

**INTERDISCIPLINARY APPROACH TO THE PREDIABETIC STATE:
BONE – ENERGY HOMEOSTASIS AXIS, CLINICAL AND
ETIOLOGICAL ASSOCIATIONS OF NON-ALCOHOLIC FATTY
LIVER DISEASE AND INSULIN RESISTANCE**

Barbara Buday MD

PhD Thesis

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- I. **Buday B**, Izsóné Katz M, Nagy E, Papp Zs, Korányi L: A metabolikus szindróma, 2-es típusú diabetes és a csontvesztés epidemiológiai összefüggései a Balaton felvidék felnőtt lakossága körében [Relationship of cardiovascular risk factors and bone status in a large adult population of the Balaton Region]. *Ca & Csont* 2007; 10(04): 132-137.

- II. **Buday B**, Horváth T, Literáti Nagy B, Kulcsár E, Barta K, Salamon Cs, Péterfai É, Korányi L: A hagyományosan használt inzulinrezisztencia- és béta-sejt-funkciós indexek diagnosztikus értéke [The diagnostic value of traditional insulin sensitivity and beta cell function indices]. *Diabetologia Hungarica* 2007; 15: 93-105.

- III. **Buday B**, Pach FP, Literati-Nagy B, Vitai M, Vécsei Z, Korányi L: Serum osteocalcin is associated with improved metabolic state via adiponectin in females versus testosterone in males. Gender specific nature of the bone-energy homeostasis axis. *Bone* 2013; 57(1): 98-104.

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- II. **Buday B**, Kulcsár E, Lieráti-Nagy B, Horváth T, Vitai M, Vecsei I, Bezzegh K, Kiss J, Péterfai É, Koltay L, Korányi L: Az osteocalcin helye a human cukor- és csontanyagcsere kapcsolatában. [The role of osteocalcin in the connection of bone and glucose metabolism in humans]. *Orvosi hetilap* 2008; 149(52):2453-61.

- III. Literáti-Nagy B, Péterfai É., Kulcsár E, Literáti-Nagy Zs, **Buday B**, Tory K, Mandl J, Sümegi B, Fleming A, Roth J, Korányi L: Beneficial effect of the insulin sensitizer (HSP inducer) BGP-15 on olanzapine-induced metabolic disorders. *Brain Research Bulletin* 2010;83(6): 340–344.
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- IV. Literáti-Nagy B, Paragh Gy, Szilvássy Z, Kolonics A, Tory K, Literáti-Nagy Zs, Barta K, Horváth T, **Buday B**, Kulcsár E, Péterfai E and Korányi L: Improvement of insulin sensitivity by a novel drug, BGP-15, in insulin-resistant patients: a proof of concept randomized double-blind clinical trial. *Horm Metab Res* 2010; 41(5):374-380.
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- V.** Vitai M, **Buday B**, Kulcsár E, Literáti-Nagy B, Vecsei I, Bezzegh K, Péterfai É, Kurucz I, Korányi L: A GRB10 gén (+11275G>A) polimorfizmusának hazai előfordulása és kapcsolata a cukoranyagcserével. [Occurrence of GRB10 (+11275G > A) polymorphism in Hungarian population and its relationship to glucose metabolism]. *Orvosi Hetilap* 2009;150(40):1845-51.
- VI.** Vitai M, Kocsordi K, **Buday B**, Literáti Nagy B, Enikő Kulcsár E, Bezzegh K, Péterfai, E, Koltay L and Korányi L: Nemhez kötött a katalázgén-polimorfizmus (RS769217) hatása az energia-háztartásra és a csontok állapotára. [Effects of catalase gene (RS769217) polymorphism on energy homeostasis and bone status are gender specific]. *Orvosi Hetilap* 2010;151(23): 923-931.
- VII.** Kiss J, **Barbara B**, Literáti-Nagy B, Dr. Faluközi J, Fogarassy Gy, Apró D, Vecsei I, Fék A, Veress G, Korányi L: A koszorúérbetegség és a csontállapot kapcsolata másképp: a lumbalis csigolyadenitáz a koszorúérbetegség pozitív prediktora nőkben? *LAM KID* 2011;1(3):43-47.
- VIII.** **Buday B**, Pach FP, Literáti-Nagy B, Vecsei Zs, Korányi L: A csontátépülés és az energia háztartás nőkre jellemző kapcsolatai. *LAM-KID* 2011;1(2):30-35.
- IX.** Pauer J, Fék A, **Buday B**, Literáti-Nagy B, Pach P, Vitai M, Péterfai É, Korányi L: Anyagcsere-eltérések a 2-es típusú cukorbetegség egészséges, első fokú férfi rokonaiban. [Metabolic alterations in healthy men with first degree type 2 diabetic relatives]. *Orvosi Hetilap* 2013; 154 (5): 178-186

X. Kovács Gy, **Buday B**, Fék A, Literáti-Nagy B, Pauer J, Pach P, Vitai M, Péterfai E, Korányi L: Anyagcsere-eltérések a 2-es típusú cukorbetegség egészséges, elsőfokú nőrokonaiban. [Metabolic differences in healthy first-degree female relatives of type 2 diabetic patients]. *Orvosi Hetilap* 2013; 154(44):1747-53.

XI. Fék A, **Buday B**, Kovács Gy, Vitai M, Vecsei Z, Pauer J, Literáti-Nagy B, Bezzegh K, Péterfai É, Korányi L: A genetikai diabeteskockázat hatása a csontanyagcsere-energiaháztartás kapcsolatokra. [The effect of genetic risk of diabetes on the correlations in bone and energy homeostasis]. *Orvosi Hetilap* 2015;156(25):1007-13.

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Abbreviations

AC:	Abdominal circumference
cAMP:	Cyclic adenosine monophosphate
ADA:	American Diabetes Association
AIR:	Acute insulin response
ALP:	Alkaline phosphatase
ALT:	Alanine aminotransferase
AST:	Aspartate aminotransferase
ATP:	Adult Treatment Panel
AUC:	Area under the curve
BCF:	Beta cell function
BFP:	Body fat percent
BMD:	Bone mineral density
BMI:	Body mass index
CI:	Confidence interval
CV:	Coefficient of variations
DEXA:	Dual-energy X-ray absorptiometry
DI:	Disposition index
FBG:	Fasting plasma glucose
FFA:	Free fatty acid
FSH:	Follicular stimulating hormone
GGT:	Gamma-glutamyl transferase
GI:	Glucose intolerant
GIR:	Glucose infusion rate
HbA1c:	Hemoglobin A1c
HDL:	High density lipoprotein

HIRI:	Hepatic insulin resistance index
HOMA:	Homeostatic Model Assessment
IFCC:	International Federation of Clinical Chemistry
IFG:	Impaired fasting glucose
IGI:	Insulogenic index
IGT:	Impaired glucose tolerance
IL-6:	Interleukin-6
IR:	Insulin resistance
IS:	Insulin sensitivity
ISI _{Cederholm} :	Cederholm's insulin sensitivity index
ISI _{comp} :	Composite insulin sensitivity index
ISI _{est} :	Estimated insulin sensitivity index
IVGTT:	Intravenous glucose tolerance test
LDL:	Low density lipoprotein
MAD:	Median absolute deviation
MCR _{est} :	Estimated metabolic clearance rate
NAFLD:	Non-alcoholic fatty liver disease
NGT:	Normal glucose tolerance
OCN:	Osteocalcin
OGIS:	Oral Glucose Insulin Sensitivity
OGTT:	Oral glucose tolerance test
OR:	Odds ratio
OPG:	Osteoprotegerin
P1NP:	Procollagen type 1 amino-terminal propeptide

RANKL:	Soluble receptor activator NF- κ B ligand
RR_Dias:	Diastolic blood pressure
SD:	Standard deviation
SHBG:	Sex hormone binding protein
T1DM:	Type 1 diabetes mellitus
T2DM:	Type 2 diabetes mellitus
TNF- α :	Tumor necrosis factor - alpha
VLDL:	Very low density lipoprotein
WHO :	World Health Organization

1. Introduction and aims of our study

Major challenges of the 21st century health care of the developed world include type 2 diabetes mellitus (T2DM) and the bone loss epidemic. The parallel increase of the two diseases poses some contradiction. Both are associated with body weight, however increasing obesity and insulin resistance have a causal role in the pathogenesis of T2DM. Conversely, in osteoporosis the higher body fat content has a bone protecting effect. Type 1 diabetic patients have decreased bone mineral density (BMD) and increased fracture risk while in T2DM this association is less strong, data exist about both increased or decreased BMD in T2DM patients [1, 2].

In order to investigate the relationship between insulin resistance (IR) or insulin sensitivity (IS) and other diseases / symptoms, like osteopenia and osteoporosis that may be associated with insulin resistance we need an easily accessible simple IR measuring method that is cheap, can be used in a large number of patients, reproducible and is validated via more sophisticated studies. Precise measurement of IR is also important for the prevention, diagnosis and the therapeutic follow up of T2DM. For measuring IS, today the “gold standard” is still the “hyperinsulinemic normoglycemic clamp” developed by DeFronzo et al. [3]. However, it is an expensive and time consuming method which cannot be used in a large number of patients in clinical setting, so there have been a number of attempts to develop methods replacing the clamp, e.g. the Homeostasis Model Assessment (HOMA) indices which use data from fasting blood samples, or so are a number of indices derived from the oral glucose tolerance test (OGTT).

The first part of my work deals with the diagnostic evaluation of IS in terms of simple fasting and OGTT derived indices which still seem to be a hurdle in IS estimation since the most widely used HOMA indices in clinical practice do not correlate well with the gold standard clamp methods. We aimed to gain further insights into the pathophysiology and diagnosis of IR and related complications by studying the association between transaminase levels and clamp measured insulin sensitivity, moreover we sought to explore a unique side of the gender specific aspect of insulin homeostasis / energy metabolism, with special regard on the non-alcoholic fatty liver disease (NAFLD) which is one of the major link between insulin resistance and cardiovascular disease. By exploring the pathophysiology of the IR related steatohepatosis often associated with the ‘unexplained’ elevation of transaminase levels in overweight insulin resistant / T2DM patients, we might be able to improve the value

of the HOMA model with no extra costs, although possible gender related differences will have to be taken into account.

To explore the relationship between T2DM / IR and bone homeostasis first we analyzed data from a large epidemiological study which included the screening results of more than 6000 people at Balaton Upper-lands. Based on the results of this epidemiological study, we conducted a cross over analysis on our existing clamp database where we measured markers of bone turnover, i.e. total non-carboxylated osteocalcin (OCN) levels and other metabolic-hormonal factors, like adipocytokines, lipids, lipoproteins, sex hormones. Previous human studies have shown that serum OCN concentration is negatively associated with the plasma glucose level and body fat mass [4-7] and positively associated with insulin secretion [8, 9], lower insulin resistance [5, 6, 10] and higher serum adiponectin concentration [4, 10]. In most of this work, the HOMA model has mainly been used to assess β -cell function, insulin sensitivity and the involvement of OCN on glucose metabolism, although we and others [11] have shown that fasting indices do not always correlate well with the real insulin resistance, therefore insulin sensitivity was measured by the gold standard clamp method. Recently, it has been demonstrated that osteoblasts are able to induce testosterone production by the testes, though they fail to influence estrogen production by the ovaries [12]. The role of testosterone in the bone–energy homeostasis is presumably gender-specific, as the effects of OCN were only demonstrated in Leydig cells and not in the ovaries; moreover, low testosterone levels are only associated with a metabolic syndrome in men [13].

Based on previous data and our preliminary assumptions discussed above our main goals were:

1. To explore the diagnostic value of the most frequently used simple fasting and OGTT derived insulin sensitivity indices compared to the gold standard clamp method.
2. To try to justify possible new approaches / directions in the simple diagnostics of insulin resistance.
3. To explore epidemiological characteristics of the relationship between bone loss and diabetes / insulin resistance.
4. To explore molecular background of the bone – energy homeostasis axis, by analyzing the associations between clamp measured insulin sensitivity, total-OCN and other metabolic biomarkers.

5. Since our previous data suggested that basic sex differences exist in the pathogenesis and manifestations of insulin resistance and associated diseases, male and female populations were separately analyzed in most of our studies to address this issue.

2. Patients and methods - General considerations

2. 1. Epidemiologic study

In our study we have analyzed the results of general screening tests of the adult population in Balaton Upper-lands which was performed between 2003 and 2006. Screenings were done on a voluntary basis advertised as primarily bone density screening measurements in local surgeries or at working places (Offices, Factories). Since the screening examinations involved bone density measurements / osteoporosis screening, there were more women than men who attended the screenings.

During screening examinations anthropometric assessments (body weight and height, abdominal circumference [AC]), sitting blood pressure (Omron 705CP digital equipment), blood sugar (Personal DCont and Optimum, 77 Electronics, Hungary), total cholesterol measurements (Accutrend GCT 1537962 and Accutrend GC 1418246, Roche, Germany) from capillary blood and calcaneus bone ultrasound density ultrasound measurements (GE-Lunar, Achilles Plus, USA) were carried out. Questions about medical history, concomitant medications and life style were also raised. Screenings were not done always in fasting state although data were available about the time of last meal besides previous concomitant treatment for diabetes. We have analyzed the following data: sex, age, systolic and diastolic blood pressure, plasma glucose, total-cholesterol, AC, BMI and T score calculated by the calcaneus ultrasound density data of young and healthy population of the same sex.

Based on available data and the results of the screening examinations we defined certain diseases / syndromes to be present in the population, like diabetes, hypertension, metabolic syndrome, osteopenia and osteoporosis. 'Bone density' measured by ultrasound is not identical to BMD measured by 'dual-energy X-ray absorptiometry' (DEXA), due to the effect of bone structure, albeit they are closely associated and as such it is widely accepted in both the diagnosis and the follow up of osteoporotic treatment [14]. Data were analyzed by sex and age groups.

2.2. Clamp studies

2.2.1. OGTT, IVGTT and clamp

All clamp studies were carried out after receiving signed informed consents from the subjects. Study was approved by the Hungarian Central Ethical Committee (A12988-2/2003-1018-EKU, ad.8-311/2009-1018EKU). Patients and healthy volunteers were recruited from our own diabetes outpatient clinic and by referral from regional primary care physicians. All subjects underwent a standard 75 mg oral glucose tolerance test (OGTT) which determined the subject's basal glucose tolerance (i.e. normal or impaired glucose tolerance, impaired fasting glucose or T2DM) during the screening period. Within 3 weeks after the OGTT patients were hospitalized for the clamp. Subjects fasted on the night prior to the clamp examination. They first underwent an intravenous glucose tolerance testing (IVGTT) to assess insulin secretion (0.3g/body weight kg iv. glucose injection). Following the IVGTT, a hyperinsulinaemic normoglycaemic clamp examination was carried out, as described by DeFronzo et al. [3]. During a continuous infusion of insulin ($45 \text{ mU} \times \text{min} \times \text{m}^{-2}$) and glucose (20 %), the steady state was set at the constant glucose infusion rate (earliest from the 120th minute of clamp), where blood sugar level stayed between 5.0 and 5.9 mM / l for at least 30 min after the beginning of steady state. Glucose and insulin levels were measured from venous blood at 0-, 3-, 5-, 10-, 20-, 30-, 40-, 50th- and 60th min samples of IVGTT, before the beginning, and at the 0-, 10- 20-, 30th min samples of the steady state of clamp. Insulin secretion was determined from IVGTT by the insulogenic index [$\text{IGI} = \Delta (\text{insulin}_{5'} - \text{insulin}_{3'}) / \Delta (\text{glucose}_{5'} - \text{glucose}_{3'})$] and the acute insulin response [$\text{AIR} = (\text{insulin}_{5'} + \text{insulin}_{3'}) / 2 - \text{insulin}_{0'}$]], both being sensitive indicators of the first phase insulin response, and hence the real beta cell function. For the liver function test – IS relationship study hepatic insulin resistance index (HIRI) was estimated from the OGTT 0th and 30th min glucose and insulin values [$\text{HIRI} = (\text{GLU-AUC}_{0-30'}) \times (\text{Ins-AUC}_{0-30'})$] described by Muhammad et al. [15]. Glucose and insulin area under the curve (AUC) values were calculated using the trapezoidal rule, both from OGTT and IVGTT. We used lean body (= muscle)-adjusted glucose uptake (M3 value, mg/min/kg) calculated from the glucose infusion rates during clamp, to measure peripheral (muscle) glucose utilization rates. Formula for calculation of serum glucose levels from mmol/l to mg/dl for the clamp M3 value: $\text{mg/dl} = 18 \times \text{mmol/l}$. Body composition was determined by dual-energy X-ray absorptiometry (DPXMD+, GE-Lunar, USA, Florida). In the insulin sensitivity diagnostic evaluation study we used fasting and OGTT derived insulin sensitivity indices.

2.2.2. Calculations of insulin sensitivity indices

2.2.2.1. Data derived from the clamp examinations [3]

Glucose infusion rate (GIR), ($\text{mg} \times \text{kg}^{-1} \times \text{min}^{-1}$): the glucose infusion rate necessary to keep a steady state between the 120th and 150th minutes of the clamp.

M1: whole body glucose uptake ($\text{mg} \times \text{kg}^{-1} \times \text{min}^{-1}$)

M2: glucose uptake adjusted for body surface ($\text{mg} \times \text{m}^{-2} \times \text{min}^{-1}$)

M3: glucose uptake adjusted for lean body (muscle) mass ($\text{mg} \times \text{kg} \text{ muscle}^{-1} \times \text{min}^{-1}$)

For clamp indices we used the whole body glucose uptake (M1) and muscle mass adjusted glucose uptake (M3) to evaluate the individual OGTT / IVGTT derived IS indices.

2.2.2.2. OGTT insulin sensitivity indices

MCR_{est} (Estimated Metabolic Clearance Rate by Stumvoll) = $18.8 - 0.271 \times \text{BMI} - 0.0052 \times \text{Ins}_{120} - 0.27 \times \text{Glucose}_{90}$ ($\text{ml} \times \text{kg}^{-1} \times \text{min}^{-1}$),

where Ins_{120} is the 120th minute insulin level, and Gluc_{90} is the 90th minute glucose level of the OGTT [16].

$$\text{ISI}_{\text{cederholm}} = \frac{\{75.000 + (\text{Gluc}_0 - \text{Gluc}_{120}) \times 1.15 \times 180 \times 0.19 \times \text{kg-bodyweight}\}}{(120 \times \log(\text{Ins}_m) \times \text{Gluc}_m)}$$

($\text{ml} \times \text{kg}^{-1} \times \text{min}^{-1}$), where Gluc_0 is the basal, Gluc_{120} is the 120th minute, Gluc_m is the mean of all OGTT glucose levels, and Ins_m is the mean of all OGTT insulin levels [17].

ISI_{est} (Estimated Insulin Sensitivity Index) = $0.226 - 0.0032 \times \text{BMI} - 0.0000645 \times \text{Ins}_{120} - 0.0037 \times \text{Gluc}_{90}$

($\mu\text{mol} \times \text{kg}^{-1} \times \text{min}^{-1} \times \text{pM}^{-1}$), where Ins_{120} is the 120th minute insulin level, Gluc_{90} is the 90th minute glucose level of the OGTT [16].

OGIS = $f(G_0, G_{90}, G_{120}, I_0, I_{90}, D_0)$, is a function of the 0th, 90th, 120th minutes glucose levels and the 0th, 90th minute insulin levels of the OGTT ($\text{ml} \times \text{min}^{-1} \times \text{m}^{-2}$), calculated by the OGIS calculator accessible from <http://webmet.pd.cnr.it/ogis/ogis.php> where D is the glucose dose employed [18].

$$ISI_{\text{comp}} (\text{Composite Insulin Sensitivity Index, Matsuda}) = \frac{10.0000}{\sqrt{(\text{Gluc}_0 \times \text{Ins}_0 \times \text{Gluc}_m \times \text{Ins}_m)}}$$

where Gluc_0 is the 0th minute glucose, Gluc_m is the mean of all OGTT glucose levels, Ins_0 is the 0th minute insulin, Ins_m is the mean of all OGTT insulin levels [19].

2.2.2.3. Fasting indices

HOMA (Homeostasis Model Assessment) [20]:

HOMA-R (for insulin resistance) = $G_0 \times I_0 / 22.5$, (HOMA-1), where G_0 is the basal glucose level (mM/L), I_0 is the basal insulin level (uU/ml).

HOMA-S (for insulin sensitivity) = $1 / \text{HOMA-R}$

QUICKI = $1 / \{\log(I_0) + \log(G_0)\}$ (logarithmic transformation of HOMA) [21].

HOMA-B (for insulin release) = $I_0 \times 20 / (\text{Gluc}_0 - 3.5)$

HOMA-S% and B% (HOMA-2): indices of insulin sensitivity (S%) and beta cell function (B%), calculated by the HOMA calculator V2.2 from the 0th minute IVGTT samples, downloaded from <https://www.dtu.ox.ac.uk/homacalculator> ('the Oxford Model') [22, 23].

FFA-QUICKI = $1 / \{\log(I_0) + \log(G_0) + \log(\text{FFA}_0)\}$, where G_0 is the basal glucose, I_0 is the basal insulin, FFA is the basal fatty acid levels (mmol/l) [24].

2.2.3. Biochemical measurements

Routine biochemical parameters were measured on Cobas Mira and Hitachi 912 laboratory automats with the same method (according to IFCC recommendations) during the recruitment period (2003 – 2008). Reference ranges, detection limits and test principles were unchanged during this test period. Alanine aminotransferase (ALT), aspartate aminotransferase (AST), gamma-glutamyl transferase (GGT), alkaline phosphatase (ALP), serum bilirubin, free fatty acid (FFA), insulin, glucose, HbA1c levels and conventional lipid parameters were determined using Roche reagents (Roche Diagnostics, Germany). HbA1c levels were measured by the IFCC reference method. Total (non-carboxylated) OCN, estradiol, testosterone, follicular stimulating hormone (FSH) and serum insulin levels were

measured with an Elecsys 2010 electrochemiluminescence automat (Roche Diagnostic, Germany). Coefficient of variation (CV) for osteocalcin test varies between 1.8 and 6.5% respectively for the kits used in our study. Serum leptin, adiponectin, interleukin-6 (IL-6) and tumor necrosis factor- α (TNF- α) levels were measured by the enzyme-linked immunosorbent method (Quantikine DLP00, Quantikine DRP300, Quantikine HS600B and Quantikine HSTA00D kits respectively; R&D Systems, Minneapolis, MN, USA). Lipid fractionation was done by the Lipoprint System® Quantimetrix, USA). Lipid subfractions (very low density lipoproteins [VLDL], intermediate-density lipoproteins [IDL-A, -B and -C], and low-density lipoproteins [LDL1-4 subfractions, LDL 2-4 subfractions = small-dense LDL]), total LDL and high-density lipoprotein [HDL] were separated by acrylamide gel electrophoresis.

2.2.4. Statistics

2.2.4.1. Epidemiological study

Data analysis for the epidemiologic study was carried out with an SPSS 10.0 Program (Statistics for Windows). Numeric data were indicated as mean and standard deviations. Mean data between the groups were compared with two-sided tests. For comparing disease prevalence within groups of different genders and ages we used the χ^2 test (in small sample numbers χ^2 was calculated by continuity correction), additionally we have applied the Fisher test. Significance level was considered at $p < 0.05$ values, with indicating odds ratios (OR) and 95% confidence intervals (CI). The relationships between T score and the individual metabolic parameters were analyzed by bivariate correlations (Pearson).

2.2.4.2. Evaluation of fasting and OGTT indices

Coefficients of variations of fasting and OGTT indices were calculated according to the equation $CV = (SD / \sqrt{2}) \times 100/\bar{X}$, where SD is the standard deviation of the intra-subject changes of indices, and \bar{X} is the mean of all values. For evaluating the simple insulin sensitivity indices we used a score system to evaluate the individual indices and all of the fasting or OGTT indices within the groups divided by several aspects (Table 1.). It determined the value of the indices based on the correlation coefficients and their significance levels between the M1 and the specific index. We considered 12 to be the

maximum points (100%) within each group ($r = 0.8-1.0$ very strong correlation, $p < 0.001$), the final score was given by the percent calculated from the ratio of the reached and maximum scores.

Points	Correlation coefficient	p	points	Correlation coefficients	p
12	Very strong (0.8-1.0)	< 0.001	5	Moderate	< 0.05
11	Strong (0.6-0.79)	< 0.001	4	Weak (0.2-0.39)	< 0.001
10	Strong	< 0.01	3	Weak	< 0.01
9	Strong	< 0.02	2	Weak	< 0.02
8	Moderate (0.4-0.59)	< 0.001	1	Weak	< 0.05
7	Moderate	< 0.01	0		
6	Moderate	< 0.02			

Table 1.: Score system used for the evaluation of IS indices. The final score (%) is the ratio of the reached and maximum scores multiplied by 100.

When calculating sensitivity and specificity, values of the “gold standard” parameters under 25 percentile (M1 for IS) were considered to be the abnormal range (“real” IR), namely the “worst” quartiles:

$$\text{Sensitivity (\%)} = \frac{\text{Number of real (M) IR cases} - \text{number of false negative cases}}{\text{number of real (M) IR cases}} \times 100$$

$$\text{Specificity (\%)} = \frac{\text{Number of real (M) IS cases} - \text{number of false positive cases}}{\text{number of real (M) IS cases}} \times 100$$

2.2.4.3. General statistical considerations in clamp studies

All statistical analyses were performed with R Statistical Software (version 2.15.0). Data points are expressed as mean \pm standard deviation, if data were not normally distributed we used mean, standard deviation, median and mean absolute deviation (MAD) for each value presented. The Wilcoxon test (or in case of normally distributed parameters two-sided T test) were used to assess group differences. Spearman's correlation coefficients were calculated to test the association between biochemical and other variables, since non-linearity characterized these associations. Partial correlation coefficients were used to assess the influence of possible confounding factors such as age, body mass index (BMI), body fat percent (BFP), HbA1c levels (as being a mixed diabetic and non-diabetic population), genetic predisposition. Further adjustment with adiponectin and FSH in females and testosterone in males was used in the OCN - clamp study to exclude the effect of menopausal state in women, and the possible role of adiponectin / testosterone in mediating the metabolic effects of OCN in males/females. A p value of < 0.05 was considered significant.

Sample size determination was done empirically based on other clamp studies in the original protocol. For the liver – IR analysis we have used boot strap analysis (Monte-Carlo simulation) to test the minimal sample number to determine statistical differences between groups.

Feature selection analysis (Boruta algorithm) was used to find the most important attributes that are related to the M3 value in all of our clamp studies. This algorithm is a wrapper built around the randomForest classification algorithm (implemented in the R package randomForest) [25]. The randomForest algorithm is an ensemble approach (divide and conquer approach); it grows many decision trees and it gives a numerical estimate of the importance of a feature. A Z score is used as the importance measure since it takes into account the fluctuations of the mean accuracy loss among trees in the forest. To avoid random fluctuations in determining the importance of any given attribute, a reference set of ‘shadow attributes’ is used for deciding which attributes are truly important, since the importance of a shadow attribute can be non-zero only due to random fluctuations [26].

Multiple regression analysis was used in the NAFLD - IR study in men and women in order to determine the ability of metabolic parameters selected as ‘important attributes’ by feature selection analysis to predict clamp measured insulin sensitivity:

Model for women: $y \sim b_0 + b_1x_1 + b_2x_2 + \dots + b_5x_5$, where response variable y is M3, and explanatory variables x_1, \dots, x_5 are BMI, AC, insulin, fasting FFA, ALT, respectively and coefficients are in Table 6. The intercept b_0 is the expected mean value of M3 when all $x_i = 0$.

Model for men: $y \sim b_0 + b_1x_1 + b_2x_2 + \dots + b_9x_9$, where response variable y is M3, and explanatory variables x_1, \dots, x_9 are AC, leptin, BMI, insulin, TG, FFA, glucose, diastolic blood pressure (RR_Dias) and age, respectively and coefficients $b_1 \dots b_9$ are in Table 7. The intercept b_0 is the expected mean value of M3 when all $x_i = 0$.

2.2.5. Ethical considerations

For the epidemiological study we used database of an adult population screening between 2003 and 2006 in Balaton Upper-lands, all screenings approved by the local Hungarian Public Health and Medical Officer Service. Screenings were carried out on a voluntary basis, most of them were advertised as general screening assessments at General Practitioners' outpatient clinics, some of them were screening examinations taken place at working environment (Factories and Offices).

For clamp studies, data were retrospectively analyzed from a scientific study approved by the Hungarian Central Ethical Committee (A12988-2/2003-1018-EKU and ad.8-311/2009-1018EKU) titled "Diagnostic investigation for the early recognition of insulin resistance syndrome and its complications" (granted by Hungarian National Research and Innovation Program: NKFB -1B/0007/2002) and "Development and application of protein microarray technics to characterize insulin resistance syndrome and autoimmune processes" (granted by KMOP-1.1.1-08/1-2008-0028).

Recruitment started in 2004 and ended in 2010. After obtaining signed informed consent, in total 306 IVGTTs were carried out followed by a normoglycemic hyperinsulinemic clamp. Subjects were classified as having normal / impaired glucose tolerance or type 2 diabetes based on a standard 75 g OGTT at screening. All details about medical history, concomitant medications and life style were available for the data analysis.

3. Results

3.1. Diagnostic evaluation of simple insulin sensitivity indices

3.1.1. Study population

We performed OGTT examinations on 317 subjects during the screening period. (From the OGTT samples we determined glucose and insulin levels). Patients were categorized according to the ADA criteria (normal glucose tolerance (NGT): fasting plasma glucose (FBG) < 5.6 mM/l, impaired fasting glucose (IFG): FBG 5.6 – 6.99 mM/l, impaired glucose tolerance (IGT): FBG < 7.0 mM/l and 120th minute glucose: 7.77 – 11.0 mM/l, T2DM: fasting glucose \geq 7.0 mM/l, and/or 120th minute \geq 11.1 mM/l) [27]. Those, taking any antidiabetic medications, were treated for hormonal disease or received hormone substitution, were excluded from the study. Baseline characteristics of the subjects are listed in Table 2.

	N	m	f	age	BMI (kg/m ²)	HbA1c (%)	M1 (mg/minxkg ⁻¹)	AIR (uE/ml)
NGT	45	25	20	40.02±15.07	28.37±6.82	5.53±0.82	11.29±4.87	55.61±49.15
IFG/IGT	67	29	38	49.13±9.24*	30.85±5.21*	6.02±0.74*	8.37±3.26*	28.07±31.56*
DM	24	11	13	52.66±5.63*	32.13±6.10*	6.48±0.88*	6.46±3.38*	14.24±24.18*

Table 2.: Basic characteristics of the examined population. *: Significant ($p < 0.05$) difference compared to the NGT group, N: number of subjects m: number of males, f: number of females, M1: whole body glucose uptake, AIR: acute insulin response of IVGTT, BMI: body mass index, HbA1c: Hemoglobin A1c, NGT: normal glucose tolerant, IFG: impaired fasting glucose, IGT: impaired glucose tolerant, DM: diabetic subjects

3.1.2. Diagnostic evaluation of simple fasting and OGTT derived insulin sensitivity indices

The cumulative evaluation is shown on Figures 1. and 2. based on the earlier described score system (Table 2.). Most of the fasting indices show moderate ($R = 0.4-0.6$) correlation with M1 within most groups (correlation matrix not shown here). There was no significant correlation between fasting indices derived from the OGTT basal samples and M1 in the elder (age above median), normal glucose tolerant (NGT), and less severe (HbA1c under median) glucose intolerant groups. No correlation found in the elder NGT group between fasting indices derived from the mean of OGTT and IVGTT basal samples and M1. In HOMA-2 model we only found good correlations ($r = 0.6-0.8$) in the young NGT, diabetic and male groups. Cumulative evaluation of the fasting and OGTT derived indices based on group subanalyses are shown on Figure 3. and Figure 4. The sensitivity of fasting indices (average $50.4 \pm 4.6\%$) is low, while their specificity ($83.3 \pm 1.6\%$) is high, i.e. there are a relatively large number of false negative but less of false positive cases (Table 3.).

	Mean of fasting IS indices	Maximum value (HOMA _{igt})	Mean of OGTT IS indices	Maximum value (ISI _{cederh} , ISI _{est})
Sensitivity	50.92±4.43%	58.82%	60.25±8.06%	66.6%
Specificity	83.56±1.79%	85.7%	87.29±.87%	89.1%

Table 3.: Mean sensitivity and specificity of IS indices, considering the 25 percentile of M1 values as the cut-off for “insulin resistant” and “insulin sensitive” (less insulin resistant) groups. IS: Insulin sensitivity, HOMA_{igt}: HOMA index from IVGTT 0th minute glucose and insulin values, OGTT: oral glucose tolerance test, ISI_{cederh}: Cederholm’s IS index, ISI_{est}: estimated Stumvoll’s IS index.

ISI_{cederholm}, MCR_{est}, and ISI_{est}, including body weight in their equations, derived from the OGTT, show the strongest correlations with M ($r > 0.6$ correlations in most groups). On Figure 3. and Figure 4. it is clearly shown that all OGTT indices show better cumulative score than the average (mean of 0th minute OGTT and IVGTT insulin and glucose values) of

fasting indices. The only non-significant correlation was observed with the Matsuda equation (ISI_{comp}) in the young NGT group. The mean sensitivity and specificity of OGTT indices were higher than that of the HOMA model ($60.2 \pm 8.0\%$, and $87.2 \pm 1.8\%$ respectively, Table 3.).

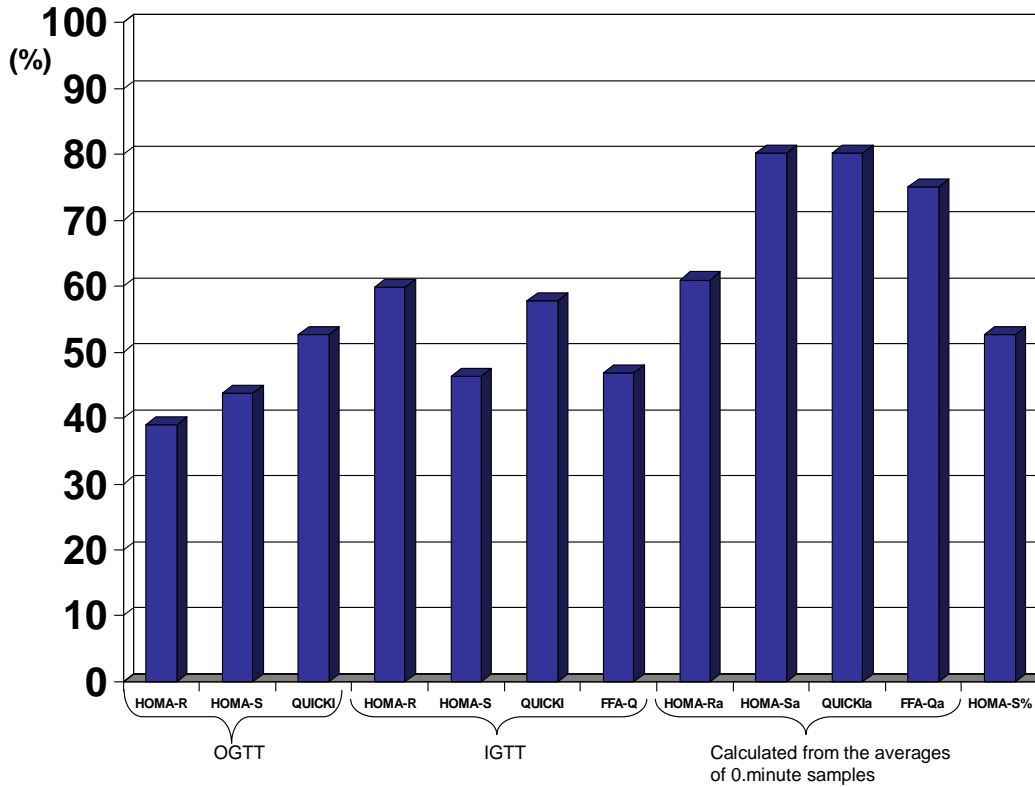


Figure 1.: Evaluation of fasting IS indices calculated from the 0th minute OGTT and IGTT values and the mean of these values, based on the correlation coefficients and significance levels with M1. “OGTT”: indices derived from 0th minute OGTT samples, “IGTT”: indices derived from 0th minute IVGTT samples, “a”: indices derived from the average of the two samples.

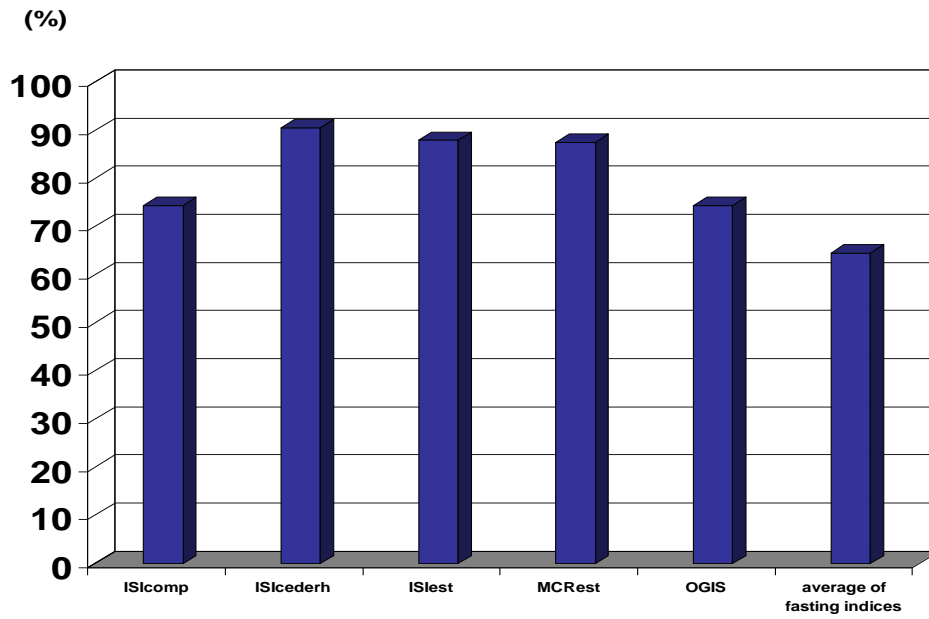


Figure 2.: Evaluation of the individual OGTT indices based on the correlation coefficients and significance levels with M1 in different subgroups. The mean evaluation of fasting indices is also indicated on the right side of the graph. ISI_{comp} : Composite insulin sensitivity index (Matsuda), $ISI_{Cederholm}$: Cederholm's insulin sensitivity index, MCR_{est} : estimated metabolic clearance rate (Stumvoll), OGIS: Oral Glucose Insulin Sensitivity Index.

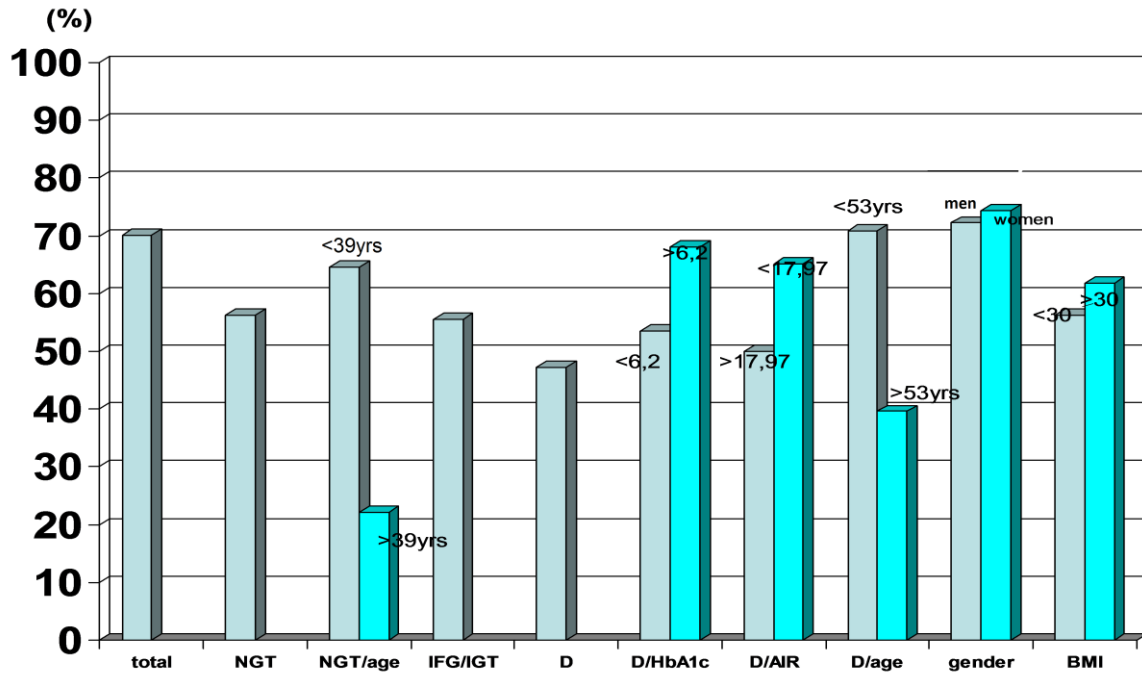


Figure 3.: Evaluation of fasting IS indices in different subgroups. From left to right: total: whole population, NGT: normal glucose tolerant, NGT/age: NGT group divided by age (below and above median), IFG/IGT: all IFG or IGT subjects , D: diabetic subjects, D/HbA1c: all IFG/IGT/D subjects divided by HbA1c (% , below and above median), D/AIR: all IFG/IGT/D subjects divided by beta cell function (acute insulin response = AIR), D/age: all IFG/IGT/D subjects divided by age (years, below and above median), gender: all subjects grouped by gender, BMI: all subjects divided by the existence of obesity (BMI < or $\geq 30 \text{ kg x m}^{-2}$).

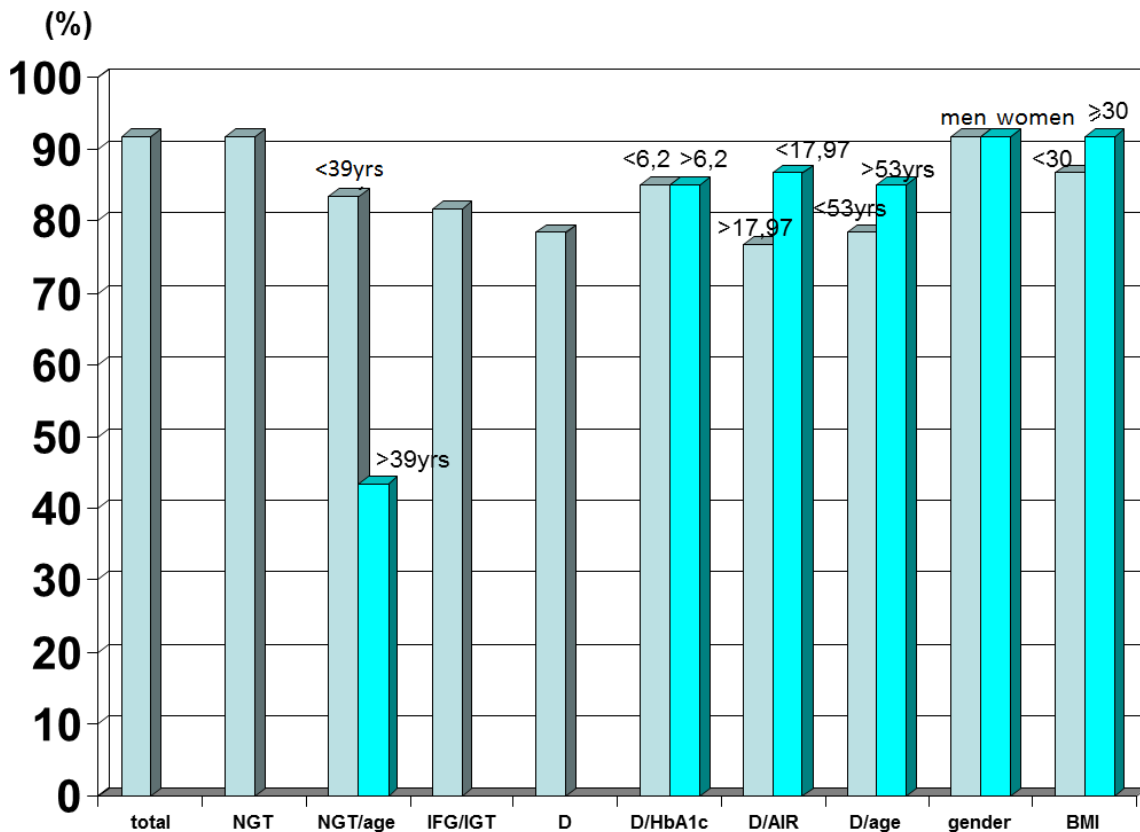


Figure 4.: Evaluation of the OGTT indices in different groups. For abbreviations of each group see Figure 3.

3.2. Clamp study on ALT – insulin sensitivity connections

3.2.1. Study population

General characteristics of the population are shown in Table 4. Mean HbA1c values were under 6.1% in all groups, i.e. the population consisted of either normal glucose tolerant or mostly prediabetic (IGT/IFG or fresh T2DM) subjects, both slightly overweight and obese individuals. Men tended to be younger and slightly more insulin sensitive than women in both the NGT and glucose intolerant (GI = IFG/IGT/T2DM) groups, although there were a lower prevalence of genetic predisposition in the NGT male group. The prevalence of genetic predisposition (the presence of diabetes in 1st degree relatives vs. genetically non-predisposed groups) were between 20 and 40%, lowest in the male NGT group, as indicated. Significant differences were found between age and metabolic parameters in NGT vs. GI groups in both

genders as expected. ALT and GGT levels were higher in the GI vs. NGT groups in both sexes, AST levels differed significantly only in the male group between NGT and GI subjects.

3.2.2. Associations between metabolic parameters, insulin sensitivity and liver function tests

Simple bivariate and partial correlation coefficients are listed in Table 4. between liver enzymes (AST, ALT and GGT) and metabolic parameters (including M3, HIRI, blood sugar level, insulin secretion, lipids and adipocytokines), after correcting for age, BMI, alcohol consumption, HbA1c, abdominal circumference (and FSH in females). In males triglyceride, HDL-cholesterol, free fatty acid and Acute Insulin Response (AIR) show significant correlations with ALT (and AST) after adjusting with the above confounding factors, while in females it is the clamp measured glucose uptake per se along with blood sugar values that stay significantly related after correction is done (see on Table 4). GGT is rather non sex-specific, i.e. corrected associations with GGT show a similar pattern in both genders.

Multiple regression analysis was carried out in order to determine the ability of the ‘important’ attributes selected by Feature selection analysis (Boruta Algorithm, applied as a pre-regression analysis here) to predict clamp measured M3 values, separately in the male and female populations. The results in Table 6. show that $F = 29.95$ ($p < 2.2e-16$) *for women*, indicating that the ‘important’ variables (BMI, AC, FFA, insulin and ALT) collectively have a significant effect on M3, ALT and BMI being significant independent predictors in women. *In men* (Table 7.) ‘important’ attributes (AC, leptin, BMI, insulin, TG, FFA, glucose, and diastolic blood pressure) have a significant effect on M3 [$F = 14.71$ ($p < 2.36e-16$)], serum leptin and insulin levels being independent predictors of clamp M3 values.

The ability of the ‘important’ attributes to predict measured M3 is indicated in Figure 5. for women, and in Figure 6. for men, where linear regression scatter plots for fitted vs. measured M3 values are shown. The regression model gave an excellent estimation of M3 in women, less so in men.

	NGT males(n=74)			GI (IFG, IGT, T2DM) males(n=74)		
	mean±SD	median	MAD	mean±SD	median	MAD
Age (years)	33.43± 11.60	30.00	10.46	48.72±9.33**	51.40	7.75
BMI (kg/m ²)	26.66±5.01	25.10	2.42	29.99±4.30**	29.41	4.01
Abdominal circumference(cm)	94.54± 13.11	90.50	6.67	105.07±14.55**	105.00	9.64
HbA1C (%)	5.41± 0.43	5.40	0.44	5.90± 0.68**	5.80	0.59
Fasting glucose (mmol/L)	4.89± 0.72	4.83	0.56	6.09±1.04**	5.96	0.96
M3(mg/min/kg)	8.83±3.10	8.81	2.82	5.88±2.77**	5.74	2.38
Hepatic insulin resistance index	54.42±33.06	48.86	24.98	63.86±33.84**	60.24	32.09
Triglyceride (mmol/l)	1.46±1.14	1.07	0.55	2.65±2.18**	2.00	1.33
HDL-C(mmol/L)	1.35±0.40	1.36	0.36	1.13±0.42**	1.08	0.29
LDL-C(mmol/L)	2.48±0.83	2.31	0.80	3.00± 1.05**	3.03	0.79
AST(U/L)	21.68±5.44	20.00	4.45	27.57±12.93**	25.00	8.90
ALT(U/L)	24.81±10.56	22.00	7.41	36.24±26.98**	29.50	15.57
GGT(U/L)	26.93±15.73	22.00	10.38	48.44±33.50**	39.00	28.17
Alcohol (g/day)	0.09±0.38	0.00	0.00	0.30±0.67**	0.00	0.00
Hypertension(%)	10.81	NA	NA	43.24	NA	NA
Smoking(%)	14.86	NA	NA	20.22	NA	NA
Genetic predisposition(%)	21.62	NA	NA	32.89	NA	NA
	NGT females (n=47)			GI (IFG, IGT, T2DM) females(n=111)		
Age (years)	45.10±10.39	46.00	10.43	50.80±8.54**	53.00	7.41
BMI (kg/m ²)	26.85± 4.25	26.57	4.74	31.49±5.25**	31.57	4.96
Abdominal circumference(cm)	91.95±12.18	92.00	14.08	104.42±12.49**	103.00	11.86
HbA1C (%)	5.62± 0.50	5.60	0.59	6.06± 0.63**	6.02	0.62
Fasting glucose (mmol/L)	5.08± 0.49	5.08	0.44	5.75±0.77**	5.65	0.76
M3(mg/min/kg)	6.64± 3.24	6.29	2.79	4.36±2.08**	3.92	1.69
Hepatic insulin resistance index	63.07±32.51	54.76	27.80	75.23±49.42**	60.77	36.46
Triglyceride (mmol/l)	1.43±0.83	1.24	0.43	1.79± 0.81**	1.57	0.74
HDL-C(mmol/L)	1.48±0.55	1.49	0.61	1.33±0.51	1.27	0.36
LDL-C(mmol/L)	2.65±0.81	2.51	0.61	3.20±1.06**	3.17	0.87
AST(U/L)	23.00±9.76	21.00	5.93	23.75±10.37	20.00	4.45
ALT(U/L)	21.79±14.21	20.00	10.38	25.33±12.98*	22.00	8.90
GGT(U/L)	25.95±28.13	19.00	11.86	31.76±25.29**	25.00	11.86
Alcohol (g/day)	0.14±0.55	0.00	0.00	0.02±0.15	0.00	0.00
Hypertension(%)	23.40	NA	NA	43.24	NA	NA
Smoking(%)	14.86	NA	NA	18.18	NA	NA
Genetic predisposition(%)	21.62	NA	NA	36.36.	NA	NA

Table 4. : Baseline biochemical and clinical characteristics of male and female populations. All values are means, medians and mean absolute deviation (MAD). NGT: normal glucose tolerant, GI= glucose intolerant, IFG=impaired fasting glucose, IGT=impaired glucose tolerant, T2DM= type 2 diabetes. M3: muscle mass adjusted glucose uptake. p<0.05*, p<0.01**, p<0.001***, p<0.0001****

	AST(U/L)				ALT(U/L)				GGT(U/L)			
	Males		Females		Males		Females		Males		Females	
	R	Partial R	R	Partial R	R	Partial R	R	Partial R	R	Partial R	R	Partial R
HbA1c(%)	n.s.	n.s.	n.s.	n.s.	n.s.	n.s.	0.204*	n.s.	n.s.	n.s.	0.229**	n.s.
Glu-0(mmol/L)	0.244**	n.s.	n.s.	n.s.	0.268***	n.s.	0.31****	0.177*	0.360****	0.25**	0.335****	0.214***
AIR(uU/mL)	-0.181*	n.s.	n.s.	n.s.	-0.189*	-0.201*	n.s.	n.s.	-0.181*	n.s.	n.s.	n.s.
FFA-0(mmol/L)	0.326****	0.234*	0.206*	n.s.	0.295***	0.203*	0.276**	n.s.	0.385****	0.185*	0.277***	0.198*
M3(mg/min/kg)	-0.167*	n.s.	-0.311****	0.216***	-0.324****	n.s.	-0.430****	-0.270***	-0.323****	n.s.	-0.337****	-0.268****
HIRI	n.s.	n.s.	n.s.	n.s.	0.198*	n.s.	0.231**	n.s.	0.193*	n.s.	0.240**	n.s.
TG (mmol/L)	0.389****	0.315****	0.201*	n.s.	0.444****	0.288**	0.216**	n.s.	0.636****	0.525****	0.299****	0.200*
HDL-C (mmol/L)	n.s.	n.s.	n.s.	n.s.	-0.255**	-0.218*	n.s.	n.s.	n.s.	n.s.	n.s.	n.s.
LDL-C (mmol/L)	n.s.	n.s.	n.s.	n.s.	0.168*	n.s.	n.s.	n.s.	0.330****	0.244**	0.167*	n.s.
Leptin (ng/mL)	n.s.	n.s.	0.239**	0.198*	0.288***	n.s.	0.166*	n.s.	0.292***	n.s.	n.s.	n.s.
Adiponectin (ug/mL)	n.s.	n.s.	n.s.	n.s.	-0.219**	n.s.	n.s.	n.s.	-0.164*	n.s.	n.s.	n.s.
IL-6(ng/mL)	0.180*	n.s.	n.s.	n.s.	n.s.	n.s.	n.s.	n.s.	0.245**	n.s.	n.s.	n.s.

Table 5.
Bivariate correlation s adjusted for confounding factors between metabolic parameters and liver enzymes in male and female subjects.

Glu-0: fasting glucose,
AIR: acute insulin response,
FFA: free fatty acid,
M3: muscle adjusted glucose uptake,
HIRI: hepatic insulin sensitivity index.

p<0.05*,
p<0.01**,
p<0.001***,
p<0.0001****.

Coefficients:	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	15.81509	1.22936	12.864	< 2e-16 ***
BMI	-0.23375	0.05272	-4.434	1.9e-05 ***
AC	-0.01780	0.02081	-0.856	0.39377
Insulin	-0.04213	0.02469	-1.706	0.09028 .
IVFFA_0	-1.00410	0.56730	-1.770	0.07899 .
ALT	-0.03159	0.01208	-2.616	0.00991 **

Table 6: Multiple regression analysis for clamp M3 in women based on the ‘important attributes’ chosen by the feature selection analysis. Significance codes: 0.0001 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05. Residual standard error: 1.88 on 158 degrees of freedom. Multiple R-squared: 0.5259, adjusted R-squared: 0.5084, F-statistic: 29.95, p value: < 2.2e-16. . Significant independent predictors: BMI (***) and ALT (**). AC: abdominal circumference, IVFFA_0: 0th minute FFA of the IVGTT.

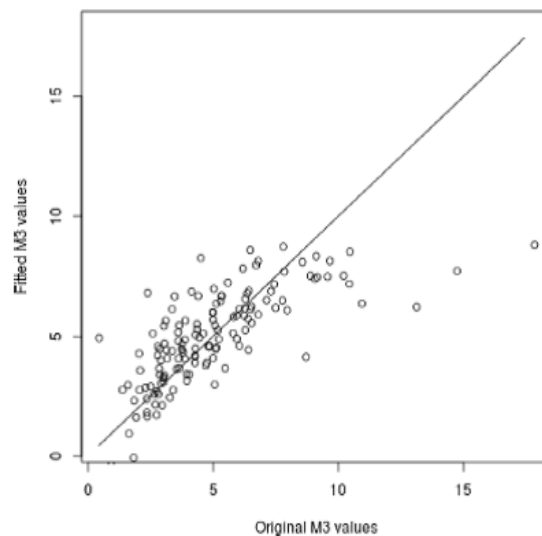


Figure 5: Linear regression for original versus fitted M3 values in women estimated by multiple regression analysis for attributes determined by feature selection. BMI (p=1.9e-05), AC (p=0.39377), serum-insulin (p=0.09028), serum-FFA (p=0.07899), ALAT (p=0.00991). Multiple R-squared: 0.5259, Adjusted R-squared: 0.5084

	Coefficients: Estimate	Std. Error	t value	Pr(> t)
(Intercept)	19.30144	2.35704	8.189	1.9e-13***
AC	-0.03122	0.02886	-1.082	0.2813
Leptin	-0.09451	0.04291	-2.202	0.0294 *
BMI	-0.07101	0.08879	-0.800	0.4253
Insulin	-0.09684	0.04147	-2.335	0.0210 *
TG	-0.07363	0.12927	-0.570	0.5699
IVFFA_0	-0.69004	0.58016	-1.189	0.2364
Glucose	-0.37848	0.23299	-1.624	0.1067
RR_Dias	-0.01234	0.02586	-0.477	0.6340
Age	-0.03591	0.01828	-1.964	0.0516 .

Table 7: Multiple regression analysis for clamp M3 in men, based on the ‘important’ attributes chosen by the feature selection analysis. Significance codes: 0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05. Residual standard error: 2.331 on 148 degrees of freedom. Multiple R-squared: 0.4989, Adjusted R-squared: 0.465, F-statistic: 14.71, p value: 2.368e-16. Significant predictors: leptin (*) and insulin (*) levels. TG = triglyceride, AC: abdominal circumference, IVFFA_0: basal FFA.

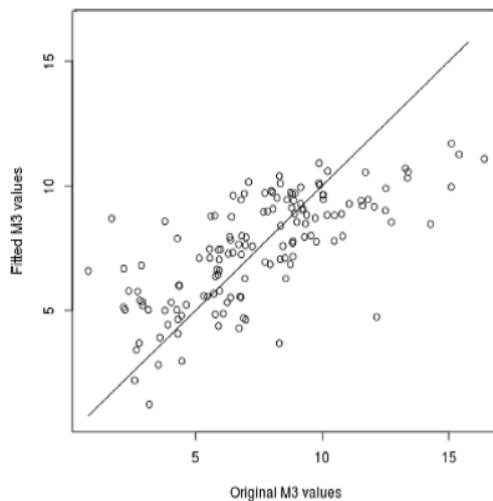


Figure 6.: Linear regression for original vs. fitted M3 values in men estimated by multiple regression analysis for attributes determined by feature selection. AC (p= 0.2813), leptin (p= **0.0294**), BMI (p = 0.4253), serum insulin (p = **0.0210**), triglyceride (p = 0.5699), serum-FFA (p=0.2364), serum-glucose (p = 0.1067), diastolic RR (p = 0.6340), age (p = 0.0516). Multiple R-squared: 0.4989, Adjusted R-squared: 0.465.

3.3. Epidemiologic association of metabolic syndrome, type 2 diabetes and bone loss in the adult population of Balaton Upper-lands

3.3.1. Study population

In our epidemiological database the results of 6282 screening tests were available, 1561 men (mean age 56 ± 13 years) and 4726 women (mean age 54 ± 13 years). The baseline characteristics (age and gender) of the screened population are shown in Table 8.

Age groups (year)	Men (n)	Women (n)
<40	163 (20.6%)	628 (79.4%)
40–50	329 (22%)	1166 (78%)
51–60	504 (27.2%)	1351 (72.8%)
61–70	356 (26.33%)	996 (73.67%)
>70	194 (26.5%)	538 (73.5%)

Table 8.: Distribution of screened subjects based on gender and age.

Based on the data that were available we have defined syndromes / diseases shown in Table 9.

<i>Diabetes</i> : presence of impaired glucose tolerance: fasting capillary plasma glucose >5.6 mmol/l and/or 2 hours capillary plasma glucose >7.8 mmol/l, any value ≥ 11.1 mmol/l and / or known / treated diabetes mellitus
<i>Hypertension</i> : untreated or treated known hypertension or systolic blood pressure > 140 And/or diastolic blood pressure >90 Hgmm
<i>Overweight</i> : $25 < \text{BMI} < 30$, <i>obesity</i> : $\text{BMI} \geq 30$, <i>severe obesity</i> : $\text{BMI} \geq 35$ kg/m ²
<i>Hypercholesterolemia</i> : total capillary plasma cholesterol >5.2 mmol/l
<i>Android obesity</i> : Abdominal circumference: men >102 cm, women >88 cm

Table 9.: Definitions of diseases / syndromes that were assessed

Categories do not always mean definitive diagnoses as one time capillary plasma measurements may not enable us to adapt internationally used criteria systems although we have tried to define categories based on the valid international / national guidelines at the

time of the publication [27-31]. In case of “Metabolic syndrome” we have defined an ‘insulin resistant’ state not necessarily identical to the disease / syndrome defined by the ATP-III or WHO criteria (Table 10.).

<i>Metabolic syndrome:</i>	Presence of DM/IGT/IFG, hypertension, hypercholesterolemia, obesity and/or android obesity in overweight patient: the presence of at least 3 of these criteria at the same time.
<i>Osteoporosis:</i>	T-score: <-2.5 and / or treated osteoporosis
<i>Osteopenia:</i>	T-score: between -1 and -2.5

Table 10.: Definition of Metabolic syndrome, osteoporosis and osteopenia

3.3.2. Results of epidemiologic data

When assessing the correlations between bone density (T score) and some of the metabolic parameters (Table 11.) we found that there is a significant positive correlation between BMI and T score in every age group in women (except in women under the age of 40), the correlation being strongest in the population > 70 years . This relationship was missing in men. Abdominal circumference shows a weak positive correlation with T score in women < 40 years and in men between 40 and 50 years. Blood sugar showed significant positive association with T score only in men over 70 years.

T score	BMI R=		Abdominal circumference R=		Blood sugar R=	
	Men	Women	Men	Women	Men	Women
<40	n.s.	n.s.	n.s.	+0.254**	n.s.	n.s.
41-50	n.s.	n.s.	+0.234*	+0.185**	n.s.	n.s.
51-60	n.s.	+0.115**	n.s.	+0.152**	n.s.	n.s.
61-70	n.s.	+0.110*	n.s.	n.s.	n.s.	n.s.
>70	n.s.	+0.230****	n.s.	n.s.	+0.359****	n.s.

Table 11.: Correlation coefficients of bivariate correlations measured between T score, BMI abdominal circumference and blood sugar levels within genders and age groups. Significant correlations are indicated: *p<0.05, **p<0.02, ***p<0.01, ****p<0.001

We have examined the prevalence of osteoporosis / osteopenia in patients with metabolic syndrome and diabetes based on gender and age groups. There was no significant

difference in the prevalence of decreased bone density between patients with metabolic syndrome and controls (those with no metabolic syndrome) in either gender or age groups. Looking at the frequency of osteoporosis / osteopenia in diabetic subjects we found that amongst 51-60 years old women the prevalence of osteopenia was significantly higher in the diabetic than in the normal glucose tolerant group (50 vs. 36.34% in diabetic vs. non-diabetic females, χ^2 : 5.237, $p < 0.022$, OR: 1.711, 95% CI: 1.076–2.722). Mean age of the two groups (diabetic and normal glucose tolerant) was not significantly different (55.7 ± 4.2 vs. 55.2 ± 4.1 years, $p = 0.7$), however the mean BMI in this diabetic group was significantly higher than in the normal glucose tolerant group (29.42 ± 5.27 vs. 27.73 ± 4.77 , $p < 0.008$) which makes our data even stronger, as high BMI is generally accepted as a protective factor against bone loss. There was no significant difference in the prevalence of osteoporosis or osteopenia in other age groups in women or in any of the age groups in men between diabetic and control subjects. Due to high BMI having presumably both diabetogenic and osteoprotective role we have separately analyzed the prevalence of different stages of bone loss in normal weight diabetic (BMI < 25), normal weight non-diabetic, all diabetic and all non-diabetic subjects (Figure 8. and Figure 9.).

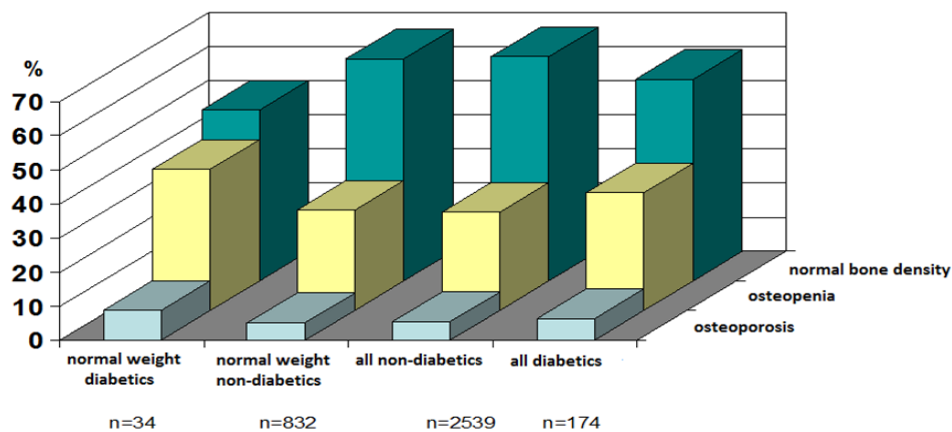


Figure 8.: Prevalence of osteoporosis, osteopenia and normal BMD in normal weight (BMI<25) diabetic, normal weight non-diabetic, all non-diabetic and all non-diabetic subjects ≤ 60 years old.

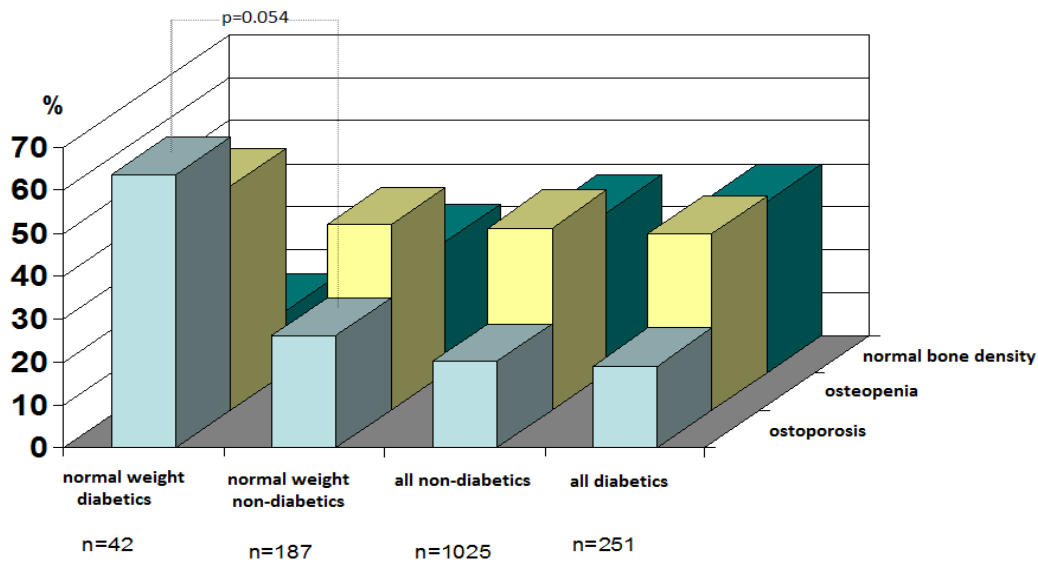


Figure 9.: Prevalence of osteoporosis, osteopenia and normal BMD in normal weight (BMI<25) diabetic, normal weight non-diabetic, all diabetic and all non-diabetic subjects > 60 years old.

Considering the small sample size in each group, for statistical rationale we only defined two age groups, below 60 and over 60 years, and genders were not separately analyzed either. Within the individual age groups there was no difference between the mean age of the normal weight and all diabetic patients (53.8 ± 10.0 vs. 56.4 ± 8.6 and 70.6 ± 6.4 vs. 69.4 ± 5.9 years). We didn't find significant difference in the prevalence of osteoporosis / osteopenia in the population under 60 years although there was a trend for increased prevalence of osteoporosis in normal weight diabetic versus all diabetic patients (Figure 8.). In the > 60 years group the frequency of osteoporosis was higher within the normal weight diabetic group than in all diabetic subjects, the difference being border significant with 'osteoporosis' (63.63 vs. 26.2%, OR: 2.71, 95% CI: 0.969–7.6, $p = 0.054$) but only a trend was seen with 'ostopenia' (53.38 vs. 43.31%, $p = 0.359$), (Figure 9.). There was no difference noted in the prevalence of osteoporosis / osteopenia in the all diabetic versus all non-diabetic group. (Figure 9.)

3.4. Clamp study; bone – energy homeostasis connections

3.4.1. Study population

After obtaining signed informed consent, we included 135 women (aged 49 ± 9 years) and 155 men (aged 42 ± 13) in our study. Subjects were classified based on results of a standard 75-g OGTT at screening (blood drawn at the 0, 30, 60, 90 and 120th minute), according to ADA criteria [27]. We included 47 normal glucose-tolerant (NGT) and 89 glucose-intolerant (GI: IFG and IGT and drug naïve T2DM) subjects in the female group; in the male group, there were 72 NGT and 83 GI subjects. The same inclusion and exclusion criteria were applied in this study as in the earlier clamp studies.

3.4.2. Baseline characteristics

General metabolic and other characteristics of male and female subjects in the study are summarized in Table 12. NGT and GI subjects are presented separately. Although OCN mean values were slightly higher in the NGT than in the GI groups in both genders, no significant difference was observed in OCN levels between NGT and GI subjects. In women this is the result of the effect of varying ages, that masks the positive association between OCN and glycemic state, in men however (see later) OCN is mainly associated with indicators of insulin sensitivity (i.e. glucose uptake rates and body composition) rather than with plasma glucose levels per se. OCN mean values were somewhat lower than described in healthy male and female population, although stayed in the normal range [32]. Subjects with extreme values were excluded from the study. Significantly higher total testosterone values were found in NGT than in GI males, while no such difference was observed between the respective female groups.

	Women		Men	
	NGT, n=47	GI, n=89	NGT, n=72	GI, n=83
BMI(kg/m ²)	27.1±5	31.6 ±5***	26.7 ±5.1	28.2 ±5.25***
Body Fat %	42.5 ±7.4	46.5 ±6**	26.8 ±7.98	28.7 ±8.11***
HbA1c (%)	5.84 ±0.8	6.06 ±0.6***	5.37 ±0.44	5.83 ±0.69***
OGTT glucose 0 min	5.1±0.45	6.1±0.83***	5.01±0.45	6.32 ±0.89***
OGTT glucose AUC (mmol/l)	38.69 ±19.3	94.96 ±45.5***	30.2 ±25.36	90.43 ±49.61***
OGTT insulin AUC (μU/ml)	6627.3 ±4017	8448 ±5617*	5402 ±3534	6796 ±4282*
M1 (mg/min/BW kg)	11.28 ±4.4	7.92 ±3.2***	12.32 ±3.78	8.76 ±3.63***
M3 (mg/min/muscle kg)	6.62±3.22	4.31±2.03***	8.95±3.45	6.11±2.71***
Leptin (ng/ml)	27.8 ±22.7	31.94 ±20.8	10.13 ±9.89	8.72 ±6.63
Adiponectin (μg/ml)	5.7 ±3.1	5.4 ±3.1	4.58 ±2.62	3.46 ±1.92
Testosterone (ng/ml)	0.74 ±	0.84 ±	15.51±5.21	13.44 ±6.24*
Osteocalcin (ng/ml)	18.89 ±7.49	18.13 ±7.66	19.87±8.27	19.08 ±9.15

Table 12.: General characteristics of the study subjects. OCN Normal range: 11–48 ng/ml for women, 14–46 ng/ml for men. Normal glucose-tolerant (NGT) and glucose-intolerant (GI) subjects, the latter consisting of impaired fasting glucose (IFG), impaired glucose-tolerant (IGT), or drug-naive type 2 diabetic (2DM) subjects, are represented separately. Significant differences between NGT and IGT groups are shown. *p < 0.05, **p < 0.01, ***p < 0.001.

3.4.3. Metabolic associations of OCN in females

Spearman's correlations and partial correlations between OCN and metabolic parameters of female subjects are shown in Table 13. Significant association was observed between OCN and adiponectin levels independent of age, HbA1c, BMI and BFP. Higher OCN values were associated with increasing age in women. Significant correlation between OCN and improving metabolic state (lower OGTT glucose values, BMI, BFP, higher M1 and M3 levels) became apparent after the adjustment for age alone. Most of these associations ceased after further adjustment with HbA1c, BMI and BFP, except for fasting glucose. Further adjustment with adiponectin resulted in lost correlation between OCN and fasting glucose value, i.e. OCN effect on improving glycemic control might be partly mediated by adiponectin in females. Further adjustment was done with FSH levels in a subset of 68 women (where FSH levels were available) in order to exclude the effect of menopausal state

on the metabolic associations of OCN (not represented in Table 13.). Positive associations between improved metabolic state and OCN became apparent after the adjustment with FSH, similar to the influence of age: $R = -0.323$, $p = 0.0062$ with HbA1c, $R = -0.349$, $p = 0.0026$ with IVGTT glucose AUC, and $R = -0.288$, $p = 0.0153$ with OGTT glucose 0th min, $R = +0.260$, $p = 0.031$ with M3. As expected, a strong positive association was found between FSH levels and OCN, independent age, BMI, body composition and HbA1c values ($R = +0.413$, $p = 0.00047$).

OCN and:	R=	R = (adjusted for age)	R=(adjusted for BMI, BFP, age and HbA1c)	R= adjusted for age, BMI, BFP, HbA1c and adiponectin)
Age	+0.231**	-	-	-
HbA1c	n.s.	-0.267**	-	-
OGTT glucose 0 min	n.s.	-0.236**	n.s.	n.s.
OGTT glucose 120 min	n.s.	-0.205*	n.s.	n.s.
OGTT glucose AUC	n.s.	n.s.	n.s.	n.s.
OGTT insulin AUC	n.s.	n.s.	n.s.	n.s.
IVGTT glucose 0 min.	n.s.	-0.269**	-0.190*	n.s.
IVGTT glucose AUC	n.s.	-0.182*	n.s.	n.s.
IVGTT insulin AUC	n.s.	n.s.	n.s.	n.s.
AIR	n.s.	n.s.	n.s.	n.s.
IGI	n.s.	n.s.	n.s.	n.s.
M1	n.s.	+0.221**	n.s.	n.s.
M3	n.s.	+0.240**	n.s.	n.s.
Leptin	n.s.	n.s.	n.s.	n.s.
Adiponectin	+0.254**	+0.254**	+0.335***	-
Triglyceride	n.s.	n.s.	n.s.	n.s.
Testosterone	n.s.	n.s.	n.s.	n.s.
BMI	n.s.	-0.192*	-	-
BFP	n.s.	-0.224**	-	-
Waist circumference	n.s.	-0.212*	n.s.	n.s.

Table 13.: Bivariate (Spearman) and partial correlation coefficients adjusted for possible confounding factors (age, BMI, BFP and HbA1c), between OCN and metabolic parameters in women. When adjusted for age alone significant associations with metabolic parameters were disclosed, i.e. these were concealed due to age. When adiponectin was added for further adjustment as possible confounding factor, the significant association between fasting glucose and OCN ceased. The simple coefficients are shown in column 1, and partial correlation coefficients in columns 2–4. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. n.s.: not significant; BMI: body mass index; BFP: body fat percent; OGTT: oral glucose tolerance test; AUC: area under the curve; IVGTT: intravenous glucose tolerance test; AIR: acute insulin response of the IVGTT. IGI: insulogenic index of the IVGTT: (insulin 5 min – insulin 3 min) / (glucose 5 min – glucose 3 min).

3.4.4. Metabolic associations of OCN in males

Spearman's and partial correlation coefficients between OCN and metabolic parameters in males are shown in Table 14.

OCN and	R=	R= (Adjusted for age, BMI, BFP and HbA1c)	R= (Adjusted for age, BMI, BFP, HbA1c and testosterone)	R= (Adjusted for testosterone alone)
Age	n.s.	-	-	n.s.
HbA1c	n.s.	-	-	n.s.
OGTT glucose 0.min	n.s.	n.s.	n.s.	n.s.
OGTT glucose 120.min.	n.s.	n.s.	n.s.	n.s.
OGTT glucose AUC	n.s.	n.s.	n.s.	n.s.
OGTT insulin AUC	-0.179*	n.s.	n.s.	n.s.
IVGTT glucose 0.min	n.s.	n.s.	n.s.	n.s.
IVGTT glucose AUC	n.s.	n.s.	n.s.	n.s.
IVGTT insulin AUC	n.s.	n.s.	n.s.	n.s.
AIR	n.s.	n.s.	n.s.	n.s.
IGI	n.s.	n.s.	n.s.	n.s.
M1	+0.229**	n.s.	n.s.	+0.174*
M3	+0.221**	n.s.	n.s.	n.s.
Triglyceride	-0.183*	n.s.	n.s.	-0.178*
Leptin	-0.168*	n.s.	n.s.	n.s.
Adiponectin	n.s.	n.s.	n.s.	n.s.
Testosterone	+0.243**	+0.193*	-	-
BMI	-0.199*	-	-	n.s.
BFP	-0.172*	-	-	-0.171*
Waist circumference	-0.224**	n.s.	n.s.	-0.221**

Table 14: Spearman and partial correlation coefficients adjusted for possible confounding factors (age, BMI, BFP, HbA1c and testosterone), between OCN and metabolic parameters in men. When adjusted only for testosterone alone, significant association between muscle glucose uptake (M3) and OCN ceased, but stayed significant between M1 and OCN. The simple coefficients are shown in column 1, and partial correlation coefficients in columns 2–4. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. See list of abbreviations below Table 13.

The OCN levels were significantly associated with improving metabolic state (decreasing OGTT insulin AUC, leptin, BMI, BFP and increasing insulin sensitivity; i.e., M1 and M3 values). These correlations disappeared after the adjustment for age, HBA1c, BMI and BFP. The significant positive correlation between OCN and testosterone levels was independent of age, HBA1c, BMI and BFP. After correction for testosterone alone, the significant positive association between OCN and M3, as well as the significant negative correlation with leptin and BMI was lost, i.e. metabolic associations of OCN were at least partly mediated by testosterone in males.

3.4.5. Feature selection

Feature selection (Boruta algorithm, Figure 10. and 11.) confirmed that age, IVGTT glucose 3, 5 and 60 minute values and adiponectin (mean Z: 9.76, 5.41, 4.09, 3.75 and 3.69, respectively) were the most important attributes in determining OCN levels in women. In men, M1, BMI, M3, leptin, BFP, OGTT 90th min glucose and insulin values, in addition to testosterone (mean Z: 5.21, 4.95, 4.41, 4.04, 3.62, 3.57 and 3.54, respectively), but not adiponectin, were confirmed as the most important parameters independently associated with OCN amongst all metabolic factors examined.

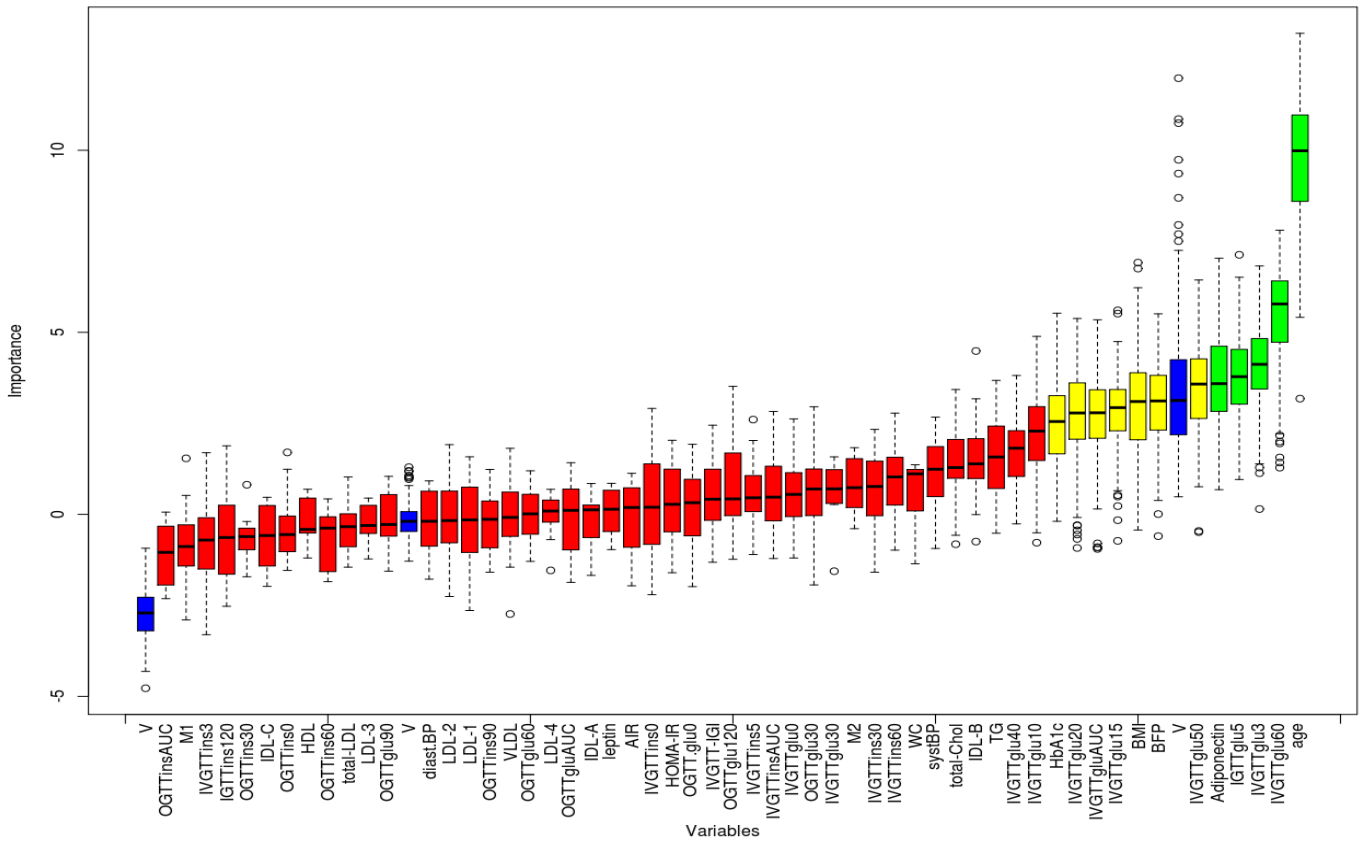


Figure 10.: Feature selection analysis (Boruta method) to identify variables most closely related to OCN values in women. All IVGTT (0-60th minutes: ‘IVGTTGlu- and Ins0-, 3-, 5-, 10-, 15-, 20-, 30-, 40-, 50- and 60’) and OGT (0-120th minutes: ‘OGTTGlu- and Ins0-, 30-, 60-, 90- and 120’) glucose and insulin values, lipids [TG, total-cholesterol (total-chol)] and lipoprotein subfractions (total-LDL, total-HDL, IDL-A, - B,- and C, LDL1-4 subfractions), adipokines (leptin, adiponectin), anthropometric parameters (BMI, body fat percent [BFP], waist circumference [WC]), testosterone, systolic- and diastolic blood pressure (systBP, diastBP), acute insulin response (AIR), age, whole- and muscle glucose uptake (M1 and M3) values were ranked for OCN association. Age, IVGTT glucose 0-, 3-, and 5th min values and adiponectin were confirmed as being ‘important’ attributes for OCN; these are represented as green columns. Yellow and red columns represent attributes that were ‘rejected’ or ‘tentative’ as being important variables for OCN. Blue columns represent ‘shadow attributes’. The Y axis represents the value of importance (mean, median, minimum, and maximum Z).

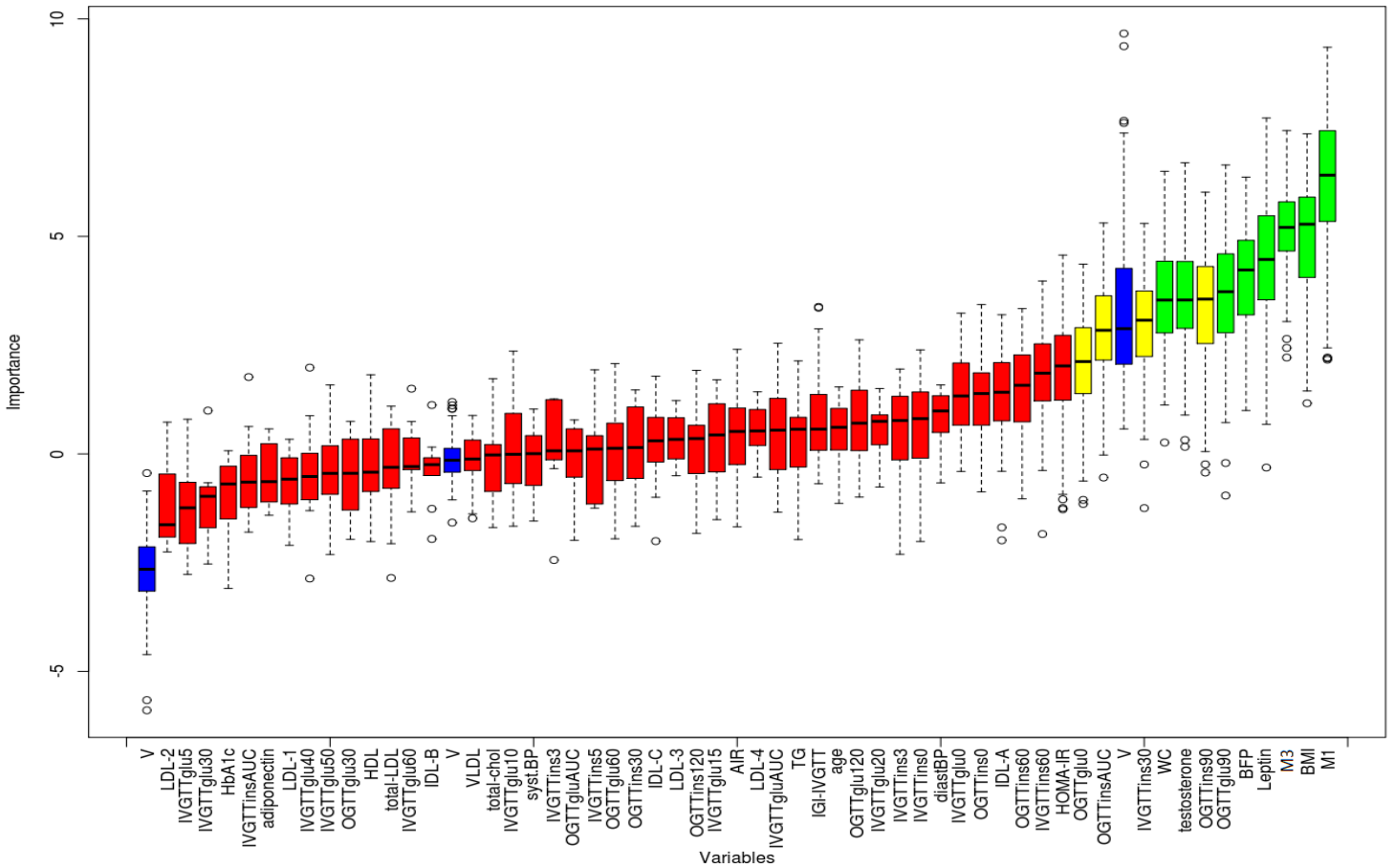


Figure 11.: Feature selection analysis (Boruta algorithm) to identify variables most closely related to OCN values in men. All IVGTT (0-60. minutes: ‘IVGTTGlu- and Ins0-, 3-, 5-, 10-, 15-, 20-, 30-, 40-, 50- and 60’) and OGTT (0-120. minutes: ‘OGTTGlu- and Ins0-, 30-, 60-, 90- and 120) glucose and insulin values, lipids [TG, total-cholesterol (total-chole)] and lipoprotein subfractions (total-LDL, total-HDL, IDL-A,- B,- and C, LDL1-4 subfractions), adipokines (leptin, adiponectin), anthropometric parameters (BMI, body fat percent [BFP], waist circumference [WC]), testosterone, systolic- and diastolic blood pressure (systBP, diastBP), acute insulin response (AIR), age, whole- and muscle glucose uptake (M1 and M3) values were ranked for OCN association. M1, BMI, M3, leptin, BFP, 90-minute glucose and insulin of the OGTT and testosterone were confirmed as ‘important’ attributes for OCN; these are represented as green columns. Yellow and red columns represent attributes that were rejected or ‘tentative’ for being important attributes for OCN. Blue columns represent ‘shadow attributes’. The Y axis represents the value of importance (Z), (mean, median, minimum, and maximum).

Discussion

4.1. Evaluation of simple indices in the estimation of clamp measured insulin sensitivity

Quantitative determination of IS and/or beta cell function (BCF) has a great significance when studying metabolic diseases, i.e. the relationship between bone metabolism and energy homeostasis. Although, a number of indices have been developed earlier [22], the comparison between the simplest fasting HOMA and clamp indices show a quite heterogeneous picture, there are those reporting about strong [33], weak [34] and non-significant [18] results. Strong correlation was found between clamp index and HOMA-R in young or middle-aged NGT subjects, or in smaller studies with T2DM patients [33, 35-37]. QUICKI, the logarithmic transformation of HOMA, improved the correlation in a small, heterogeneous population ($r = 0.78$ vs. 0.6) [21]. No significant correlation was found between either HOMA, or QUICKI and clamp indices in other reports in elderly, poorly controlled diabetic populations [38].

In our study, which included drug-naïve prediabetic or normal glucose tolerant subjects with a wide range of insulin resistance, the correlation were moderate ($r = 0.4-0.6$) between HOMA and clamp measured glucose uptake in most groups. We found weaker correlations in the NGT and in the glucose intolerant groups with higher age (above median). Correlations between fasting and clamp indices were weaker in the IGT group with lower HbA1c and higher AIR (i.e. less severe metabolic state) than in IGT subjects with higher HbA1c and lower AIR (i.e. more decreased beta cell function). According to our results the utility of HOMA-S% (Oxford Model) was far behind the expected, although today it is considered the most accepted, best fasting index [39,40]. Correlations between QUICKI and M was not improved by FFA in our study which is in contrast with previous results [24].

Amongst OGTT indices we compared the values of the five most often used indices based on empiric equations validated by clamp [16-19]. An independent study used discrepancy index (DI) to compare several OGTT indices, which showed that MCR_{est} and OGIS adjusted for body weight are structurally the closest indices to clamp, i.e. these two least under- or overestimated the correlations between basic parameters (Ins_0 , Glu_0 , BMI, Glu_{mean} , Glu_{120}) and M1 [41]. In our study we found that the three OGTT indices including body weight or BMI in their equations, MCR_{est} , ISI_{est} and $ISI_{Cederholm}$ showed the best correlation with clamp M1, as expected in a population with wide range of BMI. Based on

the correlations measured within the different groups, the cumulative scores approached 90%. We conclude that in this mostly overweight population with varying glucose tolerance, including BMI/weight into the equation improves its value compared to indices based only on glucose and insulin values. Another advantage of OGTT indices is that they show a fairly homogeneous value in most examined groups, except in elderly NGT subjects, where their value was decreased, similar to fasting indices. Individual diagnostic values of the HOMA and OGTT based IS models may still be a matter of debate, since the best sensitivity for IR diagnosis was only 50% for fasting and 67% for OGTT indices. The lower sensitivity of HOMA versus OGTT indices is shown in a study where OGTT insulin area under the curve (AUC) indicated IR in 44%, while HOMA-R only 18% of 49 normal weight women with polycystic ovary syndrome [42]. A gender (and racial) difference in the utility of insulin-based fasting and OGTT-based models has recently been described, as both gender and race, had a significant effect on explaining the predictability of clamp-measured glucose disposal rates [43]. Although we did not find significant difference between genders in the value of either fasting or OGTT indices and could not assess the effect of race as all subjects were Caucasian in our study, we did find some basic gender difference when the associations between liver enzymes and insulin sensitivity were analyzed.

4.2. ALT – a possible simple indicator of insulin sensitivity in women

One of the most important findings of our study in this healthy / prediabetic population is that after the adjustment for confounding factors such as age, BMI, abdominal circumference, body fat percent, HbA1c, alcohol consumption (and FSH levels in women), all three liver enzymes (ALT, AST and GGT) stayed significantly associated with clamp-measured insulin sensitivity (i.e. muscle glucose uptake) in women but disappeared in men. This difference was only applicable for the gold standard clamp measured peripheral insulin sensitivity, i.e. the association with the estimated OGTT derived Hepatic Insulin Resistance Index (although stronger in females than in males) disappeared in both genders after the correction was done (Table 5.). Moreover, the multiple regression model has found that ALT was a significant independent predictor of clamp insulin sensitivity besides BMI in females. In men, this was fasting insulin and leptin but none of the transaminase levels. Lee et al. described a similar gender difference in a study on adolescent population: obesity and triglyceride were the major determinants of HOMA-IR in boys, and obesity and GGT in girls, so liver function test (i.e. transaminase level) only predicted insulin resistance measured by HOMA model in females [44]. Furthermore, the independent association with IR and ALT

was stronger in girls than in boys ($p = 0.034$ vs. $p = 0.005$) [44], albeit the latter result was not confirmed by clamp studies. Those studies having found an independent association between ALT and directly (clamp or minimal model analysis) measured insulin sensitivity were carried out on mixed-gender populations (either healthy prediabetic or IGT subjects), although results stayed significant after the adjustment for sex and other confounding factors [45, 46].

Our results support the hypothesis that a very delicate sex difference exists in the progression / association of NAFLD with metabolic parameters in the adult population and this has an important clinical implication. In women, it is clearly evident that insulin resistance per se might indicate liver fat accumulation, and vice versa, elevated ALT levels might indicate decreased insulin sensitivity earlier than fasting insulin, lipoprotein or adipokine levels. In men, ALT (also AST and GGT) elevations coexist with other metabolic changes followed / caused by insulin resistance. Therefore liver enzyme elevation per se is not an indicator of decreased insulin sensitivity but a general metabolic deterioration along with insulin resistance in men, with no independent associations with the clamp M3 value. The obvious sex difference in fat distribution leads to increased susceptibility to intra-abdominal, visceral and liver fat accumulation in men, which is at least partially driven by differential sex hormone settings [47]. A further explanation and/or consequence is the sexual dimorphism displayed by liver-associated markers, such as sex hormone-binding globulin (SHBG) and adiponectin levels being much lower in men, consistent with their greater insulin resistance and greater risk of diabetes and cardiovascular disease at a younger age [48, 49] and the more severe metabolic phenotype at the diagnosis of NAFLD [50]. This is in agreement with the finding of Feitosa et al., that ALT is a significant independent predictor of coronary heart disease in men but not in women, with the association being stronger in non-diabetic men [51].

Based on the above results slightly elevated ALT may strongly indicate the presence of insulin resistance in females even without hyperinsulinemia, especially in overweight women. Hence, the use of ALT in estimating clamp measured insulin sensitivity might be more relevant in females, while that of fasting insulin-based indices (i.e. the HOMA model) physiologically seems to be more appropriate in males according to our results. A future direction could be to include transaminase levels within the HOMA index, which may increase the diagnostic value of the HOMA model, especially in females (correlations with HOMA-S in all females is improved by including ALT into the equation in women: $R = 0.495$ vs. $R = 0.588$, non-published data). As a conclusion we found that: 1. Conventional

fasting IS indices only correlate moderately with the clamp results, 2. OGTT derived IS indices correlate well with the clamp data so whenever it is possible they should be used instead of the less precise fasting indices, 3. Liver transaminase levels may increase the diagnostic value of the conventional fasting indices, especially in females, and as such they always should to be considered to be a useful tool for IR diagnosis as part of the routine laboratory assessments, either with or without HOMA model. Moreover, we must take this into consideration when trying to interpret unexplained transaminase elevation, especially in female patients.

4.3. Bone – energy homeostasis axis

4.3.1. Discussion of epidemiologic data

Numerous studies have been published about the bone effects of diabetes, most of these show higher BMD in T2DM [52-54] and lower in T1DM patients [52, 55, 56]. Diabetes is an independent risk factor for osteoporosis and bone fracture in the elderly normal weight women [57]. Patients suffering from T2DM initially even with higher BMD seem to have an accelerated bone loss and increased risk of bone fracture [58]. In most of the studies the favorable effect of T2DM on BMD and osteoporosis is decreased or ceased if data are adjusted with BMI, suggesting that the protective effect of diabetes is via increased body weight. This finding has been confirmed in our study as based on our results the increased BMD in T2DM is primarily due to increased BMI; diabetes per se rather increases the risk of than protects against osteoporosis.

Less results are available about gender differences, studies done in this field have found that the positive correlation between T2DM and bone density is weaker or lacking in men, furthermore the positive association between hyperinsulinemia and BMD appears to be stronger in women than in men [52, 54, 59, 60]. Our preliminary clamp studies have confirmed that during the initial stages of impaired carbohydrate tolerance the whole body glucose utilization rate (which is the gold standard method of insulin sensitivity) shows a significant association with bone density only in females [61]. The results of our screening tests showed that blood sugar level has a significant correlation with bone density only in elderly males and the classic positive correlation between BMI and BMD was only demonstrated in females. We found that in women in their early postmenopausal age (between 50 – 60 years) osteopenia was significantly more frequent than in the normal

glucose tolerant group despite the significantly increased BMI, although no difference was observed in the prevalence of osteoporosis.

In conclusion our results have confirmed findings which show that T2DM (similar to T1DM) increases the frequency of osteopenia and osteoporosis in certain populations (like postmenopausal women and normal weight diabetic patients), furthermore the increased BMD generally found in T2DM is a consequence of increased body mass index as such diabetes can be considered rather risk increasing than protective factor. Our results have also drawn the attention to the notion that basic gender differences may exist in the bone – energy homeostasis relationship.

4.3.2. The role of OCN; gender difference in the bone – energy homeostasis relationship

4.3.2.1. OCN and adiponectin

Between 2005 and 2010 we have measured non-carboxylated OCN levels of all either normal or impaired glucose tolerant subjects, or drug naïve type 2 diabetic patients who underwent a hyperglycemic normoglycemic clamp to measure their insulin sensitivity. OCN is believed to be one of the most important neurohumoral link between bone and energy homeostasis and as such measuring total OCN levels in patients with known insulin sensitivity in a wide range of different levels of glucose tolerance may give us important clues about the pathophysiologic background of diabetes induced bone loss and causes of possible gender differences. In our preliminary clamp studies we have first shown on human data that OCN shows a positive correlation with insulin sensitivity [62], furthermore we have also described the existence of a basic gender difference in the relationship of insulin sensitivity with bone functional metabolic unit derived from the ratio of bone formation (osteoprotegerin (OPG), procollagen type 1 N-terminal propeptide (P1NP)) and resorption markers (cathepsin-k, soluble receptor activator NF-kB ligand (RANKL) and β -crosslaps) [63].

A putative feed-forward regulatory loop ties bone turnover to energy regulation as proposed by Ferron et al. and Fulzele et al. in 2010 [64, 65]. Insulin activates skeletal remodeling (that is, increases bone formation by osteoblasts and resorption by osteoclasts), which in turn releases uncarboxylated osteocalcin from the skeletal matrix into the

circulation. This enhances insulin secretion and increases the insulin sensitivity of adipocytes. Based on the findings of Lee et al. OCN also increases insulin sensitivity by stimulating the secretion of adiponectin besides having a direct hypoglycemic effect via its secretagogue function on pancreatic beta cells [66]. We have found that even following the adjustment by confounding factors, higher OCN values were associated with higher adiponectin levels in female subjects (Table 2), similar to earlier studies [4, 10]. The same association was not found in male subjects. This is in accordance with the study of Kanazawa et al. who found a significant positive association of serum osteocalcin and adiponectin levels only in postmenopausal women, but not in men [67]. The authors suggested that the difference in adiponectin levels within sexes may have contributed to these findings but no further assessments were done to explore the possible rationale behind. In contrast it was also reported by others that the inverse relationship between carboxylated OCN and adiponectin is independent of gender and stays significant after correction with BMI [68]. Our study confirms the association of serum OCN with improved metabolic state (i.e. increased insulin sensitivity and glycemic control) but based on our findings the mechanism of action may differ significantly between sexes.

4.3.2.2. OCN in postmenopausal women

Bone turnover is accelerated with increasing age, especially in women [69, 70], which may be the reason for elevated serum OCN levels with higher age found both genders [71]. Our results confirmed that higher OCN was indeed associated with increasing age and FSH levels in female but not in male subjects. The significant positive correlation between OCN and improved metabolic parameters were only found after the correction for age and FSH, i.e. higher OCN levels due to increased bone turnover with increasing age and insulin resistance would mask its 'true' metabolic character. After further adjustment with body composition and HbA1c in addition to age, most of the significant correlations between OCN and metabolic parameters decreased or disappeared entirely, except for fasting glucose. Further adjustment with adiponectin resulted in lost correlation between OCN and fasting glucose, which supports the mediating role of adiponectin in some but not all of the metabolic effects of OCN in females. These findings disagree with recent data from Hwang et al. [9], who found that although serum OCN levels were indeed associated with improved glucose tolerance, beta cell function and OGTT measured insulin sensitivity, these were independent

of the plasma adiponectin levels. However, their data were derived from a mixed-gender population. According to our results the positive metabolic effect of OCN in women is not related to pancreatic beta cell function (as no correlation was found with AIR) but rather to adiponectin. In fact, the adiponectin mediated insulin sensitization pathways first confirmed in mice [66] were at least partly confirmed in women by our findings.

4.3.2.3. OCN in men

In contrast to female subjects, no sign of an ‘OCN–adiponectin axis’ was detected in men; i.e. no correlation was found between adiponectin and serum OCN level albeit the expected strong independent association with improved insulin sensitivity was indeed present. Recently, it was demonstrated that osteoblasts may induce testosterone production by the testes, although ovarian estrogen production in females is unaffected. Analysis of cell-specific loss- and gain-of-function models has revealed that OCN performs this endocrine function by binding to a G protein-coupled receptor expressed in Leydig cells, and OCN regulates the expression of enzymes required for testosterone synthesis in an cAMP response element-binding protein-dependent manner, thus promoting germ cell survival [12]. On the other hand, meta-analysis of cross-sectional data suggests that the metabolic syndrome is associated with male hypogonadism / hypotestosteronemia [72]. The same association was not confirmed in females [13]. Our results show that adiponectin is not associated with OCN levels in male subjects; however, increasing OCN levels were indeed associated with higher testosterone values, independent of age, weight, adiposity and HbA1c. When data were adjusted for testosterone levels, some of the significant correlation between OCN and insulin sensitivity, leptin and BMI disappeared. Correlations with whole body glucose utilization, BFP, waist circumference, and triglyceride values were not affected. These results might suggest a partial role of testosterone in the metabolic connections of OCN in males, although because these data were cross-sectional, the cause–causality relationship needs to be further clarified.

4.3.2.4. Lessons learned from feature selection analysis

Feature selection analysis clearly describes the gender difference: in women, OCN is independently associated with adiponectin, age, and three of the IVGTT glucose values; while in men, it is associated with testosterone, BMI, BFP, M1 and M3, OGTT 90th minute insulin and glucose, and leptin levels. Thus, an ‘OCN–adiponectin–glycemic state axis’

versus ‘OCN–testosterone–insulin resistance axis’ may exist in women versus men, according to our data. The sex difference might be partially explained by the difference in adiponectin concentration, as adiponectin levels are significantly higher in women at all levels of glucose tolerance (normal, prediabetes and type 2 diabetes) than in men [73]. The fact that several papers have confirmed the positive association between serum OCN and adiponectin in postmenopausal women [74, 75] but not in men [4, 67] supports the notion that OCN stimulates the expression of adiponectin only in female adipocytes in humans. In an earlier study, testosterone treatment reduced plasma adiponectin concentration in male mice, whereas testosterone also reduced adiponectin secretion in adipocytes [76]. It is therefore plausible to assume that high testosterone levels counteract with OCN in stimulating the expression of adiponectin in adipocytes in men. This could explain why we (and others) did not find any association between OCN and adiponectin in men. Furthermore, recently a large cross-over study has shown that adiponectin is inversely and independently associated with bone mass only in women but not in men [77] which result also underlines the importance of adiponectin in female over male bone physiology. The fact that in men OCN still has the earlier described positive associations with metabolic parameters proves that adiponectin independent pathways may indeed play a role in the male bone–energy homeostasis. Testosterone can only be partially responsible as our results have also confirmed this notion. Moreover, lower testosterone levels indeed coincide with worsening metabolic state with decreasing OCN, and this could alone explain numerous observations that may interfere with the interpretations of our results. Further longitudinal studies are needed to determine the real nature of the OCN–testosterone axis, particularly its long term effect on metabolic state of men.

Further data analysis of our clamp studies have shown that basic gender differences exist in other neurohumoral links of the bone – energy homeostasis relationship, i.e. the OPG/RANKL system which besides being important regulators of the bone resorption / formation cycle, have also been associated diabetes severity, diabetic micro- and macrovascular complications, including acute cardiovascular mortality and morbidity [78, 79, 80]. We have recently shown that while osteoprotegerin (OPG) seemed to be an indicator of the presence of small-dense LDL subfractions (LDL-2-5) along with decreased presence of the ‘normal’ large buoyant (LDL-1) subfraction phenotype in women, a positive association with clamp measured insulin sensitivity was shown only in men, with no clinically relevant associations with lipid subfractions [81].

The significance of this finding is still unclear, although it draws the attention to the gender specific nature of the association between bone turnover markers and metabolic diseases. Since the OPG/RANKL system is deeply involved in the OCN – insulin homeostasis axis by regulating bone resorption and so providing the acidic environment in the resorption pit for OCN undercarboxylation which results in the production of functionally active, i.e. ‘undercarboxylated’ OCN, it is plausible to presume that the OPG/RANKL system has an important role in the metabolic functions of OCN although this needs to be further assessed by in vitro / in vivo / observational and clinical studies.

4.4. Limitations of our study

Our study has several limitations. First, based on its cross sectional study design, the present findings are inherently limited in the ability to eliminate causal relationships between the investigated metabolic / bone parameters / liver markers and insulin resistance or sensitivity. Total instead of undercarboxylated OCN values were measured which latter is the metabolically active form of OCN. Some of the discrepancies found between our results and earlier data might have been resulted from this (i.e. lack of relationship between serum OCN and parameters of insulin secretion, like AIR or IGI of IVGTT), at least in part. We measured total instead of free testosterone levels, which is the bioavailable form of testosterone, although its metabolic role was indeed proven by large scale studies using both total and free testosterone measurements. The population we investigated involved both diabetic as well as non-diabetic subjects, both pre- and postmenopausal women although appropriate adjustments were made for confounding factors such as HbA1c, body composition, age or FSH. Since some of the study population had several risk factors, including hypertension, and dyslipidemia, we could not eliminate the possible confounding effect of underlying diseases on the findings, although the prevalence of controlled hypertension, smoking and dyslipidemia were similar throughout the male and female groups in our study population.

5. Summary of new observations and future implications

Between 2005 and 2010 we have carried out approximately 300 hyperinsulinemic normoglycemic clamp tests mostly on normal glucose tolerant and prediabetic subjects also including drug naïve type 2 diabetic patients at the time of the clamp. Our major purpose was to find metabolic associations in this patient population that would help us to identify potential subjects who later develop insulin resistance and type 2 diabetes and find clinical characteristics that would accompany insulin resistance although not necessarily being essential part of the ‘insulin resistance syndrome’. At first we have assessed the diagnostic value of simple fasting and OGTT derived insulin sensitivity indices on normal and impaired glucose tolerant (prediabetic) population and we found that the HOMA model used to diagnose insulin resistance in the everyday clinical practice does not precisely predict clamp measured insulin sensitivity especially in the elder populations. OGTT derived insulin sensitivity indices may be better alternative, because of the need for serial insulin measurements the cost of such investigations however may be high. In Hungary this was the first large sample sized clamp study which specifically addressed this issue. Using simple liver function test (like ALT) to improve the value of the HOMA model might be helpful, albeit because of the gender specific nature of the aethiopathogenesis of IR driven steatohepatosis described by us first in detail, it can only be a valuable approach in women according to our findings.

The results of a large sample sized epidemiologic study in ‘Balaton Upper-land’ carried out by us between 2005 and 2007 including more than 6000 subjects, drew our attention to the ‘Janus faced’ nature of the relationship between type 2 diabetes and bone metabolism, i.e. although higher BMI predicts higher BMD values, there is still an increasing prevalence of osteoporosis and osteopenia on certain diabetic / prediabetic populations, like normal weight diabetic subjects. Moreover, we have found important data about a possible gender difference in this respect which was also confirmed by our preliminary clamp data. We have measured non-carboxylated OCN in all prediabetic patients who underwent a hyperinsulinemic clamp study to assess the association between metabolic data including clamp measured insulin sensitivity and OCN values which believed to be the most important link in the bone – energy homeostasis axis. We have proved that an ‘OCN–adiponectin–glycemic state axis’ versus ‘OCN–testosterone–insulin resistance axis’ exist in women versus

men which draws the attention to the gender specific nature of the bone – energy homeostasis association but also gave insight to the potential mode of actions OCN may have on insulin and glucose metabolism.

Future directions include gender specific approach in insulin resistance studies, including the assessment of bone specific biomarkers like OCN, OPG and RANKL, their possible role in interventions to prevent or predict insulin resistance / type 2 diabetes and complications. Furthermore, studying gut microbiome in relation to bone metabolism, which is the collective metagenomic character of the overall bacterial composition of the gut (microbiota) in prediabetic state may be a promising new direction, since microbiota may be an important etiological factor and a possible promising therapeutic target for T2DM (and T1DM) [82], however only very few data exist about the association of microbiome and bone metabolism, mostly in preclinical models [83]. Moreover besides genetic, epigenetic studies are also being urged to identify possible markers of the prediabetic state, and type 2 diabetes [84]. Most importantly to find biomarkers and other characteristics in the juvenile / adolescent population in long term follow-up observational / epidemiologic studies that predict the parallel development of type 2 diabetes and bone loss later in life which could possibly make the development of these diseases / symptoms preventable. Systems medicine is a promising tool to identify factors that play a role in the etiology, disease progression and the success of therapeutic interventions of insulin resistance and type 2 diabetes [85]. Discovering more sides of the bone – energy homeostasis by using systems medicine will enable us to understand clinically meaningful associations that would help us to prevent diabetes linked bone loss, which would have a great impact on the diagnosis and prevention of the two most burdening diseases in today's health care.

6. References

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I.

II.

III.

IV.