

Type 2 Deiodinase p.Thr92Ala polymorphism does not affect the severity of obesity and weight loss after bariatric surgery

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Abstract

Introduction:

A single nucleotide polymorphism in the Type 2 deiodinase (*DIO2*) gene (p.Thr92Ala) was found to be associated with hypertension, type 2 diabetes mellitus (T2DM), insulin resistance, and body mass index (BMI). We retrospectively evaluated 182 patients to assess whether the *DIO2* p.Thr92Ala was associated with severe obesity and response to bariatric surgery.

Methods

Genomic DNA was extracted from peripheral blood leukocytes before surgery. Glycemic control parameters, liver function, cardiometabolic biomarkers and hormonal parameters were assessed at baseline and after surgery.

Results

Based on genotype evaluation, 78/182 (42.9%) patients were homozygous wild-type (Thr/Thr), 83/182 (45.6%) heterozygous (Thr/Ala), and 21/182 (11.5%) rare homozygous (Ala/Ala). Age at diagnosis was significantly lower in patients with *DIO2* p.Thr92Ala. No significant association was observed between *DIO2* p.Thr92Ala and BMI, excess weight, waist circumference, homa index and prevalence of comorbidities. After bariatric surgery, excess weight loss (EWL) % and remission from comorbidities occurred without differences according to genotypes.

Conclusions

DIO2 p.Thr92Ala does not affect the severity of obesity and its complications, but it seems to determine an earlier onset of morbid obesity. The presence of polymorphism seems not to impact on the response to bariatric surgery, both in terms of weight loss and remission of comorbidities.

Introduction

Type 2 deiodinase (D2) is an intracellular enzyme which catalyzes the conversion of thyroxine (T4) to its active form triiodothyronine (T3) controlling the intracellular T3 concentration, its availability to the nucleus and the saturation of the nuclear T3 receptor in target tissues [1]. Therefore, D2 is a very important regulator for tissue metabolic activity. A common single nucleotide polymorphism in the *DIO2* gene, resulting in a threonine change to alanine at codon 92 (p.Thr92Ala, rs225014) [2], has been identified in 12–36% of the general population [3]. Many works have evaluated how this polymorphism could participate as a disease mechanism and/or affect clinical outcomes [1] The most obvious possibility is that p.Thr92Ala substitution affects D2 catalytic activity, reducing T3 production and

causing either or both systemic/localized hypothyroidism [4]. Different studies have investigated the Ala92-D2 enzyme both *in vitro* and *in vivo* [5–7]. In a recent study, a comparison of presurgical hormonal status of LT4-treated thyroidectomized individuals with their post-surgery status revealed an association between low FT3 values and p.Thr92Ala variant [8]. In the carriers of the p.Thr92Ala-DIO2 polymorphism there is evidence of upregulation of pathways related to the mitochondria, Golgi apparatus/ER transport, oxidative stress, and apoptosis [9]. Population-based studies have suggested associations between p.Thr92Ala with hypertension [10, 5], bipolar disorder [11], mental retardation [12], low intelligence quotient (IQ) [13], recovery from lung injury [14], osteoarthritis [15], increased bone turnover [16], Grave's and Hashimoto's diseases [17]. In addition, many studies have shown that the p.Thr92Ala polymorphism is related to type 2 diabetes mellitus (T2DM) [16], insulin resistance [18; 17], and body mass index (BMI) [3, 4 16–21]. Intriguingly, most of these associations are independent of serum thyroid hormone levels, which highlight the importance of local regulation of thyroid hormones in peripheral tissues [3].

Thus, given the emerging essential physiological importance of DIO2, the aim of this study was to investigate the DIO2 p.Thr92Ala polymorphism in relation to obesity and its comorbidities in a cohort of patients with severe obesity submitted to bariatric surgery. Furthermore we investigated the role of DIO2 p.Thr92Ala polymorphism in bariatric surgery efficacy.

Materials And Methods

We retrospectively evaluated 182 patients with obesity followed at our Unit of Endocrinology and submitted to bariatric surgery from January 2011 to December 2019. The diagnosis of obesity was based on Body Mass Index (BMI) calculated using the formula: weight in kilograms divided by height in meters squared. Indication for bariatric surgery were: BMI ≥ 40 kg/m²; BMI ≥ 35 –40 kg/m² with associated comorbidities; BMI 30–35 kg/m² and type 2 diabetes with poor control despite optimal medical therapy. Different bariatric procedures were performed according to clinical comorbidities of patients: in 19,2% of patients roux-en-Y gastric bypass (RYGB), in 28,1% one anastomosis gastric bypass (OAGB), in 40,6% sleeve gastrectomy (SG) and in 12,1% adjustable gastric banding (AGB). At screening, glycaemic control parameters (fasting glucose, insulin and glycated haemoglobin), liver function and cardiometabolic biomarkers (lipid assessment, uric acid, pulse rate and blood pressure) were assessed. The index of insulin resistance, HOMA index, was calculated as the product of the fasting plasma insulin level (μ U/mL) and the fasting plasma glucose level (mmol/L), divided by 22.5. At follow-up visit (6 and 12 months after bariatric surgery) weight and excess weight loss % (EWL%) calculated using the formula: $[(\text{final weight} - \text{ideal body weight}) / \text{ideal body weight}] \times 100$ were evaluated. Metabolic biomarkers were also performed.

Thyroid hormones and TSH were evaluated using a chemiluminescent immunometric assay (Access Immunoassay Systems 2006, Beckman Coulter, Milan, Italy). Normal ranges were 2.5-4.1 pg/mL for free T3 (FT3), 5.8 -16.4 pg/mL for free thyroxine (FT4), and 0.4-4.0 mIU/L for TSH.

A written informed consent was obtained from each patient.

DNA extraction and DIO2 Thr92Ala analysis

Genomic DNA was extracted from peripheral blood leukocytes using standard procedures [22] and concentration was assessed with Nanodrop One (Thermo Scientific, Milan, Italy). DIO2 p.Thr92Ala was investigated as described [8].

Statistical analysis

Epidemiological data are presented as the mean \pm SD and median when needed. The categorical variables were compared between groups using a Chi-squared test or Kruskal Wallis test when needed. Interaction with SNP was tested by Chi-squared test at genotype and allele levels. In addition to basic tests, the association of genotype was evaluated assuming dominant and recessive models. Statistical analysis was performed using the software package SPSS v13.0. A p-value <0.05 was considered statistically significant.

Results

Baseline characteristics and prevalence of DIO2 p.Thr92Ala polymorphism in the study population

Our study population included 135 females (74.2%) and 47 males (25.8%). The mean age was 43 ± 11 years (range 18-68 years), the mean BMI was 45.3 ± 7.0 kg/m² (range 33.9-78.9 kg/m²) and the mean excess weight (EW) was 40 ± 12.7 kg (range 21.1-99.2 kg). Based on genotype evaluation, 78/182 (42.9%) patients were homozygous wild-type (Thr/Thr), 83/182 (45.6%) were heterozygous (Thr/Ala), and 21/182 (11.5%) were rare homozygous (Ala/Ala). These frequencies were similar to those observed in European and Tuscany populations in particular according with Ensembl Genome Browser Database (ENSG00000211448) [23] (Table 1).

Table 1
Genotype distribution in European, Tuscany and our obese population.

	<i>European (%)</i>	<i>Tuscany (%)</i>	<i>Study population n= 182 (%)</i>
Thr/Thr	45.1	39.3	42.9
Thr/Ala	40.8	49.5	45.6
Ala/Ala	14.1	11.2	11.5

Anthropometric and metabolic results according to p.Thr92Ala variant

The results of association between the presence/absence of DIO2 p.Thr92Ala and anthropometric characteristics, circulating free thyroid hormones, TSH, lipid and glycemic profile are shown in Table 2. Age at diagnosis was significantly lower in patients with DIO2 Thr92Ala polymorphism (p=0.03) even if we considered dominant and recessive model. BMI was similar among the three groups at genotype level but homozygous Ala/Ala patients showed a waist circumference lower than wild type and heterozygous subjects [cm 117.5±13.1 (Ala/Ala) vs. 127.4±17.1 (Thr/Ala) and 124.7±14.8 (Thr/Thr); p=0.05]. No association at genotype and allele levels was found between the DIO2 p.Thr92Ala and LDL-cholesterol, triglycerides and HOMA IR. EW was near to significance (p=0.07) being lower for homozygous Ala/Ala (36.1 ± 9.2 Kg) compared with wild type (38.7 ± 12.2 kg) and heterozygous (42.1 ± 13.6 Kg) subjects. Wild type Thr/Thr patients had greater fasting glucose levels compared with heterozygous and rare homozygous [mg/dl 118.2 ± 56.7 (Thr/Thr) vs. 115.6 ± 43.2 (Thr/Ala) and 96.4 ± 11(Ala/Ala); p = 0.04]. DIO2 Thr92Ala variant was not associated with variation in TSH, FT4 and FT3 values.

Table 2
Clinical features according with genotype in 182 obese subjects

	<i>Thr/Thr</i> (n=78) Mean ± SD	<i>Thr/Ala</i> (n=83) Mean ± SD	<i>Ala/Ala (n=21)</i> Mean ± SD	<i>P value</i>
Age (years)	45.4 ± 11,34	41.1 ± 10.8	44.6 ± 11.5	0.03
Sex (female:male)	61:17	56:27	18/3	0.1
BMI (kg/m2)	44.6 ± 6.3	46.5 ± 7.8	43.4 ± 5.8	0.1
EW (kg)	38.7 ± 12.2	42.1 ± 13.6	36.1 ± 9.2	0.07
Waist (cm)	124.7 ± 14.8	127.4 ± 17.1	117.5 ± 13.1	0.05
LDL-cholesterol (mg/dl)	111 ± 30.7	110.1 ± 32	122 ± 31.2	0.2
HOMA _{IR}	6.2 ± 6.6	6 ± 5.2	4.2 ± 2.6	0.3
TSH	2.2 ± 1.5	3.9 ± 7.8	1.9 ± 0.9	0.2
ft3	3.7 ± 1.6	3.5 ± 0.4	3.4 ± 0.4	0.3
ft4	8.7 ± 1.4	8.6 ± 1.6	8.5 ± 1.1	0.8

Prevalence of obesity comorbidities according to p.Thr92Ala variant

Obesity-related comorbidities were evaluated in a subgroup of patients (n=133) with complete 12 months follow-up. The prevalence of hypertension was 50.4% (67/133), of T2DM was 24% 32/133), of dyslipidemia was 45,9% (61/133), of metabolic syndrome was 51.9% (69/133). As shown in Table 3, the

prevalence of comorbidities was not associated with allele distribution except for hypertension that was more frequent in wild-type patients ($p=0.03$).

Table 3
Prevalence of obesity comorbidities in *Thr92Ala* subgroups

	<i>Thr/Thr</i> Prevalence (%)	<i>Thr/Ala</i> Prevalence (%)	<i>Ala/Ala</i> Prevalence (%)	<i>P value</i>
Hypertention	61,4% (35/57)	37,9% (22/58)	55,5% (10/18)	0.03
Diabetes mellitus	22,8% (13/57)	25,9% (15/58)	22,2% (4/18)	0.9
Dyslipidemia	50,9% (29/57)	37,9% (22/58)	55 5% (10/18)	0.2
Metabolic syndrome	56,1% (32/57)	48,2% (28/58)	50% (9/18)	0.6

DIO2 p.Thr92Ala and weight loss after bariatric surgery

Weight loss data were available in 150/182 (82.4%) patients at 6 months follow-up and in 133/182 (73%) patients at 12 months of follow up. The mean EWL% at 6 and 12 months was 57.4% and 49.4% in wild type patients, 62% and 50.1% in *Thr/Ala* group, 59.4% and 48% in *Ala/Ala* group, respectively, without significant differences ($p=0.7$ and $p=0.8$, respectively)(shown in Fig. 1).

p.Thr92Ala polymorphism and obesity-related comorbidities after bariatric surgery

The rate of remission of comorbidities was evaluated at 12 months follow-up (Table 4). Remission from hypertension occurred in 40% of patients (27/67), without differences between genotypes ($p=0.7$). Similarly, remission from diabetes mellitus occurred in 93.7% of patients (30/32) without differences between the *DIO2* genotypes ($p=0.8$). Remission from dyslipidemia and metabolic syndrome occurred in 57.4% (35/61) and 81.1% (56/69) of patients respectively, without differences between the *DIO2* genotypes ($p=0.7$ and $p=0.7$, respectively).

Table 4
p.Thr92Ala polymorphism and obesity-related comorbidities after bariatric surgery (12th month)

	<i>Thr/Thr</i> Prevalence (%)	<i>Thr/Ala</i> Prevalence (%)	<i>Ala/Ala</i> Prevalence (%)	<i>P value</i>
Hypertention	37,1% (13/35)	40,9% (9/22)	50% (5/10)	0,7
Diabetes mellitus	92,3% (12/13)	9 3,3% (14/15)	100% (4/4)	0,8
Dyslipidemia	55,1% (16/29)	63,6% (14/22)	50% (5/10)	0,7
Metabolic Syndrome	78,1% (25/32)	85,7% (24/28)	77,7% (7/9)	0,7

Discussion

Several previous studies suggested an association between p.Thr92Ala polymorphism and hypertension [10], bipolar disorder [11], mental retardation [12], low intelligence quotient (IQ)[13], recovery from lung injury [14], osteoarthritis [15], increased bone turnover [16], Grave's and Hashimoto's diseases [17]. To our knowledge no studies were focused on patients with obesity and, in particular on subjects affected by severe obesity (median of BMI 44.8 Kg/m²) undergoing to bariatric surgery. We investigated the DIO2 p.Thr92Ala polymorphism in relation to obesity and its comorbidities in a group of patients submitted to bariatric surgery. The prevalence of DIO2 p.Thr92Ala polymorphism observed (11.5% Ala/Ala homozygous variant) was similar to that reported in the general European population (14.1% Ala/Ala homozygous variant) and in the Tuscany population (11.2% Ala/Ala homozygous variant) [23], excluding a possible role of the p.Thr92Ala polymorphism in promoting the onset of severe obesity. Similar results were observed in a relatively large population-based cohort of whites in which DIO2 p.Thr92Ala genotype was evaluated in patients with and without obesity [19]. When a dominant or a recessive model for the penetrance of Ala92 allele were used, no association with obesity was detected [16, 18, 19]. On the contrary, other published studies reported a significant association between p.Thr92Ala polymorphism and BMI, but only when patients without obesity were evaluated [24]. Moreover a correlation was observed only in patients with both p.Thr92Ala and p.Trp64Arg ADRB3 polymorphisms [4, 18]. This interaction was not replicated in studies on the Pima Indians [16] and in another large cohort of more than 7000 Danish white subjects [19]. In our study an early-onset of severe obesity was significantly associated with DIO2 polymorphism. In details, the mean age at bariatric surgery was 45.4±11.3 years in wild type patients, 41.1±10.8 years in patients with rare homozygous variant (Ala/Ala) and 44.6±11.5 in heterozygous Thr/Ala patients (p=0.03). These results support a possible role of DIO2 p.Thr92Ala polymorphism in contributing to early-onset of severe obesity, although we do not believe that this is a major determinant for obesity among our study population. To our knowledge no published studies evaluated the association between obesity-related comorbidities and p.Thr92Ala polymorphism. We did not find any association between p.Thr92Ala polymorphism and the presence of obesity-related comorbidities such as diabetes mellitus, liver steatosis, and sleep apnoea syndrome or metabolic syndrome. Surprisingly, a higher rate of hypertension was observed in wild type patients. Published studies in which the association between the DIO2 p.Thr92Ala polymorphism and glycemic control in T2DM patients has been explored, reported contradictory results [2, 3, 19, 25]. Nevertheless, a recent meta-analysis found that people who are homozygous for p.Thr92Ala had 4.8% higher HbA1C levels, suggesting that Ala/Ala homozygosity may be associated with worse glycemic control in T2DM patients [26]. Differently to our results, Gumieniak et al reported that the Ala allele increases the risk for development of arterial hypertension [10]. In disagreement with these results no differences were observed in the prevalence of hypertension among the genotypes (Ala/Ala, Thr/Ala, Thr/Thr; 76.4%, 79.1%, 75.7% respectively, p=0.785) when 315 patients with diabetes mellitus were analyzed [27]. These apparently conflicting findings might be partially explained by differences in the studied population. In Gumieniak's report, T2DM patients were excluded and subjects were younger than those included in the study of Canani [10, 27]. Bariatric surgery is currently the most effective tool in determining an important

and lasting weight loss in patients with obesity; in addition a high rate of remission of the main obesity-related comorbidities has been reported after bariatric surgery [28–34]. In our study, bariatric surgery was effective, in terms of weight loss (EWL%) in all patients, regardless the genotype. Moreover, bariatric surgery promotes remission or a significant improvement of obesity related comorbidities without significant differences between the three groups.

Conclusion

Based on the present investigations, performed in relatively large samples of patients, we conclude that the DIO2 p.Thr92Ala polymorphism does not affect the severity of obesity and its complications, but it seems to determine an earlier onset of complicated obesity. Furthermore, the presence of polymorphism does not impact on the response to surgical treatment, both in terms of weight loss and remission of comorbidities.

Declarations

Acknowledgments: None.

Author Contributions: NB and AB contributed equally to conception and design of the study.

SC, CR, CM organized the database and performed the statistical analysis. MGC contributed to interpretation of data for the work and manuscript revision. CC, IS, AT, CV, GV contributed to manuscript revision, read, and approved the submitted version.

Conflicts of Interest Statement: The authors declared no conflict of interest.

Data Availability statements: The data that support the findings of this study are not publicly available due to their containing information that could compromise the privacy of research participants but are available from corresponding author [MGC] upon reasonable request.

Statement of Ethics: All procedure performed in studies involving human participants were in accordance with the ethical standards of the institutional research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. The study was approved by our local ethical committee “ Office of Ethical Affairs, Azienda Ospedaliero-Universitaria Senese”.

Consent to participate statement: a written informed consent was obtained from each patient.

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References

1. Bianco, A. C., Salvatore, D., Gereben, B., Berry, M. J., & Larsen, P. R. Biochemistry, cellular and molecular biology, and physiological roles of the iodothyronine selenodeiodinases. *Endocrine reviews*, 23(1), 38–89 (2002). <https://doi.org/10.1210/edrv.23.1.0455>
2. Canani, L. H. et al. The type 2 deiodinase A/G (Thr92Ala) polymorphism is associated with decreased enzyme velocity and increased insulin resistance in patients with type 2 diabetes mellitus. *The Journal of clinical endocrinology and metabolism*, 90(6), 3472–3478 (2005). <https://doi.org/10.1210/jc.2004-1977>
3. Dora, J. M., Machado, W. E., Rheinheimer, J., Crispim, D., & Maia, A. L. Association of the type 2 deiodinase Thr92Ala polymorphism with type 2 diabetes: case-control study and meta-analysis. *European journal of endocrinology*, 163(3), 427–434 (2010). <https://doi.org/10.1530/EJE-10-0419>
4. Bianco, A. C., & Kim, B. S. Pathophysiological relevance of deiodinase polymorphism. *Current opinion in endocrinology, diabetes, and obesity*, 25(5), 341–346 (2018). <https://doi.org/10.1097/MED.0000000000000428>
5. Peeters, R. P. et al. Polymorphisms in thyroid hormone pathway genes are associated with plasma TSH and iodothyronine levels in healthy subjects. *The Journal of clinical endocrinology and metabolism*, 88(6), 2880–2888 (2003). <https://doi.org/10.1210/jc.2002-021592>
6. Butler, P. W. et al. The Thr92Ala 5' type 2 deiodinase gene polymorphism is associated with a delayed triiodothyronine secretion in response to the thyrotropin-releasing hormone-stimulation test: a pharmacogenomic study. *Thyroid: official journal of the American Thyroid Association*, 20(12), 1407–1412 (2010). <https://doi.org/10.1089/thy.2010.0244>
7. Arici, M. et al. Association between genetic polymorphism and levothyroxine bioavailability in hypothyroid patients. *Endocrine journal*, 65(3), 317–323 (2018). <https://doi.org/10.1507/endocrj.EJ17-0162>
8. Castagna, M. G. et al. DIO2 Thr92Ala Reduces Deiodinase-2 Activity and Serum-T3 Levels in Thyroid-Deficient Patients. *The Journal of clinical endocrinology and metabolism*, 102(5), 1623–1630 (2017). <https://doi.org/10.1210/jc.2016-2587>
9. McAninch, E. A. et al. Prevalent polymorphism in thyroid hormone-activating enzyme leaves a genetic fingerprint that underlies associated clinical syndromes. *The Journal of clinical endocrinology and metabolism*, 100(3), 920–933 (2015). <https://doi.org/10.1210/jc.2014-4092>
10. Gumieniak, O. et al. Ala92 type 2 deiodinase allele increases risk for the development of hypertension. *Hypertension (Dallas, Tex.: 1979)*, 49(3), 461–466 (2007). <https://doi.org/10.1161/01.HYP.0000256295.72185.fd>
11. He, B. et al. Association of genetic polymorphisms in the type II deiodinase gene with bipolar disorder in a subset of Chinese population. *Progress in neuro-psychopharmacology & biological psychiatry*, 33(6), 986–990 (2009). <https://doi.org/10.1016/j.pnpbp.2009.05.003>
12. Ma, S. F. et al. Type 2 deiodinase and host responses of sepsis and acute lung injury. *American journal of respiratory cell and molecular biology*, 45(6), 1203–1211 (2011). <https://doi.org/10.1165/rcmb.2011-01790C>

13. Meulenbelt, I. et al. Identification of DIO2 as a new susceptibility locus for symptomatic osteoarthritis. *Human molecular genetics*, 17(12), 1867–1875 (2008). <https://doi.org/10.1093/hmg/ddn082>
14. Heemstra, K. A. et al. The type 2 deiodinase Thr92Ala polymorphism is associated with increased bone turnover and decreased femoral neck bone mineral density. *Journal of bone and mineral research: the official journal of the American Society for Bone and Mineral Research*, 25(6), 1385–1391 (2010). <https://doi.org/10.1002/jbmr.27>
15. Inoue, N. et al. Functional Polymorphisms of the Type 1 and Type 2 Iodothyronine Deiodinase Genes in Autoimmune Thyroid Diseases. *Immunological investigations*, 47(5), 534–542. (2018). <https://doi.org/10.1080/08820139.2018.1458861>
16. Nair, S. et al. Association analyses of variants in the DIO2 gene with early-onset type 2 diabetes mellitus in Pima Indians. *Thyroid: official journal of the American Thyroid Association*, 22(1), 80–87 (2012). <https://doi.org/10.1089/thy.2010.0455>
17. Estivalet, A. A. et al. D2 Thr92Ala and PPAR γ 2 Pro12Ala polymorphisms interact in the modulation of insulin resistance in type 2 diabetic patients. *Obesity (Silver Spring, Md.)*, 19(4), 825–832 (2011). <https://doi.org/10.1038/oby.2010.231>
18. Mentuccia, D. et al. Association between a novel variant of the human type 2 deiodinase gene Thr92Ala and insulin resistance: evidence of interaction with the Trp64Arg variant of the beta-3-adrenergic receptor. *Diabetes*, 51(3), 880–883 (2002). <https://doi.org/10.2337/diabetes.51.3.880>
19. Grarup, N. et al. Studies of the common DIO2 Thr92Ala polymorphism and metabolic phenotypes in 7342 Danish white subjects. *The Journal of clinical endocrinology and metabolism*, 92(1), 363–366 (2007). <https://doi.org/10.1210/jc.2006-1958>
20. Mentuccia, D. et al. The Thr92Ala deiodinase type 2 (DIO2) variant is not associated with type 2 diabetes or indices of insulin resistance in the old order of Amish. *Thyroid: official journal of the American Thyroid Association*, 15(11), 1223–1227 (2005). <https://doi.org/10.1089/thy.2005.15.122>
21. Dora, J. M. et al. Type 2 deiodinase Thr92Ala polymorphism is associated with disrupted placental activity but not with dysglycemia or adverse gestational outcomes: a genetic association study. *Fertility and sterility*, 101(3), 833–839 (2014). <https://doi.org/10.1016/j.fertnstert.2013.11.018>
22. Miller, S. A., Dykes, D. D., & Polesky, H. F. A simple salting out procedure for extracting DNA from human nucleated cells. *Nucleic acids research*, 16(3), 1215 (1988). <https://doi.org/10.1093/nar/16.3.1215>
23. Ensembl Release 104. Available from: <https://www.ensembl.org/index.html>. Updated May 2021. Accessed November 15, 2021.
24. Fiorito, M. et al. Interaction of DIO2 T92A and PPAR γ 2 P12A polymorphisms in the modulation of metabolic syndrome. *Obesity (Silver Spring, Md.)*, 15(12), 2889–2895 (2007). <https://doi.org/10.1038/oby.2007.343>
25. Leiria, L. B. et al. The rs225017 polymorphism in the 3'UTR of the human DIO2 gene is associated with increased insulin resistance. *PloS one*, 9(8), e103960 (2014). <https://doi.org/10.1371/journal.pone.0103960>

26. Zhang, X. et al. The *Type 2 Deiodinase Thr92Ala Polymorphism* Is Associated with Worse Glycemic Control in Patients with Type 2 Diabetes Mellitus: A Systematic Review and Meta-Analysis. *Journal of diabetes research*, 2016, 5928726 (2016). <https://doi.org/10.1155/2016/5928726>
27. Canani, L. H., Leie, M. A., Machado, W. E., Capp, C., & Maia, A. L. Type 2 deiodinase Thr92Ala polymorphism is not associated with arterial hypertension in type 2 diabetes mellitus patients. *Hypertension (Dallas, Tex.: 1979)*, 49(6), e47–e48 (2007). <https://doi.org/10.1161/HYPERTENSIONAHA.107.088278>
28. Sjöström, C. D., Peltonen, M., & Sjöström, L. Blood pressure and pulse pressure during long-term weight loss in the obese: the Swedish Obese Subjects (SOS) Intervention Study. *Obesity research*, 9(3), 188–195 (2001). <https://doi.org/10.1038/oby.2001.20>
29. Scheen, A. J. et al. L'étude clinique du mois. Chirurgie bariatrique: les résultats à 10 ans de la "Swedish Obese Subjects Study" [Bariatric surgery: 10-year results of the Swedish Obese Subjects Study]. *Revue medicale de Liege*, 60(2), 121–125 (2005).
30. Carlsson, L. M. et al. Bariatric surgery and prevention of type 2 diabetes in Swedish obese subjects. *The New England journal of medicine*, 367(8), 695–704 (2012). <https://doi.org/10.1056/NEJMoa1112082>
31. Romeo, S. et al. Cardiovascular events after bariatric surgery in obese subjects with type 2 diabetes. *Diabetes care*, 35(12), 2613–2617 (2012). <https://doi.org/10.2337/dc12-0193>
32. Burza, M. A. et al. Long-term effect of bariatric surgery on liver enzymes in the Swedish Obese Subjects (SOS) study. *PloS one*, 8(3), e60495 (2013). <https://doi.org/10.1371/journal.pone.0060495>
33. Sjöström, L. et al. Association of bariatric surgery with long-term remission of type 2 diabetes and with microvascular and macrovascular complications. *JAMA*, 311(22), 2297–2304 (2014). <https://doi.org/10.1001/jama.2014.5988>
34. Jamaly, S. et al. Bariatric Surgery and the Risk of New-Onset Atrial Fibrillation in Swedish Obese Subjects. *Journal of the American College of Cardiology*, 68(23), 2497–2504 (2016). <https://doi.org/10.1016/j.jacc.2016.09.940>

Figures

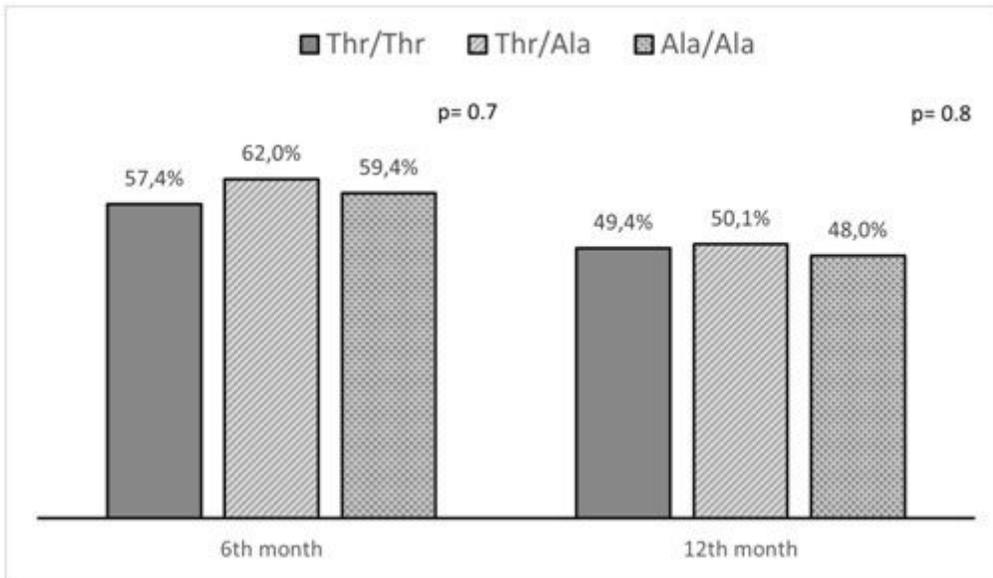


Figure 1

EWL% evaluated at six and twelve months of follow-up after bariatric surgery according to genotype (Thr/Thr, Thr/Ala, Ala/Ala), with no statistically significant differences between the three groups (p=0,7 and p=0,8) .