**Clear, Descriptive Title without Any Abbreviations
(16pt Times New Roman, Bold, Left, Title Case)**

**Given Name Family Name (in full), …, (12pt Times New Roman, bold, left)**

1*Affiliation (10pt Times New Roman, italic, left)*

*E-mail:* *author\_email@xxx.xxx* *(10pt Times New Roman, italic, left)*

2*Affiliation (10pt Times New Roman, italic, left)*

*E-mail:* *author\_email@xxx.xxx* *(10pt Times New Roman, italic, left)*

**Abstract:** An abstract of no more than 200 words should state the purposes of the study, procedures, findings and conclusions (12pt Times New Roman, justified).

**Keywords:** Keyword one, Keyword two, Keyword three... (up to 6 words) (12pt Times New Roman, justified).

**Acknowledgements**

*This section should be typed in font size 12pt Times New Roman, italic, justified.*

**References**

Up to 3-4 references allowed (examples listed below), hanging of 0.7 cm, 12pt Times New Roman, justified. The references should be listed in alphabetical order with Arabic numbers. Citations in the abstract should be with square brackets [1, 5, 10-13].

1. Edger R. C. (2009). Optimizing Substitution Matrix Choice and Gap Parameters for Sequence Alignment, BMC Bioinformatics, 10, 396-405.
2. Hompson J., D. Higgins, T. Gibson (1994). CLUSTAL W: Improving the Sensitivity of Progressive Multiple Sequence Alignment through Sequence Weighting Position Specific Gap Penalties and Weight Matrix Choice, Nucleic Acids Research, 22(22), 4673-4690.
3. Rosenblat D. (1960). On Some Aspects of Models of Complex Behavioural Systems in Information and Decision Processes (Eds., R. E. Machol, McGraw Hill), New York.
4. Murray J. D. (1995). Mathematical Biology, Springer, Verlag.
5. <http://www.mathworks.com/> (access date 20 April 2017).

**The whole submission, including title, names, affiliations, abstract, etc., should not exceed one page (A4, all margins 2.5 cm), single line spacing.**