The Ratio of High-Molecular Weight Adiponectin and Total **Adiponectin Differs in Preterm and Term Infants**

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ABSTRACT: Adiponectin consists of three subspecies (high-, middleand low-molecular weight adiponectin). Among these, high-molecular weight adiponectin (H-adn) is suggested to be an active form of this protein. To assess the relationship between H-adn and postnatal growth in preterm infants (PIs), serum H-adn and total adiponectin (T-adn) were measured in 46 PIs at birth and at corrected term, and 26 term infants (TI) at birth. T-adn and H-adn concentrations, and the ratio of H-adn to T-adn (H/T-adn) were significantly greater in TI and PI at corrected term than in PI at birth (p < 0.001). T-adn and H-adn concentrations in PI at corrected term were similar to those in TI, but H/T-adn in PI at corrected term was less than that in TI (p < 0.02). Stepwise multiple regression analysis revealed that the factors contributing to H/T-adn and serum concentrations of T- and H-adn in PI at corrected term were different from those in TI. These data suggest that quality of early postnatal growth in PIs is different from that in normally developed TI. Postnatal growth accompanying adipose tissue similar to TI may be important for PI to prevent future development of cardiovascular disease. (Pediatr Res 65: 580-583, 2009)

E pidemiologic studies have demonstrated that low birth weight is associated with an increased risk for atherosclerotic cardiovascular disease (CVD) later in life (1-3); however, the mechanisms behind this relationship have yet to be fully elucidated. Adiponectin is one of several adipocytokines secreted by the adipocytes (4) and has numerous physiologic functions. Serum levels of adiponectin are inversely associated with insulin resistance, inflammatory markers, and CVD risk factors (5-7). Taken together, these data suggest that adiponectin has antidiabetic, antiinflammatory, and antiatherogenic functions. Adiponectin consists of three multimer species (high-, middle-, and low-molecular weight adiponectins: H-adn, M-adn, and L-adn) (8-11). Among these, H-adn is suggested to be an active form of this protein, having a higher bound affinity to surface of the cell membrane and remarkably high adenosine monophosphate kinase activation compared with M-adn and L-adn (10,11). Furthermore, the ratio of H-adn and total adiponectin (T-adn) (H/Tadn) are more significantly associated with insulin resistance than T-adn, thereby suggesting the usefulness of the H/T-adn in diagnosing insulin resistance (12).

Serum levels of adiponectin are inversely associated with body weight and body mass index (BMI) in adults and schoolchildren (4,13). In contrast, serum levels of adiponectin in

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neonates were positively associated with fetal birth weight (14), with their levels being much higher than those in adults and schoolchildren (2,4,7,13,14). The physiologic significance of these findings remains to be clarified, and no information exists regarding serum multimeric adiponectin concentrations in the fetal and neonatal period. Thus, in the present study, as a first step in clarifying the relationship between low birth weight and future CVD risk, we investigated the serum concentrations of multimeric adiponectin in neonates and the difference between premature and mature infants.

SUBJECTS AND METHODS

The subjects consisted of 72 newborn infants (46 preterm born between weeks 24 and 35 of gestation and 26 term and near-term infants born after week 36 of gestation from an uncomplicated pregnancy) who had been admitted to the Ryukyu University hospital, Japan. All term and near-term infants were appropriate for gestational age (AGA). In preterm infants (PIs), 37 infants were AGA and nine infants were small for gestational age (SGA). Preterm deliveries were induced because of threatened premature delivery (21 infants), nonreassuring fetal status (four infants), viral infection of mother (one infant), premature rupture of membrane (13 infants), placenta previa (four infants), hypertension of mother (two infants), and cervical cancer of mother (one infant). Among 46 mothers, two mothers had pregnancy-induced hypertension. All PIs were fed breast milk and infant formula. Serum sample collection and anthropometric measurements were performed at birth and at the corrected term of PIs. Written informed consent was obtained from the parents. The Ethics Committee of the Ryukyu University approved the study protocol.

Anthropometric measurements of subjects. Gestational age was confirmed by ultrasound before week 20 of gestation. The umbilical cord was cut, and the placental weight was measured with a calibrated scale and recorded. The infants were weighed immediately after birth, and birth length was measured using a measuring board. Heart rate (HR) and blood pressure in the lower limb were measured with an automated oscillometric device (MP-60, Phillips Medical Systems, Eindhoven, The Netherlands). Neonatal BMI and Ponderal index were calculated as weight (kg)/body length² (m²) and weight (kg)/body length³ (m³), respectively.

Laboratory measurements. Umbilical vein samples were drawn and samples were stored at -40°C until assay. Serum adiponectin concentrations were measured by sandwich ELISA kits (Daiichi Pure Chemicals Co. Ltd., Tokyo, Japan) with a dynamic range of 0.075-4.8 ng/mL. Intraassay variations (CV) were 5.3% (T-Adn), 4.1% (M-adn + H-adn), and 3.3% (H-adn), as described previously (15).

Statistical evaluation. The differences between series of data were determined by Wilcoxon's rank sum test. Levels of T-adn, H-adn, and H/T-adn were markedly skewed. Thus, these parameters were normalized by log transformation. Pearson's correlation coefficient test was performed to assess the associations between data. To examine the relationship between adiponectin multimeric complexes and clinical data, a forward stepwise multiple regression analysis was performed. Values are reported as the mean \pm SD. Significance was set at p < 0.05. All analyses were performed with JMP 5.1 (SAS Institute Inc., Cary, NC).

Abbreviations: adn, adiponectin, CVD, cardiovascular disease, DBP, diastolic blood pressure, H-adn, high-molecular weight adiponectin, H/T-adn, the ratio of high-molecular weight adiponectin to total adiponectin, HR, heart rate, PI, preterm infant, SBP, systolic blood pressure, T-adn, total adiponectin, TI, term infants

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 Table 1. Clinical and chemical data on preterm and term infants

	At birth	р	At corrected term	р	Term infants
Number of subjects (M/F)	46 (19/28)				26 (11/15)
Postmenstrual age (wk)	31.7 ± 2.9	< 0.001	37.1 ± 1.1	ns	37.7 ± 1.2*
Placental weight (g)	371 ± 102				535 ± 138*
Body weight (g)	1609 ± 492	< 0.001	2229 ± 374	< 0.001	$2908 \pm 574*$
Body length (cm)	40.9 ± 4.2	< 0.001	43.8 ± 2.7	< 0.001	$47.3 \pm 2.1*$
BMI (kg/m ²)	9.3 ± 1.5	< 0.001	11.6 ± 1.5	< 0.001	$12.9 \pm 1.7^{*}$
Ponderal index (kg/m ³)	2.27 ± 0.29	< 0.001	2.69 ± 0.42	ns	$2.74 \pm 0.34*$
SBP (mm Hg)	61 ± 8	< 0.001	69 ± 9	< 0.001	62 ± 5
DBP (mm Hg)	35 ± 7	ns	36 ± 7	ns	36 ± 7
Heart rate	134 ± 12	< 0.01	144 ± 12	< 0.001	$123 \pm 17^{+}$
T-adn (µg/mL)	6.2 ± 3.3	< 0.001	19.9 ± 13.5	ns	$17.0 \pm 10.1*$
H-adn (µg/mL)	2.6 ± 2.0	< 0.001	9.6 ± 5.7	ns	$9.9 \pm 6.0^{*}$
M-adn (μ g/mL)	1.5 ± 1.0	< 0.001	5.3 ± 5.2	ns	$3.8 \pm 3.5 \dagger$
L-adn (µg/mL)	2.1 ± 1.2	< 0.01	5.0 ± 6.7	ns	3.3 ± 2.1
H/T-adn	0.37 ± 0.16	< 0.001	0.50 ± 0.12	< 0.02	$0.58 \pm 0.11*$

* p < 0.001, significantly different from preterm infants at birth.

 $\dagger p < 0.05$, significantly different from preterm infants at birth.

M, male; F, female; ns, not significant.

RESULTS

All data presented were not changed even after removing data of SGA infants. Therefore, we did not separate the data for SGA and AGA in analysis.

Characteristics of the subjects. As shown in Table 1, postmenstrual age of PI was significantly younger than that of term infant (TI). Postmenstrual age at corrected term of PI was similar to that of TI. Placental weight was significantly lighter in PI than that in TI. Body sizes (weight and length) were much smaller in PI at birth than those in TI. Body sizes were significantly larger in PI at corrected term than those at birth but still significantly smaller than those in TI. Systolic blood pressure (SBP) of PI at birth was similar to that of TI. SBP of PI at corrected term was significantly higher than those of PI at birth and TI. No differences in diastolic blood pressure (DBP) were found among the three groups. HR was significantly greater in PI at corrected term than in PI at birth and in TI. HR of TI was significantly less than that of PI at birth.

Serum multimeric adiponectin levels. Concentrations of adiponectin in cord serum were similar to those in serum of neonates at birth; therefore, we compared adiponectin levels of cord serum and serum levels of PI at corrected term (14). As shown in Table 1, serum concentrations of T-adn, H-adn, M-adn, and L-adn in PI at birth were significantly less than those in TI. Those of PI at corrected term were similar to those of TI. In PI at corrected term, all of these parameters were significantly higher than those of PI at birth. The ratio of H-adn to T-adn (H/T-adn) of PI at birth was significantly less than those of PI at corrected term and TI. H/T-adn of PI at corrected term was significantly less than that of TI.

Tables 2–4 show the univariate correlations between T-adn, H-adn, H/T-adn, and other parameters in PI at birth and at corrected term (n = 46) and those in TI. In PI at birth (Table 2), serum concentrations of T-adn and H-adn were positively correlated with postmenstrual age, placental weight, body weight, body length, BMI, and Ponderal index (r = 0.31– 0.79, p = 0.036–0.000). H/T-adn was positively correlated

Table 2. Univariate correlations between T-adn, H-adn, H/T-adn, and clinical variables of preterm infants at birth (n = 46)

-	T-a	ıdn	H-a	ıdn	H/T-	-adn
Independent variables	r	р	r	р	r	р
Postmenstrual age (wk)	0.75	0.000	0.75	0.000	0.48	0.001
Placental weight (g)	0.59	0.000	0.52	0.001	0.29	0.080
Body weight (g)	0.79	0.000	0.73	0.000	0.44	0.003
Body length (cm)	0.75	0.000	0.66	0.000	0.39	0.009
BMI (kg/m ²)	0.77	0.000	0.77	0.000	0.48	0.001
Ponderal index (kg/m ³)	0.31	0.036	0.47	0.001	0.36	0.018
SBP (mm Hg)	0.21	0.196	0.23	0.175	0.13	0.444
DBP (mm Hg)	-0.01	0.953	0.04	0.814	-0.02	0.886
Heart rate	-0.02	0.898	-0.09	0.590	-0.02	0.889

Bold type indicates significant correlations.

Table 3. Univariate correlations between T-adn, H-adn, H/T-adn, and clinical variables of preterm infants at corrected term (n = 46)

(n = 40)								
	T-adn		H-adn		H/T-adn			
Independent variables	r	р	r	р	r	Р		
Postmenstrual age (wk)	0.14	0.371	0.30	0.047	0.38	0.010		
Placental weight (g)	0.29	0.069	0.38	0.017	0.24	0.138		
Body weight (g)	0.34	0.021	0.41	0.005	0.19	0.209		
Body length (cm)	0.26	0.085	0.36	0.014	0.26	0.086		
BMI (kg/m ²)	0.14	0.340	0.09	0.570	-0.12	0.416		
Ponderal index (kg/m ³)	0.08	0.609	0.04	0.782	-0.08	0.619		
SBP (mm Hg)	0.32	0.030	0.18	0.241	-0.30	0.040		
DBP (mm Hg)	0.13	0.379	0.06	0.684	-0.15	0.311		
Heart rate	0.29	0.051	0.06	0.685	-0.50	0.000		

Bold type indicates significant correlations.

with postmenstrual age, body weight, body length, BMI, and Ponderal index (r = 0.36-0.48, p = 0.018-0.001). No significant correlation was found among T-adn, H-adn, H/Tadn, and other parameters. In PI at corrected term (Table 3), serum concentrations of T-adn were positively correlated with body weight and SBP (r = 0.32-0.34, p = 0.03-0.021). Serum concentration of H-adn was positively correlated with postmenstrual age, placental weight, body weight, and body

Table 4. Univariate correlations between T-adn, H-adn, H/T-adn, and clinical variables of term infants (n = 26)

	0				
T-adn		H-	H-adn		adn
r	р	r	р	r	р
0.26	0.195	0.37	0.060	0.39	0.047
0.12	0.567	0.04	0.863	-0.23	0.273
0.39	0.049	0.40	0.045	0.10	0.625
0.18	0.383	0.17	0.409	0.01	0.962
0.43	0.027	0.46	0.017	0.18	0.369
0.31	0.119	0.36	0.069	0.21	0.296
0.39	0.047	0.44	0.023	0.24	0.236
0.45	0.022	0.52	0.007	0.31	0.125
0.09	0.674	0.07	0.736	-0.03	0.871
	<pre>r 0.26 0.12 0.39 0.18 0.43 0.31 0.39 0.45</pre>	r p 0.26 0.195 0.12 0.567 0.39 0.049 0.18 0.383 0.43 0.027 0.31 0.119 0.39 0.047 0.45 0.022	r p r 0.26 0.195 0.37 0.12 0.567 0.04 0.39 0.049 0.40 0.18 0.383 0.17 0.43 0.027 0.46 0.31 0.119 0.36 0.39 0.047 0.44 0.45 0.022 0.52	r p r p 0.26 0.195 0.37 0.060 0.12 0.567 0.04 0.863 0.39 0.049 0.40 0.045 0.18 0.383 0.17 0.409 0.43 0.027 0.46 0.017 0.31 0.119 0.36 0.069 0.39 0.047 0.44 0.023 0.45 0.022 0.52 0.007	r p r p r 0.26 0.195 0.37 0.060 0.39 0.12 0.567 0.04 0.863 -0.23 0.39 0.049 0.40 0.045 0.10 0.18 0.383 0.17 0.409 0.01 0.43 0.027 0.46 0.017 0.18 0.31 0.119 0.36 0.069 0.21 0.39 0.047 0.44 0.023 0.24 0.45 0.022 0.52 0.007 0.31

Bold type indicates significant correlations.

 Table 5. Stepwise multiple regression models for predicting T-adn, H-adn, and H/T-adn

	Selected independent	r^2	
	parameters	<i>r</i> -	p
Preterm infant	s at birth $(n = 46)$		
T-adn	Body weight	0.62	0.000
H-adn	BMI	0.59	0.000
H/T-adn	Postmenstrual age	0.23	0.001
Preterm infant	s at corrected term $(n = 46)$		
T-adn	SBP, placental weight	0.22	0.011
H-adn	Body weight	0.17	0.005
H/T-adn	HR, body weight	0.34	0.000
Term infants a	t birth $(n = 26)$		
T-adn	DBP	0.20	0.022
H-adn	DBP	0.27	0.007
H/T-adn	Postmenstrual age,	0.33	0.011
	placental weight		

length (r = 0.30-0.41, p = 0.047-0.005). H/T-adn was positively correlated with postmenstrual age (r = 0.38, p = 0.010). SBP and HR showed significant negative correlations with H/T-adn (r = -0.30 to -0.50, p = 0.04-0.000). No significant correlation was found among T-adn, H-adn, H/Tadn, and other parameters. In TI (Table 4), serum concentration of T-adn was positively correlated with body weight, BMI, SBP, and DBP (r = 0.39-0.45, p = 0.049-0.022), as well as H-adn was positively correlated with body weight, BMI, SBP, and DBP (r = 0.40-0.52, p = 0.045-0.007). H/T-adn was positively correlated with postmenstrual age (r = 0.39, p = 0.047). No significant correlation was found among T-adn, H-adn, H/T-adn, and other parameters.

Because each of the above parameters can potentially contribute directly to the regulation of serum multimeric adiponectin levels, we performed stepwise multiple regression analysis with T-adn, H-adn, and H/T-adn as the dependent variables and postmenstrual age, placental weight, body weight, body length, BMI, SBP, DBP, and HR as the independent variables (Table 5). In PI at birth, the predictors of T-adn, H-adn, and H/T-adn were body weight, BMI, and postmenstrual age, respectively. In PI at corrected term, the major predictors of T-adn were SBP and placental weight. The predictor of H-adn was body weight. HR and body weight were selected as the predictors of H/T-adn. In TI at birth, the predictor of T- and H-adn was DBP. Postmenstrual age and placental weight were selected as the predictors of H/T-adn.

DISCUSSION

In the present study, we showed that i) the serum concentrations of T-adn and H-adn in PI at corrected term were three times higher than those in PI at birth, whereas serum concentrations of T-adn and H-adn of PI at corrected term were similar to those of TI; ii) the H/T-adn of TI and PI at corrected term were significantly greater than that of PI at birth, but that of PI at corrected term was still significantly less than that of TI; and iii) the factors contributing to serum concentrations of T- and H-adn differed among the three groups.

A recent study reported that concentrations of adiponectin in fetal circulation shows a 20-fold increase between week 24 of gestation and term, and serum concentrations of adiponectin were positively associated with body weight in PIs (16). In addition, Kotani et al. (14) reported that the cord serum adiponectin concentrations in full-term neonates are higher than serum adiponectin concentrations in adults (22.4 µg/mL versus 8.2 µg/ mL) and are correlated positively with fat mass-related parameters (birth weight, BMI, and leptin concentration). In the present study, serum concentrations of T- and H-adn increased by 2.5-3-fold between 32 and 38 wk of gestation, and their levels were also higher than those reported in schoolchildren (2,7,17). These data suggest that fetal development during late pregnancy contributes to serum concentrations of T- and H-adn, and that the relationship between adiposity and adiponectin levels in neonates may differ from that in adults and children. Interestingly, Inami et al. (18) recently reported a positive association between serum concentrations of adiponectin and body weight in TIs at birth, but no association was found at 1 mo of age. To date, some roles of adipocytokines such as IL-6 and TNF- α have been highlighted in the inverse correlation between adiponectin and body weight in adults (19). It has been shown that IL-6 and TNF- α reduce the secretion of adiponectin from adipocytes (20). Some types of inflammation in adipose tissue might contribute to the inverse correlation between adiponectin and adiposity. The rapid increase in fetal body weight during late pregnancy and the perinatal period is a physiologic development; thus, it is reasonable to consider that an increase in adipose tissue at this stage may not be accompanied by hypertrophy of adipocytes and the inflammation in adipose tissue. This may also cause the higher levels of adiponectin and positive correlations between adiponectin and body weight observed during the perinatal period. With respect to the higher concentrations of adiponectin in PI at corrected term and TI, Kim et al. (21) reported that serum levels of adiponectin are a starvation signal released by adipocytes, and that normalsized adipocytes do not induce insulin resistance, even in obese status. It is well known that development of adipose tissue is greatest in perinatal period (22). A recent report of Pinar et al. (23) showed that the high concentrations of H-adn found in the fetus were associated with higher insulin sensitivity. Thus, high levels of adiponectin in the perinatal period may reflect rapid growth during this period.

As shown in Table 2, T-adn, H-adn, and H/T-adn were significantly associated with adiposity-related parameters (body weight, BMI, and Ponderal index) in PI at birth. However, these associations were weakened in PI at corrected term and TI (Tables 3 and 4). Different from PI at birth, hemodynamic

parameters (blood pressure and/ or HR) were significantly associated with T-adn and H/T-adn in PI at corrected term, and with T-adn and H-adn in TI (Tables 3-5). It is of interest that expression of adiponectin in fetus was pronounced in vascular endothelium, thereby suggesting that in addition to adipose tissues, vascular endothelial cells may contribute to fetal adiponectin levels (23). It is plausible that hemodynamic parameters affect the status of vascular endothelium and lead to significant correlation of these to adiponectin in PI at corrected term and TI. Taken together, our data suggest that vascular endothelium might contribute to serum concentrations of T-adn and H-adn during late pregnancy. However, in PI at corrected term, hemodynamic parameters were not associated with H-adn. SBP and HR were inversely associated with H/T-adn. These results indicate that in PI at corrected term, ex-utero factors may influence adipose tissue development and contribution of vascular endothelium to adiponectin status. Placental weight was significantly correlated with T-adn and H-adn in PI at birth (Table 2). Serum concentrations of H-adn in PI at corrected term were still positively associated with placental weight (Table 3). A recent report showed that the adiponectin concentration in umbilical cord serum is positively associated with the weight ratio of fetus to placenta (24). In contrast, Pinar et al. proved the absence of endogenous placental adiponectin and showed its exclusive production by fetal tissues. All of these findings indicate that effect of placental weight on adiponectin status may be indirect effects; however, further studies are needed to clarify this.

Wang et al. (25) showed that ectopic fat accumulation occurred even in nonobese mice, which showed hyperinsulinemia, insulin resistance, and hypertrophy of adipocytes because of subnormal adipocyte storage capacity, whereas Ruderman et al. (26) reported metabolically obese, normal-weight patients. These previous studies, combined with the report by Kim et al. (21) strongly suggest that in neonates, normal development of adipose tissue is important to prevent future development of cardiometabolic diseases such as diabetes, hypertension, and dyslipidemia. Subnormal development of adipose tissue in neonates may reduce the number of adipocytes. If our notion is valid, the limited number of adipocytes in subjects with low birth weight may induce hypertrophy of adipocytes more readily than in subjects with normal birth weight. As a result, these subjects may be at high risk for cardiometabolic diseases in the future, even if normal weight is maintained. Ibanez et al. (27) recently reported that children born small for gestational age tend to be viscerally adipose with hypoadiponectinemia, even if they are not overweight. This may support our hypothesis.

In conclusion, our present data suggest that postnatal growth accompanying adiposity similar to TI may be important for premature and/or low birth weight infant to prevent future development of cardiometabolic diseases. Monitoring of T- and H-adn could be useful to evaluate the maturation of adipose tissue in PIs.

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