**Meta-Analysis/Systematic Review**



**Title: concisely convey the main topic(s) of the research**

[***Suggestions: No more than 16 words.*** ***No abbreviations except for standardized ones e.g., DNA, RNA, etc.***]

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**How to Use This Template**

***This template shows the manuscript structure that can be used in a meta-analysis/systematic review: Abstract, Keywords, Introduction, Methods, Results, Discussion, Declarations and References. Please note that each part has a corresponding style, that authors should follow. “Please note that the fonts in gray show writing requirements”. For any questions, you may contact the [editorial office](mailto:editorialoffice@mtodjournal.net).***

**Abstract**

[***Suggestions: No more than 250 words. No citations. Define abbreviations at their first mention.***]

**Aim:** The purpose of this study, i.e., the reason why authors write this manuscript.

**Methods:** The detailed methods applied to this study, for example, authors may describe how they compare, summarize or analysis data.

**Results:** The main results of this study, including conclusive description, analysis, and comparison with other related research results, *etc*.

**Conclusion:** The conclusion of this study. We suggest that authors may highlight its significance, emphasize the value of this study and state expectation on future studies that may need to be carried out.

**Keywords:** Morbidity, *Foxp3,* transforming growth factor-β1, *Helicobacter pylori,* DNA, contrast, meta-analysis, Notch

***[Please suggest 3-8 keywords which can be used for describing the content of the manuscript and will enable the full text of the manuscript to be searchable online]***

**INTRODUCTION**

The introduction is a beginning section of a manuscript which states the purpose of the study, overviews or summarizes previous findings and progress related to this study, and indicates its significance in this research field. It is generally followed by the body and discussion.

**METHODS**

In this section, we suggest that authors may set headings and provide all the details of how you conducted your study and what you did for it. In detail, it may contain search strategy, study selection, data extraction, synthesis and analysis, *etc*. All the information should be given in sufficient detail so that other scholars are able to know more.

**Level 2 heading**

*[*e.g.,**Search strategy***]*

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**RESULTS**

This section shows the main findings of your study. It may contain description of statistics, analysis, summary and comparison with other related research results, *etc*. Authors may set headings to separate the results of different experiments in this section. **Table 1** (other forms: Tables 1 and 2; Tables 1-3),**Equation (1)** [other forms: Equations (2) and (3); Equations (4-6)] and **Figure 1** (other forms: Figure 1A and B; Figure 2A-C; Figures 1 and 2A; Figures 1, 2A and 3-5) show the examples of diagrams. All the tables, equations and figures should be cited in sequence in the main content near to the first time they appear. For supplementary material, authors may cite table, equation and figure like **Supplementary Table 1**, **Supplementary Equation (1)** and **Supplementary Figure 1**. For details, you may refer to [Supplementary material template](http://www.oaepublish.com/files/tpl/mtod/Template_for_Supplementary_Material_mtod.docx).

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**Table 1. This is a table caption. A summary description of this table should be written here**

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Ref.** | **Country** | **Study type** | **Years of study** | **Group** | **Sample size** | **Age** |
| Boyle *et al.*[1], 1983 | United States | Retrospective | 1977-1981 | Control | 540 | 54 ± 2\* |
| CKD (mix) | 720 | 59 ± 4\* |
| Tsai *et al.*[2], 1996 | Taiwan | Prospective | 1991-1994 | Control | 640 | 55.7 ± 16.2# |
| ESRD | 5658 | 64.1 ± 11.4# |
| Cheung *et al.*[3], 2010 | Canada | Retrospective | 2000-2006 | Control | 850 | 67 ± 13 |
| CKD | 250 | 71 ± 13 |
| ESRD | 1650 | 68 ± 12 |

This part is footer. \*Data expressed as mean ± SEM (standard error of mean), #Data expressed as mean ± SD (standard deviation). CKD: chronic kidney disease; ESRD: end-stage renal disease. This table is cited with permission from Li *et al*.[1] published in xxx

***Table notes:***

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|  |  |
| --- | --- |
|  | (1) |

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|  |  |
| --- | --- |
| 图2  **A** | 图1  **B** |
| 20150519-10  **C** | |

**Figure 1.** We present examples of flow diagram, forest plot and funnel plot in Figure 1A-C. A: description of what the Figure 1A is; B: description of what the Figure 1B is; C: description of what the Figure 1C is. CI: confidence interval; SE: standard error; OR: odds ratio. This figure is quoted with permission from XX *et al*.[2]

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* ***Contributors should avoid making statements on economic benefits and costs except for economic data and analyses that serve as an integral part of the manuscript.***

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**Acknowledgments (optional)**

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***Standard journal articles (organization as author)***

Diabetes Prevention Program Research Group. Hypertension, insulin, and proinsulin in participants with impaired glucose tolerance. Hypertension 2002;40:679-86. [PMID: 12411462]

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Vallancien G, Emberton M, Harving N, van Moorselaar RJ; Alf-One Study Group. Sexual dysfunction in 1,274 European men suffering from lower urinary tract symptoms. J Urol 2003;169:2257-61. [PMID: 12771764 DOI: 10.1097/01.ju.0000067940.76090.73]

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Sherlock S, Dooley J. Diseases of the liver and billiary system. 9th ed. Oxford: Blackwell Sci Pub; 1993. pp. 258-96.

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Meltzer PS, Kallioniemi A, Trent JM. Chromosome alterations in human solid tumors. In: Vogelstein B, Kinzler KW, editors. The genetic basis of human cancer. New York: McGraw-Hill; 2002. pp. 93-113.

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FDA News Release. FDA approval brings first gene therapy to the United States. Available from: <https://www.fda.gov/NewsEvents/Newsroom/PressAnnouncements/ucm574058.htm>. [Last accessed on 30 Oct 2017]

***Conference proceedings***

Harnden P, Joffe JK, Jones WG, editors. Germ cell tumours V. Proceedings of the 5th Germ Cell Tumour Conference; 2001 Sep 13-15; Leeds, UK. New York: Springer; 2002.

***Conference paper***

Christensen S, Oppacher F. An analysis of Koza’s computational effort statistic for genetic programming. In: Foster JA, Lutton E, Miller J, Ryan C, Tettamanzi AG, editors. Genetic programming. EuroGP 2002: Proceedings of the 5th European Conference on Genetic Programming; 2002 Apr 3-5; Kinsdale, Ireland. Berlin: Springer; 2002. pp. 182-91.

***Unpublished material***

Tian D, Araki H, Stahl E, Bergelson J, Kreitman M. Signature of balancing selection in Arabidopsis. Proc Natl Acad Sci U S A Forthcoming 2002.

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