

# Serum endotrophin identifies optimal responders to PPAR $\gamma$ agonists in type 2 diabetes

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## Abstract

**Aims/hypothesis** The treatment of type 2 diabetes with full peroxisome proliferator-activated receptor gamma (PPAR $\gamma$ ) agonists improves insulin sensitivity, but is associated with weight gain, heart failure, peripheral oedema and bone loss. Endotrophin, the C-terminal fragment of the  $\alpha 3$  chain of procollagen type VI (also called Pro-C6), is involved in both adipose tissue matrix remodelling and metabolic control. We established a serum assay for endotrophin to assess if this novel adipokine could identify type 2 diabetic patients who respond optimally to PPAR $\gamma$  agonists, improving the risk-to-benefit ratio. **Methods** The BALLET trial (NCT00515632) compared the glucose-lowering effects and safety of the partial PPAR $\gamma$  agonist balaglitazone with those of pioglitazone in individuals with type 2 diabetes on stable insulin therapy. The per protocol population ( $n = 297$ ) was stratified into tertiles based on baseline endotrophin levels. Participants were followed-up after 26 weeks, after which correlational analysis was carried out between endotrophin levels and measures of glucose control. This is a secondary post hoc analysis. **Results** Endotrophin was significantly associated with therapeutic response to balaglitazone and pioglitazone. At week 26, only

individuals in the upper two tertiles showed significant reductions in HbA<sub>1c</sub> and fasting serum glucose compared with baseline. The OR for a 1% and a 0.5% reduction in HbA<sub>1c</sub> for individuals in the upper two tertiles were 3.83 (95% CI 1.62, 9.04)  $p < 0.01$ , and 3.85 (95% CI 1.94, 7.61)  $p < 0.001$ , respectively. Endotrophin levels correlated with adipose tissue mass, insulin resistance and fatty liver index. Notably, PPAR $\gamma$ -associated adverse effects, such as moderate-to-severe lower extremity oedema, only occurred in the lower tertile.

**Conclusions/interpretation** Elevated endotrophin serum levels predict response to two insulin sensitisers and reduce the risk of associated adverse effects, thereby, identifying patients with type 2 diabetes who may profit from PPAR $\gamma$  agonist treatment.

**Keywords** Balaglitazone · Efficacy · PPAR $\gamma$  · Safety · Type 2 diabetes

## Abbreviations

|               |   |
|---------------|---|
| AE            | Adverse effect  |
| ALP           | Alkaline phosphatase  |
| ALT           | Alanine aminotransferase  |
| AST           | Aspartate aminotransferase                                      |
| DXA           | Dual-energy x-ray absorptiometry                                |
| ECM           | Extracellular matrix  |
| GGT           | Gamma-glutamyl transferase                                      |
| GLP-1         | Glucagon-like peptide 1   |
| NASH          | Non-alcoholic steatohepatitis                                   |
| PHC           | Personalised healthcare   |
| PPAR $\gamma$ | Peroxisome proliferator-activated receptor gamma                |
| Pro-C6        | C-terminal pro-peptide of the $\alpha 3$ type IV collagen chain |
| SAE           | Severe adverse effect   |
| TZDs          | Thiazolidinediones  |

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## Introduction

Type 2 diabetes is a major cause of morbidity and mortality in the industrialised and developing world [1]. There are numerous treatment options including metformin, insulin, glucagon-like peptide 1 (GLP-1) agonists, dipeptidyl peptidase IV (DPP-IV) inhibitors, partial and full agonists of peroxisome proliferator-activated receptor gamma (PPAR $\gamma$ ), and drugs with alternative molecular targets [2]. However, there is a need for personalised healthcare (PHC) to identify those individuals who are likely to respond best to certain treatment options in order to increase the safety and benefits of a given intervention.

Recent research has revealed that the extracellular matrix (ECM) harbours properties of an endocrine organ. Its structural proteins generate signalling molecules that can modulate cellular processes at distant sites, including cell migration, differentiation and angiogenesis. These molecules include the potent anti-angiogenic peptide, endostatin, which is derived from type XVIII collagen, and tumstatin, vastatin and restin, which are released from types IV, VII and XV collagens, respectively [3].

The microfamentous interstitial type VI collagen, a triple helical molecule composed of the constituent chains  $\alpha 1(\text{VI})$ ,  $\alpha 2(\text{VI})$  and  $\alpha 3(\text{VI})$ , is expressed in most connective tissues and prominently in adipose tissue [4], where it anchors cells through its interconnections with other ECM proteins [5]. During formation of the microfilaments, the triple helical core of this type VI collagen is proteolytically released from its propeptide [6, 7]. Here, further cleavage of the C-terminal propeptide of the  $\alpha 3(\text{VI})$  chain generates endotrophin (also known as Pro-C6), a newly identified adipokine. Endotrophin is predominantly produced by adipose tissue and induces upregulation of TGF- $\beta$ , adipose tissue fibrosis, angiogenesis and inflammation [8]. In animal models, it has also been shown to unfavourably modulate several metabolic functions, such as insulin sensitivity, food intake, energy balance [8–10] and adipose tissue inflammation [11, 12]. These findings suggest that levels of endotrophin in the blood may be useful for classifying and/or monitoring patients with metabolic dysfunction, especially those with type 2 diabetes.

Thiazolidinediones (TZDs) are PPAR $\gamma$  agonists and have been used widely to treat type 2 diabetes because of their ability to improve insulin sensitivity, lower glucose levels and reduce the need for insulin [13, 14]. However, the use of TZDs, such as pioglitazone, has been limited substantially by associated adverse effects (AEs), such as heart failure [15], weight gain [16], peripheral oedema [17] and bone loss in women [18]. In an attempt to minimise the AEs associated with PPAR $\gamma$  agonists, partial activators of PPAR $\gamma$ , such as balaglitazone, which trigger only a subset of PPAR $\gamma$  downstream signals, have been developed [19, 20]. Such partial agonists achieve good glycaemic control with a reduced risk of AEs [21]. A serum biomarker that would optimally define

treatment responders could further improve the efficacy and safety associated with such glitazones.

Using our recently developed serum assay for the assessment of endotrophin [22], we hypothesised that endotrophin levels may predict insulin resistance and possibly identify those patients with an optimal response to insulin sensitisers, such as balaglitazone or pioglitazone. In the present work, data from participants of the BALLEET study were used to make a head-to-head comparison of balaglitazone and pioglitazone in late-stage insulin-dependent type 2 diabetes.

## Methods

**Study design** The BALLEET study was a phase III, randomised, double-blind, parallel-group, placebo- and active comparator-controlled clinical study designed to determine the efficacy and safety of 6 months of balaglitazone or pioglitazone treatment in individuals with type 2 diabetes on stable insulin therapy. The baseline demographics, CONSORT diagram and efficacy and safety data have previously been published [23]. The current serological assessment is a secondary post-hoc analysis in which we used the per protocol population of the BALLEET study (ClinicalTrials.gov registration no. NCT00515632), which consisted of 308 individuals (from whom 297 serum samples had been procured), randomised 1:1:1:1 across four groups (placebo, balaglitazone 10 mg, balaglitazone 20 mg and pioglitazone 45 mg) as previously described [23]. Existing serum samples from the BALLEET study were used and reanalysed specifically for endotrophin for the current investigation. All other measures were collected previously [23] and used in the reanalysis. Study participants gave informed consent, allowing for re-analysis of their samples, and the study was approved by the local ethics committee and carried out in accordance with the Declaration of Helsinki as revised in 2008.

**Endotrophin/Pro-C6 assay protocol** An endotrophin/Pro-C6 ELISA, targeting the C-terminus of the  $\alpha 3$  chain of endotrophin, was previously designed [22] and employed for use in the present study. This is a standard competitive ELISA, using a mono-clonal mouse detection antibody. It has previously been shown to have acceptable dilution recovery, response to spiking, and intra- and inter-individual variation [22].

**Statistical analysis** The analysis included individuals from the per protocol population, from whom a baseline measurement of serum endotrophin had been obtained. Participants were grouped into tertiles based on their endotrophin level at baseline (tertile 1,  $\leq 6.2$  ng/ml; tertile 2, 6.3–7.7 ng/ml; tertile 3,  $\geq 7.8$  ng/ml). The baseline characteristics of the three subgroups were compared by ANOVA and sex was compared by Fisher's exact test.

Spearman's ranked correlation was conducted on baseline levels of serum endotrophin, fasting serum glucose (FSG), blood HbA<sub>1c</sub>, BMI and the derived variables of HOMA-IR and the fatty liver index (FLI). The HOMA-IR was calculated according to the homeostasis model assessment

including serum glucose and insulin [24], and FLI was calculated as described by Bedogni et al [25] using the equation (triacylglycerides [mmol/l], BMI [kg/m<sup>2</sup>], gamma-glutamyl transferase [GGT; U/l] and waist circumference [cm]):

$$FLI = \frac{(e^{0.953 \times \log_e[\text{triacylglycerols}] + 0.139 \times \text{BMI} + 0.718 \times \log_e[\text{GGT}] + 0.053 \times \text{waist circumference} - 15.745})}{(1 + e^{0.953 \times \log_e[\text{triacylglycerols}] + 0.139 \times \text{BMI} + 0.718 \times \log_e[\text{GGT}] + 0.053 \times \text{waist circumference} - 15.745})} \times 100$$

Changes from baseline in FSG, blood HbA<sub>1c</sub> and serum endotrophin were studied as a function of time and treatment in each tertile. The least squares (LS) means  $\pm$ SE were estimated from a mixed-effect repeated measure

model (using baseline level, 'visit' (after 12 weeks of treatment) and end of treatment (after 26 weeks of treatment) with change from baseline as the dependent variable, and the baseline level vs visit, and end of treatment

**Table 1** Demographic and baseline characteristics of participants grouped according to baseline endotrophin level tertile

| Characteristics                    | Baseline endotrophin level tertile       |  |   | <i>p</i> value |
|------------------------------------|--|--|---|----------------|
|                                    | Tertile 1<br>≤6.2 ng/ml<br><i>n</i> = 96 | Tertile 2<br>6.3–7.7 ng/ml<br><i>n</i> = 101 | Tertile 3<br>≥7.8 ng/ml<br><i>n</i> = 100 |                |
| Treatment, <i>n</i>                |  |  |   | –              |
| Balaglitazone 10 mg                | 27                                       | 21   | 25  |                |
| Balaglitazone 20 mg                | 22                                       | 21   | 25  |                |
| Pioglitazone 45 mg                 | 24                                       | 29   | 31  |                |
| Placebo                            | 23                                       | 30   | 19  |                |
| Age (years)                        | 57.6 (8.1)                               | 60.6 (8.3)                                   | 63.4 (8.0)                                | <0.0001        |
| Sex, <i>n</i> (%)                  |  |  |   | 0.007          |
| Female                             | 21 (22)                                  | 32 (32)                                      | 43 (43)                                   |                |
| Male                               | 75 (78)                                  | 69 (68)                                      | 57 (57)                                   |                |
| BMI (kg/m <sup>2</sup> )           | 32.0 (3.9)                               | 33.6 (4.7)                                   | 34.9 (6.3)                                | 0.0005         |
| Waist circumference (cm)           | 110 (10)                                 | 114 (12)                                     | 117 (14)                                  | 0.001          |
| Hip circumference (cm)             | 109 (8)                                  | 111 (10)                                     | 115 (12)                                  | 0.0002         |
| DXA total body fat mass (kg)       | 30.8 (8.4)                               | 33.86 (8.9)                                  | 36.1 (9.8)                                | 0.0006         |
| DXA trunk fat mass (kg)            | 18.3 (5.2)                               | 20.0 (5.0)                                   | 21.7 (5.6)                                | 0.0001         |
| Blood HbA <sub>1c</sub> (%)        | 8.7 (1.4)                                | 8.4 (1.3)                                    | 8.8 (1.5)                                 | NS             |
| Blood HbA <sub>1c</sub> (mmol/mol) | 71 (15)                                  | 68 (14)                                      | 72 (17)                                   | NS             |
| FSG (mmol/l)                       | 9.4 (3.3)                                | 9.2 (3.2)                                    | 9.8 (3.4)                                 | NS             |
| Serum AST (U/l)                    | 28 (12)                                  | 32 (13)                                      | 32 (12)                                   | NS             |
| Serum ALT (U/l)                    | 31 (15)                                  | 34 (19)                                      | 33 (17)                                   | NS             |
| Serum GGT (U/l)                    | 45 (38)                                  | 55 (56)                                      | 54 (47)                                   | NS             |
| Serum ALP (U/l)                    | 163 (49)                                 | 172 (46)                                     | 187 (56)                                  | 0.004          |
| Serum bilirubin (μmol/l)           | 9.0 (3.3)                                | 9.0 (5.1)                                    | 9.0 (3.7)                                 | NS             |
| Serum triacylglycerol (mmol/l)     | 1.52 (0.94)                              | 1.85 (1.16)                                  | 2.05 (1.07)                               | 0.002          |
| Serum cholesterol (mmol/l)         | 4.34 (0.96)                              | 4.28 (0.85)                                  | 4.45 (1.04)                               | NS             |
| Serum HDL-cholesterol (mmol/l)     | 1.31 (0.35)                              | 1.23 (0.29)                                  | 1.25 (0.27)                               | NS             |
| Serum LDL-cholesterol (mmol/l)     | 2.61 (0.90)                              | 2.54 (0.76)                                  | 2.61 (0.97)                               | NS             |

Data are mean ( $\pm$  SE) unless otherwise stated

Endotrophin was measured in the current study. All other measures were obtained previously [22]

ALP: Alkaline phosphatase; ALT: Alanine aminotransferase; AST: Aspartate aminotransferase; DXA: Dual-energy x-ray absorptiometry; GGT: Gamma-glutamyl transferase; NS, non-significant

**Table 2** Spearman correlation coefficient (rho) at baseline

| Variable                   | Endotrophin | FSG  | Baseline HbA <sub>1c</sub> | HOMA-IR | FLI     | BMI     |
|----------------------------|-------------|------|----------------------------|---------|---------|---------|
| Endotrophin                | 1           | 0.07 | 0.06                       | 0.16**  | 0.32*** | 0.24*** |
| FSG                        | –           | 1    | 0.47***                    | 0.27*** | 0.20*** | 0.17**  |
| Baseline HbA <sub>1c</sub> | –           | –    | 1                          | 0.15**  | 0.17**  | 0.10    |
| HOMA-IR                    | –           | –    | –                          | 1       | 0.42*** | 0.33*** |
| FLI                        | –           | –    | –                          | –       | 1       | 0.86*** |
| BMI                        | –           | –    | –                          | –       | –       | 1       |

Data are from the placebo group only

\* $p < 0.05$ ; \*\* $p < 0.01$ ; \*\*\* $p < 0.001$

vs visit interactions as fixed effects, with an unstructured covariance structure for participants.

For each individual the mean change from baseline was calculated as AUC by the trapezoidal method, and the LS means  $\pm$  SE were estimated from an ANCOVA model with mean change as the dependent variable, baseline level as the covariate and treatment as a fixed effect. Each tertile within each of the active treatment groups was compared with the placebo group, with the level of significance adjusted for multiple comparisons using the Dunnett method. Assessment of difference between mean change from baseline compared with zero was based on the LS mean  $\pm$  SE.

All statistical calculations were performed using the SAS software package version 9.3 or higher for Windows (SAS Institute, Cary, NC, USA). Graphing was performed using

GraphPad Prism version 7.01 for Windows (GraphPad, La Jolla, CA, USA).

## Results

### Serum endotrophin is correlated with metabolic measures

The efficacy (as assessed by metabolic measures) and safety data of the treatments used in the BALLEET trial have been published elsewhere [23]. The correlations of baseline levels of endotrophin with variables associated with the metabolic syndrome are presented in Table 1. Endotrophin levels were significantly correlated with HOMA-IR, FLI, triacylglycerol and BMI, but not with FSG and HbA<sub>1c</sub> (Table 1), supporting findings that endotrophin is indeed an adipokine, related to

**Table 3** Spearman correlation coefficients at week 26

| Treatment group/variable   | Endotrophin | FSG     | HbA <sub>1c</sub> | HOMA-IR | FLI     | BMI     |
|----------------------------|-------------|---------|-------------------|---------|---------|---------|
| Placebo                    |             |         |                   |         |         |         |
| Endotrophin                | 1           | 0.05    | –0.07             | 0.28*   | 0.34**  | 0.26*   |
| FSG                        | –           | 1       | 0.24*             | 0.23*   | 0.18    | 0.26*   |
| HbA <sub>1c</sub>          | –           | –       | 1                 | 0.12    | 0.16    | 0.11    |
| HOMA-IR                    | –           | –       | –                 | 1       | 0.35**  | 0.23    |
| FLI                        | –           | –       | –                 | –       | 1       | 0.87*** |
| BMI                        | –           | –       | –                 | –       | –       | 1       |
| Pioglitazone 45 mg         |             |         |                   |         |         |         |
| Endotrophin                | 1           | –0.21   | –0.31**           | 0.02    | 0.39*** | 0.31**  |
| FSG                        | –           | 1       | 0.48***           | 0.02    | –0.11   | –0.14   |
| HbA <sub>1c</sub>          | –           | –       | 1                 | –0.06   | –0.13   | –0.12   |
| HOMA-IR                    | –           | –       | –                 | 1       | 0.30*   | 0.25*   |
| FLI                        | –           | –       | –                 | –       | 1       | 0.84*** |
| BMI                        | –           | –       | –                 | –       | –       | 1       |
| Balaglitazone 10 and 20 mg |             |         |                   |         |         |         |
| Endotrophin                | 1           | –0.22** | –0.16             | 0.07    | 0.36*** | 0.44*** |
| FSG                        | –           | 1       | 0.40***           | 0.35*** | 0.09    | –0.07   |
| HbA <sub>1c</sub>          | –           | –       | 1                 | 0.05    | 0.01    | –0.14   |
| HOMA-IR                    | –           | –       | –                 | 1       | 0.37*** | 0.26*** |
| FLI                        | –           | –       | –                 | –       | 1       | 0.85*** |
| BMI                        | –           | –       | –                 | –       | –       | 1       |

adipocyte function, fat mass and some aspects of insulin sensitivity. Endotrophin levels were not correlated with cholesterol.

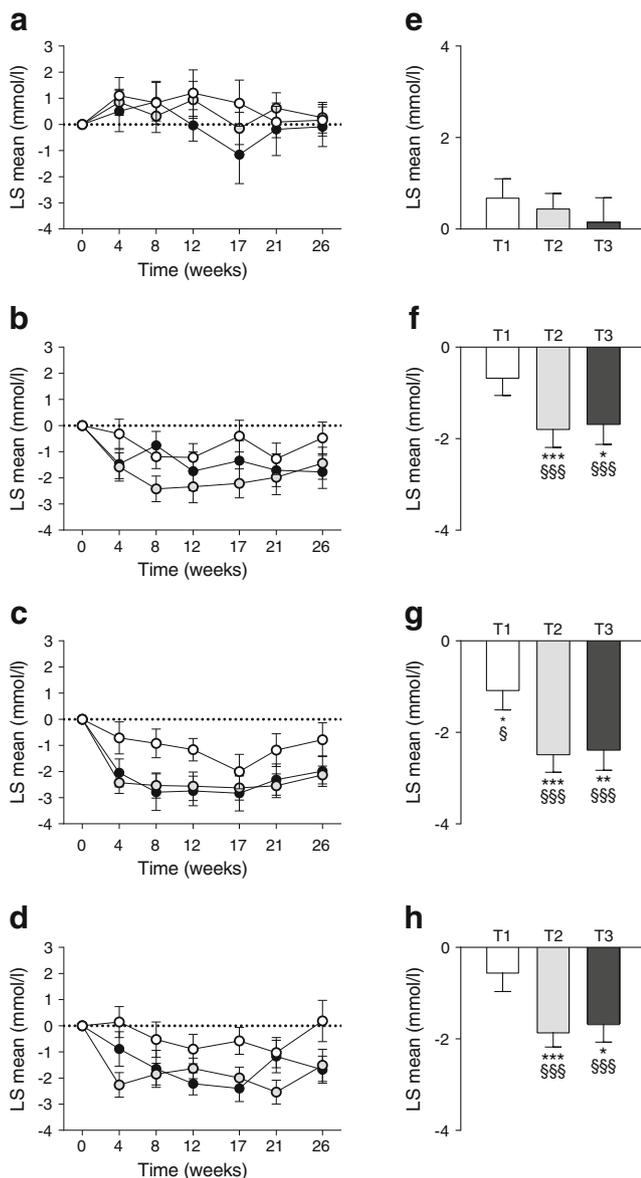
In the placebo group, the correlations observed at baseline between endotrophin and the metabolic variables, FSG, HbA<sub>1c</sub>, HOMA-IR, FLI and BMI (see Table 2), were maintained at the end of the 26 week treatment period (Table 3). However, in the

PPAR $\gamma$  agonist treatment groups, the correlation between HOMA-IR and endotrophin was eliminated, while the correlation between endotrophin and BMI or FLI persisted and even showed a trend towards being stronger (Table 3).

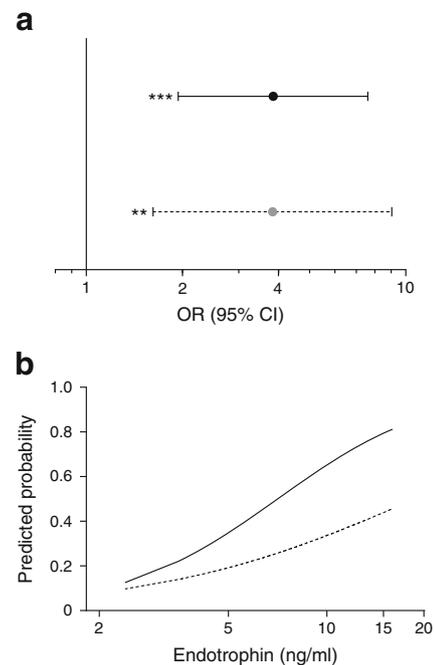
### Endotrophin identifies responders to glitazone therapy

When assessing the change from baseline in FSG over time (Fig. 1a–d) and mean per cent change in FSG over time (Fig. 1e–h), from baseline to end of treatment (week 26), it was apparent that FSG was significantly reduced by ~2.5 mmol/l compared with baseline and placebo for participants with endotrophin in the upper two tertiles in all three glitazone treatment groups. A less marked reduction in FSG, compared with baseline and placebo, was observed for participants in the lower tertile.

When response to therapy was investigated, patients in the upper two tertiles for baseline serum endotrophin were more likely to show a clinically significant response to glitazone therapy at week 26, compared with patients in the lower tertile. Furthermore, compared with participants in the lower tertile, those in the upper two tertiles had an OR of 3.83 (95% CI 1.62, 9.04) ( $p < 0.01$ ) and 3.85 (95% CI 1.94, 7.61) ( $p < 0.001$ )



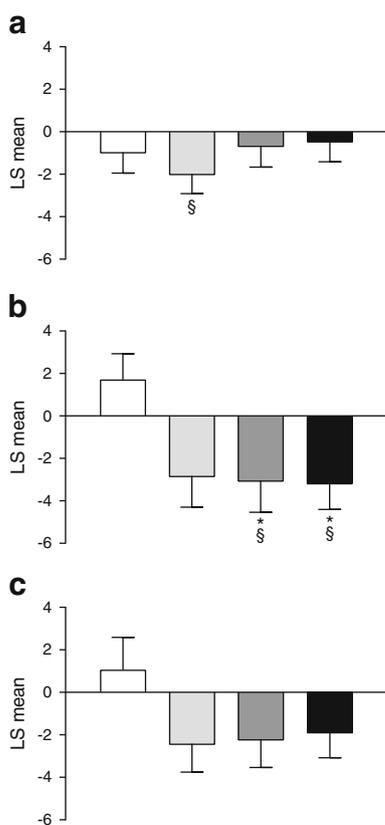
**Fig. 1** Mean change in FSG over time from baseline to end of treatment (week 26) in the (a) placebo, (b) balaglitazone 10 mg, (c) balaglitazone 20 mg and (d) pioglitazone 45 mg groups. Data were analysed using a mixed-effect repeated measures model for each tertile of baseline endotrophin. Mean absolute change in FSG from baseline to end of treatment (week 26) in the (e) placebo, (f) balaglitazone 10 mg, (g) balaglitazone 20 mg and (h) pioglitazone 45 mg groups. White circles, tertile 1; grey circles, tertile 2; black circles, tertile 3. T1, tertile 1; T2, tertile 2; T3, tertile 3. All data are LS estimates ( $\pm$  SE). \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ , Dunnett-adjusted level of significance of treatment effect against placebo; § $p < 0.05$ , §§§ $p < 0.001$ , level of significance of change from baseline



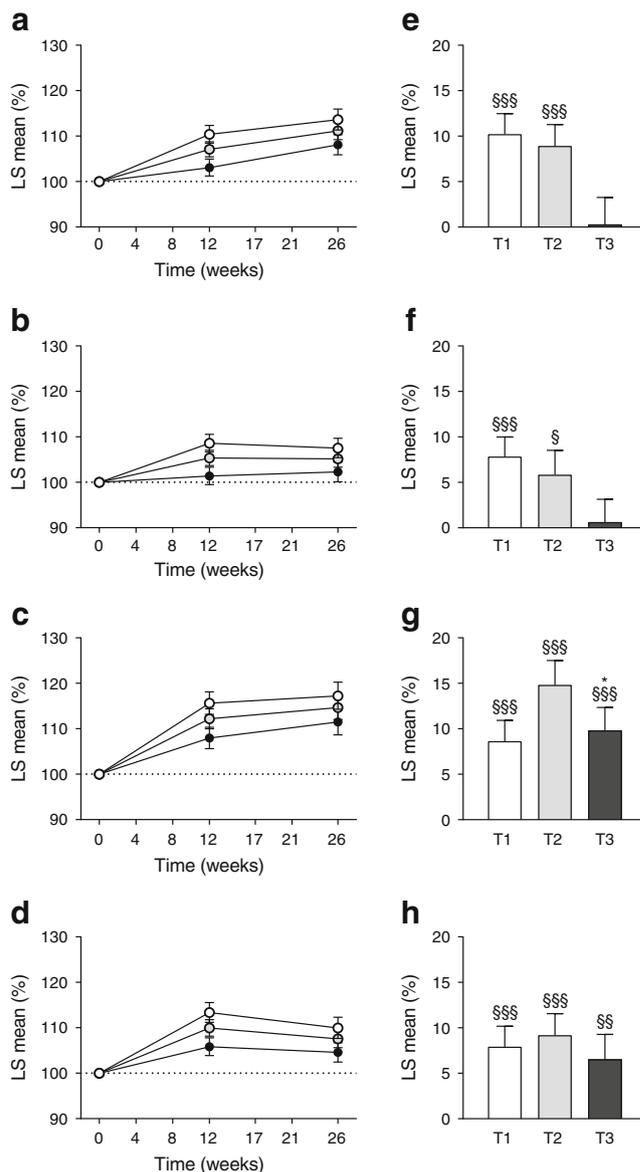
**Fig. 2** (a) Responder analysis at week 26 based on a reduction (compared to baseline) in blood HbA<sub>1c</sub> of  $\geq 0.5\%$  (black circle, solid line), or  $\geq 1.0\%$  (grey circle, dashed line). ORs (95% CI) were calculated for response to therapy based on baseline endotrophin level. The analysis compares the upper two tertiles vs the lowest tertile for treatment groups combined. \*\* $p < 0.01$ , \*\*\* $p < 0.001$ . (b) Power of baseline endotrophin levels for predicting the probability of treatment response. Data from treatment groups were combined and analysed in a logistic regression model including baseline HbA<sub>1c</sub> and baseline endotrophin as predictors. Based on regression estimates the predicted probability of endotrophin was calculated at a mean baseline level of HbA<sub>1c</sub> at 8.64% (70.9 mmol/mol)

for achieving a reduction in HbA<sub>1c</sub> of  $\geq 1\%$  or  $\geq 0.5\%$ , respectively (Fig. 2a). The power of baseline serum endotrophin for predicting the probability of responding to treatment is illustrated in Fig. 2b. For example, at a baseline serum endotrophin level of 4 ng/ml, the predicted probability of achieving a treatment response of a reduction in HbA<sub>1c</sub>  $\geq 0.5\%$  is 26%, whereas an endotrophin level of 10 ng/ml at baseline is associated with a predicted probability of treatment response of  $\sim 65\%$ . For changes in HOMA-IR during therapy, individuals in the upper tertiles again showed the strongest numerical improvement (Fig. 3a–c), albeit non-significant. Interestingly, despite the variable efficacy of therapy between tertiles there were no marked differences in weight gain (data not shown).

The effect on serum endotrophin as a function of treatment and time expressed as percentage change relative to baseline is shown in Fig. 4a–d. Figure 4e–h shows the mean change from baseline at 26-weeks. Overall, an increase in serum endotrophin was observed in the placebo group and in all three glitazone treatment groups, with the exceptions of the placebo and balaglitazone 10 mg groups within the uppermost tertile.

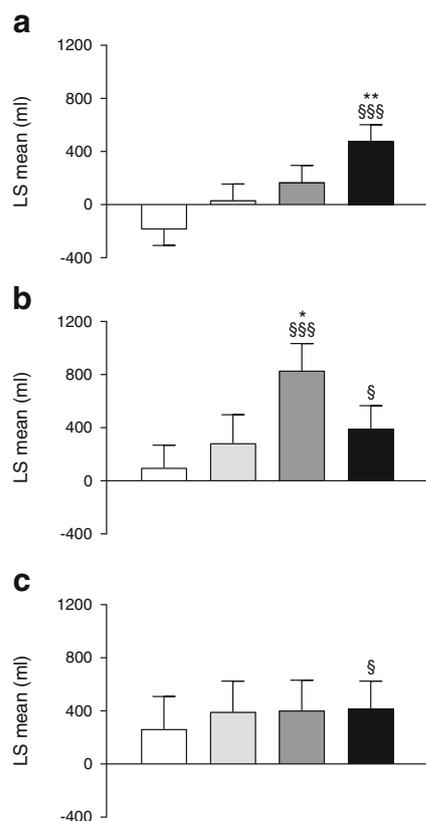


**Fig. 3** Mean absolute change from baseline in HOMA-IR during the 26 week treatment period in participants stratified into (a) tertile 1, (b) tertile 2 or (c) tertile 3 based on baseline endotrophin levels. White bars, placebo; light grey bars, balaglitazone 10 mg; dark grey bars, balaglitazone 20 mg; black bars, pioglitazone 45 mg. All data are LS estimates ( $\pm$  SE). \* $p < 0.05$ , Dunnett-adjusted level of significance of treatment effect against placebo; § $p < 0.05$ , level of significance of change from baseline



**Fig. 4** Values relative to baseline serum endotrophin level during the 26 week treatment period. (a) Relative level in the placebo, (b) balaglitazone 10 mg, (c) balaglitazone 20 mg and (d) pioglitazone 45 mg groups. Data were analysed in a mixed-effect repeated measures model. Data are presented as LS estimates ( $\pm$  SE), calculated at the geometric mean level of baseline endotrophin within each tertile: white circles, 5.2 ng/ml; grey circles, 6.9 ng/ml; black circles, 9.8 ng/ml. Mean change in endotrophin relative to baseline in the (e) placebo, (f) balaglitazone 10 mg, (g) balaglitazone 20 mg and (h) pioglitazone 45 mg groups. T1, tertile 1; T2, tertile 2; T3, tertile 3. All data are LS estimates ( $\pm$  SE). \* $p < 0.05$ , Dunnett-adjusted level of significance of treatment effect against placebo; § $p < 0.05$ , §§ $p < 0.01$ , §§§ $p < 0.001$ , level of significance of change from baseline

**Adverse effects** Lower leg oedema, measured as volume increase due to water displacement, was correlated with baseline serum endotrophin level. Certain glitazone therapies led to increased lower leg volume in the lower and middle tertiles, while there were no differences between the treatment and placebo groups in the upper tertile (Fig. 5). The AEs and



**Fig. 5** Absolute mean change in lower leg volume from baseline to 26 weeks following treatment in participants stratified into (a) tertile 1, (b) tertile 2 or (c) tertile 3 based on baseline endotrophin levels. White bars, placebo; light grey bars, balaglitazone 10 mg; dark grey bars, balaglitazone 20 mg; black bars, pioglitazone 45 mg. All data are LS estimates ( $\pm$  SE). \* $p < 0.05$ , \*\* $p < 0.01$ , Dunnett-adjusted level of significance of treatment effect against placebo; § $p < 0.05$ , §§§ $p < 0.001$ , level of significance of change from baseline

severe AEs (SAEs) observed within the different tertiles are presented in Table 4. There were no significant differences in the occurrence of AEs or SAEs between the three active treatment groups when stratified according to endotrophin level. The SAEs presented in Table 4 differ from lower leg oedema (reported in Fig. 5) since lower leg volume is a quantitative measure and oedema is a patient-reported outcome.

## Discussion

The ECM is increasingly recognised as a dynamic structure that does not only lend stability and spatial organisation to multicellular tissue and organisms, but that also transmits signals of differentiation, growth and migration to neighbouring cells. Moreover, the ECM can serve as an endocrine organ due to the signalling potential of certain fragments released by proteolysis, creating ligands with paracrine and potential endocrine functions [3]. Endotrophin (Pro-C6) is a soluble fragment of the C-terminal  $\alpha 3$  chain of type VI procollagen that is

released by naturally occurring proteolysis [4, 8, 22, 26]. Here we demonstrate that serum endotrophin levels are predictive of a response to the insulin sensitisers, pioglitazone and balaglitazone, in patients with type 2 diabetes. Individuals within the upper two tertiles for baseline endotrophin serum levels (tertile two, 6.3–7.7 ng/ml and tertile three,  $\geq 7.8$  ng/ml) were four times more likely to respond to treatment, compared with patients in the lower tertile ( $\leq 6.2$  ng/ml baseline serum endotrophin). Since the glitazones are associated with safety concerns, such as non-fatal heart failure and bone fractures [15, 27–31], identification of individuals who will gain the most benefit from treatment and also the fewest AEs is crucial for the continued use of these drugs, which are still considered highly effective insulin sensitisers. Participants in the upper two tertiles who responded to treatment with a reduction in FPG and HbA<sub>1c</sub> levels were at a reduced risk of developing lower leg oedema, a major AE associated with glitazone treatment. These efficacy and safety data are highly relevant for predicting benefit-to-risk ratios for patients treated with glitazones; this should also apply when the medications are considered for other indications, such as liver disease [32–34].

In animal models, endotrophin levels have been shown to reflect insulin sensitivity, food intake and energy balance, as well as angiogenesis [4, 5, 8]. Accordingly, suppression of endotrophin improves insulin sensitivity and attenuated adipose tissue inflammation in animals [8]. This correlates well with our findings that elevated serum endotrophin levels are indicative of a response to PPAR $\gamma$  agonists. Furthermore, mRNA levels of the endotrophin precursor procollagen  $\alpha 3(\text{VI})$  are upregulated in obese adipose tissue, paralleling adipose tissue inflammation and fibrosis [9]. The ECM and its products, especially procollagen type VI and endotrophin, may be of particular relevance in fatty liver, including in the severe form of the disease, non-alcoholic steatohepatitis (NASH). NASH is a metabolic-fibrotic disorder of the liver that shows at least a partial overlap with type 2 diabetes [32–34]. Accordingly, we expect that this novel biomarker should assist in the diagnosis and management of NASH patients where insulin sensitisers may be beneficial for subpopulations, both for the treatment of insulin resistance and liver fibrosis. Here, ECM components, in particular collagens/collagen type VI, and their functional role in the transition of fatty liver to overt fibrotic NASH needs to be further investigated.

In line with this, in the current study we observed a strong correlation between serum triacylglycerol and FLI, an index that correlates with NASH inflammatory activity and predicts more severe liver fibrosis [25]. In support of a role for type VI collagen in NASH-related fibrosis, studies have demonstrated its prominent expression in areas of active scar formation [35, 36], and elevated serum levels of the collagen VI core structure (which lacks the endotrophin domain) have been shown to be associated with advanced liver fibrosis in rodents [37], humans [38–41], and with elevated portal pressure [42]. The expression

**Table 4** AE profile for tertiles/treatment group

| AE type                | Placebo     | Balaglitazone 10 mg | Balaglitazone 20 mg | Pioglitazone 45 mg |
|------------------------|-------------|---------------------|---------------------|--------------------|
| AEs                    |             |                     |                     |                    |
| Tertile 1              |             |                     |                     |                    |
| Number of participants | 23          | 27                  | 22                  | 24                 |
| All AEs                | 16 (70%) 30 | 20 (74%) 38         | 17 (77%) 33         | 19 (79%) 45        |
| SAEs                   | 0 (0%) 0    | 1 (4%) 1            | 1 (5%) 1            | 1 (4%) 1           |
| Tertile 2              |             |                     |                     |                    |
| Number of subjects     | 30          | 21                  | 21                  | 29                 |
| All AEs                | 23 (77%) 51 | 17 (81%) 35         | 15 (71%) 36         | 17 (59%) 37        |
| SAEs                   | 3 (10%) 3   | 1 (5%) 1            | 0 (0%) 0            | 2 (7%) 3           |
| Tertile 3              |             |                     |                     |                    |
| Number of subjects     | 19          | 25                  | 25                  | 31                 |
| All AEs                | 14 (74%) 41 | 20 (80%) 55         | 20 (80%) 43         | 25 (81%) 66        |
| SAEs                   | 0 (0%) 0    | 1 (4%) 1            | 6 (24%) 6           | 4 (13%) 6          |
| SAEs                   |             |                     |                     |                    |
| Tertile 1              |             |                     |                     |                    |
| Heart failure          | 0 (0%) 0    | 0 (0%) 0            | 0 (0%) 0            | 0 (0%) 0           |
| Cardiac ischaemia      | 1 (4%) 1    | 0 (0%) 0            | 2 (9%) 2            | 0 (0%) 0           |
| Peripheral oedema      | 0 (0%) 0    | 2 (7%) 2            | 2 (9%) 2            | 5 (21%) 5          |
| Total SAEs             | 1 (4%) 1    | 2 (7%) 2            | 4 (18%) 4           | 5 (21%) 5          |
| Tertile 2              |             |                     |                     |                    |
| Heart failure          | 0 (0%) 0    | 0 (0%) 0            | 0 (0%) 0            | 0 (0%) 0           |
| Cardiac ischaemia      | 1 (3%) 1    | 1 (5%) 1            | 0 (0%) 0            | 1 (3%) 1           |
| Peripheral oedema      | 1 (3%) 1    | 3 (14%) 3           | 2 (10%) 2           | 5 (17%) 5          |
| Total SAEs             | 2 (7%) 2    | 4 (19%) 4           | 2 (10%) 2           | 5 (17%) 6          |
| Tertile 3              |             |                     |                     |                    |
| Heart failure          | 0 (0%) 0    | 0 (0%) 0            | 0 (0%) 0            | 1 (3%) 1           |
| Cardiac ischaemia      | 1 (5%) 1    | 0 (0%) 0            | 1 (4%) 1            | 3 (10%) 4          |
| Peripheral oedema      | 1 (5%) 2    | 1 (4%) 1            | 1 (4%) 1            | 2 (6%) 2           |
| Total SAEs             | 2 (11%) 3   | 1 (4%) 1            | 2 (8%) 2            | 6 (19%) 7          |

Data are presented as *n* (%) number of events

of procollagen  $\alpha 3(\text{VI})$  is regulated by PPAR $\gamma$  and our findings are in alignment with this. In fact, procollagen  $\alpha 3(\text{VI})$  mRNA is suppressed by PPAR $\gamma$ , as demonstrated by an increase in procollagen  $\alpha 3(\text{VI})$  mRNA expression in adipocyte cultures treated with siRNA against PPAR $\gamma$  [9] and by a decrease in its transcripts in subcutaneous adipose tissue of individuals with type 2 diabetes treated with the PPAR $\gamma$  agonist pioglitazone, especially in patients with high baseline tissue levels of procollagen  $\alpha 3(\text{VI})$  mRNA [12]. These data may in part explain the change in correlation, from baseline to the end of glitazone treatment, between endotrophin serum levels and HbA $_{1c}$  or HOMA-IR, and in particular the lack of a correlation between endotrophin and the metabolic variables. Another clinical study provides further supportive evidence by showing that tissue endotrophin levels in obese individuals correlated with chronic inflammation and systemic insulin resistance [10, 26].

Generally, the ECM has, until now, been considered to be a passive scaffold. Type VI collagen has mostly been recognised

through mutations in the genes *COL6A1*, *COL6A2* and *COL6A3* that encode its three constituent chains. These mutations cause muscle disorders, such as Bethlem myopathy, Ullrich congenital muscular dystrophy, limb-girdle muscular dystrophy and autosomal recessive myosclerosis [43–46]. This provides an interesting link to metabolic dysfunction, since type VI collagen mutations are associated with muscle abnormalities and muscle represents an important regulator of insulin resistance. Therefore, the available evidence strongly suggests that collagen type VI is more than a passive ECM component, but an important mediator of adipose (and liver) metabolic dysfunction related to insulin resistance, type 2 diabetes and NASH.

Of note, the measurement of only one standard biochemical variable is a weakness of the current study; because of limited sample availability we were only able to assess endotrophin levels in serum for comparisons with HbA $_{1c}$ , limiting the possibility of investigating other potential biomarkers for response to treatment.

In conclusion, elevated circulating levels of endotrophin, predominantly derived from adipocytes and adipose tissue, are correlated with insulin resistance and are predictive of the response to insulin sensitisers. This enables the identification and monitoring of patients who will respond optimally to an insulin sensitiser, improving the benefit-to-risk ratio for PPAR $\gamma$  agonists in the treatment of type 2 diabetes and possibly NASH.

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