



# Role of *TERT* Gene in Relationship between Body Fat Percentage with the Modifiable Risk Factors of Cardiometabolic Disease in Minangkabau Premenopausal

Yuniar Lestari<sup>1</sup>, Delmi Sulastr<sup>2</sup>, Desmawati Desmawati<sup>2\*</sup>

<sup>1</sup>Department of Public Health, Medical Faculty, Andalas University, Padang, West Sumatera 25163, Indonesia; <sup>2</sup>Department of Nutrition, Medical Faculty, Andalas University, Padang, West Sumatera 25163, Indonesia

## Abstract

**Edited by:** Slavica Hristomanova-Mitkovska  
**Citation:** Lestari Y, Sulastr D, Desmawati D. Role of *TERT* Gene in Relationship between Body Fat Percentage with the Modifiable Risk Factors of Cardiometabolic Disease in Minangkabau Premenopausal. Open Access Maced J Med Sci. 2020 Apr 15; 8(B):414-417. https://doi.org/10.3889/oamjms.2020.3551

**Keywords:** Body fat percentage; Cardiometabolic; Lipid profile; Telomerase reverse transcriptase  
**\*Correspondence:** Desmawati Desmawati, Department of Nutrition, Medical Faculty, Andalas University, Perintis Kemerdekaan Street, 94<sup>th</sup> Padang, West Sumatera, 2516325163 Indonesia. Tel.: +62-852-74467797. E-mail: desmawati@med.unand.ac.id

**Received:** 19-Aug-2019

**Revised:** 20-Feb-2020

**Accepted:** 18-Mar-2020

**Copyright:** © 2020 Yuniar Lestari, Delmi Sulastr, Desmawati Desmawati

**Funding:** This research was financially supported by the Andalas University, Indonesia research project (Grant no 33/UN.16/HKRGB/LPPM/2018)

**Competing Interests:** The authors have declared that no competing interests exist

**Open Access:** This is an open-access article distributed under the terms of the Creative Commons Attribution-NonCommercial 4.0 International License (CC BY-NC 4.0)

**BACKGROUND:** Cardiometabolic disease risk is increasing in the premenopausal phase of women life. The *telomerase reverse transcriptase (TERT)* gene is a gene that plays a role in protecting telomeres from progressive shortening. Telomere shortening is associated with the incidence of cardiometabolic disease.

**AIM:** This study aims to elucidate the relationship body fat percentage with the modifiable risk factors of cardiometabolic disease in Minangkabau premenopausal people.

**METHODS:** A cross-sectional study was conducted in 111 Minangkabau premenopausal people aged 40–55 years old at Padang City in West Sumatera, Indonesia. Body fat percentage analyzed by bioelectrical impedance analyzer. Fasting blood glucose and lipid profile were examined from venous blood. Polymorphism of the *TERT rs2736098* gene was analyzed by a polymerase chain reaction. Data were analyzed using Spearman's rank correlation test with significant levels  $p < 0.05$ .

**RESULTS:** Average body fat percentage of subjects was  $36.23 \pm 6.9$ , fasting blood glucose was  $101.52 \pm 12.3$  mg/dL; total cholesterol was  $193.99 \pm 41.5$  mg/dL; triglyceride was  $113.76 \pm 37.9$  mg/dL; LDL-cholesterol was  $120.59 \pm 45.7$  mg/dL and HDL-cholesterol was  $53.52 \pm 15.6$  mg/dL. Body fat percentage has a significant correlation with triglyceride ( $r = 0.368$ ;  $p = 0.001$ ) and HDL-cholesterol levels ( $r = -0.307$ ;  $p = 0.006$ ) just in subject with polymorphism but does not have a significant correlation with another biomarker.

**CONCLUSION:** Body fat percentage correlates with triglyceride and HDL cholesterol in Minangkabau premenopausal women who have polymorphism of *TERT* gene.

## Introduction

Obesity is a multifactorial disease that developed from an unbalanced energy and energy expenditure associated with cardiometabolic disorder. Obesity occurs as an effect of the interaction between genetic, environmental, and physiological factors. Body fat percentage has emerged as an essential indicator of cardiometabolic risk than body mass index. Body fat percentage is related to the biochemical marker of cardiometabolic risk factors such as blood glucose and lipid profile [1]. Cardiometabolic risk is increased in premenopausal women. Sen's study in India stated that cardiometabolic risk increased with age in women, including the premenopausal phase [2]. Another study also said that cardiometabolic risk in the premenopausal phase increased after 10-year follow-up [3]. Decreasing of estrogen levels in premenopausal women can lead the increased cardiometabolic risk in women, due to change in a hormonal milieu with declining estrogen and alteration of its ratio with testosterone [4]. The

*telomerase reverse transcriptase (TERT) rs2736098* gene is a gene that plays a role in protecting telomeres from progressive shortening. Telomere shortening is associated with the incidence of cardiometabolic disease. The prevalence is a high in Europe and the world.

This study aims to elucidate the role of the *TERT rs2736098* gene in the relationship between body fat percentage with the modifiable risk factors of cardiometabolic disease in Minangkabau premenopausal people.

## Methods

A cross-sectional study was conducted in Minangkabau premenopausal people at Padang City in West Sumatera, Indonesia. A total of 112 Minangkabau Premenopausal people aged 40-55 years were selected randomly. The research was conducted

from July to December 2017. Respondents have to sign an informed consent form. All subjects are the Minangkabau ethnicity (all of grandparent and parent are the original Minangkabau ethnicity, and there was no cross marriage) and women in the premenopausal phase (menorrhagia and/or metrorrhagia more than 3 months, but a menorrhagia <12 months). Body fat percentage analyzed by bioelectrical impedance analyzer. Fasting blood glucose and lipid profile were examined from venous blood after fasting for 8 to 10 h.

Polymorphism of the *TERT* rs2736098 gene was analyzed by polymerase chain reaction (PCR) using a self-composed primer. We used *TERT* ex F primer (5'-GAACCATAGCGTCAGGGAG-3') and *TERT* ex R (5'-TCCCAAGCAGCTCCAGAAACA-3'). The PCR amplification products were analyzed using electrophoresis techniques in agarose 1.5% and electrophoresis at 120 volts for 50 minutes. The location of DNA contained in the gel can be observed by staining using red gel, then documented in the form of an electropherogram using the Doc Gel from Biorad. After it was confirmed that the DNA in the PCR product was good enough for sequencing, the PCR product was sent for sequencing.

This study has had ethical approval from the Ethical Committee Medical Faculty of Andalas University with registration number 279/KEP/FK/2017. Data were analyzed using non-parametric Spearman's rank correlation test with significant levels  $p < 0.05$ . All data analyzed using the SPSS software.

## Results

The frequency distribution of *TERT* gene polymorphisms in the research subject is shown in Table 1.

**Table 1: Frequency distribution of TERT rs2736098 gene polymorphisms in Minangkabau premenopausal people (n=111)**

Variables	N	(%)
Wild type/polymorphism (-)	33	29.73
Polymorphism (+)	78	70.27

TERT: Telomerase reverse transcriptase.

This research got a research subject with characteristic, as shown in Table 2.

**Table 2: Distribution of body fat percentage and biochemical marker of cardiometabolic disease based on polymorphism gen of TERT in Minangkabau premenopausal people (n=111)**

Variable	Mean			p-value
	Total	Wild type/ polymorphism (-)	Polymorphism (+)	
Body fat percentage	36.23 ± 6.9	35.23	36.64	0.330
Blood glucose	101.52 ± 12.3	101.75	101.43	0.903
Total cholesterol	193.99 ± 41.5	194.10	193.95	0.986
Triglyceride	113.76 ± 37.9	193.95	193.95	0.825
LDL-cholesterol	120.59 ± 45.7	120.81	120.49	0.975
HDL-cholesterol	53.52 ± 15.6	53.33	53.59	0.935

TERT: Telomerase reverse transcriptase, LDL: Low-density lipoprotein, HDL: High-density lipoprotein.

Average body fat percentage of subjects was  $36.23 \pm 6.9$ , blood glucose was  $101.52 \pm 12.3$

mg/dL, total cholesterol was  $193.99 \pm 41.5$  mg/dL, triglyceride was  $113.76 \pm 37.9$  mg/dL, LDL-cholesterol was  $120.59 \pm 45.7$  mg/dL, and HDL-cholesterol was  $53.52 \pm 15.6$  mg/dL.

From Table 3, we can see that body fat percentage has a significant correlation with triglyceride ( $r = 0.368$ ;  $p = 0.001$ ) and high-density lipoprotein (HDL)-cholesterol levels ( $r = -0.307$ ;  $p = 0.006$ ) just in subject with polymorphism but does not have a significant correlation with another biomarker.

**Table 3: Correlation between a biochemical marker of cardiometabolic disease with body fat percentage in Minangkabau premenopausal people**

Variable	Polymorphism gene of TERT	Body fat percentage	
		r	p-value
Blood glucose	Wild type	0.027	0.884
	Polymorphism (+)	0.240	0.033
	Total	0.176	0.065
Total cholesterol	Wild type	-0.162	0.933
	Polymorphism (+)	0.095	0.405
	Total	0.063	0.514
Triglyceride	Wild type	0.113	0.538
	Polymorphism (+)	0.368	0.001*
	Total	0.297	0.002*
LDL-cholesterol	Wild type	-0.193	0.290
	Polymorphism (+)	0.113	0.323
	Total	0.028	0.772
HDL-cholesterol	Wild type	-0.120	0.513
	Polymorphism (+)	-0.307	0.006*
	Total	-0.243	0.010*

\*Significant level at  $p < 0.05$ . TERT: Telomerase reverse transcriptase, LDL: Low-density lipoprotein, HDL: High-density lipoprotein.

## Discussion

Average body fat of subjects was  $36.23 \pm 6.9$ . It means that the subject of this study was obese. This result is similar to the results of Flint *et al.* (2014) study of 4065 women aged  $44 \pm 12.9$  years, who received a body fat percentage of  $36 \pm 8.4\%$  [5]. Dos Anjos *et al.* (2013) research on 352 women in Brazil showed an average percentage of body fat that increased at an older age. The percentage of body fat with the age range of 40–50 years is  $39.6 \pm 0.59\%$  and the age range of 50–60 years is  $40.3 \pm 0.65\%$  [6]. The results of other studies in China also obtained a body fat percentage of  $33.12 \pm 5.85\%$  in 154 women aged  $40.46 \pm 13.28$  years. The results in the study also relate to the previous research which states that Indonesia is a country with a high percentage of body fat, besides China, Japan, Ethiopia, Thailand, and Mexico [7].

Based on the U.S., the National Institutes of Health Criteria standards for percent body fat and the American Dietetic Association/Canadian Dietetic Association, body fat percentage values defined as obesity when body fat percentage in men  $\geq 25\%$  and women  $\geq 30\%$ . Differences in the percentage values are found in the study of Zhao *et al.* (2013), which define obesity with a number  $>25\%$  in men and  $>35\%$  who rely on the previous studies [6], [7], [8].

From the biochemical marker of cardiometabolic disease, such as blood glucose and lipid profile, its

average is a normal value. Dyslipidemia is strongly associated with an increase in cardiovascular disease. However, only a small number of research subjects were enlarged prediabetes and dyslipidemia. Dyslipidemia plays a role in the pathogenesis of the cardiovascular disease, namely, in the formation of atherosclerosis. The management of dyslipidemia must be carried out comprehensively through pharmacological and non-pharmacologic aspects. Pharmacological management is using drugs, especially statins, while non-pharmacological control through lifestyle modification, low-fat and high-fiber diets, increase physical activity and stress management [9].

The *TERT* gene is located on chromosome 5p15.33 in base pairs to 1,253,167 to 1,295,047, having 16 exons with a length of about 35 kb [10], [11]. A total of 69.9% of subjects had *TERT* gene polymorphism rs2736098, where 50.4% heterozygotes and 19.5% were homozygous recessive.

In this study, we can see that there is no significant difference in body fat percentage, blood glucose, and lipid profile between wild type and mutation groups. Although as more than half of subjects have polymorphism of *TERT rs2736098* gene. The number of *TERT rs2736098* polymorphisms is higher than single-nucleotide polymorphism research in the Asian population in China which ranges from 42% to 66% [12], [13], [14], [15], [16], [17], in India 58.28% [18] and in Korea 54.16% [19]. The percentage of polymorphism in this study was lower than that in other Asian populations in Iran, which amounted to 74.05% and the Caucasian population in Turkey (83.26%), and Austria (76.03%) in Serbia at 76.11% [20], [21], [22], [23].

Spearman's rank correlation test showed that body fat percentage has a significant correlation with triglyceride ( $r = 0.368$ ;  $p = 0.001$ ) and HDL-cholesterol levels ( $r = -0.307$ ;  $p = 0.006$ ) just in subject with polymorphism of *TERT* gene, but does not have a significant correlation with another biomarker. A research conducted on 234 Thai adults found that body fat percentage is a good predictor of insulin resistance, hyperglycemia, hypertriglyceridemia, and hyperleptinemia, and also atherogenic lipoprotein particles [24]. Likewise, research in China suggests that body fat percentage cut off for predicting the risk of cardiac abnormalities to be 24% and 33% in Chinese men and women, respectively [25].

In this study, a significant relationship between the percentage of body fat and biomarkers of cardiometabolic disease only occurred in subjects who had polymorphisms. It has been caused by the reduced expression of the *TERT* gene which causes a change in the function of the *TERT* gene itself. *TERT*'s function is to express telomerase which serves to keep telomeres from fast restriction. If there is a change in the function of the *TERT* gene that causes its expression to be disrupted, there will be a decrease

in telomerase production. This will cause telomere shortening to occur faster which disrupts chromosomal stability. Chromosomal stability disorders will eventually lead to various health problems, including an increased risk of cardiometabolic disease [12], [26], [22], [27]. The research about the other mechanisms of relationship the *TERT* gene polymorphisms on the risk of cardiometabolic disease in Minangkabau ethnic is needed.

## Conclusion

From this study, we can conclude that body fat percentage correlates with triglyceride and HDL cholesterol in Minangkabau premenopausal people who have *TERT* polymorphism. It is recommended for Minangkabau ethnicity who have *TERT rs2736098* gene polymorphism to maintain body weight and body fat percentage under normal conditions.

## Acknowledgment

We are grateful to Andalas University for funding this research project (Grant no 33/UN.16/HKRGB/LPPM/2018) and special thank for all respondents who participated in this study.

## References

1. Bennasar-Veny M, Lopez-Gonzalez AA, Tauler P, Cespedes ML, Vicente-Herrero T, Yañez A, *et al.* Body adiposity index and cardiovascular health risk factors in caucasians: A comparison with the body mass index and others. *PLoS One.* 2013;8(5):e63999. <https://doi.org/10.1371/journal.pone.0063999>  
PMid:23734182
2. Sen P, Das S, Hore S, Bhattacharjee S, Choudhuri D. Obesity and associated cardiometabolic risk among women from tripura-a Northeastern state of India. *J Midlife Health.* 2017;8(3):110-7. [https://doi.org/10.4103/jmh.jmh\\_116\\_15](https://doi.org/10.4103/jmh.jmh_116_15)  
PMid:28983157
3. Razmjou S, Abdunour J, Bastard JP, Fellahi S, Doucet É, Brochu M, *et al.* Body composition, cardiometabolic risk factors, physical activity, and inflammatory markers in premenopausal women after a 10-year follow-up: A MONET study. *Menopause.* 2018;25(1):89-97. <https://doi.org/10.1097/gme.0000000000000951>  
PMid:28763400
4. Choudhuri S, Aithal M, Choudhuri D. Screening for cardiometabolic risk profile in middle aged premenopausal Indian women. *J Cardiovasc Dis Res.* 2015;6(2):91-6. <https://doi.org/10.4103/jcd.116.15>

- doi.org/10.5530/jcdr.2015.2.8
5. Flint E, Cummins S, Sacker A. Association between active commuting, body fat, and body mass index: Population based, cross sectional study in the United Kingdom. *BMJ*. 2014;349:1-9. <https://doi.org/10.1136/bmj.g4887>  
PMid:25139861
  6. Dos Anjos LA, Teixeira FC, Wahrlich V, Vasconcellos MT, Going SB. Body fat percentage and body mass index in a probability sample of an adult Urban population in Brazil. *Cad Saude Publica*. 2013;29(1):73-81.
  7. Zhao D, Li Y, Zheng L, Yu K. Brief communication: Body mass index, body adiposity, and percent body fat in asians. *Am J Phys Anthropol*. 2013;152(2):294-9. <https://doi.org/10.1002/ajpa.22341>  
PMid:23996556
  8. Zeng Q, Dong SY, Sun XN, Xie J, Cui Y. Percent body fat is a better predictor risk factors than body mass index. *Braz J Med Biol Res*. 2012;45(7):591-600. <https://doi.org/10.1590/s0100-879x2012007500059>  
PMid:22510779
  9. Nordestgaard BG, Chapman J, Ray K, Boren J, Andreotti F, Watts GF, et al. Perhimpunan dokter spesialis kardiovaskular Indonesia. Pedoman tatalaksana dislipidemia. *J Kardiologi Indones*. 2013;34(4):265.
  10. TERT Telomerase Reverse Transcriptase Homo Sapiens (Human); 2017. Available from: <https://www.ncbi.nlm.nih.gov/gene/7015>. Last accessed on 2017 Apr 05].
  11. Cong YS, Wen J, Bacchetti S. The human telomerase catalytic subunit hTERT: Organization of the gene and characterization of the promoter. *Hum Mol Genet*. 1999;8(1):137-42. <https://doi.org/10.1093/hmg/8.1.137>  
PMid:9887342
  12. Zhang C, Tian YP, Wang Y, Guo FH, Qin JF, Ni H. hTERT rs2736098 genetic variants and susceptibility of hepatocellular carcinoma in the Chinese population: A case-control study. *Hepatobiliary Pancreat Dis Int*. 2013;12(1):74-9. [https://doi.org/10.1016/s1499-3872\(13\)60009-0](https://doi.org/10.1016/s1499-3872(13)60009-0)  
PMid:23392802
  13. Yuan X, Cheng G, Yu J, Zheng S, Sun C, Sun Q, et al. The TERT promoter mutation incidence is modified by germline TERT rs2736098 and rs2736100 polymorphisms in hepatocellular carcinoma. *Oncotarget*. 2017;8(14):23120-9. <https://doi.org/10.18632/oncotarget.15498>  
PMid:28416747
  14. Xiao X, He W. Genetic polymorphisms in the TERT-CLPTM1 L region and lung cancer susceptibility in Chinese males. *Oncol Lett*. 2017;14(2):1588-94. <https://doi.org/10.3892/ol.2017.6289>  
PMid:28789383
  15. Xing Y, Liu F, Li JF, Lin JC, Zhu GD, Li M, et al. Case-control study on impact of the telomerase reverse transcriptase gene polymorphism and additional single nucleotide polymorphism (SNP)-SNP interaction on non-small cell lung cancers risk in Chinese Han population. *J Clin Lab Anal*. 2016;30(6):1071-7. <https://doi.org/10.1002/jcla.21982>
  16. Baode L, Liu D, Zhenyuan Y, Weijin F, Chen J, Haoyuan L. Association between genetic polymorphism of TERT and CLK3 with susceptibility of bladder cancer. *J Pract Med*. 2016;32(11):1806-9.
  17. Ma Z, Hu Q, Chen Z, Tao S, Macnamara L, Kim ST, et al. Systematic evaluation of bladder cancer risk-associated single-nucleotide polymorphisms in a Chinese population. *Mol Carcinog*. 2013;52(11):916-21. <https://doi.org/10.1002/mc.21932>  
PMid:22711262
  18. Singh V, Jaiswal PK, Mittal RD. Replicative study of GWAS TP63C/T, TERTC/T, and SLC14A1C/T with susceptibility to bladder cancer in North Indians. *Urol Oncol*. 2014;32(8):1209-14. <https://doi.org/10.1016/j.urolonc.2014.05.013>  
PMid:25218484
  19. Yoo SS, Do SK, Choi JE, Lee SY, Lee J, Cha SI, et al. TERT polymorphism rs2853669 influences on lung cancer risk in the Korean population. *J Korean Med Sci*. 2015;30(10):1423-8. <https://doi.org/10.3346/jkms.2015.30.10.1423>  
PMid:26425038
  20. Jannuzzi AT, Karaman E, Oztas E, Yanar HT, Özhan G. Telomerase reverse transcriptase (TERT) gene variations and susceptibility of colorectal cancer. *Genet Test Mol Biomarkers*. 2015;19(12):692-7. <https://doi.org/10.1089/gtmb.2015.0150>  
PMid:26501986
  21. Hashemi M, Amininia S, Ebrahimi M, Hashemi SM, Taheri M, Ghavami S. Association between hTERT polymorphisms and the risk of breast cancer in a sample of Southeast Iranian population. *BMC Res Notes*. 2014;7:895. <https://doi.org/10.1186/1756-0500-7-895>  
PMid:25491902
  22. de Martino M, Taus C, Lucca I, Hofbauer SL, Haitel A, Shariat SF, et al. Association of human telomerase reverse transcriptase gene polymorphisms, serum levels, and telomere length with renal cell carcinoma risk and pathology. *Mol Carcinog*. 2016;55(10):1458-66. <https://doi.org/10.1002/mc.22388>  
PMid:26294352
  23. Carkic J, Nikolic N, Radojevic-Skodric S, Kuzmanovic-Pficer J, Brajovic G, Antunovic M, et al. The role of TERT-CLPTM1 L SNPs, hTERT expression and telomere length in the pathogenesis of oral squamous cell carcinoma. *J Oral Sci*. 2016;58(4):449-58. <https://doi.org/10.2334/josnusd.16-0108>  
PMid:28025427
  24. Vanavanan S, Srisawasdi P, Rochanawutanon M, Kumproa N, Kruthkul K, Kroll MH. Performance of body mass index and percentage of body fat in predicting cardiometabolic risk factors in Thai adults. *Diabetes Metab Syndr Obes*. 2018;11:241-53. <https://doi.org/10.2147/dmso.s167294>  
PMid:29910627
  25. Jia A, Xu S, Ming J, Xing Y, Guo J, Zhao M, et al. Body fat percentage cutoffs for risk of cardiometabolic abnormalities in the Chinese adult population: A nationwide study. *Eur J Clin Nutr*. 2018;72(5):728-35. <https://doi.org/10.1038/s41430-018-0107-0>  
PMid:29410481
  26. Soerensen M, Thinggaard M, Nygaard M, Dato S, Tan Q, Hjelmborg J, et al. Genetic variation in TERT and TERC and human leukocyte telomere length and longevity: A cross-sectional and longitudinal analysis. *Aging Cell*. 2012;11(2):223-7. <https://doi.org/10.1111/j.1474-9726.2011.00775.x>  
PMid:22136229
  27. Atzmon G, Cho M, Cawthon RM, Budagov T, Katz M, Yang X, et al. Evolution in health and medicine Sackler colloquium: Genetic variation in human telomerase is associated with telomere length in Ashkenazi centenarians. *Proc Natl Acad Sci U S A*. 2010;107 Suppl 1:1710-7. <https://doi.org/10.1073/pnas.0906191106>  
PMid:19915151