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**The effect of growth hormone on ovarian  
function recovery in ovarian insufficiency in a  
mouse model**

**난소부전 마우스 모델에서 성장호르몬에 의한  
난소 기능회복 효과**

February 2023

Graduate School of Medicine

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# **The effect of growth hormone on ovarian function recovery in ovarian insufficiency in a mouse model**

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October 2022

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## **Abstract**

# **The effect of growth hormone on ovarian function recovery in ovarian insufficiency in a mouse model**

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### **Objectives**

The present study was designed to evaluate the effects and the mechanisms of action of growth hormone (GH) on ovarian function recovery in ovarian insufficiency induced by the administration of cyclophosphamide (CP) in a mouse model.

### **Materials and Methods**

After inducing ovarian insufficiency by administering a single intraperitoneal injection of 400 mg/kg of CP to six-week-old ICR mice, mice were divided into four groups (control group, CP administration group, GH 1 mg/kg administration group, and GH 2 mg/kg administration group). There were 10 mice in each group. GH was administered a week later for 7 days. Five mice from each group were sacrificed the next day, and their ovaries were collected for histological examination. The remaining mice were superovulated for *in vitro* fertilization (IVF). The following experiments were conducted to explore the mechanism of action of GH on ovarian

function recovery. The terminal deoxynucleotidyl transferase dUTP-nick end labeling (TUNEL) assay was used to detect apoptosis. Masson's trichrome staining was used to analyze the degree of fibrosis. To quantify angiogenesis, CD31 immunohistochemistry was performed. Angiogenesis-related gene expression profile was assessed using quantitative reverse transcription polymerase chain reaction (RT-qPCR).

## **Results**

Administration of CP induced the loss of non-growing (primordial and primary) follicles. GH significantly protected primordial follicles and increased follicular quality. The CP group showed a decrease in fertilization rate and blastocyst formation rate in IVF. In contrast, the GH treatment group enhanced IVF outcomes in a dose-dependent manner. GH treatment also decreased apoptosis, stromal fibrosis, and increased angiogenesis. Many genes involved in angiogenesis are upregulated, especially the upregulation of leptin (*Lep*), platelet endothelial cell adhesion molecule 1 (*Pecam-1*), and angiogenin (*Ang*) have been observed in GH treatment groups.

## **Conclusion**

GH treatment may promote ovarian function recovery in ovarian insufficiency induced by administration of CP via decreasing apoptosis, stromal fibrosis, and upregulation of *Lep*, *Pecam-1*, and *Ang* genes involved in angiogenesis.

**Keywords:** Growth hormone, Ovarian insufficiency, Cyclophosphamide, Breast cancer, Ovarian regeneration, Fertility preservation

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# 1. Introduction

## 1.1. Background

Breast cancer is the most commonly diagnosed cancer in women worldwide (1), and approximately 10% of breast cancer cases occur in women younger than 45 years (2). With the improved survival rate among breast cancer patients, the demand for improving the quality of life is also increasing. Fertility preservation is one of the important issues for breast cancer survivors (3). Considering the rising trend in the postponement of marriage and childbearing, the demand for fertility preservation among breast cancer survivors is expected to increase.

Cyclophosphamide (CP) is an alkylating agent widely used as adjuvant chemotherapy for breast cancer (4). CP is ranked as a high risk of gonadotoxicity as it can cause apoptosis, overactivation of primordial follicles, and atresia of growing follicles (5). Administration of CP could induce premature ovarian insufficiency, and it causes not only infertility but cognitive dysfunction, osteopenia, and cardiovascular disease (6).

Oocyte and embryo cryopreservation are considered the standard practice of fertility preservation, but those methods cannot recover the ovarian endocrine function (7). Ovarian tissue cryopreservation is an option that could resume the ovarian endocrine function (8), but there is concern about the possibility of re-entrance of tumor cells after transplantation of cryopreserved tissue in cancer patients (9). Those cryopreservation methods are not feasible in places where assisted reproductive centers do not exist. Gonadotropin-releasing hormone agonists (GnRHa) could be offered to young women with breast cancer when proven fertility preservation

methods are not feasible. The evidence for using GnRHa to preserve function is conflicting, so it cannot replace established methods (10). Research on therapeutic agents to restore damaged ovarian function has been conducted recently. A representative therapeutic agent is a stem cell. However, the safety of stem cells has not yet been proven, and the methods for injecting stem cells are not standardized (11).

Growth hormone (GH) is a monomeric peptide hormone secreted mainly by the anterior part of the pituitary gland (12). In the classical view of GH, pituitary GH acts on its hepatic receptors to produce hepatic insulin-like growth hormone I (IGF-I). However, this view has been revised because not only the liver but also the peripheral organs, including the ovary, can produce IGF-I by GH stimulation. Moreover, it has been found that reproductive cells can produce GH itself to control the signaling pathway by paracrine or autocrine actions (13). GH plays essential roles in cell growth, development, and metabolism throughout the body. In female reproduction, GH is required for the onset of puberty, follicular development, steroidogenesis, and oogenesis in the ovary (14). Since GH is involved in the regulation of female infertility, GH as adjuvant therapy in *in vitro* fertilization (IVF) has received the most attention, especially in patients with a poor ovarian response (POR) (15). In a recent systematic review and meta-analysis, GH supplementation to POR increased clinical pregnancy rate, number of oocytes retrieved, number of MII oocytes, and live birth rate (16, 17).

GH has already been administered during IVF clinically, so it seems safer than other therapeutic agents, such as stem cell therapy; however, it is unclear what effect it has in restoring ovarian function. Only a few studies were conducted to explore the effects of GH on ovarian function recovery. Mahran et al. demonstrated that GH

showed a radioprotective effect and rescued ovarian function (18). Liu et al. showed that GH promoted ovarian tissue repair, estrogen release, and oocyte maturation via activation of the Notch-1 signaling pathway (19). Nevertheless, little information was found on the mechanisms by which GH restores ovarian function. It is necessary to evaluate the effects and the mechanisms of GH on ovarian function recovery more diversely.

## **1.2. Objectives**

The present study is designed to evaluate the effects and mechanisms of action of GH on ovarian function recovery in ovarian insufficiency induced by administration of CP in a mouse model.

## 2. Materials and Methods

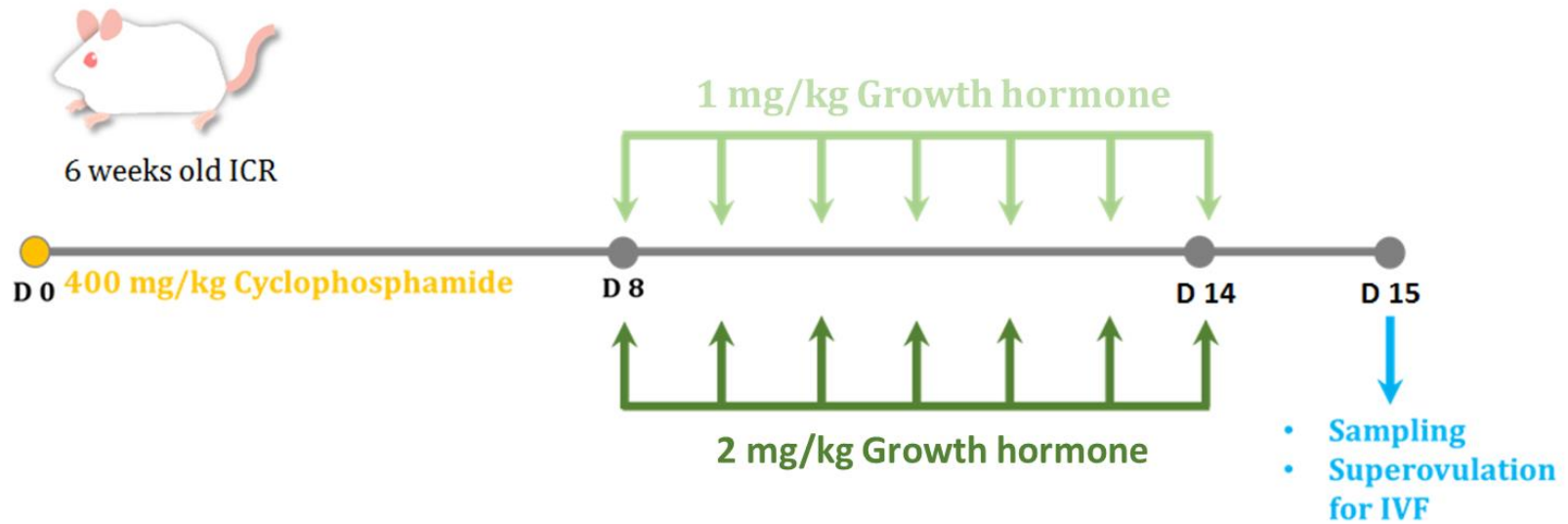
### 2.1 Animals

The experimental protocols were approved by the Institutional Animal Care and Use Committee (IACUC) of Seoul National University Bundang Hospital (BA-2009-304-087-01). Six-week-old female ICR mice (Orient Bio Inc., South Korea) were acclimated for one week before experimentation and housed in ventilated cages at room temperature, humidity, and a light-controlled environment with 12-hour light-dark cycles. They were provided food and water ad libitum.

### 2.2 Study Design

**Figure 1** shows the experimental scheme of the present study. Cyclophosphamide (CP) (Sigma-Aldrich, Burlington, MA, USA) was administered to induce ovarian insufficiency (OI). A preliminary study was conducted to determine the CP dose to be administered (**Supplementary Figures 1~3**). The mice were divided into three groups (4 mice per group) as follows: 1) control group: mice only receiving 200  $\mu$ L of 0.9% normal saline injection; 2) low-dose CP group: mice receiving 400 mg/kg of CP by a single intraperitoneal injection; 3) high-dose CP group: mice receiving 400 mg/kg of CP by an intraperitoneal injection every two days for three times. A total of 1200 mg/kg of CP was injected. Three days after the last injection, ovaries were collected to perform hematoxylin and eosin (H&E) staining, terminal deoxynucleotidyl transferase dUTP nick-end labeling (TUNEL) assay, and Masson's trichrome staining. The high-dose CP group showed a high mortality rate (50%). Furthermore, the primordial follicle damage (**Supplementary Figure 1**), apoptosis rate (**Supplementary Figure 2**), and fibrosis rate (**Supplementary Figure 3**) were

too high to be analyzed. We concluded that the low-dose CP group achieved the desired degree of ovarian damage with no harm to animal life but the deterioration of ovarian function. The mice were randomly divided into four groups (10 mice per group) as follows: 1) control group: mice only receiving 0.9% normal saline injection 200  $\mu$ L; 2) CP group: OI mice receiving 200  $\mu$ L of 0.9% normal saline injection; 3) low-dose GH treatment group: OI mice injected with 1 mg/kg GH (Growthropin, DONG-A ST, Seoul, Korea); 4) high-dose GH treatment group: OI mice injected with 2mg/kg GH. Normal saline or GH was administered intraperitoneally a week after induction of OI for 7 days. The dose (18, 19) and route of GH administration (20, 21) were determined after reviewing published studies. Five mice from each group were utilized for histological examination, and five mice per group were superovulated for IVF. Subsequently, all mice were sacrificed a day after the last injection of GH.



**Figure 1.** The experimental scheme for the present research. To induce ovarian insufficiency (OI), 400mg/kg of cyclophosphamide (CP) was administered by a single intraperitoneal injection. The mice were randomly divided into four groups: the control, CP, GH 1 mg/kg treated, and GH 2 mg/kg treated groups. Normal saline or GH was administered for a week after induction of OI for 7 days. All mice were sacrificed a day after the last injection of GH.



## 2.3 Histological analysis

Ovaries were fixed in formalin and embedded with paraffin. To count the number of follicles and evaluate follicular development and quality, serial sections at 5  $\mu\text{m}$  thickness were stained with H&E solution (Cancer Diagnostics, Durham, NC, USA) and visualized by light microscopy (Nikon, Tokyo, Japan). The total number of follicles less than 48  $\mu\text{m}$  in diameter was counted in at least eight selected non-overlapping fields. The developmental stages of follicles were classified by the following categories (22): (1) primordial follicles: a single layer of flattened granulosa cells; (2) primary follicles: a single layer of granulosa cells, one or more of which are being cuboidal; (3) secondary follicles: two or more layers of cuboidal granulosa cells lacking an antral space; (4) antral follicles: multiple layers of cuboidal granulosa cells with the antrum present. The follicular quality was assessed by morphological criteria as follows (23, 24): (1) primordial and primary follicles: grade 1, spherical with even distribution of granulosa cells; grade 2, granulosa cells pulled away from the edge of the follicle but with the oocyte still spherical; grade 3, granulosa cells with pyknotic nuclei and misshapen oocyte; (2) secondary and antral follicles: grade 1, evenly distributed granulosa and theca cells with small spaces between cells and a spherical oocyte; grade 2, intact theca cells, apparent loss of granulosa cells but with a spherical oocyte; grade 3, disrupted granulosa cells with theca cells pulled away from the edge with pyknotic nuclei and misshapen vacuolated oocyte.

## **2.4 In Vitro Fertilization**

Five mice from each group were hyperstimulated by intraperitoneal injection of 7.5 IU pregnant mare's serum gonadotropin, followed by 7.5 IU human chorionic gonadotropin after 48 hours. After 16 hours, the mice were sacrificed, and cumulus-oocytes complexes (COCs) were harvested by fallopian tube puncture. The epididymal sperm were retrieved from 10-week-old male ICR mice, and the sperm suspensions were incubated at 37°C in humidified 5% CO<sub>2</sub> in air for an hour. The metaphase II oocytes were inseminated with sperms and incubated at 37°C in humidified 5% CO<sub>2</sub> in air for 4 hours. Inseminated oocytes were washed with HTF medium (FUJIFILM Irvine Scientific, Santa Ana, CA, USA) and incubated in a fresh HTF medium. Fertilization was assessed by the formation of two cells 24 hours after insemination. Then, cleaved embryos were transferred to KSOM media (MilliporeSigma, Burlington, MA, USA). The blastocyst development was assessed 72 hours after insemination.

## **2.5 Analysis of follicular apoptosis**

Apoptosis of the ovarian follicles was analyzed using the TUNEL assay (Promega, Madison, WI, USA). Briefly, after deparaffinization and rehydration, the slides were briefly treated with proteinase K solution (Promega) for 20 minutes at room temperature, followed by incubation with a TUNEL reaction mixture for an hour at 37°C. After washing with D-PBS, the slides were mounted with Vectashield Mounting Medium with 4',6-diamidino-2-phenylindole (DAPI, ImmunoBioScience Corp., Mukilteo, WA, USA) and examined under an inverted Zeiss AX10 microscope. Apoptotic cells were visualized as green fluorescence and DAPI-stained

cells were visualized as blue fluorescence. Follicles containing more than 30% apoptotic cells were regarded as apoptotic follicles (25). Quantification of the apoptotic area was analyzed by the Image J software (National Institutes of Health, Bethesda, MD, USA).

## **2.6 Analysis of follicular fibrosis**

Masson's trichrome staining was performed to analyze follicular fibrosis using the Roche Trichrome III Blue Staining Kit (Roche, Basel, Switzerland). After deparaffinization and rehydration, the slides were treated with Bouin solution (Sigma-Aldrich) for an hour at 60°C. After washing with distilled water, the slides were stained in Weigert iron Hematoxylin solution for 10 minutes. After washing with distilled water, the slides were placed in Biebrich-Scarlet Acid fuchsin solution for 5 minutes. After washing with distilled water, the slides were placed in Aniline Blue Solution for 8 minutes. After washing with distilled water, the slides were treated with 0.5% acetic acid for a minute. The slides were dehydrated in ethanol and treated with xylene for a minute. Finally, the slides were mounted with the Mounting Medium (Dako). Stained slides were analyzed using the Image J software (National Institutes of Health).

## **2.7 Immunohistochemistry for CD 31**

The density of the blood vessel was evaluated by immunohistochemistry with anti-CD 31 antibodies. After deparaffinization and rehydration, target antigen retrieval was performed using citrate buffer. The slides were put in the microwave oven and heated for 20 minutes at 700W, followed by cooling for 30 minutes at room

temperature. The slides were placed in the peroxidase-blocking solution for 10 minutes and incubated overnight at 4 °C with anti-31 antibody (1:200 dilution, Abcam, Cambridge, UK). After washing, the slides were treated with EnVision+HRP solution (Dako) for 60 minutes and substrate-chromogen solution (1:50, Dako) for 10 minutes. The slides were counterstained with hematoxylin and dehydrated with ethanol and xylene. The CD31- positive areas were determined using the Image J software (National Institutes of Health).

## **2.8 Western blot for CD 31**

Protein samples obtained from mouse ovaries were incubated with PRO-PREP™ Protein extraction solution (iNtRON Biotechnology, Seongnam, South Korea). Twenty micrograms of proteins were separated by 10 % SDS-PAGE and transferred to a polyvinylidene difluoride membrane using a Mini Trans-Blot cell and criterion blotter (Bio-rad, Hercules, CA, USA). After blocking in 5% scheme milk in 1x TBST with 0.1% Tween 20 (Sigma-Aldrich) at room temperature for 1 hour, the blots were probed at 4°C overnight with primary antibodies 1:300 dilution of anti-CD 31 (BS-0195R; Bioss, Woburn, MA, USA) and 1:1000 dilution of anti-β-tubulin (s2128; Cell Signaling Technology, Danvers, MA, USA). Secondary incubation of a 1:2000 dilution of mouse anti-rabbit IgG (sc-2357; Santa Cruz Biotechnology, Dallas, TX, USA) was performed at room temperature for 1 hour. Band intensity was quantified using Image J and normalized by β-tubulin.

## 2.9 Quantitative reverse transcription polymerase chain reaction and data analysis

Angiogenesis-related gene expression profile was assessed using quantitative reverse transcription polymerase chain reaction (RT-qPCR). Total RNA was extracted from the ovarian tissues using a TRizol reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer's protocol. cDNA synthesis and the subsequent real-time PCR were carried out using AccuTarget™ qPCR Screening Kit (Bioneer, Daejeon, South Korea). The real-time PCR reaction contained a final volume of 20  $\mu$ L, 1  $\mu$ g reverse-transcribed total RNA, and a 2 $\times$  PCR master mix. All qRT-PCR amplification and analyses were conducted using an Accupower qPCR premix kit in an Exicycler 96 real-time quantitative thermal block (Bioneer). The PCR conditions were as follows: initial denaturation of the template DNA at 95°C for 10 minutes, followed by 40 cycles of amplification of the template DNA, in which each cycle consisted of denaturation at 95°C for 5 seconds and primer annealing and extension at 58°C for 25 seconds. All reactions were done in triplicate. The details of the analyzed genes are listed in **Table 1**. Data analysis was based on the relative quantitative method, and the  $\Delta\Delta$ CT value was used to determine the relative fold change in expression. Scatterplots of the mRNA expressions that were up or down-regulated by 1-fold compared to the control group were created. Heatmap results were generated using the R program. All the data were normalized to the reference beta-actin (ACTB) gene expression level.

## **2.10 Statistical analysis**

The data were presented as mean  $\pm$  standard error of the mean (SEM). Multiple comparisons were performed using one-way analysis of variance (ANOVA), followed by Tukey's methods as a post-hoc test by GraphPad Prism, version 8.4.0 (Graph-Pad Software Inc., San Diego, CA, USA).  $P < 0.05$  was considered statistically significant.

**Table 1.** List of genes for Real-time Polymerase Chain Reaction

<b>Gene Symbol</b>	<b>Gene ID (NCBI)</b>	<b>Gene full name</b>
Adgrb1	107831	adhesion G protein-coupled receptor B1
Akt1	11651	thymoma viral proto-oncogene 1
Ang	11727	angiogenin
Angpt1	11600	angiopoietin 1
Angpt2	26360	angiopoietin 2
Anpep	16790	Alanyl aminopeptidase, membrane
Ccl11	20292	chemokine (C-C motif) ligand 11
Ccl2	20296	chemokine (C-C motif) ligand 2
Cdh5	12562	cadherin 5
Col18a1	12822	collagen, type XVIII, alpha 1
Col4a3	12828	collagen, type IV, alpha 3
Csf3	12985	colony stimulating factor 3 (granulocyte)
Ctgf	14219	cellular communication network factor 2
Cxcl1	14825	chemokine (C-X-C motif) ligand 1
Cxcl2	20310	chemokine (C-X-C motif) ligand 2
Cxcl5	20311	chemokine (C-X-C motif) ligand 5
Edn1	13614	endothelin 1
Efna1	13636	ephrin A1
Efnb2	13642	ephrin B2
Egf	13645	epidermal growth factor
Eng	13805	endoglin
Epas1	13819	endothelial PAS domain protein 1
Ephb4	13846	Eph receptor B4
ErbB2	13866	erb-b2 receptor tyrosine kinase 2
F2	14061	coagulation factor II
F3	14066	coagulation factor III
Fgf2	14173	fibroblast growth factor 2
Fgf6	14177	fibroblast growth factor 6

Fgf1	14164	Fibroblast growth factor 1
Fgfr3	14184	fibroblast growth factor receptor 3
Figf	14205	vascular endothelial growth factor D
Flt1	14254	fms related receptor tyrosine kinase 1
Fn1	14268	fibronectin 1
Hgf	15234	hepatocyte growth factor
Hif1a	15251	hypoxia inducible factor 1, alpha subunit
Ifng	15978	interferon gamma
Igf1	16000	insulin-like growth factor 1
Il1b	16176	Interleukin 1 beta
Il6	16193	Interleukin 6
Itgav	16410	integrin alpha V
Itgb3	16416	integrin beta 3
Jag1	16449	jagged 1
Kdr	16542	kinase insert domain protein receptor
Lect1	16840	chondromodulin
Lep	16846	leptin
Mapk14	26416	mitogen-activated protein kinase 14
Mdk	17242	midkine
Mmp14	17387	matrix metalloproteinase 14 (membrane-inserted)
Mmp19	58223	matrix metalloproteinase 19
Mmp2	17390	matrix metalloproteinase 2
Mmp9	17395	matrix metalloproteinase 9
Nos3	18127	nitric oxide synthase 3, endothelial cell
Nrp1	18186	neuropilin 1
Nrp2	18187	neuropilin 2
Pdgfa	18590	platelet derived growth factor, alpha
Pecam1	18613	platelet/endothelial cell adhesion molecule 1
Pgf	18654	Placenta growth factor
Plau	18792	plasminogen activator, urokinase
Plg	18815	plasminogen
Ptgs1	19224	prostaglandin-endoperoxide synthase 1



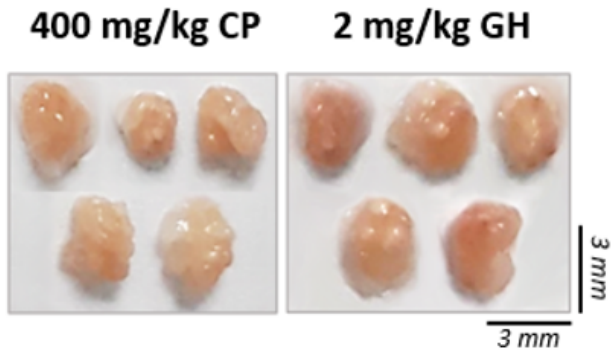
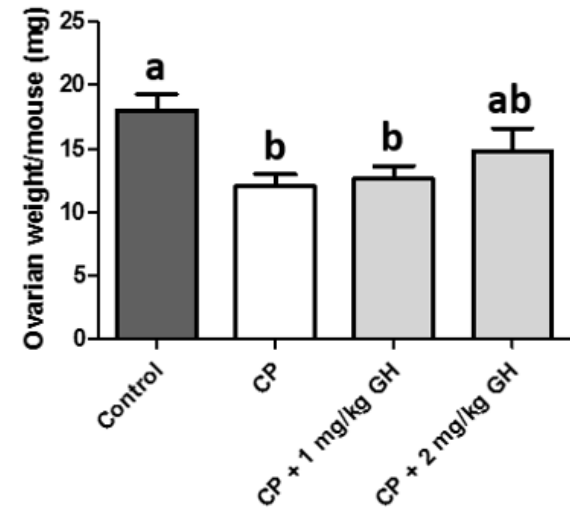
Ptk2	14083	protein tyrosine kinase 2
S1pr1	13609	sphingosine-1-phosphate receptor 1
Serpine1	18787	serine (or cysteine) peptidase inhibitor, clade E, member 1
Serpinf1	20317	serine (or cysteine) peptidase inhibitor, clade F, member 1
Smad5	17129	SMAD family member 5
Sphk1	20698	Sphingosine kinase 1
Tbx1	21380	T-box 1
Tek	21687	TEK receptor tyrosine kinase
Tgfa	21802	transforming growth factor alpha
Tgfb1	21803	transforming growth factor, beta 1
Tgfb2	21808	transforming growth factor, beta 2
Tgfb3	21809	transforming growth factor, beta 3
Tgfb1	21812	transforming growth factor, beta receptor I
Thbs1	21825	thrombospondin 1
Thbs2	21826	thrombospondin 2
Tie1	21846	tyrosine kinase with immunoglobulin-like and EGF-like domains 1
Timp1	21857	tissue inhibitor of metalloproteinase 1
Timp2	21858	tissue inhibitor of metalloproteinase 2
Tnf	21926	Tumor necrosis factor
Tnfsf12	21944	tumor necrosis factor (ligand) superfamily, member 12
Tymp	72962	thymidine phosphorylase
Vegfa	22339	vascular endothelial growth factor A
Vegfb	22340	vascular endothelial growth factor B
Vegfc	22341	vascular endothelial growth factor C

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## 3. Results

### 3.1 Gross observation and ovarian weight changes

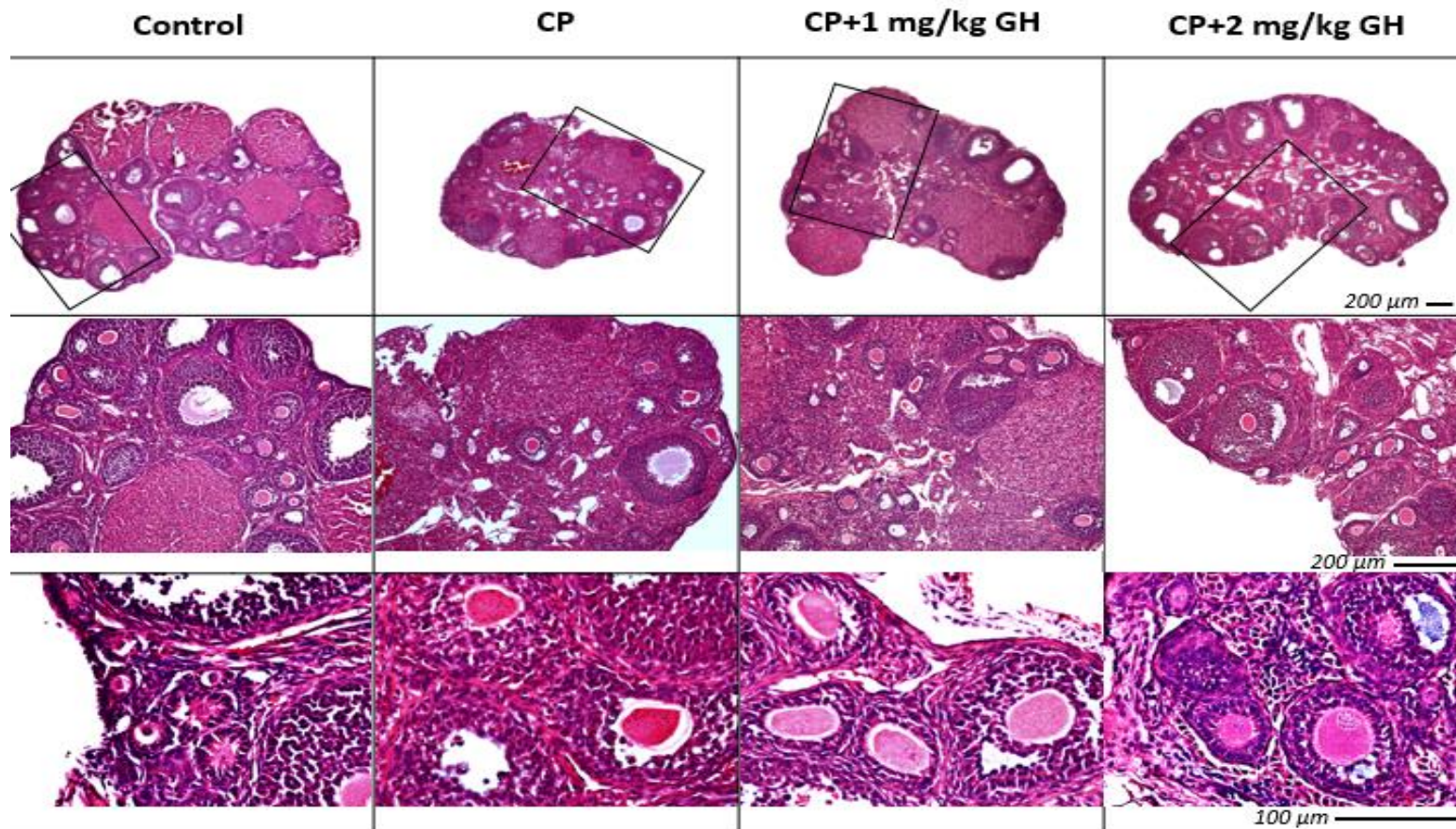
A day after the GH injection, ovaries were harvested, and the weight of the ovaries was determined. When grossly observed, the size of ovaries appeared to be increased in the GH-treated group when compared to the CP group (**Figure 2A**). As indicated in **Figure 2B**, ovarian weight was significantly decreased after CP administration. In the GH-treated group, ovarian weight increased but did not reach statistical significance.

**A****B**

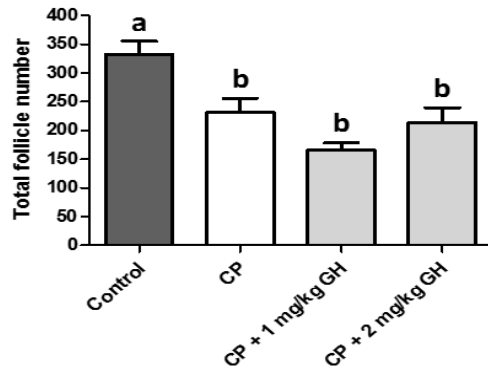
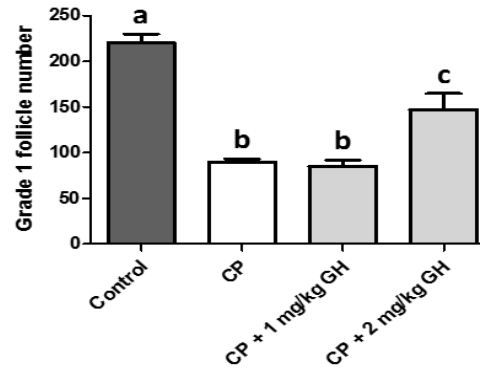
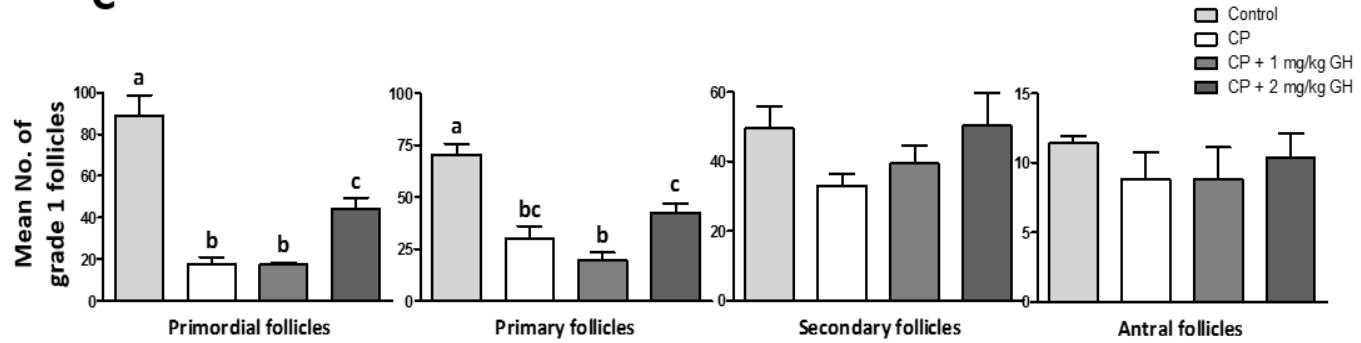
**Figure 2.** (A) The gross appearance of the ovaries. (B) The average ovarian weight. Graphs are presented as mean  $\pm$  SEM, and different superscript letters indicate statistically significant differences ( $p < 0.05$ ). CP: cyclophosphamide; GH: growth hormone.

### 3.2 Evaluation of ovarian follicle morphology

The number and structural quality of follicles were evaluated using H&E stained ovaries (**Figure 3**). The stromal density and normality of ovarian follicles were decreased in the CP group. However, the density of stromal cells in GH-treated groups appeared higher than that in the CP group, and relatively high-graded follicles were observed in the GH-treated groups. The total number of follicles and grade 1 (G1) follicles decreased in the CP and GH-treated groups compared to the control group (**Figure 4A, B**). A statistical significance was not observed in the total number of follicles between the CP and GH-treated groups (**Figure 4A**), but the number of G1 follicles increased significantly in the GH 2 mg/kg treated group than in the CP group ( $p < 0.05$ ) (**Figure 4B**). **Figure 4C** represents the mean G1 follicle number according to the follicular development stages. The primordial and primary follicles were dramatically decreased after CP administration ( $p < 0.05$ ), indicating that CP induces non-growing follicle loss. The proportion of G1 primordial follicles was significantly increased in the GH 2 mg/kg treated group compared to the CP group. The G1 follicle ratio was increased in the primary and secondary follicles in the GH-treated groups compared to the CP group, but with no statistical significance. Thus, GH treatment prevented primordial follicle loss induced by CP administration and improved the follicular quality.



**Figure 3.** Representative images of hematoxylin and eosin-stained ovarian tissue. CP: cyclophosphamide; GH: growth hormone.

**A****B****C**

**Figure 4.** Histological analysis of ovaries. (A) Total follicle number (B) Grade 1 follicle number, and (C) Mean numbers of grade 1 follicles according to the follicular development stages of ovarian tissue. Graphs are presented as mean  $\pm$  SEM; n=5 and different superscript letters indicate statistically significant differences ( $p < 0.05$ ). CP: cyclophosphamide; GH: growth hormone.

### **3.3 *In Vitro* Fertilization (IVF)**

**Table 2** shows data for IVF. MII oocyte rate was 88.9% in the control group but declined to 78.8% in the CP group. It increased to 81.1% and 89.4% in the GH 1 mg/kg and GH 2 mg/kg treated groups, respectively. After insemination, the fertilization rate (81.7%) and blastocyst formation rate (26.9%) also decreased in the CP group, whereas those values were 88.3% and 81.3% in the control group. The fertilization and blastocyst formation rates were increased in the GH-treated groups: 91.0% and 40.5% in the GH 1 mg/kg treated group and 92.3% and 50.0% in the GH 2 mg/kg treated group. The blastocyst formation rate was far more decreased in the CP group, but it increased in the GH-treated group in a dose-dependent manner.



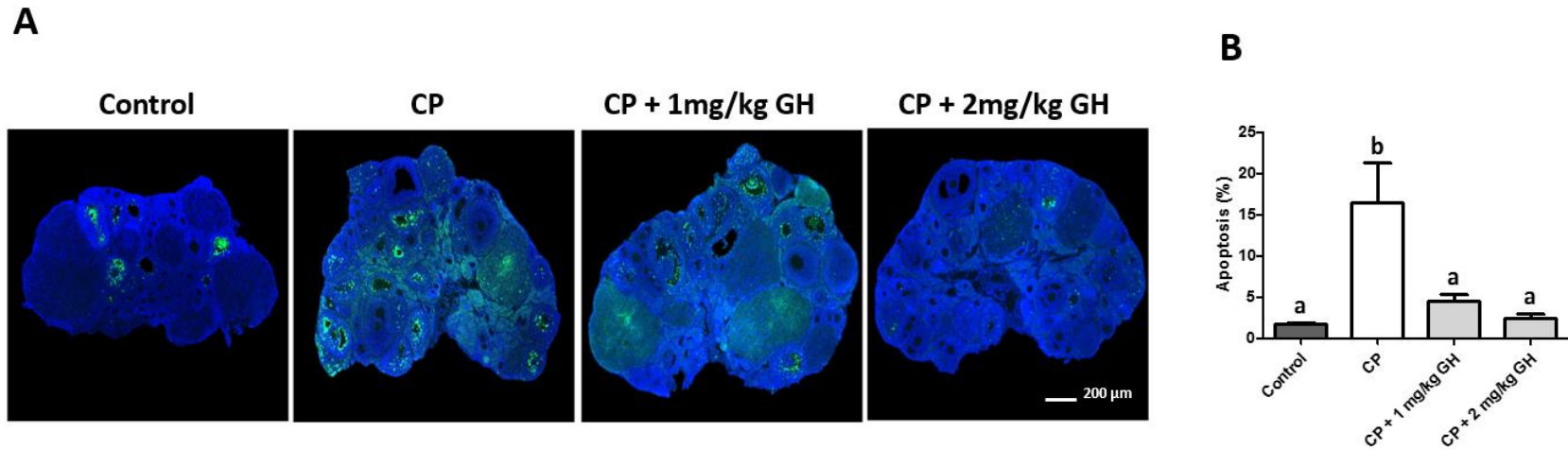
**Table 2. IVF outcomes with the control group, CP group, and GH-treated groups.**

Group	No. of mouse	No. of normal oocyte	Mean No. of normal oocyte	No. of MII oocyte	Maturation rate (%)	Fertilization rate (%)	Blastocyst formation rate (%)
Control	5	135	27	120	88.9	88.3	81.3
CP	5	104	20.8	82	78.8	81.7	26.9
CP+GH 1mg/kg	5	164	32.8	133	81.1	91.0	40.5
CP+ GH 2mg/kg	5	160	32	143	89.4	92.3	50.0

CP: cyclophosphamide; GH: growth hormone.

### **3.4 Evaluation of apoptosis**

**Figure 5A** shows a representative image of TUNEL-positive areas in the ovary of each group. Apoptosis was significantly increased in the CP groups and remarkably decreased in the GH-treated group in a dose-dependent manner (**Figure 5B**).

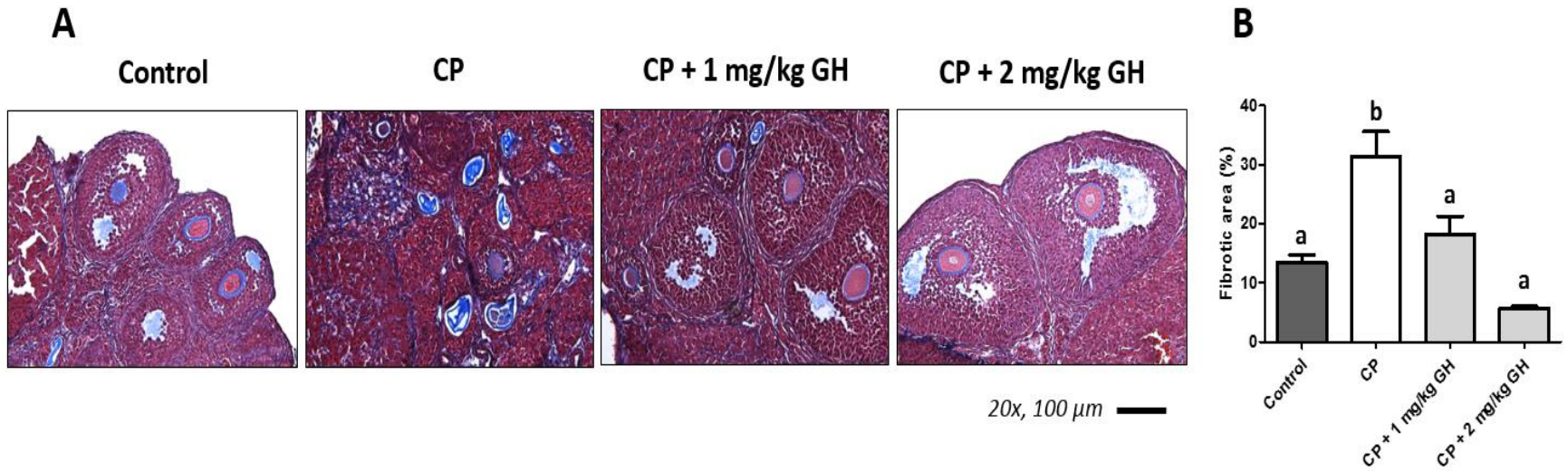


**Figure 5.** Terminal deoxynucleotidyl transferase dUTP nick-end labeling (TUNEL) assay. (A) Green fluorescence indicates apoptotic cells and blue fluorescence (DAPI) indicates intact cell nuclei. (B) Quantification of the TUNEL positive area. Graphs are presented as mean  $\pm$  SEM;  $n=5$  and different superscript letters indicate statistically significant differences ( $p < 0.05$ ). DAPI: 4',6-diamidino-2-phenylindole CP: cyclophosphamide; GH: growth hormone.

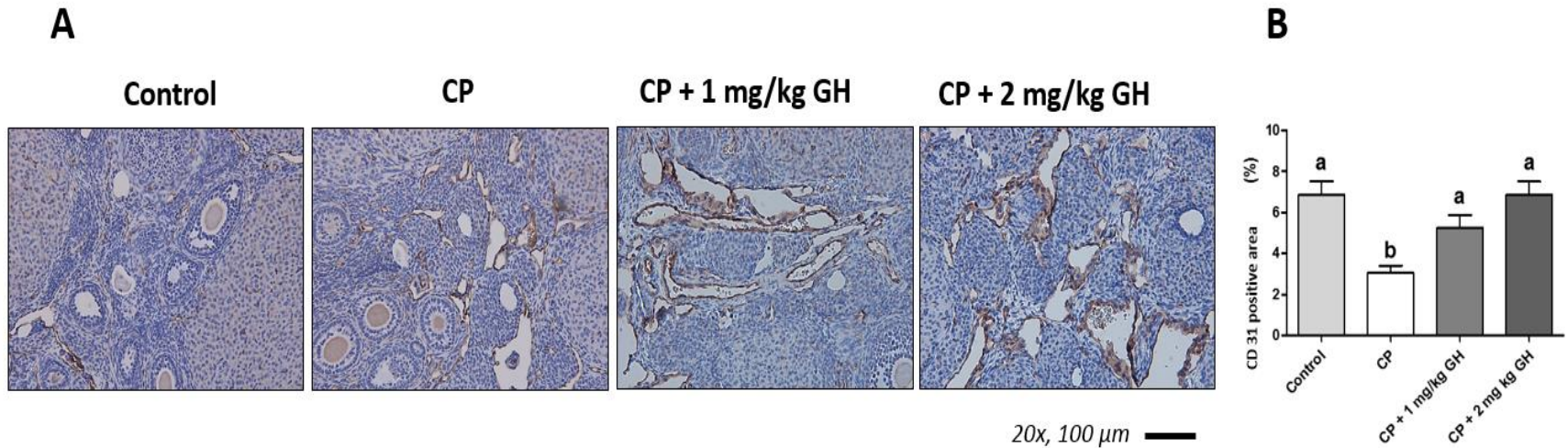
### **3.5 Evaluation of fibrosis and angiogenesis**

**Figure 6A** shows the representative image of Masson's trichrome stain. This staining is used to detect collagen fibers and is widely used to measure tissue fibrosis (26). The CP group shows an extensive fibrotic area, and its proportion was significantly reduced in the GH-treated groups (**Figure 6B**).

A representative figure of CD-31 immunostained ovaries is given in Figure 7A. CD31 is expressed on the surface of endothelial cells and is well established as the maker of vessel density (27). The CD-31 positive area was significantly decreased in the CP group compared to the control group (**Figure 7B**). The CD-31 positive area was increased in the GH-treated group with significant differences compared to the CP group, and there were no significant differences from that of the control group, indicating restoration of blood vessel density as much as the control group (**Figure 7B**).



**Figure 6.** Masson's trichrome stain. (A) The fibrotic surface, nuclei, and cytoplasm were stained blue, black, and red, respectively. (B) Quantification of the fibrotic area. Graphs are presented as mean  $\pm$  SEM;  $n=5$  and different superscript letters indicate statistically significant differences ( $p < 0.05$ ). CP: cyclophosphamide; GH: growth hormone.

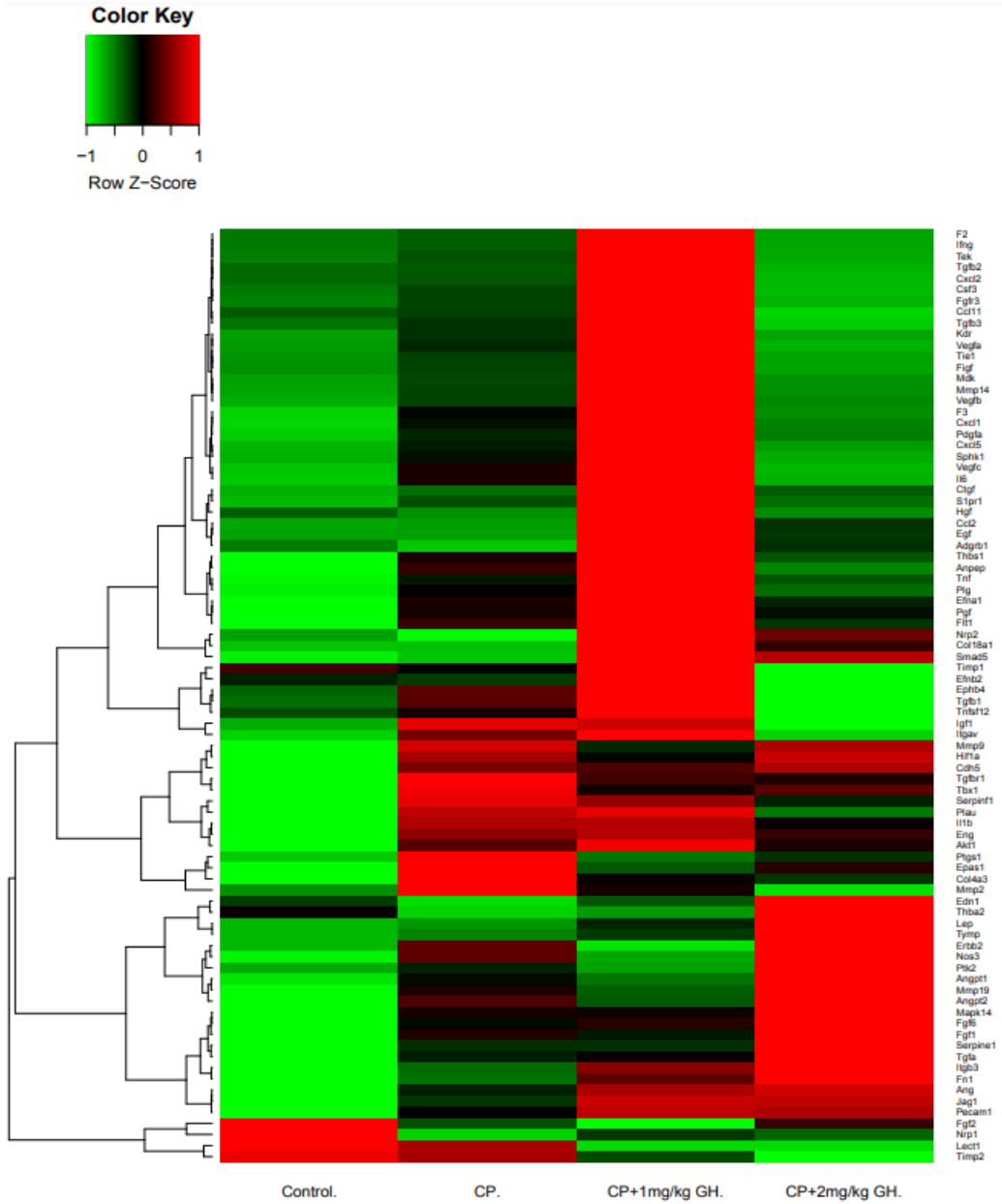


**Figure 7.** Immunohistochemical staining of ovarian tissue with CD 31. (A) Brown-stained cells are CD 31-positive cells. (B) Quantification of CD 31-positive area in ovarian tissues. Graphs are presented as mean  $\pm$  SEM; n=5 and different superscript letters indicate statistically significant differences ( $p < 0.05$ ). CP: cyclophosphamide; GH: growth hormone.

### 3.6 Angiogenesis-related gene expression

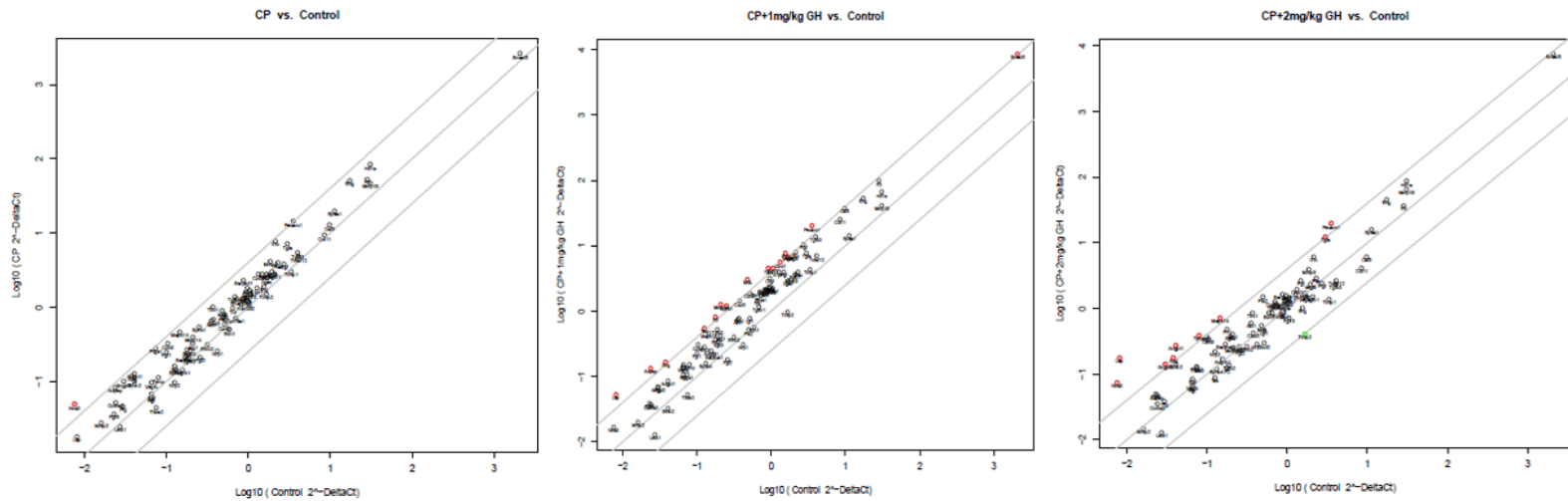
**Figure 8** shows the expression of 84 angiogenesis-related genes compared to the control group. **Table 3** demonstrates up-regulated genes with a > 4-fold change. In the GH-treated groups, three genes were up-regulated with a > 4-fold change, as shown in **Table 4** and **Figure 9** [leptin (*Lep*) (**Figure 9A**), platelet endothelial cell adhesion molecule 1 (*Pecam-1*) (**Figure 9B**), and angiogenin (*Ang*) (**Figure 9C**)].

# A





**B**



**Figure 8.** Angiogenesis-related gene expression by quantitative reverse transcription polymerase chain reaction (RT-qPCR). (A) Heatmap image, in which up-regulated and down-regulated genes are depicted in green and red colors, respectively. (B) Scatter plot. The red dots indicate up-regulation, and the green dots indicate down-regulation. The scatter graph depicts a log transformation plot of the relative expression level of each gene (2-Delta Ct) between control (x-axis) and CP (y-axis) groups. CP: cyclophosphamide; GH: growth hormone.

**Table 3.** Up-regulated genes with a fold change value > 4.0

<b>Gene</b>	<b>Description</b>	<b>Fold Change</b>
<b>CP vs. Control</b>		
Nos3	Nitric oxide synthase 3	6.42
<b>CP+1mg/kg GH vs. Control</b>		
Lep	Leptin	6.52
Mdk	Midkine	6.39
Mmp14	Matrix metalloproteinase 14	5.98
Anpep	Alanyl (membrane) aminopeptidase	5.64
Pecam1	Platelet endothelial cell adhesion molecule 1	5.61
Ifng	Interferon Gamma	4.91
Sphk1	Sphingosine kinase 1	4.81
Tie1	Tyrosine kinase with immunoglobulin-like and EGF-like domain 1	4.81
Ang	Angiogenin	4.35
F2	Coagulation factor ii (thrombin)	4.56
Tek	Tyrosin-protein kinase	4.29
Fgfr3	Fibroblast growth factor receptor 3	4.21
Cxcl1	Chemokine (c-x-c motif) ligand 1	4.18
Vegfa	Vascular endothelial growth factor-A	4.08
<b>CP+2mg/kg GH vs. Control</b>		
Lep	Leptin	21.78
Nos3	Nitric oxide synthase 3	9.93
Angpt1	Angiopoietin-1	6.71
Pecam1	Platelet endothelial cell adhesion molecule 1	5.44
Tymp	Thymidine phosphorylase	4.94
Mapk14	Mitogen-activated protein kinase 14	4.82
Angpt2	Angiopoietin-2	4.63
Ang	Angiogenin	4.61
Tgfa	Transforming growth factor-alpha	4.12

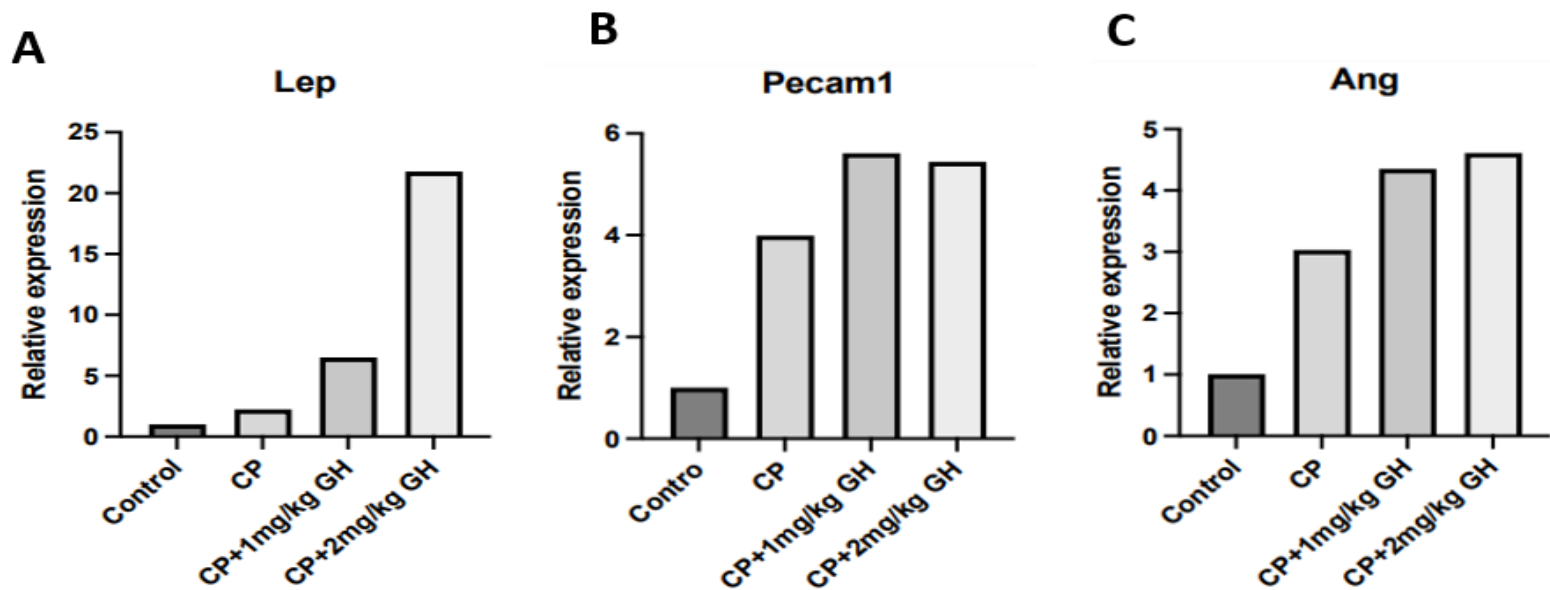
CP: cyclophosphamide; GH: Growth hormone.

**Table 4.** Up-regulated genes with a fold change value > 4.0 in the GH-treated group

	Control	CP	CP+1mg/kg GH	CP+2mg/kg GH
Lep	1.00	2.24	6.52	21.78
Pecam1	1.00	3.99	5.61	5.44
Ang	1.00	3.03	4.35	4.61

CP: cyclophosphamide; GH: Growth hormone; Lep: Leptin;

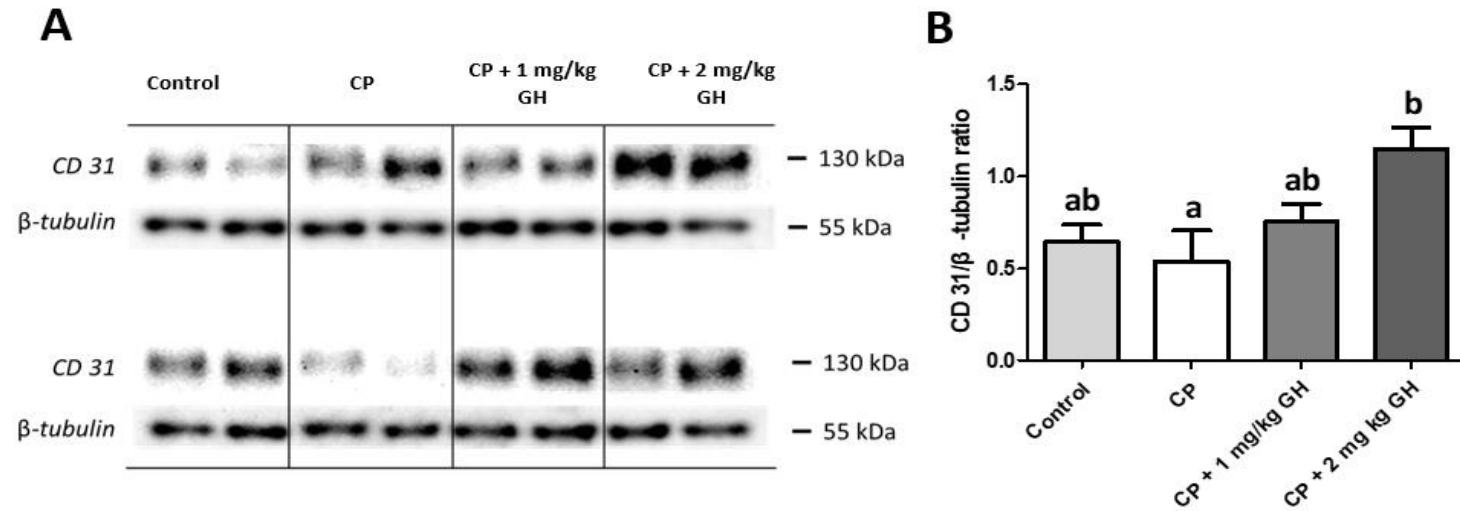
Pecam1: Platelet endothelial cell adhesion molecule; Ang: Angiogenin



**Figure 9.** Bar graphs of up-regulated genes with a >4-fold change in the GH-treated group. The fold changes represent gene expressions compared to those of the control group. (A) Leptin (*Lep*) (B) Platelet endothelial cell adhesion molecule 1 (*Pecam1*) (C) Angiogenin (*Ang*) CP: cyclophosphamide; GH: growth hormone.

### **3.7 Western blot of CD 31**

CD 31, previously a marker for determining vessel density, is a protein encoded by the *Pecam-1* gene. Western blot revealed that CD 31 was significantly increased in the GH 2 mg/kg treated group compared to the CP group (**Figure 10A, 10B**).



**Figure 10.** Evaluation of CD31 protein expression by western blot (A) Representative western blot image of CD31 (B) Relative quantification of CD31. Graphs are presented as mean  $\pm$  SEM; n=5 and different superscript letters indicate statistically significant differences ( $p < 0.05$ ). CP: cyclophosphamide; GH: growth hormone.

## 4. Discussion

This study showed the effects of GH on ovarian function recovery in ovarian insufficiency induced by the administration of CP. **Figure 11** shows the schematic results of the present study. CP damaged the non-growing follicles, such as the primordial and primary follicles. GH recovered ovarian function by decreasing apoptosis and fibrosis. GH also enhanced angiogenesis by up-regulation of Leptin (*Lep*), Platelet endothelial cell adhesion molecule 1 (*Pecam-1*), and Angiogenin (*Ang*). As a result, primordial and grade I follicles were increased. Furthermore, the fertilization rate and blastocyst formation rate were increased in IVF.

Anticancer treatment can impact ovarian reserve so that it could induce ovarian insufficiency. Ovarian insufficiency increases the risk of infertility, bone loss, and cardiovascular disease. Beyond these medical comorbidities, ovarian insufficiency impairs self-esteem and causes emotional distress (28). The highest risk of ovarian insufficiency in breast cancer is associated with the use of CP and its dose (29). CP-based chemotherapy significantly decreases the anti-mullerian hormone (AMH) level and causes treatment-related amenorrhea (3). Various strategies are being developed to enhance ovarian function to overcome these medical hurdles. Thus, this present study investigated the potential of GH as one of the therapeutic agents capable of ovarian function regeneration.

GH is a 191 amino acid protein that binds to the growth hormone receptor (GHR). Human oocytes, granulosa, and stromal cells express GHR and can be directly influenced by GH (30).

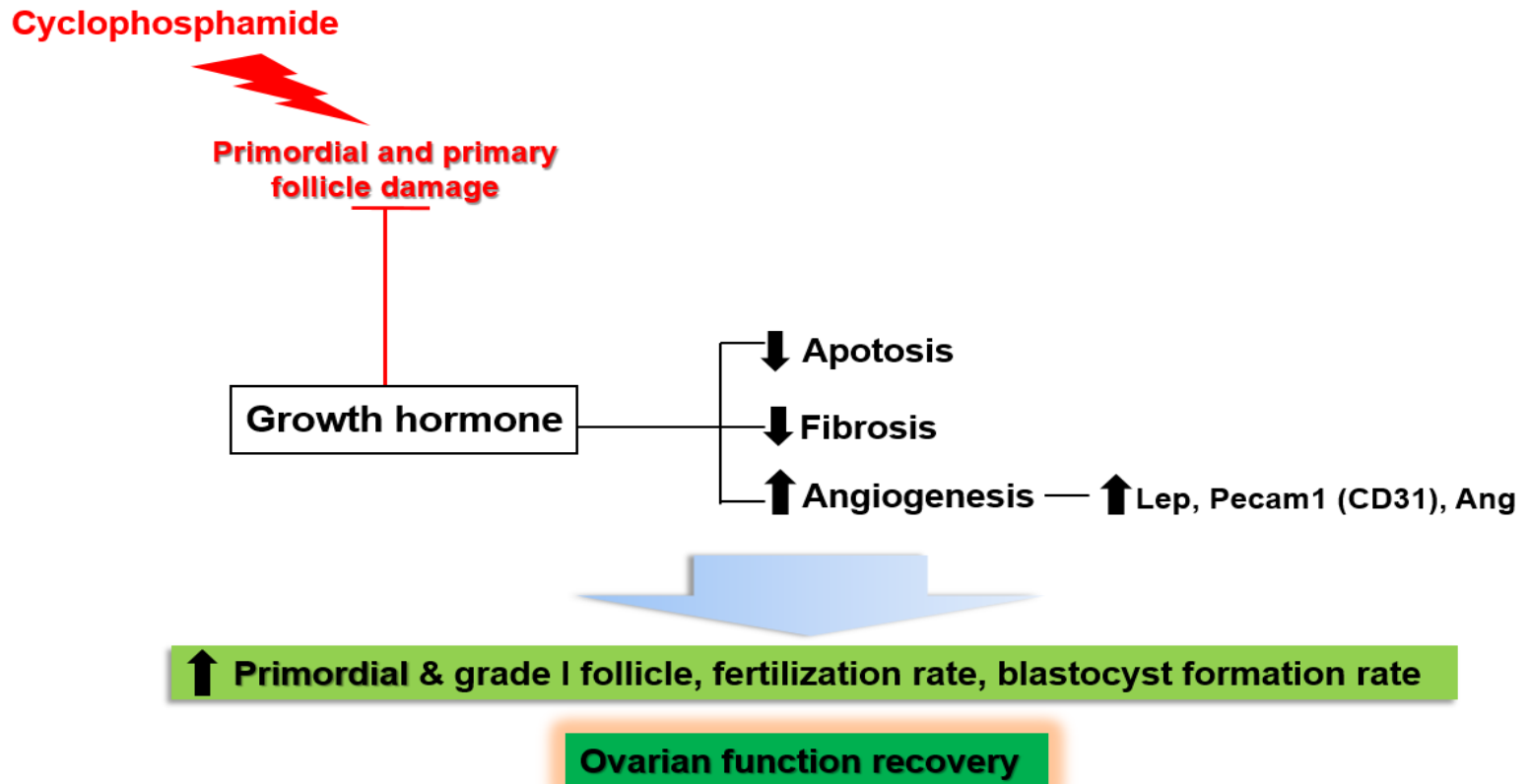


Figure 11. Schematic results of the current study.



GH indirectly affects ovarian function through autocrine or paracrine signaling mediated by the local production of secondary factors, particularly IGF-I (14). Through these direct and indirect pathways, GH influences folliculogenesis, oocyte maturation, and steroid synthesis (13). GH plays an essential role in the proliferation of ovarian follicles and is necessary for follicular maturation and survival (12). Locally released GH has a major role affecting ovarian function, but systemic GH produced by the anterior pituitary gland or exogenous administration also has an important role in female reproduction (13). GH replacement treatment recovered ovarian function with successful pregnancies in infertile patients with GH deficiency (31). This present study demonstrated that GH administration inhibited primordial follicle atresia and increased grade I follicles. The IVF outcome was also improved. Although statistical significance was not observed, the total number of follicles and grade 1 follicles were not increased as we expected in the GH 1 mg/kg treated group compared to the CP group. We speculated that these results could be explained by the small sample size with variation between objects. Moreover, these results suggest that high-dose GH might be needed to restore ovarian function.

The mechanism by which GH restores ovarian function, suggested by this study, is that GH reduces apoptosis and stromal fibrosis and enhances angiogenesis. Mahran et al. evaluated the radioprotective effects of GH. They injected GH 1 mg/kg in rats for seven days, starting three days before irradiation and lasting three days post-irradiation. GH significantly enhanced follicular development and ameliorated oxidative stress-mediated apoptosis by lowering cytochrome c and caspase-3 (18). Liu et al. injected GH 1 mg/kg in aged mice (8 months old) every two days for two months. GH reversed the age-associated depletion of ovarian reserve and the decline of oocyte quality. GH also reduced apoptosis by inhibiting the activation of Fos and

Jun signaling (32). Although those experiments were not in the same setting as the present study, GH may ameliorate apoptosis through various pathways.

To our knowledge, no study has explored the anti-fibrosis effects of growth hormones against ovarian fibrosis. Fibrosis is characterized by excessive proliferation of fibroblasts and deposition of extracellular matrix (ECM). Fibrosis develops in response to repeated tissue injury and inflammation (33). Ovarian fibrosis could be a reason for diminished ovarian function (34). Meiorow et al. conducted a case-control study to evaluate the ovarian tissue harvested for cryopreservation (35). The case group (n=17) comprised young cancer patients previously administered chemotherapy, including CP. The control group (n=18) was not exposed to chemotherapy. Over 75% of study group patients showed ovarian cortical fibrosis with the disappearance of primordial follicles. Meiorow et al. also found vascular damage showing severe narrowing and obliteration due to intimal fibrosis and hyalinization. These findings were consistent with this present study, which markedly increased fibrosis and decreased vessel density in the CP group. Meiorow et al. suggested a mechanism of ovarian damage by chemotherapy as follows: exposure to chemotherapy caused injury and obstruction of blood vessels and ischemia of ovarian tissue. Finally, intact ovarian tissue was replaced with collagenous fibrosis lacking primordial follicles. In this study, fibrosis was significantly reduced in the GH-treated groups, hence we could speculate that ovarian fibrosis is reversible by GH and it would be a therapeutic strategy to enhance ovarian function.

As previously mentioned, we identified that GH promoted angiogenesis and decreased fibrosis in the damaged ovary. We focused on the angiogenesis effects of GH because, in any way, the supply of GH to the targeted organ needs adequate

vascular flow. Angiogenesis, the formation of new blood vessels, plays a pivotal role in maintaining normal ovarian function (36). Adequate blood supply to the ovaries is essential for receiving nutrients and delivering hormones and many factors. The female reproductive organs, specifically ovarian follicles, must experience a regulated angiogenesis process emerging dominant follicles and early corpus luteum in each menstrual cycle until menopause.

To understand the mechanism of GH on angiogenesis, we analyzed gene expression by RT- qPCR. We confirmed that GH increased in the expression of angiogenesis-related genes, including *Lep*, *Pecam-1*, and *Ang*.

*Lep* is an adipocyte-derived peptide hormone that regulates food intake, body mass, and reproductive functions, such as puberty (37). Capillary endothelial cells crosstalk with adipocytes and extracellular components via paracrine pathways and direct cell-to-cell interactions (38). Surprisingly, *Lep* is also regarded as a potent angiogenic factor. Endothelial cells express the *Lep* receptor and *Lep* promotes capillary-like tube formation in vitro. Furthermore, *Lep* increases corneal neovascularization in rats (39). In addition, ovarian follicular cells express *Lep* receptors. *Lep* promotes follicular development and oocyte maturation in synergistic effects with GH and IGF-1 (40).

*Pecam-1* (CD31) is a member of the Ig superfamily and is a single-chain transmembrane glycoprotein expressed on endothelial cells, platelets, and leukocytes (41). Angiogenesis refers to the formation of new blood vessels from pre-existing vasculatures by intussusception or sprouting (42). Angiogenesis requires cascade events such as enzymatic degradation of the basal membrane of the pre-existing vessels, migration of endothelial cells, and endothelial cell proliferation (43). *Pecam-1* has been implicated in the adhesive and signaling cascades required for endothelial

cell migration, which plays an essential role in angiogenesis (44, 45). *Pecam-1* is also involved in the endothelial cell-to-cell associations required for the organization of endothelial cells into tubular networks (46).

*Ang*, a member of the ribonuclease A superfamily, is a potent angiogenic molecule (47). For inducing angiogenesis, *Ang* is taken up by endothelial cells through receptor-mediated endocytosis and rapidly translocates to the nucleus, promoting ribonuclease activity, basement membrane degradation, signaling transduction, and nuclear translocation (48). In addition, the secreted *Ang* binds to the cell surface actin to induce basement membrane and extracellular matrix degradation, promoting endothelial cell invasion and migration into the surrounding tissue (49). Angiogenesis occurs as an orderly cascade of molecular events and starts with angiogenic factor secretion. In this study, GH increased angiogenesis by promoting the secretion of those important angiogenic factors including *Lep*, *Pecam-1*, and *Ang*.

The strength of the present study was that experiments were conducted in various aspects on the mechanism by which the GH restored ovarian function, including evaluation of apoptosis, fibrosis, and angiogenesis. In addition, RT-qPCR was also conducted to explore the mechanisms of GH on angiogenesis. Our findings will provide the basis for further studies on the pharmacological recovery of ovarian function. This present study has several limitations. First, this mouse model only provided relative short-term effects of GH on recovery of ovarian function. Second, it is not clear whether GH delayed declining ovarian function or restored it. It is necessary to compare the ovarian function before GH administration and on the day of ovarian sampling to determine if there is a difference in ovarian function decline. Third, although the GH concentration was determined after reviewing previous studies, it is a very high dose compared to the actual GH concentration administered

to humans. Fourth, GH could interfere with the activity of chemotherapy. Therefore, further studies on the long-term effects of GH on ovarian function and the elaboration of the protocol applied in human trials are recommended. In addition, GnRHa might be offered to patients when cryopreservation methods are not feasible in actual clinical practice. It would be helpful in clinical practice if the effects and mechanisms of GnRHa and GH are compared in further investigation.

In conclusion, this present study demonstrated the effect of GH on ovarian function recovery in ovarian insufficiency induced by the administration of CP. The mechanisms by which GH achieved these effects might be through decreasing apoptosis, stromal fibrosis, and increasing angiogenesis by up-regulating *Lep*, *Pecam-1*, and *Ang*.

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국문 초록

# 난소부전 마우스 모델에서 성장호르몬에 의한 난소 기능회복 효과

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## 목적

이 연구는 항암제인 cyclophosphamide (CP)를 투여하여 난소부전 마우스 모델을 구축한 후 성장호르몬을 투여하여 난소 기능 회복에 대한 효과를 확인하고 그 기전을 탐색하고자 하였다.

## 연구 방법

6주령 ICR 마우스에 CP를 400 mg/kg를 복강 내로 투여하여 난소부전을 유도한 후 네 군 (대조군, CP 투여군, 성장호르몬 1 mg/kg 투여군, 성장호르몬 2 mg/kg 투여군) 으로 나누었고 각 군당 마우스 10마리를 배정하였다. CP를 투여한 후 7일 뒤 성장호르몬을 7일간 투여하였고 투여 완료한 다음 날 각 군당 5마리의 마우스의 난소를 채취하여 조직학적 검사를 진행하였다. 나머지 5마리는 체외수정 (in vitro fertilization, IVF)을 진행하였다. 조직학적 검사와 체외수정을 통해 성장호르몬의 효

과를 확인 한 뒤 난소 기능을 회복시키는 효과를 확인하기 위하여 다음의 실험을 진행하였다. 세포사멸 (apoptosis)의 정도를 비교하기 위하여 Terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) assay를 시행하였다. 난소 조직의 섬유화 정도를 알아보기 위하여 Masson's trichrome staining을 시행하였다. 혈관신생 (angiogenesis)을 정량화하기 위하여 CD31 면역조직화학염색 (CD31 immunohistochemistry)을 시행하였다. 역전사 중합효소 연쇄반응 (reverse transcription polymerase chain reaction, RT-qPCR)을 통하여 혈관신생에 관여하는 유전자의 발현을 알아보았다.

## 결과

CP 투여는 원시난포와 1차 난포의 소멸을 유발하였다. 성장호르몬의 투여는 원시난포의 소멸을 방지하였고 난포의 질을 향상시켰다. 체외수정 결과를 살펴보면, CP 투여군은 수정률 및 배반포 형성률 (blastocyst formation rate)이 저하되었다. 그 반면, 성장호르몬 투여군은 성장호르몬을 투여한 용량에 비례하여 체외수정 결과가 향상되었다. 그 밖에 성장호르몬은 세포사멸 및 난소조직의 섬유화를 감소시키고 혈관신생을 향상시켰다. 혈관신생과 연관된 여러 유전자의 발현이 증가하였는데 그 중에서 특히 Leptin (*Lep*), Platelet endothelial cell adhesion molecule 1 (*Pecam-1*), Angiogenin (*Ang*) 유전자가 성장호르몬 투여군에서 발현이 증가하였다.

## 결론

CP를 투여하여 난소 부전을 유도한 뒤 성장호르몬을 투여하면 세포자멸사 및 난소조직 섬유화가 감소되고 혈관신생과 연관된 *Lep*, *Pecam-1*, *Ang* 유전자 발현이 증가하여 난소 기능 회복이 촉진된다.

**주요어:** 성장호르몬, 난소 부전, 시클로포스파미드, 난소암, 난소 재생,

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