlation between the expression of TNF α and the IVD degeneration grades ($r = 0.301$; $p < 0.05$) (Fig. 3). Despite the fact that the expression of the IL1 β mRNA showed no significant differences with the control group, the abundance of this gene in the disc tissue correlated with the disc degeneration severity ($r = 0.37$; $p < 0.05$) (Fig. 4).

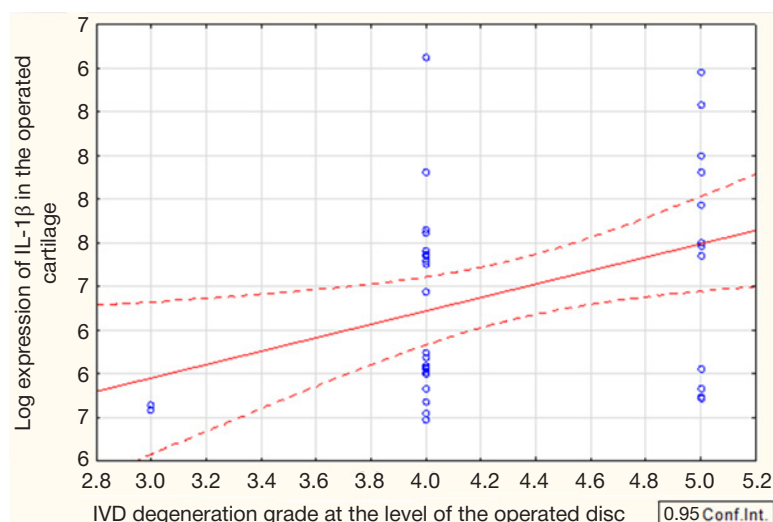


Fig. 4. Correlation between the IL1 β mRNA expression levels in the IVD tissues and the Pfirrmann degeneration grades in the index group patients

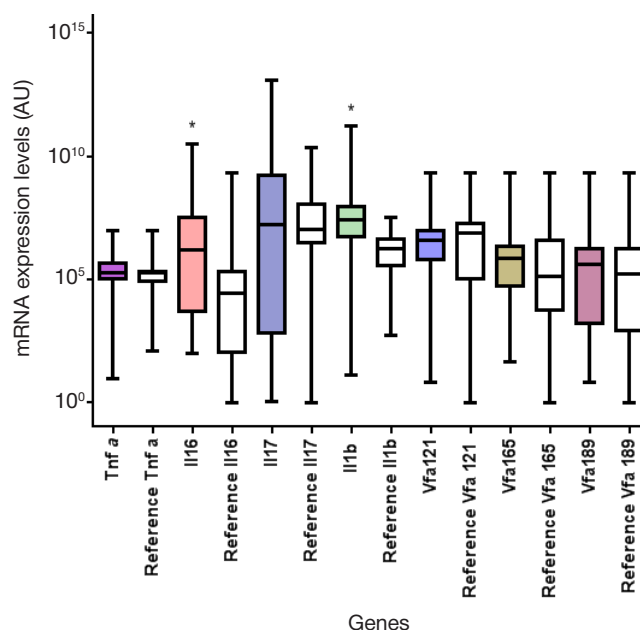


Fig. 5. Expression levels of mRNAs of the studied genes in relation to normalization genes (B2m, GUSB) in the white blood cells of the index group patients and controls ($p < 0.01$)

From the perspective of the IVD matter disorganization progression, it was interesting to assess the cartilage vascularization markers in IVDs with grade 4 or 5 degeneration. The significant negative correlation between the levels of VEGF121 and the IVD degeneration grades ($r = -0.85$; $p < 0.001$) was identified for vascular endothelial growth factors, which could indicate the decreased angiogenesis intensity in the discs of patients with the advanced stage DDD.

Analysis of gene abundance in blood cells

Further analysis was aimed to identify mRNAs of cytokines and growth factors in peripheral white blood cells. Our study revealed the significantly increased abundance of the genes, encoding interleukins IL1 β , IL6 ($p < 0.01$) (Fig. 5). Analysis of the relationship between the expression of the IL6 cytokine mRNA in peripheral blood and the IVD degeneration grade showed the decrease in the cytokine levels in the discs with the advanced IVD Pfirrmann degeneration grades ($r = -0.347$; $p < 0.05$), which was indicative of the cytokine significant contribution to degeneration at earlier stages.

Taking into account the high prevalence of Modic changes in the vertebral bodies (70%) of the index group patients, we assessed the relationship between the cytokine expression levels and the presence of Modic changes. A significant negative correlation between the IL6 levels in blood cells and Modic type 1 and 2 changes was revealed ($r = -0.31$; $p < 0.05$), which was indicative of the elevated cytokine expression at the stage of inflammatory bone marrow edema compared to the stage of fatty transformation within the adjacent vertebral bodies. Individuals with a more severe lesion in their functional spinal units, the triple combination (herniation + erosions of the vertebral body endplates + Modic) ($r = 0.401$; $p < 0.05$), had a higher expression of IL17 in their peripheral blood compared to individuals having herniation only. Regardless of the low abundance of angiogenic biomarkers in blood of patients with BP, we decided to see whether the vascular endothelial growth factor (isoforms 121, 165, 189) expression levels changed depending on the IVD degeneration grades: significant negative correlations were obtained for all three isoforms ($r = -0.44$; -0.33 , and -0.45 , respectively; $p < 0.05$).

DISCUSSION

Studying the BP pathophysiology in young patients confirmed the involvement of immune and inflammatory mechanisms in disc degeneration. High expression of TNF α and TNF α receptors, especially in the AF tissue, in patients with DDD, was also reported by other authors [17, 18]. This specific cytokine was described in experiments as an apparent inducer of matrix degradation, especially at the early stages, compared to other mediators [19]. The coupled expression of TNF α and IL17, identified during our study, was found in two studies, conducted by Chinese researchers, and confirmed by the effects of etanercept (inhibitor of TNF α receptor), which was capable of quenching the entire inflammatory cascade in the disc NP tissue [20, 21]. Pathogenetic relationships between the immune inflammation and the disc vascularization, observed during our study, were shown in the overseas human and animal studies [22, 23].

Association of the elevated IL6 gene expression in peripheral blood with Modic changes, and specifically with the inflammatory bone marrow edema grade, was also shown within the framework of the study, performed by Chinese researchers: the expression of IL6 was significantly increased in elderly patients with DDD and Modic type 1 changes compared to patients with Modic type 2 changes [24]. The assessment of changes confirmed the fact of the reactive aseptic spondylitis in patients with severe DDD. Detection of the elevated IL17 expression in individuals with the triple combination of lesions in their functional spinal units suggests the existence of the adverse clinical and instrumental phenotype of BP with the immune marker at the systemic level.

It is known that, in accordance with the Pfirrmann grading system, grade 5 of the IVD degeneration is characterized by the dramatically decreased IVD height, hypointense and nonhomogeneous signal from NP, which corresponds to severe, almost total dehydration of NR together with the extracellular matrix disintegration, replacement of the disc central space with type I collagen, and blurring between NP and AF [25]. When distinguishing between the natural age-related disc degeneration process and the abnormal disc degeneration, we wish to emphasize the key role of inflammatory markers in the second scenario, being particularly evident and manifesting

as severe degeneration in young adults [26]. By the age of 4, blood vessels and capillaries of the disc vanish, and the disc becomes a completely avascular structure [27]. As the “low-immune” inflammation develops in the IVD tissues, neovascularization becomes the way of the immunocompetent cells delivery from the systemic blood flow to the cartilage tissue with subsequent activation of catabolic pathways [28] and resorption of the tissue fragment that has fallen out of AF. The vascular endothelial growth factor A (VEGF) is one of the major regulators of angiogenesis. VEGF plays an important role in physiological and pathological neovascularization [29]. The VEGF expression is promoted by the activity of chondrocytes, which form clusters within the NP. In grade 5 DDD, the disc might no longer have the NP along with the rest of the NP cellular content, which is likely to disrupt the angiogenesis stimulation pathway in this settings. These data are consistent with the study, which has shown that the VEGF angiogenic factor expression levels in IVD samples with mild degenerative changes are significantly higher compared to advanced degeneration grades [12]. Taking into account the low reparative capacity of the disc with grade 5 degenerative changes, the use of therapeutic bioengineering, involving the application of tissue engineering and 3D IVD scaffold fabrication with subsequent cellular therapy, is the most promising.

CONCLUSIONS

The study sheds light on the range of activated genes, which express cytokines, and shows the inflammatory profiles of those in IVD tissues and peripheral blood depending on the DDD severity and the area of the functional spinal unit lesion. The abundance of mRNAs of the studied cytokines and vascular endothelial growth factor isoforms in the IVD tissues (TNF α , IL17, VEGF121, VEGF165) was higher in individuals with BP and DDD. The vascular endothelial growth factor expression levels, reflecting the possible processes of neovascularization, dramatically decreased in patients with the terminal grade DDD, both in the IVD tissues and in peripheral blood. However, the levels of TNF α and IL1 β cytokines in the cartilage tissue positively correlated with the severity of the IVD degeneration, which was in line with the concept of immune inflammation, associated with DDD. The IL6 gene expression levels in white blood cells turned out to be increased in patients with Modic changes and were to a greater extent associated with inflammatory bone marrow edema in adjacent vertebral bodies at the level of the compromised segments, and the levels of IL17 turned out to be increased in patients with a combination of herniation, erosions of the adjacent vertebral body endplates, and Modic changes. The findings would help to identify the molecular targets and new directions for the anti-inflammatory and reparative therapy of DDD.

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