Author’s response to reviews

Title: Spectrum of CFTR mutations in Chechen cystic fibrosis patients: high frequency of c.1545_1546delTA (p.Tyr515X; 1677delTA) and c.274G>A (p.Glu92Lys, E92K) mutations in North Caucasus.

Authors:

Nika Petrova (npetrova63@mail.ru)
Nataliya Kashirskaya (kashirskayanj@mail.ru)
Dzhamilya Saydaeva (saydaeva78@mail.ru)
Alexander Polyakov (polyakov@med-gen.ru)
Tagui Adyan (tagui.adyan@yandex.ru)
Olga Simonova (oisimonova@mail.ru)
Yulia Gorinova (ygorinova@yandex.ru)
Elena Kondratyeva (elenafpk@mail.ru)
Victoria Sherman (tovika@yandex.ru)
Olga Novoselova (lgnovoselova@gmail.com)
Tatiyana Vasilyeva (vasilyeva_debriv@mail.ru)
Andrey Marakhonov (marakhonov@gmail.com)
Milan Macek Jr. (milan.macek.jr@lfmotol.cuni.cz)
Evgeny Ginter (ekginter@mail.ru)
Rena Zinchenko (renazinchenko@mail.ru)

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Dear Editor,

Editor Comments:
A more detailed description of the methodologies (AFLP and RFLP), used to identify mutations, is needed and will be helpful for the readers. In alternative, I suggest to cite a paper previously published by the authors, in which the method is detailed (PMID: 26948992 DOI: 10.1016/j.jcf.2016.02.003 for example?).

Answer: On page 2 line 30 we sited our previous work (doi: 10.1016/j.jcf.2016.02.003.).

In-house molecular genetic methods previously described [11], including amplified fragment length (AFLP) and restriction fragment length (RFLP) polymorphism techniques were utilized to detect insertion/deletion variants and nucleotide substitutions, respectively.

Several further language corrections are also needed prior to publication.

Answer: Our text was once more reviewed by the English speaking editor and all the correction are added in the revised text.

Thank you for your cooperation,

Nataliya