

Serum adipocyte-fatty acid binding protein level is elevated in severe OSA and correlates with insulin resistance

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ABSTRACT: Obstructive sleep apnoea (OSA) is associated with insulin resistance and metabolic syndrome. There is evidence that adipocyte-fatty acid binding protein (A-FABP) may be involved in the development of cardiometabolic dysfunction. The present authors hypothesise that A-FABP is upregulated in OSA.

A total of 124 males without hypertension, diabetes mellitus, hyperlipidaemia or cardiovascular disease were recruited and underwent polysomnography. Serum A-FABP levels showed significant positive correlations with duration of oxygen desaturation and minimal oxygen saturation, fasting insulin and insulin resistance index by homeostasis model assessment. When subjects were divided into tertiles according to apnoea/hypopnoea index (AHI), serum A-FABP levels were significantly higher in the group with AHI \geqslant 34.4 events·h⁻¹ than the groups with AHI 13.2=34.4 events·h⁻¹ or with AHI <13.2 events·h⁻¹. Serum A-FABP levels were significantly higher in the AHI \geqslant 34.4 group than obesity-matched subjects with AHI <34.4 events·h⁻¹.

Serum adipocyte-fatty acid binding protein levels correlated with obstructive sleep apnoea and insulin resistance, independently of obesity, and were significantly higher in severe obstructive sleep apnoea. Adipocyte-fatty acid binding protein may play a role in obstructive sleep apnoea and metabolic dysfunction.

KEYWORDS: Adipocyte-fatty acid binding protein, obstructive sleep apnoea

bstructive sleep apnoea (OSA) is highly associated with obesity and other components of the metabolic syndrome [1–3]. This clustering of OSA with cardiometabolic risk factors may imply that there is a common molecular pathway or mechanism underlying these disorders which has not been completely worked out. Visceral fat acts as a reservoir for various mediators, which are released into the circulation, and may play pathogenetic roles in metabolic or vascular dysfunction; OSA may act as a trigger to initiate or maintain the expression and release of these mediators [4–6].

Fatty acid binding proteins (FABPs) are a family of small intracellular lipid-binding proteins that reversibly bind hydrophobic ligands, which influence energy metabolism and inflammation, such as long chain fatty acids and eicosanoids [7]. Adipocyte-FABP (A-FABP), also known as aP2 or FABP4, is a cytosolic protein present in an abundant amount in adipocytes and macrophages [8]. A growing body of evidence from both animal and clinical studies suggests that A-FABP plays an

important role in the interaction between lipid metabolism and inflammation, and may be involved in the development of the metabolic syndrome [9, 10]. The pro-inflammatory role of A-FABP expression is supported by the observation that its expression in macrophages is induced by oxidised low-density lipoprotein (LDL) [11], as well as toll-like receptor activators [8]. In animal experiments, it has been shown that mice lacking A-FABP and mal1 (a minor form of FABP in adipose tissue) were protected from the development of insulin resistance in the context of dietinduced obesity [12], type 2 diabetes [13], fatty liver disease [14] and hypercholesterolaemia atherosclerosis [10]. Furthermore, obese mice (ob/ob) lacking A-FABP and mal1 (aP2/mal1-/-) were able to maintain normal blood glucose levels and peripheral insulin sensitivity despite being severely obese, implying that the A-FABP deficiency genotype is associated with improved systemic insulin sensitivity, which is independent of leptin action [15]. A recent report on BMS309403, a potent inhibitor of A-FABP, showed that treatment of obese mice (ob/ob) with this selective inhibitor led AFFILIATIONS

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STATEMENT OF INTEREST
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to reduced hyperglycaemia and hyperinsulinaemia [16]. A-FABP has also been shown to be involved in the mediation of allergic airway inflammation, implicating its role in asthma and possible action in bronchial epithelial cells [17].

Although traditionally thought to be a cytoplasmic protein, A-FABP is also detectable in the human bloodstream. Recent evidence demonstrates circulating A-FABP level to be an independent predictor of the development of metabolic syndrome and diabetes after adjustment for the effects of adiposity [18, 19]. In addition, serum A-FABP level has been shown to be closely associated with carotid atherosclerosis in Chinese females [20]. These epidemiological observations were also corroborated by a recent large population-based genetic study showing a reduced risk for hypertriglyceridaemia, type 2 diabetes and coronary artery disease in subjects carrying a functional genetic variant of the A-FABP gene that resulted in reduced adipose tissue A-FABP gene expression [21].

In light of the close association between OSA, adiposity and insulin resistance, the aim of the present study was to evaluate the role of A-FABP as a potential mediator in the development of metabolic dysfunction in OSA. The present authors hypothesise that severe OSA may lead to upregulation of A-FABP production, which correlates with insulin resistance, promoting the development of cardiometabolic dysfunction.

METHODS AND MATERIALS

Subject characteristics

Consecutive male subjects (aged ≥18 yrs) undergoing overnight polysomnography (PSG) at the Sleep Laboratory of Queen Mary Hospital (Hong Kong) were recruited from the period commencing October 2002 until March 2006 (excluding the period March-September 2003 when sleep laboratory services were suspended) in order to explore various cardiometabolic aspects in relation to OSA; subjects without other prevailing comorbidities were targeted (Obstructive Sleep Apnoea Cardio-Metabolic Study (OSACMS) cohort). The present authors investigated the relationship between serum A-FABP levels and sleep disordered breathing. The exclusion criteria were as follows: known history of hypertension or diabetes mellitus (fasting plasma glucose ≥7.0 mmol·L⁻¹); hyperlipidaemia on lipid-lowering therapy; cardiovascular diseases or other major chronic medical illness; use of regular medication.

All subjects gave written informed consent. The study protocol was approved by the Hong Kong University/Hong Kong Hospital Authority (Hong Kong) Institutional Review Board Ethics Committee.

Anthropometric and biochemical measurements

Clinical and anthropometric measurements (body height, body weight, waist circumference, body mass index (BMI)) were collected as previously reported [2]. Plasma glucose was measured using the glucose oxidase method with a Beckman Autoanalyzer (Beckman Instruments, Bream, CA, USA). Serum insulin was determined with a microparticle enzyme immunoassay on the Abbott IMx system (Abbott, Abbott Park, IL, USA). The insulin resistance index estimated by the homeostasis model assessment method (Homeostasis Model

Assessment – Insulin Resistance (HOMA-IR)) was calculated as follows [22]:

fasting serum insulin ($\mu U \cdot mL^{-1}$) \times fasting plasma glucose (mmol·L⁻¹)/22.5

Plasma total cholesterol and triglyceride were measured enzymatically using a Hitachi 912 Analyzer (Roche Diagnostics, GmbH, Mannheim, Germany). High-density lipoprotein cholesterol was measured using a homogenous method with polyethylene glycol-modified enzymes and alpha-cyclodextrin. LDL cholesterol was calculated using the Friedewald equation.

PSG

PSGs were performed using a computerised diagnostic system (Alice 3 or 5; Respironics, Pittsburgh, PA, USA) in the Sleep Laboratory [2]. Manual scoring of sleep and respiratory events was performed according to standard criteria [23, 24].

ELISA for quantitative measurement of human A-FABP

Serum A-FABP levels were measured using a human A-FABP ELISA kit (Biovendor Laboratory Medicine Inc., Modrice, Czech Republic). Serum samples were diluted 1:10 before assay, then 100 μL of diluted sera, calibrators and quality-control samples were applied to 96-well microtitre plates coated with an affinity-purified goat anti-human FABP antibody. A calibration curve was constructed by plotting the absorbance values at 450 nm *versus* the A-FABP concentrations of the calibrators, and concentrations of unknown samples were found using this calibration curve. The intra- and interassay variations were 3.7/6.4% and 2.6/5.3% for high/low quality controls, respectively [25].

Data analysis

Sleep parameters, including apnoea/hypopnoea index (AHI), duration of desaturation, arousal index and minimum oxygen saturation, were log or logit-transformed as appropriate for further statistical analysis. Pearson correlation coefficients and partial correlation coefficients with adjustment for age, waist and BMI, were used to examine the association between serum A-FABP levels with various anthropometric, biochemical and sleep parameters. Comparisons of group means were performed using unpaired t-tests, with assumption of unequal variance (for comparison between any two groups) and using one-way ANOVA (for multiple group comparison) where appropriate. Matrix contrast examination was performed where significant variation in univariate ANOVA was seen, to examine for between-group effects. All data are expressed as mean ± SD and p-values < 0.05 were considered to be statistically significant.

RESULTS

A total of 128 males, recruited between October 2002 and March 2006, were analysed. Four subjects were excluded due to either extreme hyperlipidaemia (total cholesterol 9.0 mmol·L⁻¹ and triglyceride 25.6 mmol·L⁻¹) or insufficient total sleep time (<240 mins) at PSG. The demographic characteristics of the subjects are summarised in table 1. Of the subjects, 96 (77.4%) had OSA with an AHI ≥ 5 events·h⁻¹.

Serum A-FABP levels showed significant correlations with waist circumference and BMI (table 2). On controlling for the effects of age and obesity (BMI and waist circumference),



EUROPEAN RESPIRATORY JOURNAL VOLUME 33 NUMBER 2 347

SERUM A-FABP IN OSA D.C-L. LAM ET AL.

TABLE 1

Clinical characteristics, sleep parameters and adipocyte-fatty acid binding protein (A-FABP) of the recruited patients and between-group comparisons of the tertile groups

	Overall	AHI events·h ⁻¹			p-values#
		<13.2	13.2–34.4	≥34.4	
Subjects n	124	41	42	41	
Age yrs	44.5 ± 9.7	40.2 ± 8.2	47.4 ± 9.3	45.9 ± 10.2	0.001
Waist circumference cm	93.8 ± 9.5	92.5 ± 9.0	91.9 ± 9.2	97.2 ± 9.5	0.021
BMI kg·m ⁻²	27.0 ± 4.0	26.5 ± 3.4	26.1 ± 3.8	28.4 ± 4.4	0.020
Current/ex-smokers n	56	16	19	21	0.540
Current/ex-alcohol drinkers n	64	20	25	19	0.440
AHI [¶] events⋅h ⁻¹	27.6 ± 24.3	4.3 ± 3.7	20.4 ± 5.7	58.2 ± 13.9	<0.001§
Duration of Sa,O ₂ <90% mins [¶]	45.7 ± 69.3	3.2 ± 4.2	20.8 ± 23.0	113.6 ± 83.6	<0.001§
Arousal index [¶]	30.5 ± 20.0	14.4 ± 6.9	26.9 ± 9.3	50.3 ± 20.4	<0.001§
Minimum Sa,O ₂ % ⁺	74.4 ± 14.0	84.7 ± 7.4	76.8 ± 8.6	62.0 ± 14.2	<0.001§
A-FABP level ng·mL ⁻¹	16.6±6.9	15.6 ± 5.6	15.1 ± 5.4	19.3 ± 8.7	0.030§

Data are presented in mean ±sp, unless otherwise stated. AHI: apnoea/hypopnoea index; BMI: body mass index; Sa,o2: oxygen saturation. #: between-group comparisons were performed using ANOVA; *1: log-transformed before analysis, *1: logit transformed before analysis. *5: significance taken at p<0.05 for between-group comparisons after adjustment for age, waist and BMI.

serum A-FABP levels still demonstrated significant correlations with the duration (mins) of oxygen desaturation (Sa,O2) <90% during sleep (r=0.293, p=0.001) and the minimum S_{a,O_2} (r= -0.255, p=0.004). Serum A-FABP also had significant correlations with fasting serum insulin levels (r=0.343, p<0.001) and HOMA-IR (r=0.333, p<0.001; table 2). There was no significant correlation between serum A-FABP levels and serum lipid levels, with the exception of serum triglycerides (r=0.284, p=0.001; table 2).

The whole group of study subjects was divided into three groups according to their AHI in tertiles, as follows. AHI Group 1: <13.2 events·h⁻¹; AHI Group 2: 13.2–34.4 events·h⁻¹; AHI Group 3: $\geqslant 34.4$ events·h⁻¹ (table 1). After adjustment for age, waist and BMI, between-group comparisons showed significant differences in AHI (p<0.001), duration of Sa,O2 <90% (p<0.001), arousal index (p<0.001), minimum S_{a,O_2} level (p<0.001) and serum A-FABP levels (p=0.030; table 1). Contrast examination (K matrix) to test for between-group

TABLE 2 Correlations of adipocyte-fatty acid binding protein (A-FABP) with studied parameters, adjusted for waist and body mass index (BMI)

	A-FABP ng⋅mL ⁻¹				
	Pearson correlation		Partial correlation#		
	r-values	p-values	r-values	p-values	
Age yrs	-0.104	0.252			
Waist circumference cm	0.304 [§]	0.001 §			
BMI kg·m ⁻²	0.244 [§]	0.006§			
AHI [¶] events·h ⁻¹	0.154	0.087	0.139	0.130	
Duration of Sa,O ₂ <90% mins [¶]	0.293 [§]	0.001§	0.254	0.005\$	
Arousal index [¶]	0.104	0.248	0.157	0.085	
Minimum Sa,O ₂ % ⁺	-0.255 [§]	0.004 [§]	-0.230	0.012 [§]	
Cholesterol mmol·L ⁻¹	0.146	0.105	0.131	0.152	
Triglyceride [¶] mmol·L ⁻¹	0.284 [§]	0.001 §	0.241	0.008§	
LDL mmol·L ⁻¹	0.013	0.889	-0.027	0.765	
HDL [¶] mmol·L ⁻¹	0.024	0.794	0.112	0.222	
Fasting glucose [¶] mmol·L ⁻¹	0.095	0.295	0.098	0.285	
Fasting serum insulin level pmol·L ⁻¹	0.343 [§]	<0.001 [§]	0.210	0.021 §	
HOMA-IR [¶]	0.333 [§]	<0.001 [§]	0.207	0.022§	

AHI: apnoea/hypopnoea index; Sa,O2: oxygen saturation; LDL: low-density lipoprotein; HDL: high-density lipoprotein; HOMA-IR: Homeostasis Model Assessment – Insulin Resistance. #: adjusted for age, BMI and waist circumference; 1: log-transformed before analysis; 1: logit transformed before analysis. 5: significance taken at p<0.05.

D.C-L. LAM ET AL. SERUM A-FABP IN OSA

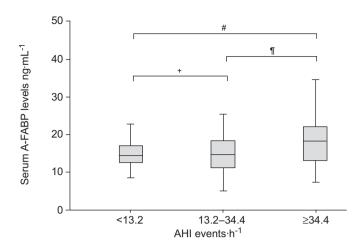


FIGURE 1. Significant difference in serum adipocyte-fatty acid binding protein (A-FABP) levels between tertile groups divided according to apnoea/hypopnoea index (AHI). Boxes show the median and the whiskers show the range. $^{\#}$: p=0.018; $^{\$}$: p=0.026; $^{+}$: p=0.812.

differences showed that there were significant differences in serum A-FABP levels between AHI Group 3 and Group 1 (p=0.018), and between AHI Group 3 and Group 2 (p=0.026), but not between AHI Group 1 and Group 2 (p=0.812; fig. 1).

Based on the difference between AHI Group 3 and the other AHI groups, an AHI of $34.4 \text{ events} \cdot \text{h}^{-1}$ was chosen as a cut-off level for further comparison. When comparing AHI Group 3 subjects (n=40) with waist-matched subjects with an AHI $<34.4 \text{ events} \cdot \text{h}^{-1}$ (n=40), serum A-FABP levels in AHI Group 3 subjects were significantly higher (p=0.036; table 3).

DISCUSSION

The present study demonstrates that serum A-FABP levels were significantly raised in otherwise healthy males with severe OSA compared with subjects with no or less severe OSA. A-FABP levels also showed significant correlations with sleep hypoxaemia parameters, including duration of oxygen

TABLE 3

Serum adipocyte-fatty acid binding protein (A-FABP) level in subjects with severe obstructive sleep apnoea (apnoea/hypopnoea index (AHI) ≥34.4 events·h⁻¹) compared with waist-matched subjects with an AHI <34.4 events·h⁻¹

	AHI eve	p-values	
	≽34.4	<34.4	
Subjects n	40	40	
Age yrs	44.4 ± 8.8	46 ± 10.2	0.447
Waist circumference cm#	95.9 ± 8.7	95.6 ± 8.6	0.867
BMI kg·m ⁻²	27.8 ± 4.0	27.4 ± 3.4	0.462
Serum A-FABP levels ng·mL ⁻¹	19.2 ± 8.9	15.9 ± 4.7	0.036 [¶]

Data presented as mean \pm sp. #: waist circumferences were matched to ± 5 cm. \P : significance taken at p<0.05 with between-group comparisons after adjustment for age, body mass index (BMI) and waist circumference.

desaturation and minimal oxygen saturation, independent of age and obesity. Increasing serum A-FABP levels were also associated with increased insulin resistance, independent of age and adiposity.

There is increasing evidence in cellular and animal experiments that A-FABP might play a central role in insulin resistance and glucose intolerance [10, 26]. In animal experiments, the apparent function of A-FABP was mediation of insulin resistance, independent of the effects of obesity and the promotion of atherosclerosis [9, 27]. A-FABP appeared to assume an important position in influencing metabolic responses, and hence lipid and glucose metabolism, and the mediation of subsequent development of atherosclerosis. The present results confirmed that in this cohort of males, the majority of whom had OSA of varying degrees of severity, serum A-FABP level also correlated with waist circumference/ BMI, as previously reported in the general population [18]. The subsequent partial correlation analyses were therefore performed with adjustment for waist circumference and BMI (table 2). The comparisons between groups of different OSA severity, determined according to AHI tertiles, suggest that the difference in serum A-FABP levels mainly existed between those with high AHI of $\geqslant 34.4$ events·h⁻¹ and those with AHI <34.4 events·h⁻¹, while there was no difference between subjects whose OSA would be considered as moderate compared to those with mild/no OSA. There was also significant positive linear correlation between FABP levels and the severity of oxygen desaturation, independent of obesity. These findings suggests that there may be a threshold effect of OSA on FABP levels and that FABP levels may be upregulated mainly in those subjects with high AHI and a more severe degree of hypoxaemia. There is scanty information regarding the regulation of FABP, although there has been previous evidence of FABP expression being regulated by hypoxia in human trophoblasts in vitro [28]. It is possible that the different intensity of intermittent hypoxia in OSA subjects results in different degrees of heightened expression of A-FABP, which then assumes a mediating role between OSA and metabolic dysfunction.

The significant correlations between serum A-FABP levels and insulin resistance remained after adjustment for obesity, simulating the finding in animal experiments that serum A-FABP deficiency protected against development of insulin resistance, independent of the effects of obesity [14]. An independent association between OSA and insulin resistance has been reported and multiple mechanisms may be involved, in particular intermittent hypoxia, sleep fragmentation, sympathetic overactivity or even activation of the hypothalamic pituitary axis [3, 29, 30]. Multiple biomediators may be involved and A-FABP may well be one of them. As yet, little is known about the regulation of A-FABP. However, the relationships between these pathophysiological mechanisms in OSA and the regulation of A-FABP, and their respective contributions towards the development of insulin resistance, are beyond the scope of the present study's design.

Several strengths and limitations of the current study warrant further discussion. The results of the current study were based on male subjects and should not be generalised to female subjects. Furthermore, only subjects without cardiometabolic



SERUM A-FABP IN OSA

D.C-L. LAM ET AL.

comorbidities were recruited to avoid the confounding effects of these diseases; this is considered one of the strengths of the present study, given that the major objective was to investigate for any independent relationship between OSA and A-FABP. However, in clinical practice, many OSA subjects will have overt metabolic syndrome or cardiovascular diseases, and the current findings cannot be entirely extrapolated. Another potential limitation is the use of a sleep clinic-based cohort, which may introduce potential sample bias, though this would have the advantage of including a significant number of subjects with a wide range of OSA severity within a not-toolarge sample size. Although the results from the present study were being controlled for age, BMI and waist circumference, it would be more definitive to assess body fat content with more sophisticated and detailed measurements. Given the finding of an independent association between A-FABP levels and severity of OSA in the present study, further definitive information on the cause-effect relationship and interactive pathways may be derived by studying the effects of continuous positive airway pressure treatment of OSA on serum A-FABP levels, as well as in vitro studies using cell culture or animal models.

Conclusion

In summary, the present authors have demonstrated significant differences in serum adipocyte-fatty acid binding protein levels between subjects with severe obstructive sleep apnoea compared to those with no or mild obstructive sleep apnoea in terms of apnoea/hypopnoea index. Serum adipocyte-fatty acid binding protein levels were also found to correlate with nocturnal hypoxaemia and insulin resistance, independent of adiposity. The findings support the hypothesis that adipocyte-fatty acid binding protein could be a link in the association between obstructive sleep apnoea and cardiometabolic dysfunction.

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350 VOLUME 33 NUMBER 2 EUROPEAN RESPIRATORY JOURNAL

D.C-L. LAM ET AL. SERUM A-FABP IN OSA

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EUROPEAN RESPIRATORY JOURNAL VOLUME 33 NUMBER 2 351