| 1 | Original Research Article |
|----|---|
| 2 | |
| 3 | Association of T-786C and 27 bp (4b/4a) polymorphism of |
| 4 | endothelial nitric oxide synthase gene with obesity in Egypt |
| 5 | |
| 6 | Abstract: |
| 7 | Background and objective: Endothelial nitric oxide synthase gene polymorphism (e |
| 8 | NOS) is one of three isoforms that synthesize nitric oxide (NO), that participates in |
| 9 | several biological processes have been associated with obesity. This study was |
| 10 | undertaken to determine if eNOS gene (T786C) and 27bp (4b/4a) were associated with |
| 11 | susceptibility of obesity. Materials and Methods: The study was carried out on 200 |
| 12 | cases divided into 100 obese patient and 100 healthy as control. The mean age cases was |
| 13 | (35.3 ± 11.58) they include 79 female and 21 males. All participants were subjected to an |
| 14 | estimation of their body mass index (BMI), weight hip ratio (WHR), in addition to |
| 15 | random blood sugar (RBS), total cholesterol, triglyceride (TG), and lactate |
| 16 | dehydrogenase enzyme (LDH). DNA was amplified using PCR-SSP for detection of |
| 17 | relation between polymorphism and endothelial nitric oxide synthase gene in two parts |
| 18 | T786C and 27bp (4b/4a). Results: All cases showed that there were no significant |

- 19 difference between cases and controls regarding to their chemical lab's analysis (TG,
- 20 Cholesterol, LDL and HDL). All cases showed significant frequency of T786C TT, CC,
- TC vs. controls (p < 0.001) these was considered risk factor for disease. On the other hand
- there no significant difference between 27bp aa, bb, and ab (p=0.618) vs. controls.
- **Conclusion:** The polymorphism T786C not the 27bp in eNOS was associated with
- obesity. However it didn't affect their chemical lab's analysis parameter.
- 25
- 26 Key words: Endothelia Nitric oxide, gene polymorphism, obesity
- Abbreviations: Endothelial nitric oxide (eNOS), polymerase chain reaction with
 sequence specific primers PCR-SSP. Nitric oxide NO.
- 29
- 30
- 31

32 Introduction

Obesity is a medical condition in which surplus body fat accumulated to the range that it might had a negative effect on health. (1), People are generally considered obese when their body mass index (BMI), A measurement obtained by dividing a person's weight by the square of the person's height, is over 30 kg/m2, with

- the range 25–30 kg/m2 defined as overweight (1), Some East Asian countries use lower
- values.(2) Obesity increases the incidence of various diseases and conditions, specially
- cardiovascular diseases, type 2 diabetes, obstructive sleep apnea, definite types of
- 40 cancer, osteoarthritis and depression. (3), (4).
- 41 Obesity is most commonly caused by a mixing of excessive food intake, lack of physical
- 42 activity, and genetic susceptibility.(1),(5) A few cases are caused firstly
- 43 by genes, endocrine disorders, medications, or mental disorder. (6) on the other hand
- obese people eat little next to gain weight because of a slow metabolism is not medically
- 45 supported.(7) .On average, obese people have a greater energy usage than their normal
- 46 people because of the energy required to maintain an increased body mass.(7), (8).
- 47 Obesity might be a cause of death which can be preventable worldwide, with increasing
- rates in adults and children (1). In 2015, 600 million adults (12%) and 100 million
- 49 children were obese in 195 countries. (9) Obesity is more common in women than men
- 50 (1). Several studies viewed that obesity is one of the most dangerous public health
- problems of the 21st century. (10) In 2013, obesity is classified as a disease by
- 52 the American Medical Association. (11), (12).
- 53 Impaired NO. Production is involved in the pathogenesis of several diseases such as
- 54 hypertension, diabetes mellitus, obesity, erectile dysfunction, and migraine. A large
- number of studies showed that polymorphisms in NOS3 gene affect the susceptibility to
- these diseases. Although NOS3 is a highly polymorphic gene, three genetic
- 57 polymorphisms in this gene have been widely studied: the single nucleotide
- polymorphisms (SNPs) g.-786T>C located in NOS3 promoter and in exon 7,
- respectively, and the variable number of tandem repeats 4b/4a (<u>VNTR</u>) characterized by
- 27 bp repeat in intron 4. (13). The C allele for the T786C polymorphism, which
- results in reduced eNOS expression and NO.Production was associated with
- 62 increased risk for hypertension, (14). The VNTR in intron 4 affects eNOS
- expression, (15). And the susceptibility to hypertension, (14). Obesity, (16).
- 64

65 Materials and methods

- 66 **Study group:** This study includes 200 cases 100 obese patients they were recruited from
- 67 the department of diabetes and endocrine unit in specialized medical hospital Mansoura
- 68 University, Egypt, as well as Ministry of Health Hospitals of Dakahlia, Egypt, (New
- 69 General Hospital).during the period September 2016 to May 2018, as 100 patients obese.
- The mean age of cases were 35.30 ± 11.58 years
- 71 **Control group:** for comparison 100 healthy controls were selected.
- 72 **Biochemical analysis:** after obtaining 5ml blood were collected from each case and
- control in an empty tube blood sample for biochemical analysis (random blood sugar
- 74 (RBS), total cholesterol, triglyceride (TG), LDL and HDL.
- 75 Biochemical measurements were carried according to validated method.

76 And Another Venous blood samples were collected from each case and control in

ethylenediamine tetra acetate (EDTA) as an anticoagulant containing tubes., DNA was

extracted from peripheral blood using promptly using DNA extraction and purification

kit (Gentra system, USA) according to manufacturer's instructions and then stored at - 20

80 c till use .

81 Capture column kit extraction and purification:

82 The generation DNA purification capture column kit (Gentra System, USA) is based on a

- proprietary system that uses two reagents, a DNA purification solution and a DNA
- 84 elution solution, along with a specially formulated purification matrix. In this kit, a
- sample is applied directly to the purification matrix contained a spin column .the cells
- contained in sample lyse upon contact with the matrix .once the cells were lysed, DNA
- 87 was captured by the matrix material which make it possible to efficiently wash away

contaminants, leaving the DNA bound to the matrix. Contaminants, including protein

89 heme and RNA were removed from the matrix by washing with DNA purification

90 solution.

91 Following removal contaminants, the DNA released from the matrix using DNA elution

92 solution and heat .samples of purified DNA were ready for analysis and not require93 precipitation.

94 PCR amplifications of each eNOS studied: Single nucleotide polymorphism (SNPs) for

nitric oxide synthase gene (eNOS) were genotyped in this case-control study C786Tand

27bp polymorphism using polymerase chain reaction PCR. Amplification were

97 performed in sequence-specific primer polymerase chain reaction (SSP-PCR) employing

a forward and reverse primer for each part. The region containing one (Restriction

99 Fragment Length Polymorphisms) RFLPs within the eNOS gene was amplified with tag

- 100 DNA polymerase, PCR buffer, Mgcl2 and dNTPs.
- The entire reaction volume plus 5 micro L of bromophenol blue track dye were
 loaded into 2% agarose gel (Bohringer Mannheim) containing ethidium bromide. And for
- 103 30 minutes at 100V Gels were electrophoresed, then photographed under UV light (320
- 104 nm) and then detect the presence or absence of an allele specific bands.

105 Statistical analysis:

- 106 Data were processed and analyzed using the statistical package of social science (SPSS
- 107 program version 17). The student t-test was used to compare the numerical values related
- to genotype polymorphism, cholesterol, TG, LDL and HDL, where CHI square test and
- 109 One-Way ANOVA test were used to compare frequencies of different genotypes and
- alleles between cases and controls
- 111 Results

112 On studying general characteristics of cases of obesity and healthy controls (table1):

the mean age (\pm SD) and the gender frequency in cases of obesity were significant

different from the control as the mean standers for age were $(35.3 \pm 11.58, 28.81 \pm 0.02)$

9.83) for patient and controls with P value 0.001* (males were 21while females were
79 in obese cases and male were 6 while females 94 in healthy controls), however the

age onset were non significantly between cases of obesity and controls with p value 0.744

- as the mean standers for age onset were (27.02 ± 10.90) for patient and 27.51 ± 10.26
- 119 for controls).

121 (p <0.001 for each) (246.32
$$\pm$$
 60.23), (140.76 \pm 95.91), (49.94 \pm 15.60), and (168.85

122 \pm 64.86) for cases of obesity respectively than in healthy controls (181.16 \pm 44.48),(

123 101.74 ± 47.85), (37.54 ± 13.48) and (124.10 ± 40.89) respectively.

124

125 Table 1: Descriptive data of studied cases of obesity and healthy controls.

126

| | Patients (N=100) | Control (N=100) | t | Р |
|-------------|--------------------|--------------------|--------|---------|
| age | 35.3±11.58 | 28.81 ± 9.83 | 4.273 | <0.001* |
| Нір | 122.69 ± 12.96 | 89.26 ± 17.18 | 15.536 | <0.001* |
| Weight | 106.03 ± 16.95 | 68.66 ± 17.77 | 15.216 | <0.001* |
| Height | 162.47 ± 8.26 | 166.38 ± 7.55 | 3.495 | 0.001* |
| BMI | 40.13 ± 6.40 | 25.02 ± 7.67 | 15.132 | <0.001* |
| WHR | 0.95 ± 0.14 | 0.82 ± 0.12 | 7.268 | <0.001* |
| waist | 116.16 ± 15.47 | 74.57 ± 24.76 | 14.245 | <0.001* |
| Age onset | 27.02 ± 10.90 | 27.51 ± 10.26 | 0.327 | 0.744 |
| Cholesterol | 246.32 ± 60.23 | 181.16 ± 44.48 | 8.703 | <0.001* |
| TG | 140.76 ± 95.91 | 101.74 ± 47.85 | 3.640 | <0.001* |
| HDL-C | 49.94 ± 15.60 | 37.54 ± 13.48 | 6.014 | <0.001* |
| LDL-C | 168.85 ± 64.86 | 124.10 ± 40.89 | 5.835 | <0.001* |

127 N = number of cases, t =Student t-test

128 *p <0.001 (significant).

129

131 Table 2: descriptive data of studied cases of obesity.

| | Pa | tients | C | Control | ~? | |
|-------------------------|----|--------|--------|---------|---------|---------|
| | N | % | Ν | % | ×2 | þ |
| Family History | | | | | | |
| there is a family | 60 | 60.09/ | 0 | 0.09/ | | |
| history(+ve) | 00 | 00.0% | 0 | 0.070 | 85 714 | <0.001* |
| no family history (-ve) | 40 | 40.0% | 100 | 100.0% | | |
| BMI gp. | | | | | 10 | |
| <35 | 17 | 17.0% | 100 | 100.0% | 1/1 880 | <0.001* |
| 35-> | 83 | 83.0% | 0 | 0.0% | 141.000 | ~0.001 |
| disease | | | \sim | | | |
| obesity | 53 | 53.0% | 0 | 0.0% | | |
| obesity+D.M | 21 | 21.0% | 0 | 0.0% | - | |
| obesity+HTN | 12 | 12.0% | 0 | 0.0% | | |
| obesity+D.M+ HTN | 14 | 14.0% | 0 | 0.0% | 200.000 | <0.001* |
| normal, no disease | 0 | 0.0% | 100 | 100.0% | - | |
| Sex | | | | | | |
| Male | 21 | 21.0% | 6 | 6.0% | 9.634 | 0.001* |
| Female | 79 | 79.0% | 94 | 94.0% | 9.034 | 0.001 |
| M.SYN | | | | | | |
| Metab syndrome | 46 | 46.0% | 46 | 46.0% | 0.0 | 1.0 |
| Simple obesity | 54 | 54.0% | 54 | 54.0% | 0.0 | 1.0 |

132 N= number of cases, % = percentage of cases, $\chi 2$: Chi-square test

133 *p <0.001 (significant).

134

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139 Comparing all cases with obesity and healthy controls regarding their 140 genotype distribution of e NOS gene polymorphism (in T786C), (table 3): all genotypes 141 (TT), (TC), and (CC) were highly significant (p<0.001) vs. controls. While on alleles 142 analysis both (T) and (C) were significantly. (P<0.001)

143

144

145 Tablet 3: comparison between all cases with obesity and healthy controls regarding

146 their genotype distribution of eNOS gene polymorphism in (T786C).

147

| T786C | | Pa | Patients | | ontrol | ~~~ | n |
|----------|-----|-----|----------|-----|--------|--------|---------|
| | | Ν | % | Ν | % | λ2 | Р |
| Genotype | TT | 33 | 33.0% | 84 | 84.0% | | |
| | TC | 55 | 55.0% | 14 | 14.0% | 53.736 | <0.001* |
| | CC | 12 | 12.0% | 2 | 2.0% | | |
| Alleles | (T) | 121 | 60.5% | 182 | 91% | 50 641 | <0.001* |
| | (C) | 79 | 39.5% | 18 | 9% | 50.041 | ~0.001 |

148

| 149 | N= number of cases, | % = percentage | of cases, TT | T = thymine t | thymine, TO | C =thymine |
|-----|---------------------------------------|----------------|--------------|---------------|-------------|------------|
| | · · · · · · · · · · · · · · · · · · · | 1 0 | , | 5 | <i>J</i> | 5 |

- 150 cytosine, CC= cytosine cytosine, T=thymine, C=cytosine
- 151 Significance using $\chi 2$: Chi-square test:
- 152 *p<0.001 (significant)

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157 Comparing all cases with obesity and healthy controls regarding their 158 genotype distribution of e NOS gene polymorphism **(27 bp) repetition**), (table 4): all 159 genotypes (aa), (ab), and (bb) were non significantly (p=0.618) vs. controls. While on 160 alleles analysis (a) and (b) did not show any significant difference between both, (P 161 =0.482).

163 Tablet 4: comparison between all cases with obesity and healthy controls regarding 164 their genotype distribution of eNOS gene polymorphism in (27 bp) repetition

| 27bp | | Pa | Patients | | ontrol | | n |
|----------|-----|-----|----------|-----|--------|-------|-------|
| | | N | % | Ν | % | ×2 | h |
| Genotype | aa | 15 | 15.0% | 14 | 14.0% | | |
| | ab | 63 | 63.0% | 58 | 58.0% | 0.961 | 0.618 |
| | bb | 22 | 22.0% | 28 | 28.0% | | |
| Alleles | (a) | 93 | 46.5 | 86 | 43% | 0.495 | 0.482 |
| | (b) | 107 | 53.5 | 114 | 57% | 0.495 | 0.402 |

¹⁶⁵ N= number of cases, % = percentage of cases, a=allele a, b= allele b

| 166 | Significance | using χ2: | Chi-square t | est. |
|-----|--------------|-----------|--------------|------|
|-----|--------------|-----------|--------------|------|

Electrophoresis result of PCR showing enzymatic digestion of T786C polymorphism of eNOS gene:

Wild type TT is found which appear at 236bp in lanes 1, 2, 4 and 5, digestion of PCR producted for T786C polymorphism of eNOS gene using NgoMIV enzyme. Which digest the 236-bp fragments into 203 and 33-bp fragments (heterozygous mutated genotype TC which has 236, 203, 3381p fragments lanes 6 only) but (homozygous mutated genotype CC is found which has 203, 3381p fragments lanes 3, 7) by using DNA size marker 50bp 183

| | bp | M | 1 | 2 | 3 | 4 | 5 | 6 | 7 | |
|--------------------------|--|--|--|------------------------------------|---------------------------------------|--|---------------------------------------|--------------------------------------|--------------------------------|-------------|
| | 450 400 350 300 250 200 150 100 | | | | | | | | | 236 203 |
| 405 | 50 | | | | | | | | | < 33 |
| 185 | | | Contract (State) | | | 0 | | | | |
| 100 | Fig 1. | Enzymatic | direction | of T794 | C nolum | ornhism | of aNOS | gano | | |
| 100 | гıg I. | Enzymatic | urgestion | 01 1/80 | л рогуп | norphism | or enus | gene. | | |
| 100 | | | | | | 1 | | | | |
| 100 | | | | | | | | | | |
| 101 | | | | | | | | | | |
| 102 | | | | | | | | | | |
| 102 | | | \sim | 1 | | | | | | |
| 193 | | 1 | | | | | | | | |
| 194 | | | | | | | | | | |
| 196 197 | Electro polymo | ophoresis r orphism of | esult of P NOS g | CR shov gene: | wing PCR | amplific | ation of i | Intron 4 | b/a (27b | op) |
| 198 199 200 201 | PCR p homoz carrier marker | broduct of in ygous lanes heterozygo 50 bp. | ntron 4b/a s 2, 3, 6 ar us which | polymor nd 7 and l has (220, | phism hav have ban s 193 bp fra | ve ban size ize (193) agments la | e (220) by in aa hon anes 1 and | p in bb ca nozygous d 5) by us | arrier 5 lanes 4 sing DN | and ba A |
| 202 | | | | | | | | | | |

| | bp 450 400 350 | M | 1 | 2 | 3 | 4 | 5 | 6 | 7 | |
|-------------------|--------------------------------|-----------------------------------|-------------------------------------|----------------------------------|--------------------------------------|-----------------------------------|---|---------------------------------|----------------------------------|---------------------|
| | 250 | | | - | | | | | | ← 220 |
| | 150 100 | | | | | | | | | < <u>193</u> |
| | 50 | | | | | | | | | |
| 203 | | | | | | | | | | |
| 204 | | | | | | | | | | |
| 205 | Fig 2: I | PCR amp | lification | of intron | 4b/a poly | morphisn | n of eNOS | S gene | | |
| 206 | | | | | \sim | | | | | |
| 207 | | | | | | ́с` | | | | |
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| 215 | | | | | | | | | | |
| 216 | Discuss | ion | | | | | | | | |
| 217 218 219 | Obesity accumu are gen | and Over lated to erally co | rweight i the exten onsidered | s a med nt that it obese w | ical cond might hat then their | ition in y ad a nega body m | which exc trive effection ass index | ess body et on hea (BMI), | fat lth. <i>(1)</i> a meas | .People surement |

are generally considered obese when their body mass index (BMI), a measurementobtained by dividing a person's weight by the square of the person's height, is

over 30 kg/m², with the range 25–30 kg/m² defined as overweight. (1). Overweight and obesity are major risk factors for a number of chronic diseases, including diabetes,

- cardiovascular diseases and cancer. Once considered a problem only in high income
- countries, overweight and obesity are now dramatically on the rise in low- and middle-
- income countries, particularly in urban settings (1). Obesity is one of the
- leading preventable causes of death worldwide. (17),(18).
- 227 Growing evidence supports the association of diseases with
- 228 NOS3 haplotypes (combination of alleles in close proximity, within a DNA block). This
- approach may be more informative than the analysis of genetic polymorphisms one by
- one. (19). Haplotypes including the SNPs g.-786T>C and Glu298Asp, g- G894T and the
- 231 VNTR in intron 4 affected the susceptibility to hypertension, (20). And there is
- association between NOS3 and the susceptibility to obesity (16). And diabetes
 mellitus (21).
- The present study was carried to determine genotype distribution of the eNOS (T786 C,
- and 27bp) gene region in obesity cases and their allele frequencies, the present study aim
- mainly to investigate the association of T786C and 27 bp Polymorphism of eNOS gene with
- 237 Possibility to occur of obesity, the present study results showed that:
- 238 Homozygous mutated TT and homozygous mutated CC genotypes, mutant T and C
- allele of T786C polymorphism had significant frequency among cases of obesity
- compared with controls,
- But Homozygous mutated bb and homozygous mutated aa genotypes, mutant b and a
- allele of 27 bp polymorphism had NO significant frequency among cases of obesity
- compared with controls.
- In T786C polymorphism the result showed that homozygous mutated CC and
- homozygous mutated TT genotypes , mutant T and C allele of T786Cpolymorphism had
 significant frequency among cases of obesity compared with controls which considered
 as genetic risk factors for obesity
- 248
- Our results are in agreement with results of a study In BRAZIL (22) suggest that the
 biological variations associated with the T786C polymorphism predispose to MetS in
 both obese children and adolescents.
- Another study in japan also agree with our study which found that cases of obesity and controls was significantly in genotypes and allele frequencies of T786C POLY
- 255 MORPHISM OF e NOS gene (23).
- 256

- Our results are in agreement with a study in Brazil its data findings that the eNOS haplotype, T786C, 4b is associated with hypertension in obese children and adolescents (16).
- 260
- Another agreement with our study data obtained from study on north Indian population
- found association between T786C polymorphism and *Type 2 Diabetes with obese patient*

,But this study not effective enough to confirm that the relation between e NOS gene andobese patient in contrast type 2 diabetes (24).

- Another study in Ukraine (25) show that there is a Changes of lipid metabolism in patients with rheumatoid arthritis and, abdominal obesity and diabetes mellitus type 2
- 267 depending on the gene polymorphism T-786C of endothelial nitric oxide synthase.
- 268

Another study carried on Brazilian obese women From the local community in brazil
agree with our results finding that in obesity an impairment of NO endotheliumdependent relaxation is observed and it seems that the association of decreased NO.
production and increased NO.scavenging might compromise NO. bioavailability. And
they evaluated whether the T-786C polymorphism of eNOS may modulate the

- simvastatin effect on nitrite levels (26).
- 275 276
- On the other hand, our study disagrees with a study in Hungary (27) suggest that no
 significant differences were seen in the case of the eNOS promoter polymorphism
 (T786C) and the eNOS 4th intron 27-bp repeat polymorphism.
- 280
- The present study disagrees with the study in taxes (USA) *(28)* which failed to find significant association between the T-786C with Type 2 diabetes in Mexican Americans obese patients.
- 284
- Another disagreements with us is a study carried in Italy *(29)* where there is lack of association with eNOS variants and cardiovascular damage onset.in obese and non-obese.
- Another disagreement in Poland *(30)* was not found association with increased risk for both obese and non-obese subjects especially in patient with hypertension and nitric oxide synthase gene T786C polymorphism.
- In 27 bp polymorphism the results showed that homozygous mutated bb and homozygous
 mutated aa genotypes, mutant b and a allele of 27 bp polymorphism had NO significant
- frequency among cases of obesity compared with controls.
- A study in Hungary (27) agreed with our results which suggest that no significant differences were seen in the case of the eNOS 4th intron 27-bp repeat polymorphism and
- the eNOS promoter polymorphism.
- Our result are in agreement with a study in USA by *(28)* obtained that 27bp-VNTR appears to be a minor contributor to the variation in T2DM-related traits in obese
- 300 Mexican Americans.
- 301
- 302 On the other hand, a disagreement with us is a study carried between Tunisian people by
- 303 (31) this study provides the first evidence for the association 4a/b (27bp) variants with
- 304 body mass index and the risk of obesity in Tunisians. These polymorphisms did not

- 305 exhibit, however any significant association with both metabolic traits and vascular
- 306 function.
- 307
- A Brazilian study (16) also disagreed with our results where it suggest that the eNOS
- 309 gene polymorphism is associated with hypertension in obese children and adolescents.
- Further studies examining the possible interactions of eNOS haplotypes with
- environmental factors and other genetic markers might cause the development of obesity
- 312 and its complications are warranted.

313 CONCLUSIONS

- The C786T polymorphism of e NOS gene was found to be significantly associated with
- development of obesity .and T, C alleles, (CC and TT genotypes of C786T) might significantly
- considered genetic risk factor for development of obesity.

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