

The Effect of Secretome Hypoxia Mesenchymal Stem Cells on PDGF and IL-1b Gene Expression (Experimental Study on Wistar Rats Hyperglycemic Wound Models)



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ABSTRACT: Hyperglycemic wounds take longer to heal, and high glucose results in an increase in inflammatory cells, a decrease in the angiogenesis process, and the production of growth factors. Antibiotic treatment, surgery, or debridement can increase the risk of amputation. This study aimed to determine the effect of administering gel secretome hypoxia mesenchymal stem cells (SH-MSCs) on the expression of PDGF and IL-1b in Wistar rats with a hyperglycemic wound model. In vivo laboratory experimental research with a Post Test Only Control Group Design. The samples consisted of 30 Wistar rats, divided into 5 groups, namely healthy rats (P1), negative control (P2), positive control with gentamicin (P3), gel secretome dose of 20 μ L/rat (P4), and gel secretome dose of 40 μ L/rat (P5). MSC secretome hypoxia gel treatment for 10 days, then skin tissue samples were examined using the RTq-PCR method to analyze PDGF and IL-1b gene expression. Analysis of PDGF gene expression showed significant differences between treatment groups using the Kruskal-Wallis test with a result of 0.018 ($p < 0.05$), but there were no significant differences between groups using the Mann-Whitney test ($p < 0.05$). Analysis of IL-1b gene expression showed significant differences between treatment groups using the Kruskal Wallis test with a result of 0.001 ($p < 0.05$), various doses of secretome gel affected reducing IL-1b gene expression using the Mann Whitney test with a result of 0.004 ($p < 0, 05$). The most significant decrease was at a secretome dose of 40 μ L/mouse. MSC secretome hypoxia gel at a dose of 40 μ L/rat effectively reduces IL-1b gene expression in Wistar rats with a hyperglycemic wound model. However, various doses of MSC secretome hypoxia gel do not significantly increase PDGF gene expression.

KEYWORDS: PDGF, IL-1b, SH-MSCs, hyperglycemic wounds

I. INTRODUCTION

Poor glycemic control in the long term negatively affects wound recovery.¹ Hyperglycemic wound conditions take longer to heal.² The difficulty of wound healing is caused by high glucose in the microenvironment and various biological factors. Wounds are easily damaged, difficult to heal, and eventually develop into chronic wounds.¹ Wound healing becomes inhibited due to an increase in inflammatory cells, a decrease in the process of angiogenesis, and a decrease in growth factor production.³ Hyperglycemia is known to induce the expression of interleukin 1b (IL-1b) in several different cell types in each injury event. The IL-1b pathway can be part of a positive feedback loop that maintains inflammation in chronic wounds and contributes to impaired healing.⁴ Platelet-derived growth factor (PDGF) is a growth factor that regulates cell growth and division, PDGF promotes wound healing by strengthening chemotactic migration, mitosis, and proliferation of inflammatory cells and fibroblasts to promote the formation of granulation tissue in wounds.⁵

Treatment of hyperglycemic wounds with antibiotics, surgical or debridement, can still increase the risk of amputation with frequent recurrences.⁶ Growth factor therapy is an interesting strategy to accelerate wound closure, MSCs are multipotent stromal cells that can regenerate tissue damage.⁷ Therefore, an effective and safe adjunct therapeutic approach is needed for wounds, one of which is mesenchymal stem cells (MSC) secretome hypoxia that can polarize into anti-inflammatory cytokines in a hypoxic environment and produce various anti-inflammatory cytokines and growth factors.⁸

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Wound healing is disrupted due to complications of the disease that reduce physical activity and cause chronic wounds up to amputation.⁹ Hyperglycemia in patients with diabetes, vascular disease, and neuropathy is a condition that hinders the wound healing process.⁴ Wound conditions in hyperglycemia are linked to complications of diabetes mellitus, affecting 439 million people globally by 2030 (aged 20 to 79 years), potentially causing chronic ulcers and increasing the risk of amputation 17 to 40 times higher.¹⁰

Research shows that MSC secretome hypoxia secretes various mediators and growth factors that promote skin regeneration, namely PDGF, Hepatocyte growth factor (HGF), Vascular endothelial growth factor (VEGF), Fibroblast growth factor (FGF) and Transforming growth factor- β 1 (TGF- β 1). The hypoxia secretome has anti-inflammatory potential and has the ability to detect the presence of injury signaling molecules, migrate to the injury area and accelerate wound healing.¹¹ In Vitro research shows that MSC secretome activates the Phosphatidylinositol 3-kinase (PI3K/Akt) signaling cascade or Focal adhesion kinase mediated signaling (FAK)/ Extracellular regulated kinase (ERK1/2) and further increases the proliferation and migration ability of various skin cell types, such as fibroblasts, keratinocytes, and vascular epithelial cells thereby accelerating contraction wound, inhibition of this signaling pathway resulting in beneficial effects induced by secretions on various skin cells.¹² Other preclinical studies reported routine antioxidant and anti-inflammatory effects on wound healing in streptozotocin-induced hyperglycemic rats. Rutin increases the production of antioxidant enzymes induced by nuclear erythroid related factor 2 (NRF2), inhibits the expression of matrix metalloproteinases (MMPs) regulated by NF- κ B, and decreases the expression of vascular endothelial growth factor (VEGF), reduces oxidative stress and inflammatory response in hyperglycemic rats thereby improving wound healing and further reducing the risk of ulcer development.¹

PDGF at a concentration of 20-200 pmol accelerates the infiltration of inflammatory cells and fibroblasts, extracellular matrix deposition, and collagen formation in wounds, resulting in a rapid healing process compared to untreated models.¹³ In Vivo studies report that several factors may influence the phenotype of macrophages expressed in chronic wounds, having the potential to maintain proinflammatory environment. Macrophages in chronic wounds express high levels of proinflammatory molecules IL-1b, matrix metalloproteinase 9 (MMP-9), tumor necrosis factor-alpha (TNF- α), and decreased expression of nonhealing markers insulin-like growth factor 1 (IGF-1), transforming growth factor beta (TGF- β), and interleukin 10 (IL-10). In hyperglycemic conditions known to induce IL-1b interleukin expression in a number of different cells, the IL-1 pathway β being part of a positive feedback loop that sustains inflammation in chronic wounds and contributes to impaired healing.⁴

MSC secretome hypoxia can suppress the secretion of pro-inflammatory cytokines thereby lowering levels of IL-6, IL-1b, TNF- α , IL-4 and accelerating the improvement of wound conditions in mice.¹⁴ Research reveals MSC secretome hypoxia secretes growth factors that support skin regeneration, such as PDGF regulates cellular activity in the wound healing process, detects the presence of injured signal molecules, migrates to the injured area and accelerates wound healing.^{3,11} Various studies report the potential of the precondition MSC secretome in repairing wound tissue, enhancing regenerative properties in cell secretion. MSC secretome hypoxia as a source of tissue protective and regenerative secretions in wound healing.¹⁵ However, the potential of MSC secretome hypoxia gel in regenerating hyperglycemic wounds has never been done. Based on this background, it is necessary to conduct research to determine the effect of MSC secretome hypoxia gel on the expression of PDGF and IL-1b genes in male Wistar rat hyperglycemic wound models.

II. MATERIAL AND METHOD

Study Design and Experimental Animals

This research is a true research of laboratory experiments in vivo with the design of Post Test Only Control Group Design. The study subjects used male white rats of the Wistar strain, ranging in age from 2-3 months with a weight between 200-250gr which met the criteria of inclusion and inclusion, acclimatized for 7 days. This study used 5 treatment groups, a group of healthy mice (P1), a group of negative control rats of hyperglycemic wound model with base gel administration (P2), a positive control group of hyperglycemic wound rat models with standard therapy using gentamicin antibiotics (P3), a group of hyperglycemic wound rat model with administration of MSC secretome hypoxia gel dose of 20 μ L / rat (P4), and a group of hyperglycemic wound model with MSC gel administration secretome hypoxia dose of 40 μ L/rat (P5).

Research Materials

This research uses equipment consisting of 1 cc injection syringe, balance sheet, handsoon, gluco test easy touch, stemple mold wound 2x2 cm tweezers, Handsoon surgical scissors, sterile swab, biosafety Cabinet (BSC), Oxygen meter, beaker glass, stirring rod, micropipette, CO2 incubator, Dissecting kit, Flask 75T, Hypoxic chamber.

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Research Equipment

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MSC Isolation from Umbilical Cord

The isolation stage is carried out in biosafety cabinet class 2, with aseptic and sterile-conditioned tools. The steps are as follows: (1) Collect the Umbilical cord then place it in a sterile container containing 0.9% NaCl. (2) The Umbilical cord is placed on a petri dish, and then washed thoroughly with PBS using tweezers. (3) Cut the umbilical cord, separate it from the rat fetus, and remove the blood vessels. (4) Finally chop the umbilical cord, insert it into the 25T flask then flatten, and let stand for 3 minutes until the tissue sticks to the surface of the flask. (5) Complete media (DMEM, fungizone, penstrep, and FBS) are added slowly until the network is covered with media. (6) Incubation with an incubator at 37°C and 5% CO₂. (7) Cell growth is detected about 14 days after the culture process, then add media every 3 days by removing half of the media and completely replacing it with new media. (8) The addition of cells is repeated so that the cells reach 80% confluence.

Hypoxia and TFF Manufacturing

(1) MSC that has reached 80% confluence plus complete media up to 10 mL (2) Flask containing MSC is inserted into the hypoxic chamber. (3) Nitrogen gas is flowed through the inlet valve and the oxygen meter is placed in the sensor hole to determine the oxygen concentration. (4) Nitrogen is then added until the oxygen concentration is 5%. (5) After 24 hours, culture media will be removed and screened using TFF to obtain hypoxia MSC secretome and then will be used on P4 and P5.

Preparation Manufacturing Gel MSC Secretome Hypoxia

(1) MSC secretome hypoxia gel preparation is prepared by mixing 20 µL secretome added 20 µL in 200mg gel so as to obtain a concentration of 20 µL/rat (P4). The MSC secretome hypoxia gel preparation for (P5) was prepared by mixing 40 µL of secretome with 20 µL added in 200 mg of gel to obtain a concentration of 40 µL/rat. (2) Aseptically mix until homogeneous based on physical properties observed using a microscope.

Hyperglycemic Mouse Model Making

(1) Rats were acclimatized for 7 days, then satisfied for 8-12 hours, then anesthetized with a mixture of ketamine (60 mg/kgBW) and xylazine (20 mg/kgBW) and given intraperitoneal doses using streptozotocin 65 mg/kgBW/day dissolved in 0.1 M citrate buffer pH 4.0 for 30 days. A blood glucose test is performed on day 7 after initiation. The diagnosis of hyperglycemia is established when blood glucose levels >200 mg/dl. (2) After 30 days of Streptozotocin injection, rat hair was shaved on the back. Making wounds using a circular biopsy punch diameter of 6 mm with a depth of 2 mm.⁵⁹

Validation of Hyperglycemic Wound Examination

Microscopically, skin samples were taken from all groups to make histopathological preparations with hematoxylin-eosin (HE) staining. (1) Skin samples of mice of all groups were taken and fixed in 10% NBF solution, Rinse with 70% alcohol until free of fixative residue. (2) Skin samples were dehydrated with 70%, 80%, 90%, 96% alcohol, and absolute alcohol for 30 minutes each. (3) The sample is soaked with toluol for 1 hour until clear or transparent. (4) In the oven, the paraffin infiltration process is carried out at a temperature of 56-60°C by placing the skin sample in a mixture of toluol and paraffin with a ratio of 3: 1, 1: 1 and 1: 3 for 30 minutes. Skin samples were placed on pure paraffin I, pure paraffin II, and pure paraffin III for 30 minutes at each treatment. (5) Embedding process, a sample of pure paraffin skin is implanted into a paraffin block mold containing liquid paraffin and wait for it to harden, and then the paraffin block is cut to a thickness of 6 µm with microtomes. (6) Slices of skin tissue samples are attached to a glass object by applying Mayer's albumin and then dripped with a little distilled water and then heated using a hot plate until the slices stick firmly. (7) The histological preparation is deparafinisation by placing it on xylol for 24 hours. (8) Continue with HE staining. The xylol content is taken with filter paper, then added in alcohol 96%, 90%, 80%, 70%, 60%, 50%, 40%, 30%, and aqueous is done for 1-2 minutes. The preparation is placed on the HE dye for 5-10 seconds, then rinse under running water for 10 minutes. The next process is to soak the preparation using 30%, 40%, 50%, 60%, and 70% alcohol for 3-5 minutes. The preparation is placed in eosin for 5-10 minutes, then soaked in 70%, 80%, 90%, and 96% alcohol for 3-5 minutes, and dried with filter paper. The preparation is put into xylol for 15 minutes, then the histological preparation is given drops of Canada balsam. (9) The preparation is covered with a cover glass, given etiquette, and stored in the preparation box.⁶⁰

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Hyperglycemic Wound Care

Hyperglycemic wounds formed on the sixth day, then rats were treated for 10 days with daily administration of MSC secretome hypoxia gel, with a composition of MSC secretome hypoxia gel dose 20 $\mu\text{L}/\text{rat}$ (P4) and dose 40 $\mu\text{L}/\text{rat}$ (P5). The positive control mice (P3) received standard antibiotic therapy gentamicin, and the negative control mice received base gel treatment (P2).

RNA Isolation Process and cDNA Synthesis

(1) A 100 mg skin tissue sample is cut into small pieces and inserted in a tube containing 0.5 mL of Iso Plus RNA. (2) Skin sample pieces were smoothed with micropestle with RNase free DNA added 0.5 mL of Iso Plus RNA and then stored at room temperature for 5 minutes. Add 2 mL of chloroform and stir the solution until milky white using a vortex mixer. (3) Incubation for 2-3 minutes at room temperature and centrifugation for 15 minutes at 4°C at a speed of 15,000 rpm until three layers are obtained. (4) The topmost layer containing mRNA is transferred into the microtube and isopropanol is added in equal proportions. (5) Shake the microtube until white fibers appear, then centrifuge at 4°C at 15,000 rpm for 10 minutes. The supernatant is discarded and the remaining white pellets appear at the bottom of the tube. After drying, add 100 μL of 70% ethanol to the DEPC (Diethyl pyrocarbonate) solution flip back several times then centrifuge again at 4°C, 15,000 rpm for 5 minutes. (6) Remove the supernatant and add DEPC 30-50 μL , then as much as 5 μL of RNA sample is quantified using Nanodrop with a wavelength of 260 nm. The quantification result will be calculated to make 3000 ng. (7) The process of cDNA synthesis by mixing as much as 1 μL of RNA samples that have been calculated previously with 1 μL of OligoDT and Nuclease Free Water (NFW) primers until reaching a volume of 10 μL , then inserted in a thermal cycler for 5 minutes at a temperature of 70°C. (8) Add 5X buffer 4 μL , DEPC-Treated H_2O 5 μL , ReverTraAce 1 μL . (9) The mixture is incubated at 45-50°C for 30 min.⁶¹

Diagnosis of PDGF and IL-1b Gene Expression RTq-PCR method

(1) PDGF and IL-1b gene expression were analyzed using RTq-PCR with a mixture of 1 μL cDNA sample each, KAPA SYBR FAST qPCR Master Mix as much as 10 μL , primer forward and reverse as much as 1 μL each, and nuclease-free water up to a total of 20 μL . (2) PCR gene expression was analyzed using qRT-PCR illumine with a predenaturation temperature profile of 95°C for 2 minutes, denaturation of 95°C for 30 seconds, then aneling for 20 seconds at a temperature of 61°C with a cycle of 50 times. (3) Increased gene expression was analyzed in the ratio of improvement to gene housekeeping using EcoStudy software.⁶²

III. RESULT

Isolation of MSC Secretome from Umbilical Cord

The hypoxia MSC secretome used in this study was the result of filtration of MSC culture medium under 5% hypoxia conditions using Tangential Flow Filtration (TFF) for 24 hours then MSC was isolated from the umbilical cord of Wistar rats in a 19-day bunting state, the addition of media consisting of DMEM (Dulbeccos's modified eagle medium), fungizone, penstrep, and FBS, the results of isolation was then cultured in a plastic flap. MSC validation is performed after the 4th phase. Microscopic examination of MSC

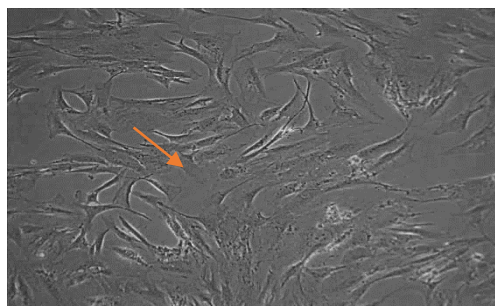


Figure 1. MSC isolation with spindle-like cells

culture results at 80% confluenc

e showed cells attached to the base of the flask with spindle-like cell shapes (Figure 1).

Examination of the capacity of MSCs to differentiate into osteogenic and adipogenic cells serves to validate the results of MSC cell isolation procedures. Oil red O staining is used in adipogenic differentiation assays to show the formation of red lipid droplets (Figure 2.a). The osteogenic differentiation assay of Alizarin red staining shows the production of calcium disposition, which is seen in red (Figure 2.b).

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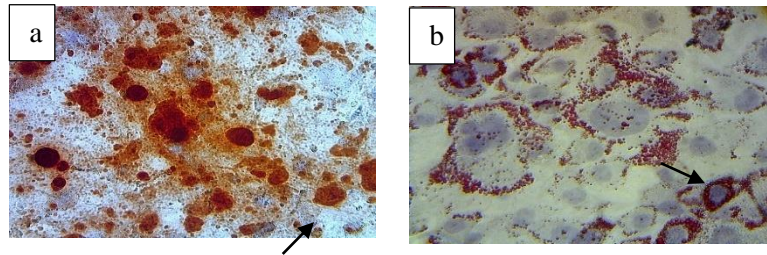


Figure 2. (a) MSCs differentiate into osteocytes and (b) MSCs differentiate into Adipocytes

Validation with flow cytometry, the results of MSC secretome hypoxia isolation show the capacity to express various special surface markers (surface markers). MSC expresses CD90 (97.6%), CD29 (96.4%) and expresses CD45 (1.7%), and CD31 (3.9%), the results of flow cytometry analysis are shown in Figure 3:

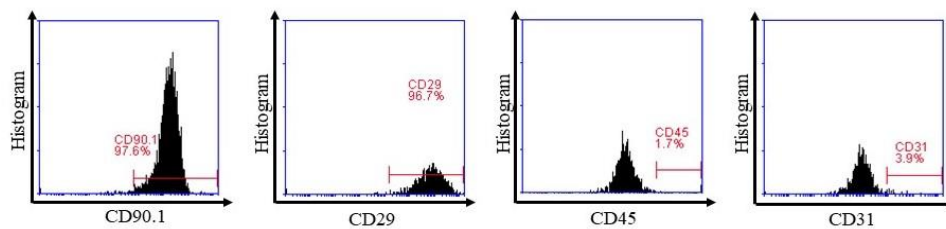


Figure 3. Flow cytometry analysis of CD90, CD29, CD45 and CD31 expression

MSC secretome hypoxia is suspected to contain anti-inflammatory cytokines and growth factors, therefore, after the filtration process is carried out analysis of the content of anti-inflammatory cytokines and growth factors, measurement of growth factor content in MSC secretome hypoxia is carried out using the ELISA test. MSC secretome hypoxia is known to produce PDGF levels of 947.26 ± 34.93 pg/mL, VEGF levels of 1137.56 ± 37.51 pg/mL, bFGF levels of 1175.54 ± 38.82 pg/mL, IL-10 levels of 525.12 ± 10.24 pg/mL, and TGF- β levels of 372.53 ± 9.18 pg/mL.

Validation of Blood Sugar Levels in Hyperglycemic Model Rats

Validation examination of glucose levels in research subjects on day 30, 36, and 46, to ensure the condition of hyperglycemia in each study subject according to the inclusion criteria for hyperglycemic wound models. The results of the glucose level examination are described in Table 1.

Table 1. Results of glucose level validation in hyperglycemic group rats

Group	P1 H0	P1 H30	P1 H36	P1 H 46	P3 46	P4	P5
Mean	105.7	419.3	388.2	382	311.8	304	294
SD	6.19	90.28	74.41	57.48	45.2	29.19	51.11

Average glucose levels in the 30th-day hyperglycemic treatment group showed a significant increase in the results of the healthy group of 105.7 ± 6.19 increased significantly to 419.3 ± 90.28 , the average level of the 36th-day hyperglycemia group obtained results of 388.2 ± 74.41 , the average levels in the 46th day hyperglycemic group obtained results of 382 ± 57.48 , compared to the gentamicin group 311.8 ± 45.2 , secretome group $20 \mu\text{L}$ 304 ± 29.2 and secretome group $40 \mu\text{L}$ 294 ± 51.1 .

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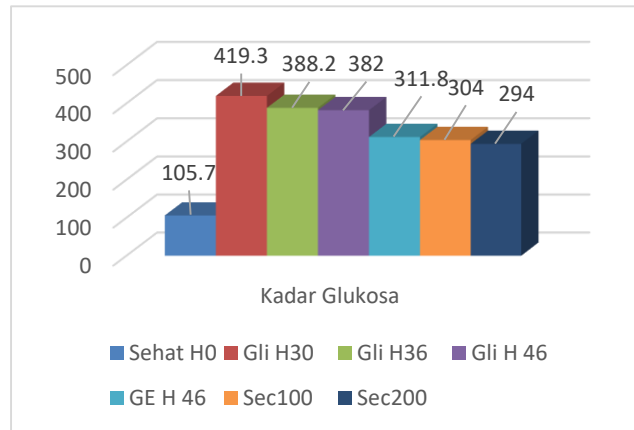


Figure 4. Results of glucose level validation in treatment group rats

The results of glucose levels showed a significant increase in the hyperglycemic group on day 30, 36, and 46 so this hyperglycemic condition can be modeled as hyperglycemic wounds.

Macroscopic Picture of Hyperglycemic Wounds

The results of macroscopic observational studies on hyperglycemic wounds on day 36, 43, and 46, showed accelerated wound healing using secretome. Using the 40 μ L/rat secretome most macroscopically significant, on day 46 wound healing was visibly camouflaged and covered with growing hair, indicating that the use of secretome is effective in accelerating wound healing, the results of macroscopic wound closure analysis are shown in Figure 5.

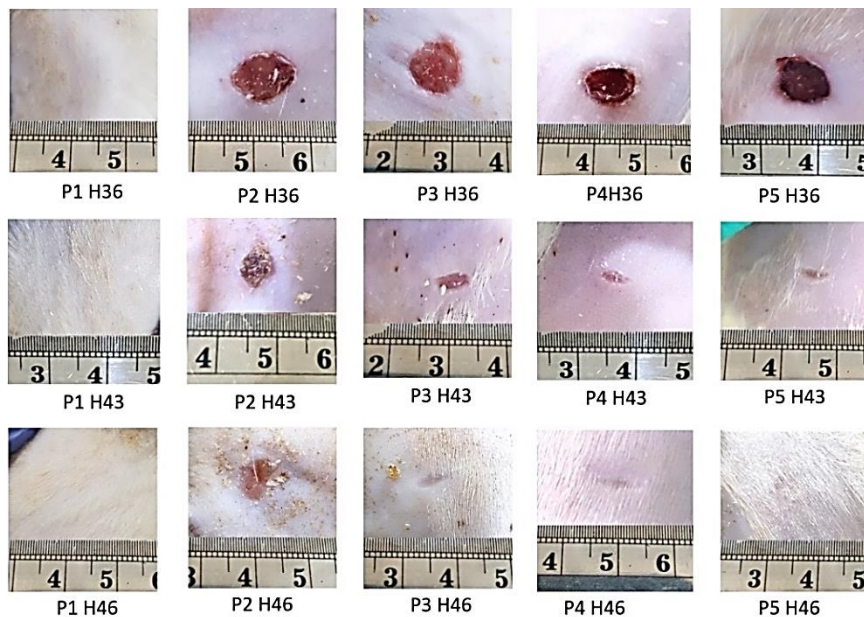


Figure 5. Macroscopic hyperglycemic wounds

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Histopathological Results of HE Staining

The readings of histopathological preparations with hematoxylin-eosin (HE) staining showed that the STZ-induced rat group of 65 mg / kgBW had thickened in the epithelium, necrosis in epithelial cells and inflammation in immune cells compared to the group of healthy mice. The results of the analysis of HE histopathological preparations are shown in Figure 6.

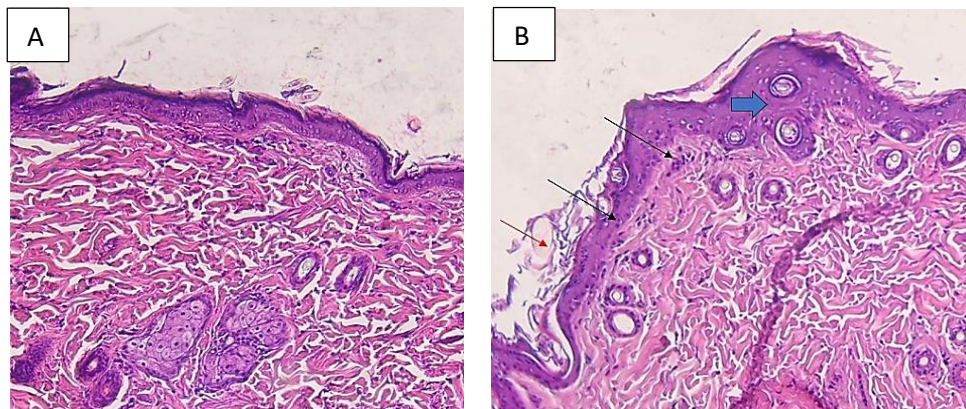


Figure 6. HE Staining Histopathology

Information

A = Group of healthy mice

B = Group of hyperglycemic rats

HE staining in the hyperglycemic group showed thickening of the epithelium (blue arrow), inflammation of immune cells (black arrow), and necrosis of epithelial cells (red arrow).

Effects of MSC Secretome Hypoxia Administration on PDGF Gene Expression

In this study, it was found that MSC secretome hypoxia was able to increase PDGF gene expression in hyperglycemic wound model mice. The results of PDGF gene expression analysis using the RTq-PCR method using the Illumina Eco Real-Time PCR tool are described in Table 2.

Table 2. Mean expression, normality and homogeneity of PDGF between treatment groups

Group	Base gel (P2)	Gentamicin (P3)	Sec 20 μL (P4)	Sec 40 μL (P5)	p value
Rat 1	0,35	0,18	0,66	0,79	
Rat 2	0,25	0,27	0,34	0,27	
Rat 3	0,40	0,12	0,26	0,77	
Rat 4	0,30	0,21	0,66	0,69	
Rat 5	0,25	0,42	0,34	0,35	
Rat 6	0,43	0,13	0,26	0,77	
Mean	0,33	0,22	0,42	0,61	
SD	0,08	0,11	0,19	0,23	
Shapiro wilk	0,42*	0,29*	0,03	0,03	
Levene test	0,007				
Kruskal Wallis	*0,018				

Information:

*Saphiro Wilk test ($p > 0.05$ = normal)

* Levene's Test ($p > 0,05$ = homogenous)

Kruskal Wallis ($p < 0.05$) = significant difference)

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Based on Table 2 above, the average results of PDGF gene expression in the base gel group (P2) were 0.33 ± 0.08 , the mean gentamicin group (P3) 0.22 ± 0.11 , the mean secretome group 20 μL (P4) 0.42 ± 0.19 and the mean secretome group 40 μL (P5) 0.61 ± 0.23 (Figure 6). The average results of PDGF gene expression were normally distributed with Shapiro Wilk test values in the P2=0.42 and P3=0.29 ($p > 0.05$ groups), while normal undistributed data were obtained in the P4=0.03 and P5=0.03 groups with normal values of $p > 0.05$ and inhomogeneous data variants with Levene's Test test results with $p = 0.007$ values with homogeneous values ($p > 0.05$). It was concluded that the data resulting from the expression of PDGF gene groups P4 and P5 were not normally distributed and were not homogeneous.

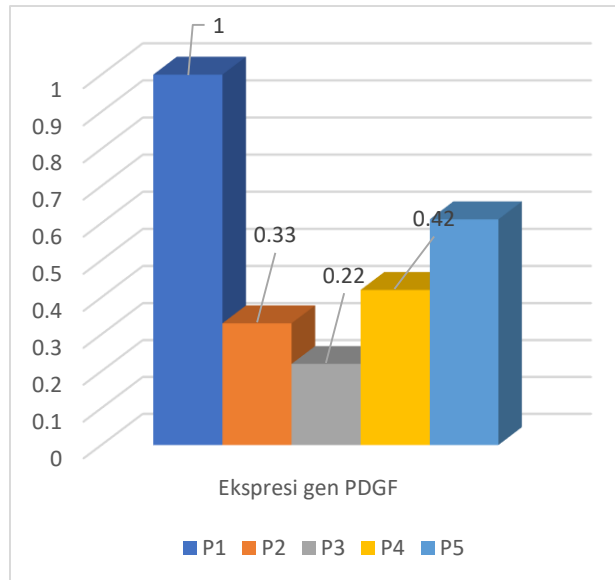


Figure 7. Average PDGF gene expression between treatment groups

Non-parametric statistical analysis using the Kruskal-Wallis test obtained results of 0.018 ($p < 0.05$), it was concluded that there were significant differences between treatment groups on PDGF expression.

Effects of MSC Secretome Hypoxia Administration on IL-1b Gene Expression

The results showed that MSC secretome hypoxia was able to reduce IL-1b gene expression in hyperglycemic wound model mice. The results of IL-1b gene expression analysis using the RTq-PCR method using Illumina's Eco Real-Time PCR tool are described in Table 3.

Table 3. Mean expression, normality, and homogeneity of IL-1b gene expression between treatment groups

Group	Base gel (P2)	Gentamicin (P3)	Sec 20 μL (P4)	Sec 40 μL (P5)	p value
Rat 1	2,14	0,26	0,25	0,34	
Rat 2	1,72	0,41	0,20	0,29	
Rat 3	3,66	0,55	0,26	0,18	
Rat 4	1,39	0,23	0,25	0,10	
Rat 5	2,14	0,41	0,20	0,14	
Rat 6	3,92	0,51	0,26	0,30	
Mean	2,49	0,39	0,24	0,22	
SD	1,04	0,13	0,03	0,09	
Shapiro wilk	0,20*	0,47*	0,17	0,44*	
Levene test	0,000				
Kruskal	*0,001				
Wallis					
Mann					
Whitney	*0,004				

Information:

*Saphiro Wilk test ($p > 0.05$ = normal)

** Levene's Test ($p > 0,05$ = homogenous)

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*Kruskal Wallis ($p < 0.05$) = significant difference)

*Mann Whitney ($p < 0,05$) = different meaning)

Based on Table 3, the average IL-1b gene expression results in the base gel group (P2) were 2.49 ± 1.04 , the average gentamicin group (P3) was 0.39 ± 0.13 , the average secretome group was 20 μL (P4) was 0.24 ± 0.03 and the average secretome group was 40 μL (P5) was 0.22 ± 0.09 .

IL-1b gene expression was highest in the base gel (P2) group and lowest in the 40 μL secretome group, showing that the 40 μL secretome was effective in reducing IL-1b gene expression in mice with hyperglycemic wound models.

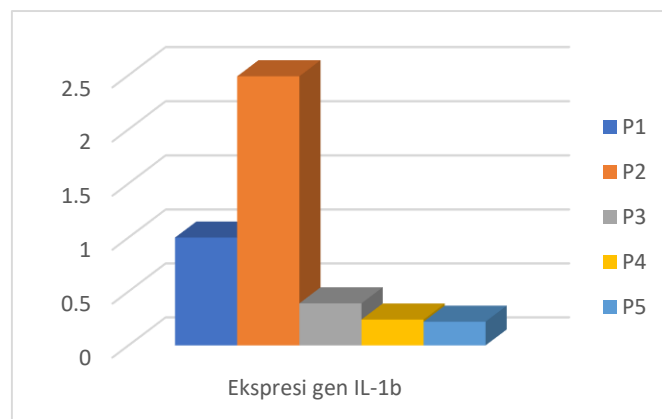


Figure 8. Mean expression, gen IL-1b gene expression between treatment groups

IL-1b gene expression data were normally distributed with Shapiro Wilk test values in the P2=0.20, P3=0.47, and P5=0.44 groups, while in the P4 group, the data were not normally distributed with values of $p=0.03$ ($p > 0.05$) and had inhomogeneous data variants with Levene's Test test results with $p = 0.000$ ($p > 0.05$). It was concluded that the distribution and variant of IL-1b gene expression data are not normally distributed and are not homogeneous. The Kruskal-Wallis non-parametric test obtained results $p= 0.001$ ($p < 0.05$), followed by the Mann-Whitney test obtained results of 0.004 ($p < 0.05$) so that it was concluded that there were significant differences between treatment groups of IL-1b gene expression in mice with hyperglycemic wound models.

IV. DISCUSSION

Inflammation is a local response to various inflammatory cells to eliminate, cleanse, build, and maintain the integrity of the tissue homeostasis system. The inflammatory process is initiated by inflammatory cells due to the stimulation of danger signal molecules produced when pathogenic bacteria damage certain tissues or substances.⁶³ Impaired wound healing due to hyperglycemia alters endothelial cell activity and causes vascular dysfunction in wounds, increasing the risk of infection up to amputation. Changes in a number of endogenous factors, including decreased growth factor production, angiogenic response mediators, macrophage activity, collagen synthesis, keratinocyte migration, and fibroblast proliferation lead to delayed wound healing.⁵⁹ Ormazabal et al, 2022 study using MSC-EC hypoxic secretome or specific protein combination enriched in MSC-EC hypoxic secretome improves the wound healing process in hyperglycemic conditions.⁵⁷ The group that received secretome treatment showed rapid wound healing and less scarring. In addition, further research using the human fibroblast secretome serves as a human protein control. When wounds are treated with stem cell secretion, epidermal and dermal thickness increases markedly faster.¹⁷

MSCs actively contribute to angiogenesis through direct differentiation, interaction of cell contact with endothelial cell derivatives, and paracrine release of pro-angiogenic molecules. Paracrine factor can increase the blood supply of damaged tissues through the activation and recruitment of stem cells and progenitor cells.⁶⁶

The results of the PDGF gene expression study in the base gel group (P2) were 0.33 ± 0.08 , showed an increase in the average gentamicin group (P3) 0.22 ± 0.11 , the average secretome group 20 μL (P4) 0.42 ± 0.19 showed an increase from the P3 group and the average secretome group of 40 μL (P5) showed the highest increase of 0.61 ± 0.23 , It can be concluded that the administration of MSC secretome hypoxia gel a dose of 40 $\mu\text{L}/\text{rat}$ was significant for increased expression of the PDGF gene in male white rats of the Wistar strain model of hyperglycemic wounds. In line with research by Chen et al, 2020 routine use can increase the production of antioxidant enzymes induced by nuclear factor erythroid 2-related factor 2 (NRF2), inhibit the expression of matrix metalloproteinases (MMPs) regulated by NF- κB , and decrease the expression of vascular endothelial growth factor (VEGF). It also promotes the expression of the associated neurogenic protein (UCH-L1) reduces oxidative stress and

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inflammatory response in hyperglycemic rats, promotes wound healing, and reduces the risk of ulcers.¹ PDGF acts as one of the growth and division factors, especially during the angiogenesis phase and the formation of blood vessels from the previous vascular network. MSC mechanism in the wound area through the process of differentiation, and vascularization by involving the release of PDGF.⁶¹

Another study by Das et al, 2016. reported a significant increase in wound healing activity with co-administration of PDGF-BB with syndecan-4 proteoliposomes (S4PL). significant translational potential in enhancing the effects of PDGF-BB for diabetic wound healing. The addition of syndecan-4 co-treatment may improve keratinocyte migration and wound closure in the context of diabetes S4PL has an immunomodulatory effect on wound macrophages, reduces inflammation, and improves phenotypic markers of M2 macrophages.⁵¹

The results of the IL-1b gene expression in the base gel group (P2) of 2.49 ± 1.04 , experienced an average decrease in the gentamicin group (P3) of 0.39 ± 0.13 , the average in the secretome group of 20 μL (P4) decreased from the P3 group to 0.24 ± 0.03 and the lowest average expression of the IL-1b gene in the secretome group of 40 μL (P5) of 0.22 ± 0.09 . Showed that administration of MSC secretome hypoxia gel dose 40 $\mu\text{L}/\text{rat}$ effectively decreased IL-1b gene expression in mice with hyperglycemic wound models. In line with research by Shiddiqi et al, 2017. IL-1 and Caspase-1 expression can be lowered in mice with pristane-induced lupus with MSC secretome administration. This is indicated by the fact that IL-1 β and Caspase-1 are expressed to a lesser extent in the pristane + secretome group compared to the pristane group.⁶⁷

The decrease in caspase-1 and IL-1b expression is caused by inhibition of NLRP3 inflammasome activation so that caspase-1 activation does not occur which causes IL-1b not to form. Inhibition of NLRP3 inflammasome activation is caused because the MSC secretome contains stanniocalcin-1 (STC-1) which also plays a role in inhibiting ROS production.⁶⁸ Administration of MSC secretome inhibits NLRP3 inflammasome by lowering ROS levels in mitochondria. ROS plays an important role in activating the NLRP3 inflammasome. NLRP3 inflammasome activates caspase-1 which converts pro IL-1b into active IL-1b which plays a role in inflammatory mechanisms.⁶⁷ Peripheral blood angiocrine MSCs can stimulate endothelial functional characteristics by activating the VEGF-A signaling pathway through a number of factors, including endothelin-1, IL-8, platelet-derived growth factor-AA (PDGF-AA), and IGF-2. MSC hypoxia expresses much higher levels of Heat Shock Protein (HSP) molecules than normoxic, capable of folding proteins that lead to optimal intracellular protein function, such as VEGF and PDGF. Hypoxic MSCs are more effective in regenerating damaged tissue than normoxic MSCs.⁶⁶

V. CONCLUSION

MSC secretome hypoxia gel administration significantly affected PDGF and IL-1b gene expression in male white rats of Wistar strain model hyperglycemic wound.

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