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Serum concentrations of resistin-like molecules β and γ are elevated in high-fat-fed and obese *db/db* mice, with increased production in the intestinal tract and bone marrow

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Abstract *Aims/hypothesis:* Resistin and the resistin-like molecules (RELMs) comprise a novel class of cysteine-rich proteins. Among the RELMs, RELM β and RELM γ are produced in non-adipocyte tissues, but the regulation of their expression and their physiological roles are largely unknown. We investigated in mice the tissue distribution and dimer formation of RELM β and RELM γ and then examined whether their serum concentrations and tissue expression levels are related to insulin resistance. *Methods:* Specific antibodies against RELM β and RELM γ were generated. Dimer formation was examined using COS cells and

the colon. RELM β and RELM γ tissue localisation and expression levels were analysed by an RNase protection assay, immunoblotting and immunohistochemical study. Serum concentrations in high-fat-fed and *db/db* mice were also measured using the specific antibodies. *Results:* The intestinal tract produces RELM β and RELM γ , and colonic epithelial cells in particular express both RELM β and RELM γ . In addition, RELM β and RELM γ were shown to form a homodimer and a heterodimer with each other, in an overexpression system using cultured cells, and in mouse colon and serum. Serum RELM β and RELM γ levels in high-fat-fed mice were markedly higher than those in mice fed normal chow. Serum RELM β and RELM γ concentrations were also clearly higher in *db/db* mice than in lean littermates. Tissue expression levels revealed that elevated serum concentrations of RELM β and RELM γ are attributable to increased production in the colon and bone marrow. *Conclusions/interpretation:* RELM β and RELM γ form homo/heterodimers, which are secreted into the circulation. Serum concentrations of RELM β and RELM γ may be a novel intestinal-tract-mediating regulator of insulin sensitivity, possibly involved in insulin resistance induced by obesity and a high-fat diet.

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Serum concentration

Abbreviations FIZZ: found in inflammatory zone · GST:
glutathione S-transferase · PPAR γ : peroxisome
proliferator-activated receptor γ · RELM: resistin-like
molecules · XCP: ten-cysteine protein

Introduction

Type 2 diabetes is characterised by insulin resistance of peripheral tissues such as the liver and muscle and adipose tissue [1–4]. Recent studies have indicated that adipose

tissue is, in addition to being a lipid storage site, an endocrine organ producing hormones, cytokines and other substances [5–7]. Recently, resistin was newly identified as an adipocyte-secreted protein [8]. Serum resistin levels are reportedly elevated in genetically obese mice and are down-regulated by administration of thiazolidinediones [9, 10], peroxisome proliferator-activated receptor γ (PPAR γ) agonists, although contradictory data also exist [11]. Resistin has been demonstrated to antagonise insulin action in cultured cells such as 3T3-L1 adipocytes, as well as in rodents [8]. Resistin knock-out mice exhibited lower blood glucose with reduced hepatic glucose production [12]. Thus resistin could be one of the important adipokines causing insulin resistance.

The presence of resistin-like molecules (RELMs) indicates that resistin belongs to a novel family of cysteine-rich secreted proteins (RELM/found in inflammatory zone [FIZZ]/ten-cysteine protein [XCP]). RELM α /FIZZ1 [13] and RELM β /FIZZ2 [14] are homologous with resistin/FIZZ3 and expressed mainly in white adipose tissue and the colon respectively [14, 15]. The administration of RELM β to rats resulted in acute impairment of hepatic insulin sensitivity and glucose metabolism [16], which suggested RELM β is a link between the intestine and hepatic insulin action. Finally, RELM γ /FIZZ4/XCP1, a fourth member of the RELM/FIZZ/XCP family, was identified as a gene with decreased expression in rat nasal respiratory epithelium exposed to cigarette smoke [17]. RELM γ was expressed in the bone marrow, spleen, pancreas and colon, and was revealed to play a role as a cytokine in haematopoiesis [18, 19].

The consensus structure of the RELMs is composed of two domains; one half is the N-terminal, including an N-terminal signal sequence and a variable middle portion, and the other half is the C-terminal domain, which has a highly conserved C-terminal signature sequence containing a unique spacing of the cysteine residues. The N-terminal domains of RELM α , RELM β and RELM γ are 15, 35 and 17% identical to resistin, whereas the C-terminal domains are 47, 54 and 52% identical to resistin. The C-terminus of RELM γ is highly homologous (84%) with RELM β . Recently, the crystal structures of resistin and RELM β were revealed to have a unique multimeric structure [20]. Each protomer is comprised of a 'head' and 'tail' segment and circulates as an assembly of hexamers and trimers, which reflect activation of resistin.

We previously identified RELM γ cDNA independently by PCR using degenerated oligonucleotide primers, and investigated its tissue distribution. Since we succeeded in preparing highly specific antibodies against RELM β and RELM γ , we were able to detect the endogenous proteins and measure serum concentrations as well as tissue contents of RELM β and RELM γ in the mouse. In this study, we show for the first time that RELM β and RELM γ form not only a homodimer but also a heterodimer with each other in both tissue and serum. Interestingly, we also found that the serum concentrations of RELM β and RELM γ are significantly elevated in high-fat-diet-induced and obese diabetic mice. These observations are probably attribut-

able to increased production in the colon and bone marrow. Thus, this report is the first to raise the possibility of a novel intestinal-tract-mediating regulatory mechanism of insulin sensitivity, which may be involved in insulin resistance induced by obesity and a high-fat diet.

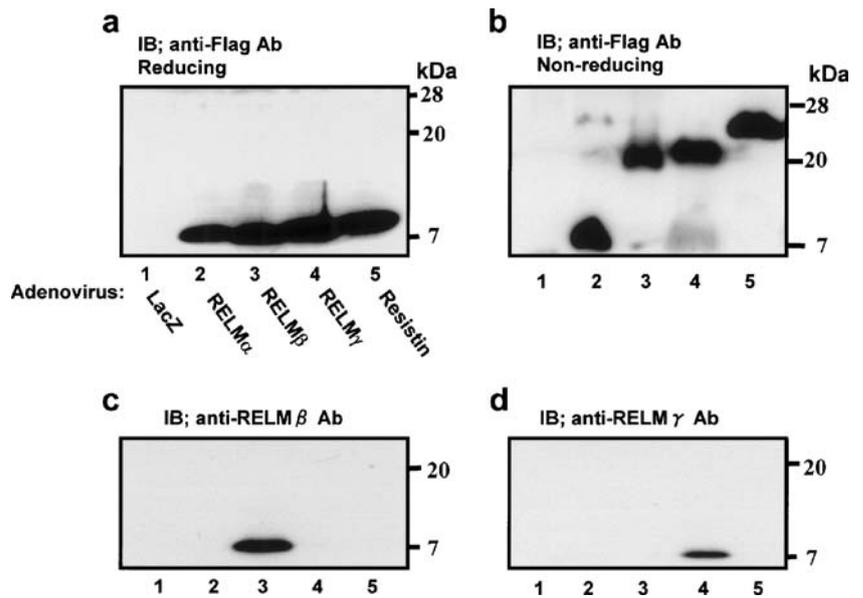
Materials and methods

cDNA cloning of a novel RELM/FIZZ isoform Two degenerate oligonucleotide primers were synthesised for PCR. These primers were 5'-ATGAAGA/CCTACAA/CC/TT/GTGTTTC/TC-3' and 5'-TTAG/AGA/CCAG/TTT/CGGC/GCAGCG-3', corresponding to amino acid residues 1–8 and 104–111 of RELM α / β , which are highly conserved among RELM/FIZZ isoforms. PCR was performed using mouse embryonic cDNA. A DNA fragment of approximately 330 bp was then separated by electrophoresis, cloned into TA vector pCRII (Invitrogen, San Diego, CA, USA), and sequenced using a DNA sequencer. We obtained two independent sequences: one, F1, turned out to be completely homologous to mouse RELM α /FIZZ1; the other, F2, encoded a protein to mouse RELM γ /FIZZ4. Then, a mouse embryonic cDNA library produced by a standard method (Stratagene, La Jolla, CA, USA) was screened under standard hybridisation conditions using a 32 P-labelled F2 cDNA fragment as a probe to obtain a full-length cDNA encoding RELM γ /FIZZ4. Positive clones were excised into pBluescript and sequenced.

RNase protection assay of various tissues Mice were killed by cervical dislocation, and various tissues (cerebrum, cerebellum, brainstem, heart, lung, liver, oesophagus, stomach, jejunum, ileum, colon, kidney, testis, spleen, pancreas, abdominal fat, epididymal fat, muscle, aorta, femoral bone marrow) were removed. Total RNA was isolated with Isogen (Nippon Gene, Japan). [α - 32 P]UTP-labelled RNA probes were prepared using nucleotides 1–200 of mouse RELM β , and 83–315 of mouse RELM γ as templates. An RNase protection assay was performed using an RPA III kit (Ambion, Austin, TX, USA) according to the manufacturer's instructions.

Preparation of the antibodies An antibody against the whole mouse resistin molecule was prepared by immunising rabbits with the recombinant mouse resistin protein, obtained as described previously [21]. Sequences corresponding to nucleotides 11–173 of RELM β /FIZZ2 and 47–227 of RELM γ /FIZZ4 were amplified by PCR and cloned into the pGEX-3T expression vector (Pharmacia, Piscataway, NJ, USA). Glutathione S-transferase (GST) fusion proteins (GST-RELM β and GST-RELM γ) were prepared according to the manufacturer's instructions (Pharmacia). The antisera were raised by immunising rabbits with GST-RELM β and GST-RELM γ . From these antisera, an antibody against GST protein was removed by filtering through Affigel-10 covalently coupled to GST proteins. Then, specific antibodies against RELM β and RELM γ were affinity purified with Affigel-10 covalently

Fig. 1 Immunoblotting of RELMs secreted by COS7 cells. The four cDNAs coding RELM α , RELM β , RELM γ and resistin, with the Flag tag at their C-termini, were expressed into COS7 cells and the medium from each cell type was subjected to SDS-PAGE under reducing (a, c and d) and non-reducing (b) conditions and immunoblotted (IB) with anti-Flag (a, b), anti-RELM β (c) and anti-RELM γ (d) antibody (Ab)



coupled to GST-RELM β and GST-RELM γ respectively. Antibodies against Flag tag were purchased from Upstate Biotech, Inc. (Lake Placid, NY, USA).

Immunohistochemistry Intestinal tissues removed from the mice were fixed in 10% phosphate-buffered formalin (pH 7.4) and embedded in paraffin. After sectioning, the tissues were dewaxed in ethanol, rehydrated in 10 mmol/l citric acid buffer, and microwaved for 13 min. The tissue sections were blocked with serial incubation in 30% H₂O₂, avidin D-blocking reagent, biotin-blocking reagent and protein-blocking reagent. The sections were then incubated with an affinity-purified polyclonal antibody for murine RELM β and RELM γ or control serum at a 1:100 dilution overnight at 4°C. After washing with PBS, the slides were incubated with a biotinylated goat anti-rabbit secondary antibody followed by detection of horseradish peroxidase.

Animal studies Nine-week-old male mice (C57BL/6J) were purchased from Jackson Laboratories (Bar Harbor, ME, USA). They were divided into two groups. One group ($n=9$) was maintained on standard rodent chow, the other ($n=9$) was fed a high-fat diet (60% fat, 25% carbohydrate and 15% protein). Genetically obese *db/db* mice ($n=6$) and lean littermates ($n=6$) were purchased from Jackson Laboratories. They were maintained on standard rodent chow. Tissues from the mice were homogenised in ice-cold lysis buffer. Insoluble materials were removed and the cell lysates were incubated for 2 h at 4°C with the indicated antibody. Blood samples were centrifuged at 3,000 rev/min for 20 min and sera containing equal amounts of protein were used for immunoprecipitation. Immunoblotting against these immunoprecipitates was performed as previously described [21]. Animal care and procedures of the experiments were approved by the Animal Care Committee of the University of Tokyo.

Statistical analysis Data are expressed as means \pm SE. Comparisons were made using unpaired *t*-tests. Serum RELM levels were compared with body weight, serum glucose and insulin levels by Pearson's correlation. Values of $p<0.05$ were considered significant.

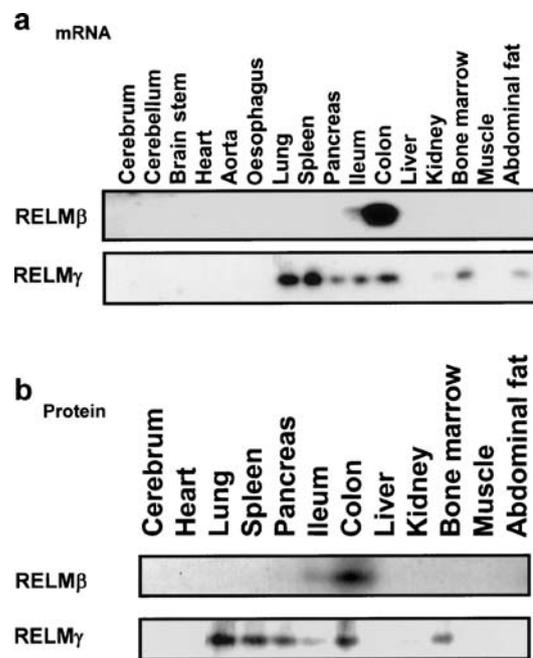
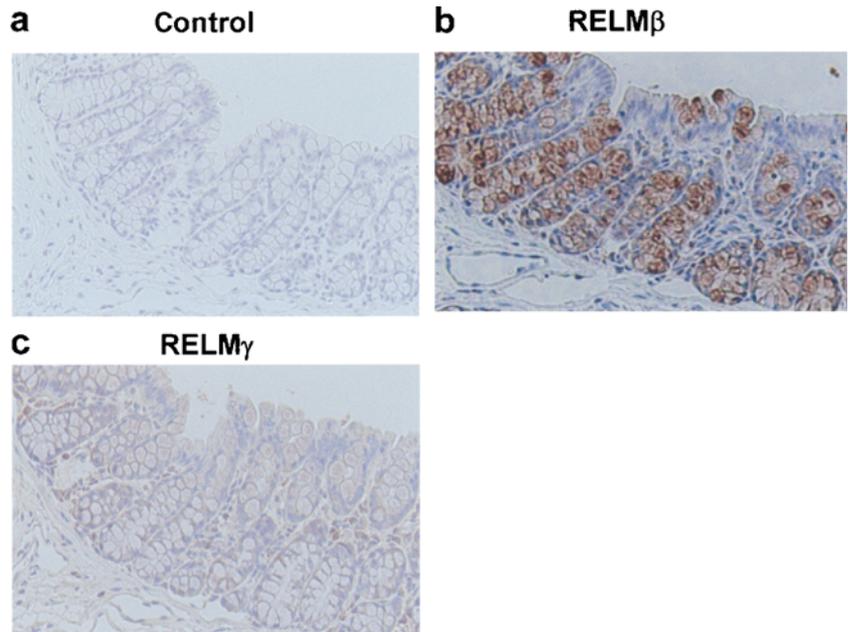


Fig. 2 RELM β and RELM γ expression in various tissues. **a** 10 μ g of RNA from various mouse tissues were prepared and hybridised with [α -³²P]UTP-labelled RNA probes for RELM β or RELM γ . The RNase protection assay was performed using an RPA III kit (Ambion) according to the manufacturer's instructions. **b** Cell lysates from various mouse tissues were prepared and then immunoprecipitated and immunoblotted with anti-RELM β antibody. Cell lysates from tissues were prepared and then immunoprecipitated and immunoblotted with anti-RELM γ antibody

Fig. 3 Immunohistochemical study of RELM β and RELM γ . Immunohistochemical study of the colon with control antibody (a) confirmed the specificities of specific antibodies ($\times 200$). The avidin–biotin–peroxidase complex method with antibodies against RELM β (b) and RELM γ (c) was used to detect cells expressing these RELMs. RELM β and RELM γ staining is brown, that of the control blue



Results

Cloning of RELM γ cDNA To obtain a cDNA fragment corresponding to a novel isoform of RELM/FIZZ, PCR was performed using degenerate oligonucleotides as primers and mouse embryonic cDNA as a substrate. PCR products, with a length of approximately 300–350 bp, were separated, subcloned and sequenced. Isolated PCR products were shown to consist of two different cDNAs. One corresponded to RELM α cDNA [14], the other to RELM γ . The full-length cDNAs encoding resistin, RELM α and RELM β were prepared by PCRs based on the reported sequences. The full-length cDNA encoding RELM γ was obtained by screening a cDNA library.

Overexpression of RELMs in COS7 cells and preparation of specific antibodies against RELM β and RELM γ Four cDNAs encoding RELM α , RELM β , RELM γ and resistin, with the Flag tag at their C-termini, were ligated into adenovirus expression vectors. COS7 cells were infected with these adenoviruses to achieve similar protein expression levels, as assessed by immunoblotting using anti-Flag antibody. The media from cells transfected with these adenoviruses were subjected to SDS-PAGE and immunoblotted with anti-Flag antibody (Fig. 1a). The expression of these proteins was observed as very similar band densities of 7–12 kDa. Thus, it was clear that RELM γ was secreted into the media, like other members of the RELM family, when expressed in COS7 cells.

Next, we investigated the electrophoretic mobilities of these RELMs under non-reducing conditions. As shown in Fig. 1b, RELM β , RELM γ and resistin migrated with apparent molecular masses twice those of the respective monomers, whereas RELM α behaved as a monomer. Taking previous [22, 23] and current results into account, it was confirmed that resistin, RELM β and RELM γ form a

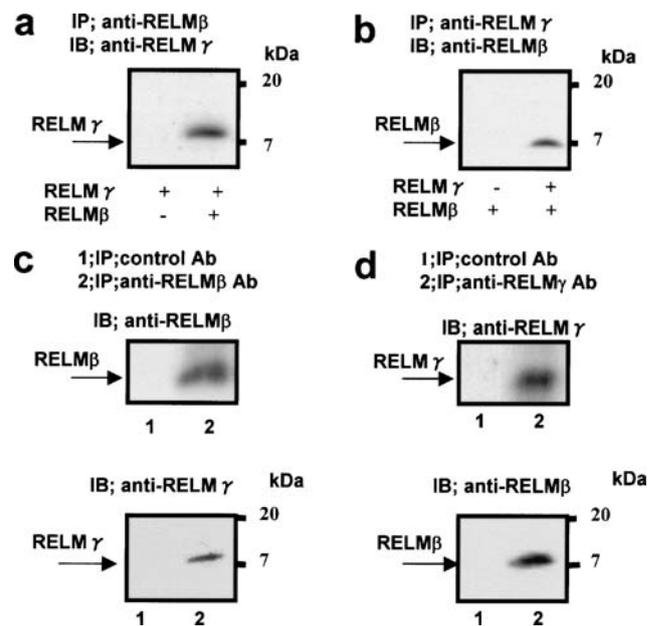


Fig. 4 Heterodimer formations of RELM β /RELM γ (a, b) in COS7 cells and endogenous RELM β /RELM γ in colon (c, d). a Secreted RELM γ , co-expressed with (right lane) or without (left lane) RELM β , was immunoprecipitated (IP) with anti-RELM β antibody (Ab) and then immunoblotted with anti-RELM γ Ab. b RELM β , co-expressed with (right lane) or without (left lane) RELM γ was immunoprecipitated with anti-RELM γ Ab and immunoblotted with anti-RELM β Ab. c Colon cell lysates were immunoprecipitated with control IgG (lane 1) and anti-RELM β Ab (lane 2) and then immunoblotted with anti-RELM β Ab (upper panel) and anti-RELM γ Ab (lower panel). d The colon cell lysates were immunoprecipitated with control IgG (lane 1) and anti-RELM γ Ab (lane 2) and immunoblotted with anti-RELM γ Ab (upper panel) or anti-RELM β Ab (lower panel)

Table 1 Serum RELM β and RELM γ in high-fat-fed mice

	4 weeks		8 weeks		12 weeks	
	Control	High-fat	Control	High-fat	Control	High-fat
RELM β	1 \pm 0.05	1.08 \pm 0.06	1 \pm 0.05	1.74 \pm 0.08 ^a	1 \pm 0.06	2.16 \pm 0.09 ^a
RELM γ	1 \pm 0.05	1.05 \pm 0.05	1 \pm 0.06	2.00 \pm 0.10 ^a	1 \pm 0.05	2.70 \pm 0.12 ^a
RELM β/γ	1 \pm 0.05	0.97 \pm 0.06	1 \pm 0.05	1.65 \pm 0.09 ^a	1 \pm 0.05	2.54 \pm 0.11 ^a
Body weight (g)	18.3 \pm 1.18	19.3 \pm 1.29	26.6 \pm 1.32	38.8 \pm 2.09 ^a	30.1 \pm 2.07	44.4 \pm 1.95 ^a
Glucose (mmol/l)	5.03 \pm 0.10	5.11 \pm 0.20	5.24 \pm 0.16	6.77 \pm 0.30	5.13 \pm 0.08	7.93 \pm 0.27 ^a
Insulin (pmol/l)	22.4 \pm 1.67	22.7 \pm 2.64	26.0 \pm 5.00	48.7 \pm 2.7 ^a	33.1 \pm 1.67	62.9 \pm 2.45 ^a

Male 12-week-old C57BL/6J mice were fed normal chow ($n=9$) or a high-fat diet ($n=9$) from 4 to 12 weeks of age. Sera were immunoprecipitated and then immunoblotted with RELM β and RELM γ antibody. The data are shown as ratios to the control diet values and expressed as means \pm SE

^a $p<0.01$ relative to control-diet-fed mice

disulphide-linked dimer, while RELM α exists mainly as a monomer.

Immunoblotting with anti-RELM β and anti-RELM γ antibodies (Fig. 1c, d) revealed that these antibodies do not recognise other members of the RELM/FIZZ family, suggesting anti-RELM β and anti-RELM γ antibodies to be highly specific for the corresponding isoforms.

Tissue distribution of RELM β and RELM γ We next evaluated RELM mRNA expression with an RNase protection assay (Fig. 2a) and protein expression using Western blotting (Fig. 2b) in various mouse tissues. RELM β mRNA and protein were abundant in the colon, and to a lesser extent in the ileum. On the other hand, RELM γ mRNA

was abundant in the colon, ileum, bone marrow, spleen, pancreas and fat, consistent with previous reports [18–20] (Fig. 2a, lower panel). RELM γ protein expression is also detectable in the colon, ileum, bone marrow, spleen and pancreas (Fig. 2b, lower panel). Thus, we found that the colon and ileum express both RELM β and RELM γ , and the localisations of RELM β and RELM γ were investigated by immunohistochemical staining of the colon (Fig. 3). It was demonstrated that epithelial cells throughout the crypt and surface of the colon express both RELM β and RELM γ , while no significant staining was observed with the control antibody. The highest level of RELM β expression was observed in goblet cells of the colon, consistent with previous reports [15, 24], and RELM γ protein

Fig. 5 Relationships between serum RELM β and RELM γ concentration and body weight, glucose and insulin in high-fat-fed mice. Male 12-week-old C57BL/6J mice were fed normal chow ($n=9$) or a high-fat diet ($n=9$) from 4 to 12 weeks of age. RELM β , RELM γ and various parameters were measured at 4, 8 and 12 weeks of age. The data are plotted as the percentage of the means of RELM β and RELM γ correlated positively with body weight (**a**, $p<0.0001$; **b**, $p<0.0005$), glucose (**c**, $p<0.0005$; **d**, $p<0.0005$) and insulin (**e**, $p<0.001$; **f**, $p<0.0005$)

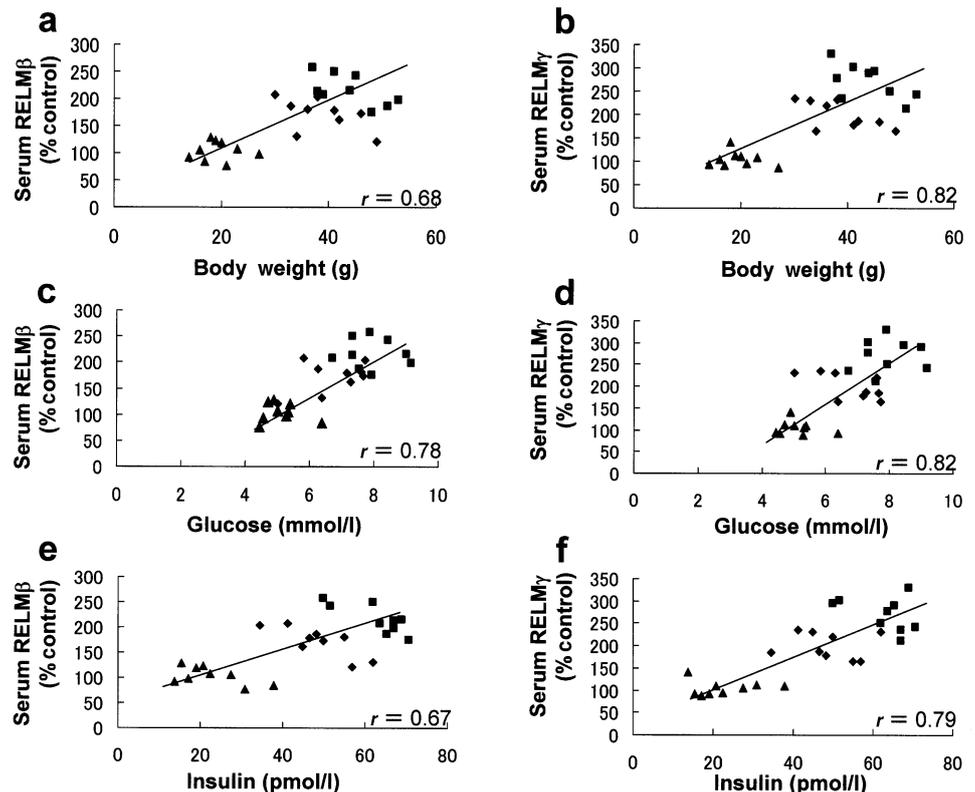


Table 2 Expression and regulation of mRNA and protein of RELM β and RELM γ in high-fat-fed mice

		Colon		Bone marrow		Spleen		Lung		Pancreas	
		Control	High-fat	Control	High-fat	Control	High-fat	Control	High-fat	Control	High-fat
RELM β	mRNA	1 \pm 0.04	1.97 \pm 0.04 ^a	–	–	–	–	–	–	–	–
	Protein	1 \pm 0.03	1.96 \pm 0.05 ^a	–	–	–	–	–	–	–	–
RELM γ	mRNA	1 \pm 0.04	1.84 \pm 0.03 ^a	1 \pm 0.04	1.89 \pm 0.05 ^a	1 \pm 0.04	1.03 \pm 0.05	1 \pm 0.04	1.05 \pm 0.05	1 \pm 0.04	0.97 \pm 0.05
	Protein	1 \pm 0.04	2.04 \pm 0.03 ^a	1 \pm 0.04	1.78 \pm 0.05 ^a	1 \pm 0.03	1.02 \pm 0.04	1 \pm 0.04	1.03 \pm 0.04	1 \pm 0.04	0.98 \pm 0.05

Male 12-week-old C57BL/6J mice were fed normal chow or a high-fat diet from 4 to 12 weeks of age. RNase protection assay and immunoblotting show RELM β and RELM γ expression in the colon, bone marrow, spleen, lung and pancreas. The data are shown as ratios to the control diet values and expressed as means \pm SE

^a p <0.01 relative to control-diet-fed mice

was also shown to be localised in goblet cells. In the bone marrow, about 30% of haematopoietic cells were stained with the anti-RELM γ antibody and these cells were myelocytes and metamyelocytes or neutrophils (data not shown), consistent with previous reports [19, 20].

Heterodimer formation between RELM β and RELM γ in COS7 cells and tissues Since colonic cells express both RELM β and RELM γ , we subsequently investigated whether these RELMs form a heterodimer. The RELM β and RELM γ antibodies were highly specific and did not immunoprecipitate the respective isoforms of RELM (data not shown). As shown in Fig. 4a, b, when RELM β and RELM γ were co-expressed, RELM γ or RELM β was detected in the RELM β or RELM γ immunoprecipitates respectively. These results suggest that RELM β and RELM γ associate with each other and form a heterodimer.

Subsequently, we investigated whether or not endogenous RELM β and RELM γ form a heterodimer using the proximal colon. RELM β was detected in RELM γ immunoprecipitates but not in those of control antibodies (Fig. 4c). RELM γ was also detected in RELM β immunoprecipitates but not in those of control antibodies (Fig. 4d). These results indicate that the heterodimerisation between RELM β and RELM γ is physiological.

Increased expression of RELM β and RELM γ in high-fat-fed mice Male 12-week-old C57BL/6J mice were fed normal chow or a high-fat diet from 4 to 12 weeks of age. A

high-fat diet resulted in body weight, serum glucose and insulin increasing time-dependently (30.1 \pm 6.21 vs 44.4 \pm 5.84 g, 5.13 \pm 0.24 vs 7.93 \pm 0.81 mmol/l, and 33.1 \pm 5.02 vs 62.9 \pm 7.35 pmol/l respectively) after 12 weeks of feeding (Table 1). Serum RELM β and RELM γ levels were increased by 116 and 170% respectively at the end of the 12-week feeding period. We detected the RELM β /RELM γ heterodimer in serum by immunoblotting of the anti-RELM β immunoprecipitate with anti-RELM γ antibody, and this heterodimer was also increased.

Serum RELM β and RELM γ and other parameters were measured at 4, 8 and 12 weeks after initiation of the high-fat diet. Close examination of these three sets of time-dependent data revealed serum RELM β and RELM γ to correlate positively with body weight (Fig. 5a, r =0.68, p <0.0001; Fig. 5b, r =0.82, p <0.0005), serum glucose concentration (Fig. 5c, r =0.78, p <0.0005; Fig. 5d, r =0.82, p <0.0005) and serum insulin (Fig. 5e, r =0.67, p <0.001; Fig. 5f, r =0.79, p <0.0005).

Subsequently, we investigated the expression levels of RELM β and RELM γ in tissues at the end of the 12-week feeding period. RELM β mRNA and protein levels in the distal colon of the high-fat-fed mice were elevated by 97 and 96% respectively (Table 2). Similarly, RELM γ mRNA and protein levels in the distal colon were elevated by 84 and 104% respectively. RELM γ mRNA and protein levels in the bone marrow were also elevated by 89 and 78% respectively. However, no significant alterations were observed in the spleen, lung or pancreas.

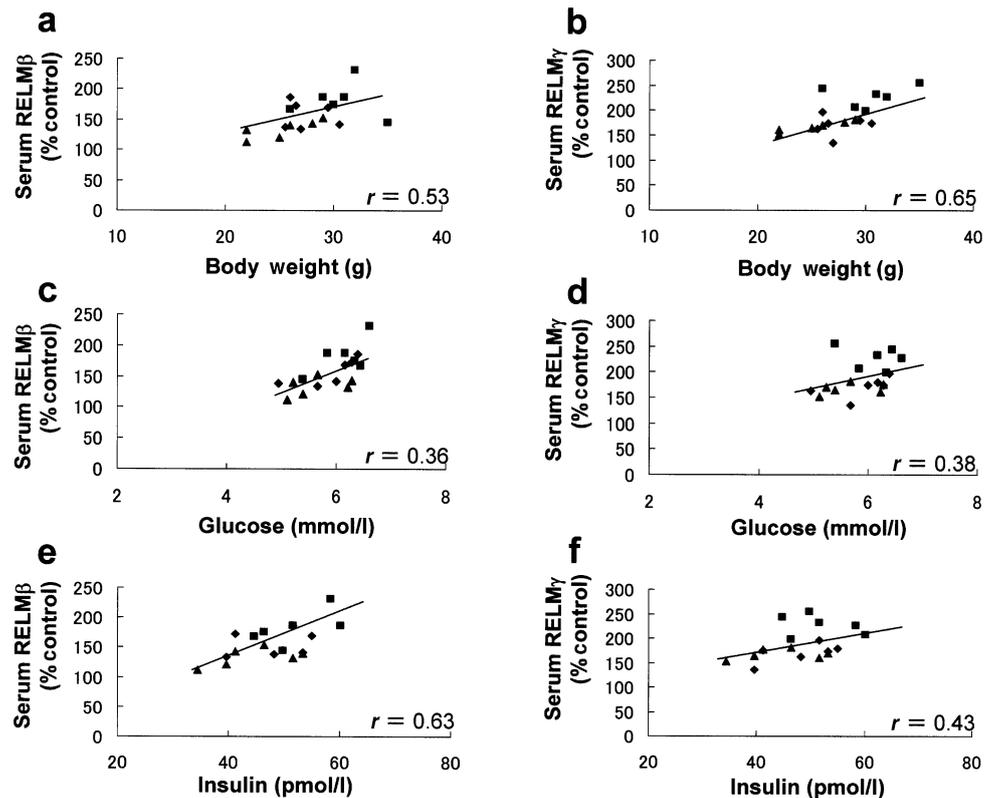
Table 3 Serum RELM β and RELM γ in *db/db* mice

	4 weeks		5 weeks		6 weeks		
	Control	<i>db/db</i>	Control	<i>db/db</i>	Control	<i>db/db</i>	
RELM β		1 \pm 0.05	1.34 \pm 0.06	1 \pm 0.06	1.57 \pm 0.09 ^a	1 \pm 0.07	1.82 \pm 0.11 ^a
RELM γ		1 \pm 0.06	1.68 \pm 0.04	1 \pm 0.04	1.70 \pm 0.08 ^a	1 \pm 0.04	2.27 \pm 0.09 ^a
RELM β / γ		1 \pm 0.04	1.48 \pm 0.05	1 \pm 0.05	1.58 \pm 0.08 ^a	1 \pm 0.05	1.79 \pm 0.06 ^a
Body weight (g)		19.8 \pm 0.90	25.3 \pm 0.92	21.0 \pm 1.00	27.5 \pm 1.13	23.2 \pm 0.92	30.1 \pm 1.24 ^a
Glucose (mmol/l)		4.71 \pm 0.12	5.65 \pm 0.20	4.94 \pm 0.12	5.91 \pm 0.22	4.62 \pm 0.07	6.13 \pm 0.18 ^a
Insulin (pmol/l)		21.4 \pm 1.64	44.4 \pm 2.99	22.0 \pm 1.74	48.2 \pm 2.63	26.3 \pm 1.68	51.9 \pm 2.57 ^a

Male lean littermates (n =6) and *db/db* mice (n =6) were fed a standard diet. Sera were immunoprecipitated and then immunoblotted with RELM β and RELM γ antibody. The data are shown as ratios to the control lean littermate values and expressed as means \pm SE

^a p <0.01 relative to lean littermates

Fig. 6 Relationships between serum RELM β and RELM γ concentrations and body weight, glucose and insulin in *db/db* mice. Male lean littermates ($n=6$) and *db/db* mice ($n=6$) were fed a standard diet. RELM β , RELM γ and other parameters were measured at 4, 5 and 6 weeks of age. The data are plotted as percentage of the means of control lean littermates. RELM β and RELM γ correlated positively with body weight (a, $p<0.05$; b, $p<0.01$). Neither RELM β nor RELM γ showed any correlation with glucose (c, d). RELM β (e, $p<0.01$), but not RELM γ (f), also positively correlated with insulin



*Increased expression of RELM β and RELM γ in *db/db* mice* Serum RELM β and RELM γ and other parameters in male *db/db* mice and their littermates were measured at 4, 5 and 6 weeks of age. Male 6-week-old *db/db* mice weighed more (30.1 ± 3.03 vs 23.2 ± 2.26 g), and had higher serum glucose (4.62 ± 0.16 vs 6.13 ± 0.45 mmol/l) and insulin (51.9 ± 6.29 vs 26.3 ± 4.11 pmol/l), than their lean littermates. Serum RELM β and RELM γ levels of *db/db* mice were higher and were increased by 82 and 127% respectively at the age of 6 weeks, as compared with the controls (Table 3). The RELM β /RELM γ heterodimer in serum was also increased. Serum RELM β and RELM γ correlated positively with body weight (Fig. 6a, $r=0.53$, $p<0.05$; Fig. 6b, $r=0.65$, $p<0.01$), but not with the serum glucose concentration (Fig. 6c, d). Serum insulin correlated positively with RELM β (Fig. 6e, $r=0.63$, $p<0.01$) but not with RELM γ (Fig. 6f).

RELM β mRNA and protein levels in the distal colons of *db/db* mice at the age of 6 weeks were elevated by 94 and 87% respectively (Table 4). Similarly, RELM γ mRNA and protein levels in the distal colon were elevated by 77 and 98% respectively (Table 4). RELM γ mRNA and protein levels in the bone marrow were elevated by 68 and 53% respectively. There were no significant changes in RELM γ levels in the spleen, lung or pancreas.

Discussion

Resistin and the three RELMs comprise a novel class of cysteine-rich proteins. Resistin is expressed exclusively in adipose tissues and reportedly causes insulin resistance [8]. RELM α was originally identified in broncho-alveolar

Table 4 Expression and regulation of mRNA and protein of RELM β and RELM γ in *db/db* mice

		Colon		Bone marrow		Spleen		Lung		Pancreas	
		Control	<i>db/db</i>	Control	<i>db/db</i>	Control	<i>db/db</i>	Control	<i>db/db</i>	Control	<i>db/db</i>
RELM β	mRNA	1 \pm 0.04	1.97 \pm 0.07 ^a	–	–	–	–	–	–	–	–
	Protein	1 \pm 0.05	1.87 \pm 0.07 ^a	–	–	–	–	–	–	–	–
RELM γ	mRNA	1 \pm 0.06	1.77 \pm 0.07 ^a	1 \pm 0.07	1.63 \pm 0.07 ^a	1 \pm 0.04	0.98 \pm 0.05	1 \pm 0.05	0.96 \pm 0.07	1 \pm 0.06	1.10 \pm 0.07
	Protein	1 \pm 0.05	1.98 \pm 0.07 ^a	1 \pm 0.05	1.53 \pm 0.07 ^a	1 \pm 0.06	1.02 \pm 0.07	1 \pm 0.05	1.05 \pm 0.07	1 \pm 0.06	1.02 \pm 0.07

Male lean littermates and *db/db* mice were fed a standard diet. RNase protection assay and immunoblotting show RELM β and RELM γ expression in the colon, bone marrow, spleen, lung and pancreas. The data are shown as ratios to the control lean littermate values and expressed as means \pm SE

^a $p<0.01$ relative to control-diet-fed mice

lavage fluid in experimentally induced pulmonary inflammation, but is most abundant in adipose tissues [13]. RELM β is specifically expressed in the intestinal tract, especially abundantly in the colon [14] and has been suggested to be related to bacterial colonisation [24]. Injecting RELM β reportedly induced insulin resistance in rats [16]. RELM γ is expressed in the colon, bone marrow, spleen and lung [17–19] and reportedly increases the proliferation rate of promyelocytic cells and modulates their differentiation [18]. RELM γ is also expressed in rat nasal respiratory epithelium, and is altered by cigarette smoke [17]. Although the physiological roles of these isoforms are still unclear, a series of previous reports seems to suggest their interaction with inflammatory processes and/or insulin resistance.

In this study we showed that RELM γ overexpressed in and secreted by COS7 cells forms a homodimer. Homodimerisation of RELM β and resistin, but not RELM α , was previously reported and conserved cysteines (Cys26 of resistin and Cys25 of RELM β) were considered to be required for disulphide bond formation [23, 24]. In RELM γ , however, the corresponding cysteine is not conserved. Therefore, Cys11 or Cys45 in RELM γ may be regarded as critical for dimer formation. Considering that Cys11 is conserved while Cys45 is missing from RELM α , which is not capable of forming a homodimer, Cys45 in RELM γ is very likely to be involved in dimerisation. Interestingly, we also found that RELM β and RELM γ secreted by COS7 cells and also endogenously expressed in colonic tissues partially heterodimerised (Fig. 4).

RELM γ reportedly binds α -defensin, a cysteine rich 3–4-kDa antimicrobial peptide stored in the cytoplasmic granules of neutrophils, some macrophages and intestinal Paneth cells [19]. Thus, it is possible that some protein(s), like α -defensin, which bind to RELM γ , also bind to RELM β and may form polymers with disulphide bonds. In this case, this heteromeric formation may modulate the antimicrobial effects of defensin. In addition, homodimerisation and heterodimerisation of RELM γ would affect binding to other molecules such as their receptor(s) and their resultant functions. Thus, whether or not RELM γ heterodimers and homodimers have different roles in insulin resistance and/or inflammatory processes is important. Further analysis, considering the crystal structure data of resistin and RELM β , is necessary to clarify the mechanism of homo/heterodimer formation of resistin family proteins [20].

This is the first study to investigate the serum concentrations and tissue contents of RELM β and RELM γ in insulin-resistant animals, while resistin expression has been analysed in several models of obesity and diabetes [8, 11, 25–30]. In high-fat-fed mice and *db/db* mice, serum levels of RELM β and RELM γ were apparently increased. Taking into consideration that RELM β causes insulin resistance by impairing hepatic glucose production [16], it is reasonable to consider the increased serum concentrations of RELM β and RELM γ to be among the molecular mechanisms underlying the insulin resistance in these diabetic mice. Furthermore, we demonstrated that elevated serum

concentrations of RELM β and RELM γ are attributable to increased production in the colon (both RELM β and RELM γ) and bone marrow (only RELM γ). Since the heterodimer consisting of serum RELM β and RELM γ was also increased, a considerable portion of the RELM β and RELM γ in serum appears to be derived from the colon and ileum.

We observed that the PPAR γ agonist rosiglitazone had no effect on expression levels ($10 \text{ mg kg}^{-1} \text{ day}^{-1}$ for 2 weeks, orally; data not shown). We can suggest two possible mechanisms of RELM β and RELM γ upregulation in the insulin-resistant diabetic mouse. One involves the expression of these isoforms being regulated by exposure of intestinal and bone marrow cells to nutrients and/or factors such as glucose, lipid, insulin or certain cytokines or hormones. The other involves signals related to inflammation triggering increased RELM β and RELM γ expression. Since inflammation has mechanistic significance in obesity and insulin resistance [8, 31–34] and some resistin family proteins have been implicated in the inflammatory response [13, 17], increased expression of RELM β and RELM γ may be involved in the mechanisms connecting inflammation and insulin resistance, which are associated with obesity and/or diabetes. In this regard, a transcriptional factor may be involved in the inflammation-induced increased expression of RELM β and RELM γ . Indeed, the promoter region of RELM β contains a binding sequence of nuclear factor κ B and signal transducer and activator of transcription 6 [24] and RELM γ is reportedly a target gene of CCAAT/enhancer binding protein ϵ mediating the role of promyelocytic cell development [19]. Our other interesting finding was that the expression of RELM γ was unchanged in the spleen, lung and pancreas in the insulin-resistant as compared with the control mice. A similar observation was that RELM α expression in adipose tissue responds to food deprivation, except in the lung [16]. Thus, it seems that organs specifically responding to nutritional or inflammatory conditions have a system which allows expression of RELM α and RELM γ to be regulated, although the underlying molecular mechanism is unclear.

In summary, adipose tissue and the intestinal tract are major tissues which produce RELM family proteins (resistin and RELM α from adipose tissue, and RELM β and RELM γ from the intestinal tract). The production of RELM β and RELM γ in the intestinal tract was clearly demonstrated to be increased in diet-induced and genetically obese mice, with elevated serum concentrations suggesting a hormonal link between the intestinal tract and insulin sensitivity. Furthermore, this link may respond to the total calorie and/or nutrient content of the food ingested. Future studies will examine whether the serum concentrations of RELM β and RELM γ are also increased in obese or insulin-resistant diabetic human subjects. If so, the measurement of serum RELM β and RELM γ concentrations may be diagnostically useful. We can also suggest the possibility that RELM β and RELM γ are potential molecular targets for future antidiabetic drugs, and additional studies are needed to investigate these possibilities.

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