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Evidence from studies in rodents and in isolated adipocytes that agonists of the chemerin receptor CMKLR1 may be beneficial in the treatment of type 2 diabetes

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ABSTRACT

It is unclear whether the adipokine chemerin has pro- or anti-inflammatory properties, plays any role in the aetiology of obesity or type 2 diabetes, or whether agonists or antagonists of the chemerin receptor CMKLR1 have potential in the treatment of these diseases. To address these questions, we investigated the metabolic phenotypes of both male and female, CMKLR1 knockout and heterozygote mice; the effects of murine chemerin and some C-terminal peptides on glucose uptake in wild-type and CMKLR1 knockout adipocytes; and plasma chemerin levels and chemerin gene mRNA content in adipose tissue in models of obesity and diabetes, and in response to fasting or administration of the insulin sensitizing drug rosiglitazone, which also has anti-inflammatory properties. Both male and female, CMKLR1 knockout and heterozygote mice displayed a mild tendency to obesity and impaired glucose homeostasis, but only when they were fed on a high fat, rather than a standard low fat diet. Obesity and impaired glucose homeostasis did not occur concurrently, suggesting that obesity was not the sole cause of impaired glucose homeostasis. Picomolar concentrations of chemerin and its C15- and C19-terminal peptides stimulated glucose uptake in the presence of insulin by rat and mouse wild-type epididymal adipocytes, but not by murine CMKLR1 knockout adipocytes. The insulin concentration-response curve was shifted to the left in the presence of 40 pM chemerin or its C-15 terminal peptide. The plasma chemerin level was raised in diet-induced obesity and ob/ob but not db/db mice, and reduced by fasting and, in ob/ob mice, by treatment with rosiglitazone. These findings suggest that an agonist of CMKLR1 is more likely than an antagonist to be of value in the treatment of type 2 diabetes and have associated anti-obesity and anti-inflammatory activities. One mechanism by which an agonist of CMKLR1 might improve glucose homeostasis is by increasing insulin-stimulated glucose uptake by adipocytes.
INTRODUCTION

Chemerin is translated as a 163 amino acid pre-proprotein and secreted as a 143 amino acid proprotein. Further C-terminal cleavage by extracellular proteases results in peptides that have chemoattractant properties. Some experimental evidence shows that these peptides have pro-inflammatory properties, but there is also evidence that some have anti-inflammatory properties (Ernst & Sinal, 2010). The C-15 peptide is reported to have potent anti-inflammatory activity and the C-19 peptide to be inflammatory but to lack any chemotactic activity for macrophages (Cash et al., 2008). We and collaborators have shown that chemerin also stimulates angiogenesis (Bozaoglu et al., 2010).

The possibility that chemerin plays a role in metabolic disease, in particular obesity and type 2 diabetes, arose when it was discovered that adipose tissue is one of the main tissues from which chemerin is secreted (i.e. it is an adipokine), and that circulating levels in humans correlate with body mass index, plasma triglycerides and blood pressure (Ernst & Sinal, 2010). Insulin sensitising thiazolidinedione drugs that activate peroxisome proliferator-activated receptor (PPAR)γ alter the expression of chemerin and circulating chemerin levels, but contrasting effects have been reported (Ernst & Sinal, 2010, Roman, Parlee & Sinal, 2012). For example, amelioration of insulin resistance and hyperglycaemia by the PPARγ agonist pioglitazone or by metformin in patients type 2 diabetes is accompanied by reduced plasma chemerin (Esteghamati et al., 2014), whereas treatment of mice with the PPARγ agonist rosiglitazone increased the chemerin mRNA level in adipose tissue and the circulating chemerin levels (Muruganandan et al., 2011).

Since chemerin levels are raised in obesity, a pro-inflammatory effect might be one mechanism by which obesity causes insulin resistance. Mechanisms that are independent of inflammation might also play a role, because in some studies chemerin has inhibited glucose uptake too rapidly to suggest a role for inflammation or inhibition of glucose uptake has been observed in isolated cell systems. In addition, chemerin promotes adipogenesis, which might exacerbate obesity (Ernst & Sinal, 2010, Roman, Parlee & Sinal, 2012). However, it is by no means certain that compounds that block the effects of chemerin would be of value in the treatment of metabolic disease. For example, inhibitory as well as stimulatory effects of chemerin on glucose uptake in 3T3-L1 adipocytes have been described (Takahashi et al., 2008; Kralisch et al., 2009). Thus elevated plasma levels of chemerin might play a causal
role in obesity and its metabolic complications, but it is also possible that they play a counter-
regulatory role. In particular, chemerin might mitigate insulin resistance by permitting 
adipose tissue expansion through its angiogenic effect and by promoting adipogenesis 
(Bozaoglu et al., 2010).

The first chemerin receptor to be identified is known as CMKLR1 or ChemR23. It is highly 
expressed in adipose tissue as well as various cell types involved in innate and adaptive 
immunity, and endothelial cells (Ernst & Sinal, 2010, Kaur et al., 2010). Adipose tissue is 
therefore both a major source of chemerin and a site of its action. Studies conducted in mice 
that lack this receptor have revealed a complex and inconsistent metabolic phenotype. Both 
beneficial and adverse age-, sex- and diet-related effects on body composition and glucose 
homeostasis have been described, as well as an absence of these effects (Ernst et al., 2012, 
Rouger et al., 2013, Gruben et al., 2014).

Here we have followed four main lines of inquiry. First, we show that in our hands both male 
and female mice that lack CMKLR1 and are fed on a high fat diet have a mild tendency to 
increased body fat and impaired glucose homeostasis, though these do not always occur at the 
same age. We are the first to describe similar results in mice that are heterozygote for 
CMKLR1. Second, we show for the first time that murine chemerin and its C-15 and C-19 
terminal peptides stimulate glucose uptake in primary adipocytes, and provide evidence that 
this effect is mediated by CMKLR1. Third, we investigate the effects of fasting and diet-
induced obesity on plasma chemerin levels and chemerin gene (RARRES2) mRNA content in 
adipose tissue, comparing for the first time the effects of diet-induced obesity on plasma 
chemerin levels in C56Bl/6 and FVB mice, the latter being relatively resistant to obesity. 
Fourth, we report that, as in wild-type mice (Muruganandan et al., 2011), rosiglitazone, 
increases the expression of the chemerin gene RARRES2 in inguinal adipose tissue and raises 
the plasma chemerin level in leptin deficient ob/ob mice. Our findings are consistent with 
chemerin, through its receptor CMKLR1, having beneficial effects in diabetes, which is 
contrary to the opinions of some other workers.
MATERIALS AND METHODS

Materials

Murine chemerin was purchased from R&D Systems Europe Ltd (Abingdon, Oxfordshire, UK). Its C-15 (amino acids 140 to 154), C-19 (amino acids 138 to 156) and stable analogue of C-9 (amino acids 148 to 156) (Shimamura et al., 2009, Cash et al., 2008) terminal peptides were custom-synthesized by Cambridge Research Biochemicals (Billingham, Cleveland, UK). Their amino acid sequences were:

C-15: AGEDPHGYFLPGQFA
C-19: AQAGEDPHGYFLPGQFAFS
C-9: yFLPsQFaTicS (This is a modification of the C-9 sequence FLPGQFAFS, with D-Tyr\(^{147}\), D-Ser\(^{151}\), D-Ala\(^{154}\), Tic\(^{155}\). Although we refer to it as C-9 analogue because it was first described as an analogue of C-9, the introduction of D-Tyr\(^{147}\) actually makes it a C-10 peptide.)

Other reagents were obtained from Sigma-Aldrich, Poole, UK, unless otherwise stated.

Animals

Mice and rats were from Charles Rivers (Maidstone, Kent, UK), except the embryos used to generate CMKLR1 mice were generated by Deltagen (see below). Mice were received at five weeks of age and rats at 150 g body weight. They were fed on standard laboratory chow (Beekay Feed, B&K Universal Ltd., Hull, UK) until used, except that for studies on diet-induced obesity male C57Bl/6 and FVB mice were fed from the age of 6 weeks on a high fat (63% by energy; Open Source D12492, Research Diets, New Brunswick, NJ, USA) diet for 6 months, Housing and procedures were conducted in accordance with the UK Government Animal (Scientific procedures) Act 1986 and approved by the University of Buckingham Ethical review Board. Animals were killed 3-4 hours after the onset of day light cycle, by a UK Government Animal Scientific Act 1986 schedule 1 method.

Whole body CMKLR1 knockout mice were originally generated by Deltagen (San Mateo, CA, USA) and supplied as embryos to AstraZeneca, Alderley Park where they were backcrossed onto the C57Bl/6 background for three generations. Heterozygote mice were supplied to the University of Buckingham and bred for a minimum of five further generations to generate the animals for the phenotyping experiments. Genotyping of experimental animals was performed in accordance with the recommendations of Deltagen.
Adipocytes were prepared from mesenteric fat pads of female wild-type C57Bl/6 mice, epididymal fat pads of male Sprague-Dawley rats (350-400 g), and epididymal fat pads of male wild-type and CMKLR1 knockout C57Bl/6 mice by a method that we have described previously (Zaibi et al., 2010). All animals were 12 to 14 weeks old.

The effect of fasting for 16 h overnight was studied in 8-week-old male wild-type C57BL/6J mice. The effect of treatment with rosiglitazone (3 mg/kg body weight, p.o., once daily; Sequoia Research Products Ltd, Japan) for three weeks and of fasting for 16 h overnight was studied in female ob/ob mice. Male db/db mice and wild-type (misty C57Bl/Ks) mice were 11 to 12 weeks old when studied.

**Phenotyping experiments**

Food intake and body weight were measured weekly. Non-fasting blood glucose was measured at 9 weeks, and 3 and 6 months of age. Blood glucose following a 5 h fast was measured at 8 weeks, 11 weeks and 6 months of age. Insulin tolerance was measured at 6 months of age. Body composition and glucose tolerance were conducted at 7 to 8 weeks and 3 and six months of age.

Body composition was measured under light gaseous anaesthetic (Isofluorane, Isoba, Shering-Plough Animal Health, UK) using dual-energy X-ray absorptiometry (Lunar PIXImus 2 mouse densitometer and version 1.46 software, GE Medical Bedford, UK).

Oral glucose tolerance was measured after mice had been fasted for five hours before being dosed with glucose (3 g/kg body weight, p.o.). Blood samples (10 μl) were taken from the tip of the tail after applying a local anesthetic (lignocaine™, Centaur Services, UK) 30 min and immediately before, and 30, 60, 90, 120 and 180 min after dosing the glucose load. Whole blood was mixed with hemolysis reagent and blood glucose was measured in duplicate using the Sigma Enzymatic (Glucose Oxidase Trinder, ThermoFisher Microgenics, UK) colorimetric method and a SpectraMax 250 (Molecular Devices Corporation, Sunnyvale, CA, USA). Areas under the curve were analyzed from 0 to 120 minutes.

Insulin tolerance was measured after mice had been fasted for five hours before being dosed with insulin (Actrapid™ Centaur Services, UK at 0.5 units/kg body weight, i.p.). Blood
samples were taken, as described for the glucose tolerance test, 15 min and immediately
before, and 15, 30, 45 and 60 min after the administration of insulin.

**Glucose uptake by adipocytes**

At least 10 mice and two rats were used for each preparation of adipocytes. Tissue was
minced and digested with collagenase type II in Krebs-Ringer HEPES buffer containing 10
mM HEPES, 1% bovine serum albumin (fraction V), 2.5 mM CaCl₂, 5.5 mM glucose and
200 nM adenosine at pH 7.4 at 37°C. The preparation was filtered through a 250–300 µm
nylon mesh. The infranatant was removed and the floating layer of adipocytes was washed
four times with fresh buffer. Adipocytes were concentrated to 40% of final volume of Krebs-
Ringer HEPES buffer containing 5% BSA and 0.3 mM glucose and pre-incubated for 45 min
under 95% O₂: 5% CO₂ before dispensing them into 300 µl polyethylene tubes for the
measurement of glucose uptake.

Glucose transport was measured as described previously (Kashiwagi, Huecksteadt & Foley,
1983). Adipocytes were incubated in Krebs-Ringer HEPES buffer containing BSA, 0.3 mM
glucose and D-[U-¹⁴C] glucose (0.2 μmol/l; 0.2 μCi/ml), for 1 h at 37°C in the absence or
presence of different concentrations of chemerin, or C-terminal chemerin peptides and
insulin. The reaction was stopped by separation of the cells through silicone oil and
radioactivity in the cells was measured. The extracellular space was measured in parallel
incubations using D-[U-¹⁴C] sucrose. Uptake is expressed relative to the weight of the cells,
or relative to uptake in the presence of insulin but absence of chemerin or its C-terminal
peptides, as indicated in the figure legends. For each preparation of adipocytes there were at
least four or five replicates of each treatment. Apart from the preliminary data shown in
Supplemental Figure 8, the data presented are means of the replicate means from four or five
preparations of adipocytes. Glucose uptake was calculated assuming that 2-deoxyglucose
and glucose are not distinguished by cellular uptake mechanisms. Further details are given in
the figure legends.

**Plasma chemerin assay**

Plasma chemerin levels were measured using a murine chemerin quantikine ELISA kit (R
and D Systems, Oxford, UK) according to the manufacturer’s recommendation. Whole blood
was collected into EDTA tubes and spun at 3000g for 5 min at 4°C and the plasma stored at -
80°C prior to analysis. Plasma samples were assayed in duplicate and the absorbance of both
unknowns and standards measured using the Spectromax at 450 nm with a sensitivity range of 1.08-3.47 pg/ml.

**Gene expression**

Gene expression analysis was performed as described previously (Wargent et al 2013). Briefly, total RNA was isolated from white adipose tissue and analysed using a NanoDrop ND1000 (Thermo Fisher Scientific, Delaware, USA). Real time PCR (StepOne™, Applied Biosystems, Paisley, UK) was carried out using Assay-on-Demand pre-designed primer and probe sets for the chemerin gene, *RARRES2*, and GAPDH (Applied Biosystems, Paisley, UK). GAPDH was chosen as the housekeeping genes because it showed consistent C<sub>T</sub> values in adipose tissue. Data were analysed using the comparative ΔC<sub>T</sub> method. All procedures were carried out in accordance to the manufacturer’s recommendation.

**Statistics**

Unless stated differently in the legends (see Figure 6), data were analyzed by one-way ANOVA followed by Fisher’s least significant difference test, using GraphPad Prism version 5 (GraphPad software, San Diego CA, USA). Two-sided significance levels are given. Differences were considered significant at *P* < 0.05. Results are expressed as means ± SEM.
RESULTS

CMKLR1 knockout mice fed on a high fat diet have impaired glucose homeostasis; only the males have increased percentage body fat

With the exception of insulin tolerance, which was measured only at 6 months of age, and food intake and body weight, which were measured weekly, measurements were conducted at 7 to 9 weeks, 11 to 13 weeks and six months of age.

When they were fed on a standard chow diet, there were no differences in body weight between genotypes (Supplementary Figure 1), but the female knockout mice had a lower body fat content than the wild-type mice at the age of 6 months (Supplementary Figure 2). Both the knockout and heterozygote female mice had a lower fasting blood glucose than the wild-type mice before the glucose tolerance test at the age of 3 months (Supplementary Figure 3). There were no significant differences at any time between wild-type, heterozygote and knockout genotypes (either males or females) in food intake (results not shown), glucose tolerance, insulin levels in an oral glucose tolerance test, or blood glucose concentration following administration of insulin (Supplementary Figures 3, 4 and 5). There were no differences between genotypes in any of these measurements when the mice were 7 to 9 weeks old.

A different picture emerged when the mice were fed on a high fat diet. These were in the direction of increased adiposity or impaired glucose tolerance in the knockout or heterozygote mice compared to the wild-type mice. Thus, although there were no significant differences between genotypes in body weight (Figure 1), both the male knockout and the male heterozygote mice had an increased percentage body fat at 3 but not 6 months of age, whilst both the female knockout and heterozygote mice had an increased percentage body fat at 6 but not 3 months of age (Figure 2). These increases were mostly due to increased body fat content, except that the 3-month-old male heterozygote mice had less lean tissue than the wild-type mice, coupled with a statistically non-significant increase in body fat content. There were no differences in body composition between genotypes when the mice were 7 to 8 weeks old (results not shown).
The male knockout mice fed on a high fat diet had a higher fasting blood glucose than the wild-type controls at 8 weeks and at 3 months of age (Figure 3), and a higher blood glucose level 10 min after administration of insulin at 6 months of age (Figure 4), despite body composition being no different from that of wild-type mice at 6 months of age. The male heterozygote mice had elevated plasma glucose at 3 months of age and an elevated fed blood glucose at both 3 and 6 months of age (Figure 3). Oral glucose tolerance and plasma insulin during the glucose tolerance test was unaffected by genotype at all ages in both the male knockout and male heterozygote mice (Figure 4; Supplementary Figures 6 and 7).

The female knockout mice fed on a high fat diet had a higher fasting plasma glucose concentration than the wild-type mice (Figure 3) and a higher blood glucose level 10 min after administration of insulin (Figure 4). At this age (6 months), they had increased body fat. Their glucose tolerance was impaired at 3 months of age, despite their body composition being no different from that of wild-type mice (Figure 2). The female heterozygote mice had a higher blood glucose level than the wild-type mice 10 min after administration of insulin at 6 months of age (Figure 4).

Chemerin and its C-15 and C-19 terminal peptides stimulate glucose uptake by adipocytes

Initial experiments (Supplementary Figure 8) using murine mesenteric adipocytes suggested that chemerin and its C-15 and C-19 terminal peptides (Cash et al., 2008), but not the C-9 analogue peptide (Wittamer et al., 2004), stimulated glucose uptake in the presence of 0.1 nM insulin, which had no effect alone, but not in the absence of insulin or in the presence of 10 nM insulin. Higher concentrations than 1 nM chemerin appeared less effective, a phenomenon that has been observed for other effects of chemerin (Bozaoglu et al., 2010, Cash et al., 2008) and the C-15 peptide (Cash, Christian & Greaves, 2010).

We were able to generate more robust data by using rat, rather than murine epididymal adipocytes. Studies were conducted on the C-9 analogue, C-15 and C-19 fragments using four separate adipocyte preparations to ensure that the results were not a peculiarity of one preparation. The C-9 analogue was ineffective, C-15 had similar potency to chemerin with a
peak effect at 40 pM, whilst C-19 had a similar peak effect but at the higher concentration of
160 pM (Figure 5A).

Four separate rat epididymal adipocyte preparations were then used to compare the effects of
40 pM chemerin and C-15 on the insulin concentration-response curve. The C-15 peptide
shifted the insulin curve further to the left than did full length recombinant chemerin (Figure
5B). Chemerin and C-15 were ineffective in the presence of high concentrations of insulin.

The stimulatory effect of the C-15 fragment of chemerin is partly mediated by
CMKLR1

Having established that the C-15 peptide had the greatest effect on glucose uptake in rat
epididymal adipocytes, the effect of this peptide (40 pM) was compared in murine
epididymal adipocytes from wild-type and knockout mice. The peptide increased insulin-
stimulated glucose uptake in adipocytes from wild-type, but not knockout mice. A similar
result was obtained with full length recombinant chemerin, but the effect of chemerin in the
wild-type adipocytes did not achieve statistical significance in this experiment (Figure 6).

The plasma chemerin concentration is raised in obesity and decreased by fasting

FVB mice are known to be less susceptible than C57Bl/6 mice to diet-induced obesity
(Glendinning et al., 2010, Kim et al., 2013), as we confirmed (Figure 7A; Supplementary
Figure 9). Plasma chemerin levels were raised by feeding male FVB or C57Bl/6 mice on a
high fat diet for 6 months (Figure 7B). The increase in plasma chemerin level in response to
the high fat diet in the FVB mice was 75% of that in the C57Bl/6 mice, whereas the increases
in body fat content and percentage body fat in FVB mice were 36% and 57% respectively of
those in C57Bl/6 mice.

The concentration of chemerin in plasma of fed male ob/ob mice was higher than in wild-
type controls (Figure 7C). We cannot exclude the possibility that this is because the ob/ob
mice were two weeks older than the wild-type mice. However, by contrast, the expression of
RARRES2, the chemerin gene, was lower in inguinal adipose tissue in ob/ob than in C57Bl/6
mice (Figure 7D). Other workers have also noted this paradox (Ernst et al., 2010). We found
no difference in the plasma chemerin level or the expression of RARRES2 in inguinal adipose tissue between male \textit{db/db} mice and wild-type mice of the background (C57Bl/Ks) strain (Supplementary Figure 10).

Plasma chemerin was reduced by fasting in both wild-type male C57Bl/6 and female \textit{ob/ob} mice (Figures 7C and 8A). The expression of RARRES2, the chemerin gene, was similarly lower in inguinal adipose tissue of fasted than fed wild-type mice (Figure 7D).

Fasting or feeding on a high fat diet did not alter the expression of RARRES2 in epididymal adipose tissue (Supplementary Figure 11).

Rosiglitazone increases the plasma chemerin concentration in \textit{ob/ob} mice

The plasma chemerin concentration was increased by treatment of female \textit{ob/ob} mice with rosiglitazone (3 mg/kg body weight, p.o.) for 18 days (following an overnight fast) or 21 days (in the fed state; Figure 8A). Treatment of female \textit{ob/ob} mice with rosiglitazone increased the expression of RARRES2 in inguinal (Figure 8B) but not parametrial white adipose tissue (Supplementary Figure 11B).
DISCUSSION

The finding that plasma chemerin levels are raised in human obesity (Bozaoglu et al., 2007, 2009) and in genetically obese ob/ob mice (Ernst et al., 2010), the adipogenic effect of chemerin (Goralski et al., 2007), and a report of decreased percentage body fat in mice that lack the chemerin receptor CMKLR1 (Ernst et al., 2012) have led to suggestions that raised plasma chemerin levels promote obesity. Other findings have led to suggestions that chemerin is in part responsible for the link between obesity and insulin resistance. Thus, raised plasma chemerin is associated with the metabolic syndrome (Stejskal et al., 2008, Bozaoglu et al., 2007) and chemerin is released from adipocytes (Goralski et al., 2007). The majority of studies find chemerin to be pro-inflammatory (Ernst & Sinal, 2010). Chemerin inhibited insulin action in skeletal muscle cells (Sell et al., 2009) and in one study in 3T3-L1 adipocytes (Kralisch et al., 2009). Administration of chemerin exacerbated glucose tolerance in mice that are obese and diabetic (Ernst et al., 2010), and mice that lack the chemerin receptor are susceptible to diet-induced insulin resistance (Ernst et al., 2012). These results support the view that an antagonist of chemerin action, in particular of CMKLR1, might be useful in the treatment of type 2 diabetes associated with obesity.

The findings that we report here lead, however, to a different conclusion – that chemerin, acting through CMKLR1, opposes diet-induced obesity and insulin resistance, and that therefore an agonist of CMKLR1 might be of value in the treatment of type 2 diabetes.

First, although our six-month-old female CMKLR1 knockout mice had a lower body fat content than wild-type mice when they were fed on a low fat diet, knockout and heterozygote mice of both sexes had a slightly higher percentage body fat than wild-type mice at three or six months of age when they were fed on a high fat diet.

Second, when fed on the high fat diet, fasting glucose was higher in our knockout than wild-type mice when males were nine weeks or three months old, and females six months old. Fasting blood glucose was also raised in three-month-old heterozygote male mice. Glucose tolerance was impaired in the female knockout mice when they were three months old and there was some impairment in insulin tolerance in both male and female six-month-old knockout mice.
Overall, our results suggest that CMKLR1 knockout and heterozygote mice tend to be mildly obese and have impaired glucose homeostasis when they are fed on a high fat diet. The impaired glucose homeostasis does not appear to be entirely a consequence of the obesity, because it sometimes occurred in the absence of obesity. Previous studies have shown an inconsistent metabolic phenotype for CMKLR1 mice. The first study reported was conducted in male mice (sex of the mice from personal communication from Professor Christopher Sinal). The mice displayed a lean phenotype when fed on either a low or high fat diet. In mice fed on a high fat diet, leanness was associated with lower fasting plasma glucose and serum insulin. However, regardless of diet, the mice displayed impaired glucose tolerance. This was associated with decreased insulin secretion, there being no alteration of insulin tolerance (Ernst et al., 2012). In another study, rather than being lean, male but not female mice that lacked CMKLR1 developed increased adipose tissue mass from the age of 8 months when fed on a standard (3.5% fat) diet, but glucose tolerance was unchanged. However, when the mice were fed on a high fat diet for 20 weeks there was no difference in body weight between the knockout and wild-type male mice (Rouger et al., 2013). In a third study, in which male mice were fed on a high fat diet, the absence of CMKLR1 had no effect on body weight, food intake or insulin sensitivity (Gruben et al., 2014).

Our results agree with reports that plasma chemerin levels are raised in obese rodents. Thus they were higher in ob/ob than in wild-type mice as reported by others (Ernst et al., 2010). We also found that they are raised in diet-induced obese mice, which is more representative of human obesity, and that they were reduced by overnight fasting. It might be argued that, since chemerin has been found to stimulate adipogenesis (Goralski et al., 2007, Roh et al., 2007), raised plasma levels would exacerbate obesity. This view is not, however, supported by comparing our results in FVB and C57Bl/6 mice. Plasma chemerin levels were raised by feeding either FVB or C57Bl/6 mice on a high fat diet. The increase in the chemerin level in the FVB mice was about 75% of that in the C57Bl/6 mice, whereas the increase in the body fat content in FVB mice was about 30% of that in C57Bl/6 mice. This might be interpreted in terms of increased plasma chemerin protecting the FVB mice from diet-induced obesity, rather than it promoting obesity through its adipogenic activity. By contrast with our results, others have reported that the plasma chemerin concentration was not raised by feeding NMRI mice on a high fat or cafeteria diet for three weeks, despite these treatments roughly doubling.
weight gain (Hansen et al., 2014). NMRI mice are less prone than C57Bl/6 mice to diet-induced obesity (Matyskova et al., 2008).

If chemerin does provide some protection against obesity, it does not seem that this is directly linked to increased expression of RARRES2 mRNA adipose tissue. We did not find any changes in the expression of RARRES2 in response to high fat feeding or fasting in perigenital adipose tissue, nor any consistent parallels between plasma chemerin levels and the expression of RARRES2 in inguinal adipose tissue. In fact, there were no changes in expression in response to diet-induced obesity in either inguinal or epididymal adipose tissue. Moreover, expression of RARRES2 was lower in inguinal adipose tissue of ob/ob mice than in wild-type mice, despite plasma chemerin concentrations being higher. Other workers have reported increased expression of RARRES2 in epididymal adipose tissue in mice fed on a high fat diet (Roh et al., 2007), and decreased expression in rat perirenal, inguinal and epididymal adipose tissue (though only perirenal results are shown) in response to fasting for 72 h or food restriction by 50% for one month (Stelmanska et al., 2013). However, our results resemble those of Hansen et al., in that cold acclimation reduced and obesogenic diets increased RARRES2 expression in inguinal but not epididymal adipose tissue. Expression in interscapular brown adipose tissue paralleled that in inguinal adipose tissue, the latter site (like the perirenal site) being one where brite/beige adipocytes are expressed (Hansen et al., 2014).

The plasma chemerin concentration has been reported to be higher in db/db mice than in wild-type C57Bl/6 mice, which is the background strain for ob/ob mice (Ernst et al., 2010). We did not find that the plasma chemerin concentration was higher in db/db mice relative to wild-type mice of its own background strain (C57Bl/Ks), though there was a slight trend in this direction.

Treatment of ob/ob mice with the insulin sensitising drug rosiglitazone increased their plasma chemerin concentration and the expression of RARRES2 in inguinal but not parametrial adipose tissue. Others have also reported that rosiglitazone increased RARRES2 mRNA in inguinal adipose tissue of normal mice and identified a PPARγ response element within the RARRES2 promoter (Muruganandan et al., 2011). It is well-established that rosiglitazone and other PPARγ agonists have anti-inflammatory properties (Ceriello, 2008). Therefore the effects of rosiglitazone in our study are more consistent with chemerin having anti-
inflammatory and anti-diabetic than pro-inflammatory and pro-diabetic activity. However, we note that other PPARγ agonists have been reported to reduce \textit{RARRES2} mRNA levels in epididymal and mesenteric adipose tissue when administered to mice (Vernochet \textit{et al.}, 2009), and plasma chemerin levels when administered to patients with type 2 diabetes (Esteghamati \textit{et al.}, 2014).

It is possible that the effect of rosiglitazone on \textit{RARRES2} mRNA levels is unrelated to its anti-inflammatory activity, but due to its promoting brown adipocyte formation, whilst possibly decreasing sympathetic activity (Sell \textit{et al.}, 2004). This seems unlikely, however, because cold acclimation and obesogenic diets, both of which increased brown adipose tissue formation, did not affect the plasma chemerin concentration. Moreover, cold acclimation reduced whereas obesogenic diets increased \textit{RARRES2} mRNA levels in interscapular brown and inguinal brite/beige adipose tissue (Hansen \textit{et al.}, 2014).

We investigated whether chemerin might have a direct insulin sensitising effect in primary adipocytes. This had been suggested by a previous study using 6 nM chemerin in 3T3-L1 adipocytes (Takahashi \textit{et al.}, 2008), although a study using an extremely high concentration (10 μM) of chemerin found reduced insulin-stimulated uptake (Kralisch \textit{et al.}, 2009). In addition to chemerin, we studied some C-terminal fragments of chemerin because we were interested in whether they might provide the basis for a drug discovery programme. The C-terminal nonapeptide has been identified as being the smallest peptide that had functional activity at low nanomolar concentration in a Chinese hamster ovary cell line that expressed \textit{RARRES2} (Wittamer \textit{et al.}, 2004). This peptide did not, however, stimulate glucose uptake by mouse mesenteric or rat epididymal adipocytes, either in the absence or presence of insulin. By contrast, the C-15 and C-19 terminal peptides (Cash \textit{et al.}, 2008) and full length chemerin itself appeared to stimulate glucose uptake by mouse mesenteric adipocytes in the presence but not the absence of a submaximal concentration of insulin, and this property was more clearly demonstrated using rat epididymal adipocytes. The peak effects of chemerin and the C-15 peptide were at 40 nM, and at this concentration both full length chemerin and the C-15 peptide shifted the chemerin concentration-response curve to the left. Using mouse epididymal adipocytes from knockout and wild-type mice, we were able to show that CMKLR1 mediated the response to the C-15 peptide and probably to chemerin as well, although unfortunately the effect of chemerin did not achieve statistical significance in that
experiment. CMKLR1 has also been shown to mediate the anti-inflammatory effect of the C-15 peptide (Cash et al., 2008).

In conclusion, knockout and heterozygote mice for the chemerin CMKLR1 receptor tend to be mildly obese and display mild impairment of glucose homeostasis. Rosiglitazone increases the plasma chemerin concentration in ob/ob mice, suggesting that chemerin might have be anti-inflammatory and anti-diabetic. Chemerin and its C-15 terminal peptide, acting via the CMKLR1 receptor, sensitise adipocytes to the stimulatory effect of insulin on glucose uptake. These findings suggest that an agonist of CMKLR1 might be useful for the treatment of type 2 diabetes associated with obesity.

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REFERENCES


Figure 1. **Body weight of wild-type, CMKLR1 knockout and CMKLR1 heterozygote mice fed on a high fat diet.** n = 17 to 21 for males, 14 to 17 for wild-type and heterozygote females, and 8 to 10 for female knockout mice. The upper three lines are for males and the lower three for females.
Figure 2. **Body composition of wild-type, CMKLR1 knockout and CMKLR1 heterozygote mice fed on a high fat diet at 12 weeks (A, B, C) and six months (D, E, F) of age.** n = 17 to 21 for males, 14 and 17 for wild-type and heterozygote females at 12 weeks and 6 months respectively, and 8 and 10 for female knockout mice at 12 weeks and 6 months respectively. *P<0.05 for knockout or heterozygote compared to wild-type mice of the same sex.
Figure 3. Blood glucose concentration in 5 h-fasted (A, C, E) and fed (B, D, F) wild-type, CMKLR1 knockout and CMKLR1 heterozygote mice fed on a high fat diet at 8 to 9 weeks (A, B), 13 weeks (C, D) and six months (E, F) of age. n = 17 to 21 at 8 to 9 weeks, and 12 to 15 at 13 weeks and six months of age, except for female knockout mice, where n = 5 to 10. *P<0.05; **P<0.01 for knockout or heterozygote compared to wild-type mice of the same sex.
Figure 4. Oral glucose tolerance and intraperitoneal insulin tolerance in wild-type, CMKLR1 knockout and CMKLR1 heterozygote mice fed on a high fat diet. Glucose tolerance curves in 12-week-old male (A) and female (B) mice. Areas under the glucose tolerance curves in 12-week-old (C) and 6-month-old (D) mice. Insulin tolerance curves in 6-month-old male (E) and female (F) mice. Further data from the oral glucose tolerance tests are given in Supplementary Figures 3 to 7. n = 17 to 21, except for female knockout mice, where n = 10. *P<0.05 for knockout or heterozygote compared to wild-type mice of the same sex.
Figure 5. **Effect of murine chemerin and C-terminal peptides on glucose uptake by rat epididymal adipocytes.** (A) Glucose uptake in the presence of 0.05 nM insulin in response to 10, 100 or 1000 pM of chemerin, C15-terminal peptide, C19-terminal peptide or the stable analogue of the C9-terminal peptide. **P<0.01; ***P<0.001 for knockout or heterozygote compared to wild-type mice. (B) Glucose uptake in the absence or presence of murine chemerin (40 pM) or its C15-terminal peptide in response to insulin. Results in each panel are means of four experiments on different preparations of adipocytes. In each experiment, each data point was the mean of five replicates. In (A) results for each experiment were expressed relative to the mean of the five replicates for uptake in the presence of insulin alone.
Figure 6. **Glucose uptake in the presence of insulin (0.3 nM) in response to murine chemerin (40 pM) or its C15-terminal peptide (40 pM) in epididymal adipocytes from wild type and CMKLR1 knockout mice.** Results in each panel are means of five experiments on different preparations of adipocytes. In each experiment, each data point was the mean of five replicates expressed relative to the mean of the five replicates for uptake in the presence of insulin alone. The effects of full length chemerin and its C15-terminal peptide in wild-type versus knockout adipocytes were compared by unpaired t-tests using Welch’s correction to allow for unequal variances. **P<0.01 for the effect of C15-terminal peptide on glucose uptake in wild-type versus CMKLR1 knockout adipocytes.** The effects of chemerin and its C15-terminal peptide in the wild-type adipocytes were analysed by repeated measures (repeated preparations of adipocytes) 2-way ANOVA. † P<0.01 for the effect of the C15-terminal peptide.
Figure 7. Effects of obesity and fasting on the plasma chemerin concentration and chemerin gene (RRARES2) expression in inguinal adipose tissue. The effect of feeding on a high fat diet for 6 months from weaning on body fat mass (A; n = 7 to 9) and plasma chemerin concentration (B; n = 5 to 8) in male C57BL6 and FVB mice. Plasma chemerin (C; n = 5 to 6) and RARRES2 mRNA concentration (D; n = 5) in inguinal tissue in fed and 5 h-fasted C57BL/6 8-week-old wild-type and 10-week-old ob/ob mice. *P < 0.05; **P < 0.01; ***P < 0.001 for the comparisons indicated by the bars.
Figure 8. The effect of fasting and rosiglitazone on plasma chemerin concentration (A) and RARRES2 expression in inguinal adipose tissue (B) in female C57Bl/6J ob/ob mice. Mice were dosed with rosiglitazone (3 mg/kg body weight, p.o. daily) for three weeks. Plasma chemerin was measured before the 19th dose following a 16 h overnight fast (n = 7) and at termination in the fed state (n = 8). RARRES2 expression was measured in the fed state (n = 5). *P < 0.05; **P < 0.01; ***P < 0.001 for the comparisons indicated by the bars.