ARTICLE





Delayed onset of obesity and glucose tolerance in interleukin 18 deficient mice by single housed condition



Boyoung Kim¹, Yoo Yeon Kim¹, Harry Jung^{1,3}, Hajin Nam¹ and Jun Gyo Suh^{1,2*}

Abstract

Interleukin 18 (IL18) is a kind of proinflammatory cytokine that belongs to the interleukin-1 family. IL18 is associated with obesity and type 2 diabetes. To discover whether body composition parameters in IL18 deficient mouse are altered in single-housed condition, body weight, glucose tolerance, lipid profiles, fat masses, and size of white adipocytes were examined. Mice were housed singly and were divided as follows: C57BL/6 J male (B6-M), IL18 deficient male (IL18-M), C57BL/6 J female (B6-F), IL18 deficient female (IL18-M). Body weight statistically significantly increased in IL18-M at 9 months (p < 0.05). Glucose tolerance occurred in IL18-M at 6 and 9 months. Total cholesterol and HDL cholesterol were statistically significantly increased in IL18-F compared with B6-F at 9 and 12 months, respectively (p < 0.05). Also, total cholesterol of IL18-M was statistically significantly increased compared with B6-F and IL18-M at 9 months (p < 0.05). The perirenal and inguinal fat masses were statistically significantly increased in IL18-M at 9 months. In single-housed condition, onset of obesity and glucose tolerance were delayed by 3 months in IL18-M. Taken together, these results suggest that housing condition is a very important factor for weight gain and onset of glucose tolerance in IL18 deficient male mouse.

Keywords: IL18 deficient mice, Single-housed condition, Obesity, Dyslipidemia, Glucose tolerance

Introduction

Environmental and social factors are very important for understanding physiological differences in laboratory mice. To enhance the replicability and reliability of animal research, these factors should be regulated precisely. Inbred strains of mice are widely used for physiological research and are regulated for body composition by environmental and genetic factors [1, 2]. In single-housed mice, variance of body fat mass is less than that in grouphoused mice [2]. Therefore, housing condition (single, group) should be considered in research into metabolic related diseases such as obesity and diabetes.

The mice's body weight is determined by the metabolic rate. Obesity is excessive accumulation of body fat,

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Interleukin 18 (IL18) is associated with obesity and type 2 diabetes. Hyperphagia and weight gain appear at 6 months of age in IL18 deficient mouse [9, 10]. In addition, IL18 deficient mouse shows abnormal lipidemia in liver at 9 months [11]. All those studies achieved their results in group-housed condition. Therefore, our hypothesis is that body composition parameters in IL18 deficient mouse will be altered in single-housed condition. In this study, body weight was statistically significantly increased in IL18 deficient male mouse compared



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with C57BL/6 J male mouse at 9 months. Lipidemia and glucose tolerance appeared in IL18 deficient male mouse at 9 months. In addition, the size of white adipocytes was increased in IL18 deficient male mice at 9 months. The phenotypes of body weight gain and diabetes were delayed in IL18 deficient male mouse by single-housed condition. Our results suggest that housing condition is a very important factor for weight gain and onset of diabetes in IL18 deficient male mouse.

Materials and methods

Animals

C57BL/6 J and interleukin 18 deficient mice were obtained from the Laboratory Animal Resources Center, Hallym University, Korea. The animal room retained at temperature (22 ± 2 °C), humidity ($55\pm10\%$) and light (8 a.m. to 8 p.m.). The animals were provided chow diet (Protein: 20%, Fat: 4.5%, Fiber: 6%, Calcium 0.5%, Phosphorus 1%, Cargill Agri Purina, Korea) and water ad libitum. The experimental groups were divided as follows: (1) C57BL/6 J-male (B6-M, n=60), (2) C57BL/6 J female (B6-F, n=48), (3) IL18 deficient male (IL18-M, n=60), and (4) IL18 deficient female (IL18-F, n=60). Mice were housed singly ($13 \times 24 \times 12$ cm, MJ Ltd, Korea). The animals were handled with Hallym University guidelines (Hallym2016-65).

Measurement of body weight, food intake, and fasting blood glucose

The body weight was measured at 3, 6, 9, and 12 months using an animal electronic scale. Food intake was measured at 3 $(2 \sim 3 \text{ M})$, 6 $(4 \sim 6 \text{ M})$, 9 $(7 \sim 9 \text{ M})$, and 12 months $(10 \sim 12 \text{ M})$. To measure fasting blood glucose, blood was collected from the retro-orbital plexus after fasting (16 h). Blood glucose monitoring meter (Accu-Chek, Roche, USA) was used to measure fasting blood glucose.

Measurement of lipid profiles and Hemoglobin A_{1C}

Blood samples (300 µl) were collected at 3, 6, 9, and 12 months using the same method as described above. Then, total cholesterol, triglyceride (TG), low density lipoprotein cholesterol (LDL), high density lipoprotein cholesterol (HDL) were measured by blood biochemistry (Kornelab20XT, Thermo, USA). Hemoglobin A_{1C} (Hb A_{1C}) was measured using whole blood by an Hb A_{1C} analyzer (HLC-723G7, Tosoh, Japan).

Oral glucose tolerance test (OGTT)

OGTT was performed at 3, 6, 9, and 12 months. All animals were fasted for 16 h. Mice were given 2 g/kg of D-glucose (Sigma, USA) by oral administration. After 0, 30, 60, and 120 min of D-glucose administration, blood were collected and measured glucose levels using a

glucose meter (Accu-Chek, Roche, USA). Then, to determine whether glucose tolerance exists, the area under the curve (AUC) was calculated.

Histological analysis

Mice were anesthetized with isoflurane (gas flow with flowmeter 100-200 ml/min and vaporizer at 3%) and were perfused transcardially with 4% paraformaldehyde. Whole adipose tissues were dissected and weighed. The interscapular, epididymal, and ovary fat were embedded in paraffin and sectioned at 5 µm for hematoxylin-eosin staining. For histological analysis, the adipose tissue sections were blocked in blocking solution (1X TBS, 1% BSA, and 10% goat serum) at room temperature for 1 h. Then the adipose tissue sections were incubated with mitochondrial uncoupling protein-1 (UCP-1) antibody (1:500, Abcam, UK) at 4 °C overnight. After that, the adipose tissue sections were incubated with anti-rabbit secondary antibody (1:200, Cell Signaling, USA) at room temperature (1 h). At least 4 mice per group were used for the histological analysis.

Statistical analysis

All data were expressed as mean \pm S.D. The significant differences were compared using one-way analysis of variance (one-way ANOVA). All results were analyzed by SPSS Statistics 22.0 (SPSS Inc., Chicago, USA).

Results

IL18 deficient male (IL18-M) mice show obesity independently of food intake at 9 months

Body weight of IL18-M was statistically significantly increased compared with that of C57BL/6 J male (B6-M) at 9 months (p < 0.05). There was no difference in body weight between IL18 deficient female (IL18-F) and C57BL/6 J female (B6-F) at all ages. Body weight of IL18-M was statistically significantly increased compared with that of IL18-F at all ages (Fig. 1a, p < 0.005). Food intake of IL18-M was statistically significantly increased compared with that of IL18-F at 6 months (Fig. 1b, p < 0.05). Fasting blood glucose (FBG) was statistically significantly decreased in IL18-F compared with B6-F at 12 months (p<0.05), but IL18-M had similar FBG levels compared with B6-M at 3, 6, 9, and 12 months (Fig. 1c). Hemoglobin A1_C (HbA1_C) was statistically significantly different in IL18-M compared with that of IL18-F at 3, 6, and 12 months (Fig. 1d, p < 0005).

IL18 deficient male (IL18-M) mice have glucose tolerance and dyslipidemia

To verify abnormal glucose metabolism, an OGTT (oral glucose tolerance test) was performed. At 3 and 12 months, there was no difference in area under the



curve (AUC) of all groups (data not shown). AUC was statistically significantly increased in IL18-M compared with the other groups at 6 and 9 months (Fig. 2a, b, p < 0.05). These findings suggest that IL18 deficient male (IL18-M) had glucose tolerance at 6 and 9 months.

Dyslipidemia is a major risk factor of abnormal glucose metabolism. At 9 months, total cholesterol was statistically significantly increased in IL18-F compared with B6-F (Table 1, p < 0.05). Total cholesterol of IL18-M was statistically significantly increased compared with B6-F and IL18-F at 9 months (p<0.05). HDL cholesterol showed no significant differences among all groups. LDL cholesterol of IL18-F was statistically significantly increased compared with that of B6-F at 12 months (p<0.05). In B6-F and IL18-F, LDL cholesterol was statistically significantly increased compared with B6-M and IL18-M at 12 months (Table 1, p < 0.05). TG in IL18-M was statistically significantly increased compared with that in B6-M at 9 months (p < 0.05), but IL18-F showed a similar TG level compared with B6-F at 6, 9, and 12 months.

IL18 deficient male (IL18-M) mice promote adipose tissue accumulation

Adipose tissue weight at 9 months is shown in Table 2. To reduce variation, the weight of adipose tissue was expressed as a ratio of body weight. The total fat mass of B6-M and IL18-M was 10.85 ± 5.43 and 12.45 ± 6.22 , respectively. The perirenal and inguinal fat masses statistically significantly increased in IL18-M compared with B6-M at 9 months (Table 2, p < 0.05). The total fat mass of B6-F and IL18-F was 9.89 ± 4.95 and 8.66 ± 4.33 , respectively. There was no difference in fat mass between B6-F and IL18-F at 9 months.

Histological analysis was performed with interscapular and epididymal (ovary) fat at 9 months (Fig. 3a). The expression of UCP-1 protein, brown adipose tissue marker, was observed in interscapular fat, but not in epididymal (ovary) fat. UCP-1 protein was expressed in the mitochondria of brown adipose tissue. The size of the adipocytes was measured because the size of adipocytes increases with obesity (Fig. 3b). The mean size of epididymal adipocytes was statistically significantly increased in IL18-M compared with that of B6-M at 9 months (Fig. 3b, p < 0.05). In IL18 F, the mean size of ovary adipocytes increased slightly compared with that of the B6-F group at 9 months. These findings indicate that IL18-M have obesity via adipose tissue accumulation at 9 months.

Discussion

This study hypothesized that body composition parameters might be changed in IL18 deficient mouse in accordance with single-housed condition. IL18 deficient male mice showed obesity at 9 months, dependent on adipose



Table 1 Lipid profiles of interleukin 18 deficient mice at 6,9 and 12 months

Group	6 M	9 M	12 M	
Total choles	sterol (mg/dL)			
B6-M	96.87 ± 6.80	103.02 ± 4.72	102.74 ± 8.86	
IL18-M	106.20 ± 11.62	115.74 ± 8.15	116.36 ± 6.87	
B6-F	81.87 ± 6.37	85.70 ± 8.23	84.08 ± 5.77	
IL18-F	91.76 ± 13.88	$100.26 \pm 5.21^{*}$	94.30 ± 8.40	
HDL choles	terol (mg/dL)			
B6-M	39.55 ± 3.85	44.40 ± 2.52	46.41 ± 4.47	
IL18-M	44.66 ± 6.74	47.83 ± 6.37	53.57 ± 4.14	
B6-F	31.52 ± 3.51	35.01 ± 4.76	34.81 ± 3.41	
IL18-F	36.82 ± 7.12	40.46 ± 2.92	39.48 ± 5.33	
LDL cholest	erol (mg/dL)			
B6-M	7.41 ± 1.08	5.29 ± 0.74	7.32 ± 1.18	
IL18-M	6.89 ± 0.71	4.18±0.79	7.05 ± 0.93	
B6-F	8.34 ± 0.96	7.08 ± 0.85	9.43 ± 0.87	
IL18-F	8.76 ± 1.71	6.85 ± 0.83	$9.52 \pm 1.46^{*}$	
TG (mg/dL)				
B6-M	69.21 ± 11.02	83.40 ± 10.91	77.97 ± 10.41	
IL18-M	79.86 ± 12.01	$111.77 \pm 12.64^{*}$	85.44 ± 11.55	
B6-F	53.69 ± 9.19	53.11 ± 5.99	63.32 ± 8.61	
IL18-F	55.40 ± 8.83	63.16±6.78	84.60±15.49	

These data are expressed as mean $\pm\,$ S.D

*p < 0.05 compared with same sex

tissue accumulation (Figs. 1, 3). These male mice also developed dyslipidemia at 9 months and glucose tolerance from 6 to 9 months (Table 1, Fig. 2). In addition, IL18 deficient male mice had more beige body fat compared with control mice (Table 2). Netea et al. reported that IL18 deficient mice showed diet-induced obesity and glucose tolerance at 6 months [9]. IL18 deficient male mice develop dyslipidemia and steatohepatitis in group condition housing [11]. There are two differences between our results and those of previous studies [9, 10]. First, the onset of obesity in our experiment was delayed by 3 months compared with Netea's results. Second, obesity in our experiment was induced by fat accumulation rather than increased food intake. There are two possible reasons for these discrepancies: (1) housing condition as group-housed or single-housed and (2) difference of microbiome composition

Mice are social animals; therefore, they are affected by an isolated environment such as single housing [12]. Group-housed mice show a considerable increase in body weight compared with single-housed mice [1]. Increased body weight in the group-housed mice is due to the differences in thermoregulation [13]. Therefore, in grouphousing, IL18 deficient male mice showed rapid increases in weight gain because of increased food intake and differences in thermoregulation [13]. Ultimately, this leads to obesity and type 2 diabetes at 6 months [9, 10, 14]. On

Type of adipose tissue (%)	B6-M	IL18-M	B6-F	IL18-F
Cervical	0.21 ± 0.11	0.26±0.13	0.22 ± 0.11	0.19±0.09
Axillary	0.73 ± 0.36	0.73 ± 0.36	0.86 ± 0.43	0.66 ± 0.33
Interscapular	2.40 ± 1.20	2.42 ± 1.21	2.47 ± 1.23	2.13 ± 1.07
Mesenteric	1.21 ± 0.60	1.53 ± 0.76	1.14 ± 0.57	0.78 ± 0.39
Mediastinic	0.13 ± 0.07	0.15 ± 0.07	0.12 ± 0.06	0.10 ± 0.05
Retroperitoneal	0.77 ± 0.39	0.52 ± 0.26	0.35 ± 0.18	0.26 ± 0.13
Epididymal and Ovary	2.77 ± 1.38	3.21 ± 1.60	1.89 ± 0.95	2.18 ± 1.09
Perirenal ^a	0.40 ± 0.20	$0.93 \pm 0.47^{*}$	0.74 ± 0.37	0.49 ± 0.25
Inguinal ^a	1.67 ± 0.84	$2.06 \pm 1.03^{*}$	1.58 ± 0.79	1.50 ± 0.75
Gluteal	0.56 ± 0.28	0.65 ± 0.32	0.53 ± 0.26	0.38 ± 0.19
Total	10.85 ± 5.43	12.45 ± 6.22	9.89 ± 4.95	8.66 ± 4.33

Table 2 Relative fat mass of interleukin 18 deficient mice at 9 months

These data are expressed as mean \pm S.D

* p < 0.05 compared with same sex

^a Beige fat



the other hand, single-housed mice show genetic effects on body composition compared with group-housed mice [2]. With single-housed mice, depression-like behavior is induced [15]. The multi-housed mice are more active compared to the single-housed mouse [16]. Thus, the single-housed mouse need more calories to keep it's body temperature compared to group housed mouse. Therefore, body weight gain and onset of diabetes were delayed in IL18 deficient male mouse by single-housed condition.

Second, recent studies demonstrated that the gut microbiome in very important in human metabolism such as obesity and diabetes [17, 18]. When the microbiota (obese microbiota) of obese mice transplant to normal mice, obesity occurs in the normal mice [19, 20]. Also, obese microbiota influence lipid metabolism such

as triglyceride [21]. In group housing, the microbiome of mice is homogenized by the cage effect [22]. IL18 deficient mice used in our study were purchased from Jackson Laboratory several years ago and were maintained at the Laboratory Animal Resources Center, Hallym University. Therefore, the microbiota of the IL18 deficient mice used in our study might have a different composition compared with that of Netea's IL18 deficient mice [9]. To confirm this speculation, microbiota analysis of the IL18 deficient mice of the two institutions is needed.

In our results, IL18 deficient female mice did not show any statistically significant differences in body weight and size of adipocytes in ovary fat (Figs. 1a, 3b). These mice also showed no statistically significant differences in the OGTT at 3, 6, 9, and 12 months (Fig. 2). The prevalence of diabetes in woman is lower compared with that in man [23]. Females are protected from β cell death in rodent diabetic models [24]. Therefore, gender differences in our study might be caused by estrogen, which modulates glucose homeostasis and insulin resistance.

In conclusion, IL18 deficient male mice showed delayed onset of obesity and glucose tolerance in the singlehoused condition. This result indicates that the housing condition and gender are key factors in metabolic studies such as body composition and glucose tolerance.

Authors' contribution

BK, YYK, HJ, and HN carried out the animal experiment. BK wrote the first draft of the manuscript. JGS initiated the project, supervised the experiment, and critically reviewed the manuscripts. All authors read and approved the final manuscript.

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Competing interest

The authors declare that they have no competing interests.

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