

## Worldwide trends in scientific publications on association of gut microbiota with obesity

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### ABSTRACT

**Objective(s):** Recent evidence has shown underlying roles of gut dysbiosis and metabolic endotoxemia in obesity and its complications. Despite the large number of experimental and clinical researches performed on gut microbiota and obesity, no bibliometrics' study has been conducted so far. We aimed to assess the trend of global scientific publications in the field of gut microbiota and obesity. **Materials and Methods:** The bibliometrics' data from January 2000 to April 2017 were retrieved based on Scopus database. The analysis of the publication year, main source, citation, subject area, co-authorship network, and geographical distribution were carried out, accordingly. The data were analyzed using the Scopus analysis tools, SPSS version 15 and Visualizing Scientific Landscapes (VOS) viewer version 1.6.5.

**Results:** Out of 4384 documents that were identified, the United States published the highest number (28.2%), followed by China and United Kingdom. The number of publications showed an increasing trend over the years of which the most productive year was 2016. The leading subject area was medicine. Most of published scientific documents were original articles and the top source was "PLOS One". The documents were cited totally 153576 times with average citations per article as 35.03, and h-index of 159. Top author in the co-authorship network assessment was "Wang J." from China. **Conclusion:** This study could provide practical sources to researchers to find highly cited studies. Moreover, the study could pave the way for researchers to be engaged in studies which potentially lead to more publication in the field.

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### Introduction

The prevalence of obesity, as one of the most important health problems influenced by environmental and genetic factors, has been increasing worldwide, both in developed and developing countries (1-3). Although obesity increases the risk of developing serious health conditions, there is no successful method for its control up to now (4). Multiple socio-economic, behavioral, and biological determinants have been known as influencing factors in the prevention and management of obesity. Current researches have focused on the basic aspects of obesity and the emerging science for its control (5). Microbiota, as one of the hot topics in the research field of obesity pathology and management has strong supports either from national institutes or pharmaceutical companies. Gut microbiota that is the microbial community inhabiting the intestine plays as an endocrine organ and its imbalance is related to numerous disorders (6). The gut microbiota might affect energy balance through several mechanisms including the fermentation of indigestible dietary compounds, secretion of complex biochemical compositions,

impacting on intestinal permeability and interference with metabolic pathways (6-8).

The rationale for ongoing efforts in investigating the relationship of gut microbiota and obesity is the development of monitoring tools and the discovery of new treatment and preventive strategies (9, 10). Despite the large number of experimental and clinical researches performed on gut microbiota and obesity, no bibliometrics' study has been yet conducted. The bibliometric analysis, which is an effective method to evaluate the current research performance, could provide practical information for basic researchers, care givers and health authorities engaged in microbiota research to conduct better studies and accelerate therapeutic innovations (11, 12).

The aim of the present study is to assess the trend of global scientific publications in field of gut microbiota and obesity with a focus on aforementioned points.

### Materials and Methods

#### Data source

A descriptive bibliometric study of scholarly products

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covering the role of gut microbiota in obesity was conducted in Scopus database available at <http://www.scopus.com>. An all-embracing coverage of the Scopus database in health and biomedical fields as well as its high coverage of citation reports and its easy access to various valid analytical tools made it a suitable choice for our study (13).

### Search strategies

Articles were searched in Scopus database across title, abstract and keywords using the following queries: (gut AND (microbiot\* OR bacteria OR microbiome) OR prebiotic OR probiotic OR (intestin\* AND (flora OR bacteria OR microbiot\* OR microbiome)) OR antibiotic OR dysbiosis) AND ((obes\* OR adipos\* OR anthropometric OR overweight OR weight OR (body AND (size OR composition OR mass)) OR BMI OR (fat AND mass)). The above key words were chosen from the list of Medical Subject Headings (MeSH) provided by the National Library of Medicine (NLM) /PubMed.

All relevant publications in the field of gut microbiota and obesity published before April 2017 were included in the analysis with no language limitation. Studies in the subject areas of veterinary, poultry science, soil biology, dentistry, engineering, material science, humanity and art science, computer and mathematics were excluded.

### Data analysis

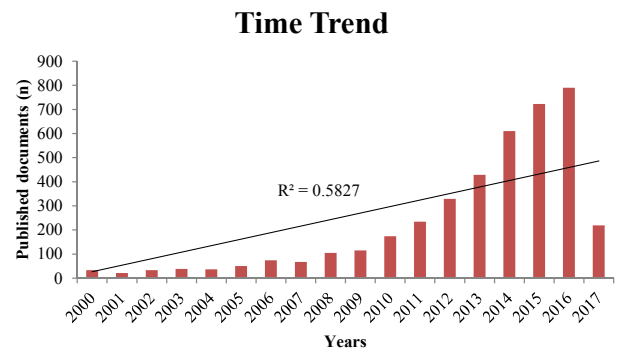
The collected data were publication year, main source (journal) with its impact factor, author's name and affiliation, geographical distribution, document's type and language, subject area, and document's citations which were retrieved and analyzed using the Scopus database. The impact factors (IF) of the journals that were retrieved from the Journal Citation Report (JCR) available at <https://jcr.incites.thomsonreuters.com> was used to compare the relative importance of journals within a specific field (14, 15). Moreover, the h-index was utilized to estimate the quality and impact of research documents. H-index could be determined based on the number of citations that each document received in other publications (16).

Analysis of the extracted data was performed through 'Analyze search results' function of Scopus web databases. SPSS software, version 15 (SPSS Inc., Chicago, IL, US) was used to assess the correlation between the number of published papers and the year of publication. By usage VOS viewer (Visualizing Scientific Landscape) software, version 1.6.5 was assessed scientific collaboration between authors with published papers in the field (17). The co-authorship mapping and clustering information could be provided by VOS viewer, available at [www.vosviewer.com](http://www.vosviewer.com).

## Results

### Time trend in publications

Overall, 4384 documents, published from January 2000 to April 2017 were analyzed. The time-trend distribution of publications is shown in Figure 1. Publications in the field of the gut microbiota and obesity have increased significantly over the recent years. The highest number of documents has published



**Figure 1.** Time-trend distribution of published documents in field of gut microbiota and obesity

in 2016 (790 documents, 18%). The overall association between the number of published documents and the year of publication is 0.806 with  $P$ -value  $<0.001$ . The R-squared value of 0.583 suggests a steady and significant increase over the same period. Details are shown in Figure 1.

### Subject area, type, language, and main source of documents

Among the subject area of the documents, top five are medicine (66.3%), biochemistry, genetics and molecular biology (31.7%), agricultural and biological sciences (22.6%), immunology and microbiology (17.4%), and nursing (12.7%).

Most of our analyzed documents are original articles (3039 documents, 69.3%) that followed by 969 review articles (22.1%), and 102 conference papers (2.3%). The remaining papers are letters, editorials, short surveys and notes (184 documents, 4.2%). Top ten languages of scientific publications are English, French, Chinese, German, Spanish, Japanese, Russian, Polish, Czech and Portuguese, respectively which most of them published in English language (96.1%).

Table 1 shows the characteristics of the main journals (sources). The "Plos One" with 186 documents has been scored as the first rank followed by "British Journal of Nutrition", "Gut Microbes", "Scientific Reports", "Journal of Nutrition", "World Journal of Gastroenterology", "Applied and Environmental Microbiology", "Proceedings of the National Academy of Sciences of the United States of America", "Nature" and "Nutrients", respectively. The impact factor of all top 10 journals, except for "Gut Microbes" and "Applied and Environmental Microbiology" is remarkable (18).

### Geographical distribution

Considering the geographical distribution of published articles, the United States is the most productive country with 1237 documents (28.2%). China is ranked as the second (9%) and the United Kingdom, France, Germany, Italy, Canada, Spain, Sweden and Belgium are ranked subsequently as the top 10 countries with a high number of published documents (3554 documents/ 81.07% of total published papers).

### Authors and Institutes' characteristics of published documents

We found 159 authors as the first-author with

**Table 1.** Characteristics of top 10 sources for the published documents in field of gut microbiota and obesity

| Title of journal  | IF (SJR)        | Documents (number/percent) | Total citations to document | Citation per document | Citation to highest cited document |
|---|-----------------|----------------------------|-----------------------------|-----------------------|------------------------------------|
| Plos One  | 3.057 (1.395)   | 186 (4.24)                 | 5399                        | 28.87                 | 624                                |
| British Journal of Nutrition  | 3.311 (1.587)   | 95 (2.17)                  | 3879                        | 40.83                 | 634                                |
| Gut Microbes  | ---- (1.473)    | 60 (1.37)                  | 849                         | 14.15                 | 135                                |
| Scientific Reports  | 5.228 (2.073)   | 53 (1.21)                  | 283                         | 5.34                  | 42                                 |
| Journal of Nutrition  | 3.740 (2.040)   | 52 (1.19)                  | 4327                        | 83.21                 | 1964                               |
| World Journal of Gastroenterology   | 2.787 (1.076)   | 45 (1.03)                  | 872                         | 19.38                 | 101                                |
| Applied and Environmental Microbiology  | ---- (1.891)    | 42 (0.96)                  | 2688                        | 64                    | 431                                |
| Proceedings of the National Academy of Sciences of the United States of America | 9.423 (6.883)   | 38 (0.87)                  | 9899                        | 260.5                 | 2007                               |
| Nature  | 38.138 (21.936) | 36 (0.82)                  | 21701                       | 602.81                | 3332                               |
| Nutrients   | 3.759 (2.275)   | 36 (0.82)                  | 749                         | 20.81                 | 219                                |

Legend: IF: Impact factor, SJR: SCImago journal rank

**Table 2.** Characteristics of top 10 institutes' affiliation for published documents in field of gut microbiota and obesity

| Rank | Name of institute  | Documents (number, percent) | Country        |
|------|--|-----------------------------|----------------|
| 1    | Universite Catholique de Louvain                                     | 101 (2.3%)                  | Belgium        |
| 2    | Kobenhavns Universitet   | 83 (1.89%)                  | Denmark        |
| 3    | Inserm   | 64 (1.46%)                  | France         |
| 4    | Goteborgs Universitet  | 64 (1.46%)                  | Sweden         |
| 5    | University College Cork  | 62 (1.41%)                  | Ireland        |
| 6    | National University of Ireland, Cork, Alimentary Pharmabiotic Centre | 59 (1.34%)                  | Ireland        |
| 7    | University of Reading  | 57 (1.3%)                   | United Kingdom |
| 8    | Imperial College London  | 57 (1.3%)                   | United Kingdom |
| 9    | Wageningen University and Research Centre                            | 54 (1.2%)                   | Netherlands    |
| 10   | VA Medical Center  | 51 (1.16%)                  | USA            |

published articles in the field of gut microbiota and obesity. Top 10 authors who have the most number of publications in this field are Cani PD with 86 documents followed by Delzenne NM (65 documents), Backhed F. (51 documents), Gordon JI and Shanahan F (each one with 34 documents), Neyrinck AM, Gibson GR, Raoult D, Everard A, and Cryan JF with 33, 31, 30, 28, and 26 documents, respectively. Out of above top 10 authors, four authors are from Belgium, two authors from Ireland, and others are from Sweden, USA, United Kingdom, and

Canada.

The first ranked institute for publications in the field of gut microbiota and obesity is “Universite Catholique de Louvain”, followed by “Kobenhavns Universitet” having the second rank and “Inserm” and “Goteborgs Universitet”, both owning the third rank. The details of findings are shown in Table 2.

**Citation number**

The trend of citation number is shown in Figure 2.



**Figure 2.** Chart of citation of published documents in our study

**Table 3.** Characteristics of top 10 highly cited published documents in field of gut microbiota and obesity

| Rank | Name of article  | Number of citations | Year | Type of article  | Country                |               | Title of journal  | IF (SJR) |
|------|--|---------------------|------|------------------|------------------------|---------------|---|----------|
|      |  |                     |      |                  | Corresponding author   | Co- Authors   |   |          |
| 1    | An obesity-associated gut microbiome with increased capacity for energy harvest  | 3312                | 2006 | Original article | USA                    | USA           | Nature  | 38.138   |
| 2    | A human gut microbial gene catalogue established by metagenomic sequencing   | 3098                | 2010 | Original article | China, France, Denmark | International | Nature  | 38.138   |
| 3    | A core gut microbiome in obese and lean twins  | 2596                | 2009 | Original article | USA                    | USA           | Nature  | 38.138   |
| 4    | Microbial ecology: Human gut microbes associated with obesity  | 2555                | 2006 | Original article | USA                    | USA           | Nature  | 38.138   |
| 5    | The gut microbiota as an environmental factor that regulates fat storage   | 1993                | 2004 | Original article | USA                    | International | Proceedings of the National Academy of Sciences of the United States of America | 9.423    |
| 6    | Obesity alters gut microbial ecology   | 1925                | 2005 | Original article | USA                    | USA           | Proceedings of the National Academy of Sciences of the United States of America | 9.423    |
| 7    | Host-bacterial mutualism in the human intestine  | 1916                | 2005 | Review           | USA                    | USA           | Science   | 34.661   |
| 8    | Metabolic endotoxemia initiates obesity and insulin resistance   | 1655                | 2007 | Original article | France                 | International | Diabetes  | 8.095    |
| 9    | Enterotypes of the human gut microbiome  | 1373                | 2011 | Original article | Germany, France        | International | Nature  | 38.138   |
| 10   | Changes in gut microbiota control metabolic endotoxemia-induced inflammation in high-fat diet-induced obesity and diabetes in mice | 1242                | 2008 | Original article | France                 | International | Diabetes  | 8.095    |

IF: Impact factor, SJR: SCImago journal rank.

Total citation number of 4384 documents is 153576 times at the time of data analysis (until the fourth April 2017). Therefore, the average number of citations per article is 35.03. Citations in the field of the gut microbiota and obesity have increased over the time with the highest number of citations being 33911 times in 2016. 16.95% of all documents (743 papers) don't receive any citation until the date of analysis.

The h-index for analyzed documents is 159 indicating that 159 documents have cited at least 159 times. Characteristics of top 10 highly cited published documents in the field of gut microbiota and obesity are summarized in Table 3.

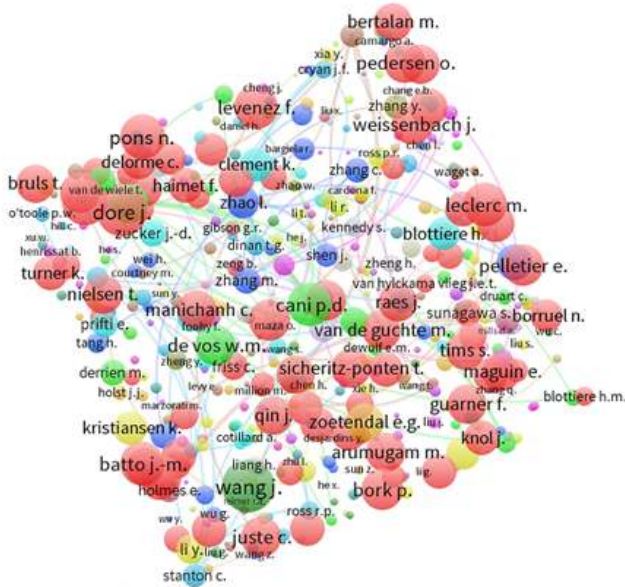
The citation number of these top 10 articles ranged from 3312 to 1242 times. The highest cited paper is an original article entitled "An obesity-associated gut microbiome with increased capacity for energy harvest", been cited 3312 times. The second ranked article entitled "A human gut microbial gene catalogue established by metagenomic sequencing" has 3098 citations. The third rank article is "A core gut microbiome in obese and lean twins" with 2596 citation numbers at the time of our

study.

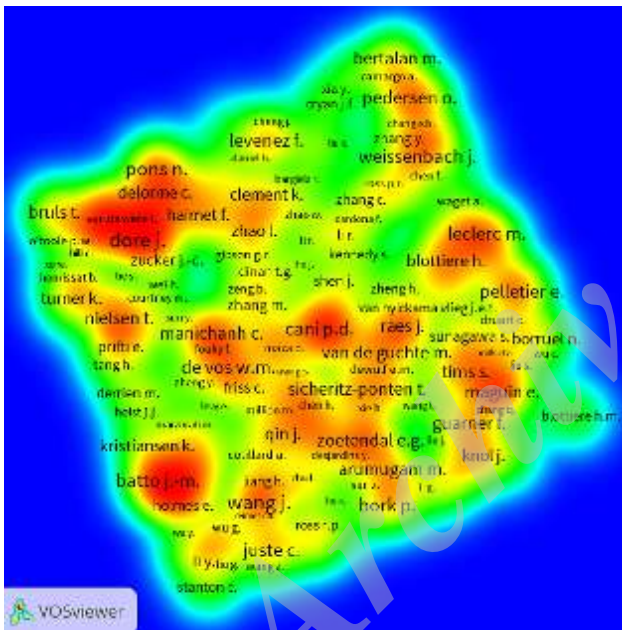
As shown in Table 3, of top 10 highly cited documents, 9 papers are original articles and 1 paper is a review, entitled "Host-bacterial mutualism in the human intestine". Of these top 10 articles, 5 documents are published in Nature, 2 documents in each of Diabetes and Proceedings of the National Academy of Sciences of the United States of America and 1 document in Science. The Washington University School of Medicine (USA) with 5 articles from 10 highly cited documents is ranked as the first.

#### **Co-authorship network mapping**

In this part of the study, VOS viewer software is used to assess the co-authorship network of authors. In order to map this network, the minimum number of documents published by an author is considered as 2 papers. Out of 16351 authors, 3561 authors meet this threshold. After excluding 101 authors with no co-authorships, the remaining authors are analyzed. The labels and density views of co-authorship network of authors in field of gut microbiota and obesity are shown in Figure 3 and Figure



**Figure 3.** Map (label view) of co-authorship network of the authors published scientific documents in field of gut microbiota and obesity



**Figure 4.** Density view of co-authorship network of the authors published scientific papers in field of gut microbiota and obesity

4, respectively. Top 10 authors in the field based on co-authorship networks of authors are “Wang J” (444 co-authorships) followed by “Dore j”, “Ehrlich S”, “Pons N”, “Cani PD”, “Levezon F”, “Batto JM”, “Leclerc M”, “Bork P” and “Pedersen O”, each one with 411, 348, 339, 329, 327, 323, 323, 297, and 297 co-authorships, respectively (Figure 3). According to Figure 4, the highest density in the network is attributed to “Wang J” from China.

**Discussion**

In the present study, the trend of global scientific publications in the field of the gut microbiota and obesity was assessed. In this view, 4384 documents were extracted from the Scopus database from January 2000 till April 2017. Although, the documents that

were indexed outside of the Scopus database were not included in this analysis, it should be mentioned that Scopus search engine covers an acceptable number of publishers and peer-reviewed journals in different fields (19). In addition, in a study that compared the citations’ reports in three essential web databases including Web of Science, Scopus, and Google Scholar, a high coverage of citation reports was shown for Scopus (20). Thus, this study could give a reliable view about the characteristics of research in the field of gut microbiota and obesity.

The number of publications in this field showed an increasing trend over the years marking 2016 as the most productive year. This progressive increase indicated an augmented interest in this topic to promote the development of novel therapeutic strategies like the application of probiotics and prebiotics for human health. It could be assumed that major financial investments including Human Microbiome Project (HMP) and Metagenomics of the Human Intestinal Tract (MetaHIT) had an impact on research output in the field of gut microbiota and obesity in the United States and Europe. The HMP was established by the United States National Institutes of Health (NIH) as a conceptual extension of the Human Genome Project in 2007 with the aim of identifying and characterizing the human microbiota in health and disease states (21-23). MetaHIT was known as a European Union Project that launched by the European Commission to explore the associations between microbial genes and human phenotypes, especially targeted obesity and inflammatory bowel diseases. This project was implemented in order to compare the gut microbiota of healthy and sick individuals, follow up patients in clinical remission, conduct nutritional interventions, and compare gut microbiota of patients responding or not to a drug (24, 25). Furthermore, the advent of next generation sequencing (NGS) technology in microbiota analysis allowed more sophisticated microbiota studies over the previous culture-based and molecular methods and led to significant growth in the number of literature in the field. To put it in another way, one reason for the upward trend in the number of articles could be attributed to the emergence of this technology and the increasing interest of scientists in this field (25-28).

Generally, a relatively small number of countries were responsible for the majority of research on gut microbiota and obesity. The United States as the leading country published the most articles and China ranked the second. It should be noted that although the United States was published nearly one third of articles in this field, “Universite Catholique de Louvain” from Belgium was found as the top institute focused on this topic. This could be well explained considering the fact that several research entities in the USA are working in this field; however, the majority of research in this area in Belgium is conducted in Universite Catholique de Louvain and Belgium have good collaboration with other countries.

The majority of published documents in this field were original articles (69.3%). This observation could be related to challenge of the global epidemic of obesity for the public health services and increasing efforts to combat this problem. The top subject area of published documents was medicine that followed by biochemistry,

genetics and molecular biology studies refer to the interests not only for better understanding the signaling pathways on the association of gut microbiota with obesity, but also on the role of gut microbiota, as a new target therapy for obesity. Dietary manipulation of the gut microbiota such as prebiotics and probiotics could affect the gut microbiota composition. However, responses of obese subjects to these manipulations are varied. Analysis of the microbiome in obese individuals could be added to future routine personalized medicine, which leads to the development of novel therapeutic strategies (29).

The majority of articles were published in high impact and high quality international journals indicating the worldwide attention to this topic. Among the top 10 sources, "Nature" with an IF of 38.138 was shown to present the importance of new findings about gut microbiota. The "PLoS One" with the highest number of papers in this field was indexed in many of the important citation databases such as PubMed, MEDLINE, Web of Science, and Scopus.

Citations in the field of gut microbiota and obesity had increased over the time with the highest number of citations in 2016, indicating the progressive appealing of the medical professionals and researchers to this topic. The h-index of 159 for analyzed documents was considered as the good quality of the articles confirming the accepted visibility of the article. The 10 top-cited articles were published from 2004 to 2011. The United States with 5 articles from 10 highly cited documents was ranked as the first. It should be noted that the other half of these studies were multinational collaboration. It could show the multidisciplinary efforts in this field. The most cited article (3312 times) was published by Turnbaugh *et al* from the Washington University in 2006, entitled "An obesity-associated gut microbiome with increased capacity for energy harvest" (30). This study published in "Nature" with an IF of 38.138 and SJR of 21.936.

It was shown that co-authorship network is the best bibliometric indicator to demonstrate different co-authorship patterns (31). Co-authorship networks could show research collaborations between different academic institutions. Top author in the co-authorship network assessment was "Wang J" from China. Analysis of the label and density views of co-authorship network of authors that showed 6 authors of the top 10 authors were French, showing that French researchers had good international collaboration in the field of obesity and gut microbiota.

Our study had some strengths and limitations. First, we focused on specific subjects on scholarly products in the field of gut microbiota and obesity. Second, we used Scopus web database that has a high multidisciplinary coverage in science. Third, we assessed the worldwide trends of scientific publications concomitant draw international collaboration and co-authorship network of authors and institutions. The potential limitation of this study was related to the database used to retrieve relevant documents. Included database could not represent all published scientific journals; in fact, some highly cited scientific publications may appear in journals other than those indexed in Scopus. In addition,

given the fact that the universities consisted of several research centers, unfortunately, it was not possible for us to distinguish these centers in the present study.

## Conclusion

The results of the present scientometric study showed an ascending trend in global publications on the relationship between gut microbiota and obesity over the years. The trend was in line with the development of novel diagnostic tools and helped identifying approaches to the modulation of the individual's microbiota to promote health and manage obesity. Evaluation of studies in this field was mentioned necessity to perform analysis of gut microbiota of different countries' population regarding the effect of geography, race and environmental factors on gut microbiota. This bibliometric analysis could be used to explore research productivity over time, map research needs in this field, and identify funding research centers to produce high-quality documents. It could also highlight the highest contributions of the academic institutions on the subject matter. Overall, this study could provide practical sources for researchers to find highly cited studies. Additionally, the study could pave the way for researchers to be engaged in studies potentially lead to more publications in this field.

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## Conflict of Interest

The authors declare that they have no conflict of interest.

## References

1. Jafari-Adli S, Jouyandeh Z, Qorbani M, Soroush A, Larijani B, Hasani-Ranjbar S. Prevalence of obesity and overweight in adults and children in Iran; a systematic review. *J Diabetes Metab Disord* 2014;13:121.
2. Tabatabaei-Malazy O, Khodaeian M, M. Amoli M. Association between genetic variants and obesity in Iranian population: review article. *Iran J Public Health* 2014;43:71-82.
3. Nguyen DM, El-Serag HB. The epidemiology of obesity. *Gastroenterol Clin North Am* 2010;39:1-7.
4. Segula D. Complications of obesity in adults: a short review of the literature. *Malawi Med J* 2014;26:20-24.
5. Palou A, Bonet ML. Challenges in obesity research. *Nutr Hosp* 2013;28 Suppl 5:144-153.
6. Ejtahed HS, Soroush AR, Angoorani P, Larijani B, Hasani-Ranjbar S. Gut Microbiota as a target in the pathogenesis of metabolic disorders: a new approach to novel therapeutic agents. *Horm Metab Res* 2016;48:349-358.
7. Homayouni-Rad A, Soroush AR, Khalili L, Norouzi-Panahi L, Kasaie Z, Ejtahed HS. Diabetes Management by Probiotics: Current knowledge and future perspective. *Int J Vitam Nutr Res* 2017:1-13.
8. Kvit KB, Kharchenko NV. Gut microbiota changes as a risk factor for obesity. *Wiad Lek* 2017;70:231-235.
9. Dore J, Multon MC, Behier JM. The human gut microbiome as source of innovation for health: Which physiological and therapeutic outcomes could we expect? *Therapie* 2017;72:21-38.
10. Dahiya DK, Renuka, Puniya M, Shandilya UK, Dhewa T, Kumar N, *et al*. Gut microbiota modulation and its relationship with obesity using prebiotic fibers and probiotics: A review. *Front Microbiol* 2017;8:563.

11. Tabatabaei-Malazy O, Atlasi R, Larijani B, Abdollahi M. Trends in publication on evidence-based antioxidative herbal medicines in management of diabetic nephropathy. *J Diabetes Metab Disord* 2015;15:1-8.
12. Tabatabaei-Malazy O, Ramezani A, Atlasi R, Larijani B, Abdollahi M. Scientometric study of academic publications on antioxidative herbal medicines in type 2 diabetes mellitus. *J Diabetes Metab Disord* 2016;15:48.
13. Bar-Ilan J. Citations to the "Introduction to informetrics" indexed by WOS, Scopus and Google Scholar. *Scientometrics* 2010;82:495-506.
14. Garfield E. The history and meaning of the journal impact factor. *JAMA* 2006;295:90-93.
15. Saha S, Saint S, Christakis DA. Impact factor: a valid measure of journal quality? *J Med Libr Assoc* 2003;91:42-46.
16. Baldock C, Ma R, Orton CG. Point/counterpoint. The h index is the best measure of a scientist's research productivity. *Med Phys* 2009;36:1043-1045.
17. Van Eck NJ, Waltman L. Text mining and visualization using VOSviewer. *Int Soc Scientometr Informetr Newslett* 2011;7:50-54.
18. Falagas ME, Kouranos VD, Arencibia-Jorge R, Karageorgopoulos DE. Comparison of SCImago journal rank indicator with journal impact factor. *FASEB J* 2008;22:2623-2628.
19. Philippe Mongeon, Ade`le Paul-Hus. The journal coverage of Web of Science and Scopus: a comparative analysis. *Scientometrics* 2016;106:213-228.
20. Kulkarni AV, Aziz B, Shams I, Busse JW. Comparisons of citations in Web of Science, Scopus, and Google Scholar for articles published in general medical journals. *JAMA* 2009;302:1092-1096.
21. Proctor LM. The human microbiome project in 2011 and beyond. *Cell Host Microbe* 2011;10:287-291.
22. Human Microbiome Project Consortium. A framework for human microbiome research. *Nature* 2012;486:215-221.
23. Robles-Alonso V, Guarner F. From basic to applied research: lessons from the human microbiome projects. *J Clin Gastroenterol* 2014;48 Suppl 1:S3-4.
24. Blanco-Miguez A, Gutierrez-Jacome A, Fdez-Riverola F, Lourenco A, Sanchez B. MAHMI database: a comprehensive MetaHit-based resource for the study of the mechanism of action of the human microbiota. *Database* 2017;2017.
25. Dusko Ehrlich S. [Metagenomics of the intestinal microbiota: potential applications]. *Gastroenterol Clin Biol* 2010;34 Suppl 1:S23-S28.
26. Finotello F, Mastrorilli E, Di Camillo B. Measuring the diversity of the human microbiota with targeted next-generation sequencing. *Brief Bioinform* 2018;19:679-692.
27. Ji B, Nielsen J. From next-generation sequencing to systematic modeling of the gut microbiome. *Front Genet* 2015;6:219-227.
28. Rogers GB, Bruce KD. Next-generation sequencing in the analysis of human microbiota: essential considerations for clinical application. *Mol Diagn Ther* 2010;14:343-350.
29. Ejtahed HS, Hasani-Ranjbar S, Larijani B. Human Microbiome as an approach to personalized medicine. *Altern Ther Health Med* 2017; 23:8-9.
30. Turnbaugh PJ, Ley RE, Mahowald MA, Magrini V, Mardis ER, Gordon JL. An obesity-associated gut microbiome with increased capacity for energy harvest. *Nature* 2006;444:1027-1031.
31. Newman ME. The structure of scientific collaboration networks. *Proc Natl Acad Sci U S A* 2001;98:404-409.