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The journal *Cytokine* has an open access mirror journal *Cytokine: X* which has the same aims and scope and peer-review process. To submit to *Cytokine: X* visit [https://www.editorialmanager.com/CYTOX/default.aspx](https://www.editorialmanager.com/CYTOX/default.aspx).

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New requirements: Authors must provide the name, position, institution, and e-mail contacts of at least 5 potential reviewers, who are not the collaborators, friends and relatives, and are working at the authors’ organization/institution.

Authors cannot use broad terminologies, such as Cytokines, Chicken, Human, etc., at the time of submission of their manuscripts. Such terms are not useful for identifying potential reviewers. They should provide at least 6 keywords based on the content of the study. For example, if an author is submitting a paper, "Interleukin-6 effects on arthritis and the study describes therapy with a new drug and the discovery of biomarkers for the action of the said drug" they must provide some keywords like the following: IL-6, arthritis, transcriptome, signal transduction, inhibition, B-cells, macrophage, therapeutics.

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Author response(rebuttal) to review comments must be complete. Responses such as, "Done", "Completed", "Changed", "Modified as suggested' are no longer acceptable. The rebuttal must indicate the details of all modifications made in response to each question/concern raised in the review. Such details must include the exact page numbers, paragraphs, figure or table numbers, subsection of a manuscript. They also must detail all data changes (deleted or added) to the revised manuscript and references (added or deleted), the subpanels of a figure, or the columns and rows of the revised manuscript where the new data are presented. Manuscripts not adhering to these instructions will not be considered.

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All primers, their sequence and the expected PCR product sizes must be presented in a tabular format. The application for which a primer set was used (for example RT-PCR, or ChIP assay, mutational analyses) should be indicated in the table legend.

Unless there are large data sets that cannot fit into a standard figure or table, all relevant supplementary data should be part of the main paper for an easier understanding.

The importance of a protein involved in a given biological process must be validated using RNAi technology. shRNA expression vectors for many genes are commercially available. A control shRNA and a gene-specific shRNA must be included when validating the data. Exogenous add-back of knocked-down gene product could be used for better validation controls of phenotypes as well. Authors must evaluate quantitatively the efficiency of knockdown in their experimental system.

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**Types of paper**

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1) All genetic polymorphism (such as Single Nucleotide Polymorphism) analyses must include at least 400 individual samples.

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**Types of paper**

**New policies following an editorial meeting at Vienna, Oct 20, 2019:**

Since many manuscripts are taking longer than the usual time to publish from the date of first submission and it is becoming increasingly difficult to recruit reviewers, the editorial board in consultation with the Editor-in-Chief, and the publisher, made the following recommendations. These guidelines will be updated at the submission website too.

Manuscripts will be reviewed to a maximum of 3 rounds (i.e., original + 2 rounds of revision). A revision of manuscript is indicated by the letter R. After the original review is completed, reviewer comments will be provided to authors. The authors must complete all major revisions in the R1 version. Major revisions include providing data from additional experiments, data presentation and analyses, and quality of the images and the correct use of scientific English. Only minor revisions such as modification of an interpretation, missing references, replotting the data or order of figure presentation (if necessary), and exclusion of a missing control will be allowed in the R2 version. If a manuscript is not suitably modified by the R2 stage, editors can reject a further review.

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Authors cannot use broad terminologies, such as Cytokines, Chicken, Human, etc., at the time of submission of their manuscripts. Such terms are not useful for identifying potential reviewers. They should provide at least 6 keywords based on the content of the study. For example, if an author is submitting a paper, "Interleukin-6 effects on arthritis and the study describes therapy with a new drug and the discovery of biomarkers for the action of the said drug" they must provide some keywords like the following: IL-6, arthritis, transcriptome, signal transduction, inhibition, B-cells, macrophage, therapeutics.

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