Author's response to reviews

Title: Burkitt lymphoma beyond MYC translocation: NMYC and DNA methyl transferases deregulation

Authors:

Giulia De Falco (defalco@unisi.it)
Fabio Fuligni (fabio.fuligni2@unibo.it)
Maria R Ambrosio (maradot@libero.it)
Anna Onnis (onnis2@unisi.it)
Cristiana Bellan (bellan@unisi.it)
Bruno J Rocca (brunojim@libero.it)
Mohsen Navari (navari@unisi.it)
Maryam Etebari (etebary@unibo.it)
Lucia Mundo (mundol@hotmail.it)
Sara Gazaneo (sara.gazaneo@gmail.com)
Fabio Facchetti (facchett@med.unibs.it)
Stefano A Pileri (stefano.pileri@unibo.it)
Lorenzo Leoncini (leoncinii@unisi.it)
Pier P Piccaluga (pierpaolo.piccaluga@unibo.it)

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Author's response to reviews: see over
Dear Editor,

on behalf of all the authors I would like to thank you for your consideration.

Please find following a point-by-point letter describing the revisions you required us to make.

**Editor's Requests:**

1.) **Requesting deposition of data:**
Nucleic acid sequences, protein sequences, and atomic coordinates should be deposited in an appropriate database in time for the accession number to be included in the published article. In computational studies where the sequence information is unacceptable for inclusion in databases because of lack of experimental validation, the sequences must be published as an additional file with the article. Where appropriate, authors should adhere to the standards proposed by the Microarray Gene Expression Data Society (http://www.mged.org) and must deposit microarray data in MIAME-compliant format in one of the public repositories, such as ArrayExpress (http://www.ebi.ac.uk/arrayexpress), Gene Expression Omnibus (GEO; http://www.ncbi.nlm.nih.gov/projects/geo/) or the Center for Information Biology Gene Expression Database (CIBEX; http://cibex.nig.ac.jp).

The requesting data have been now deposited in GEO. The accession number is the following GSE71471, and the link is [http://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE71471](http://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE71471)

2.) **Availability of Supporting Data section:**
We encourage authors whose supporting data are available in an open access repository to include an “Availability of supporting data” section in their manuscript, before the Competing interests and Authors’ contributions. The section should state the name of the repository in which your data is deposited and include a link to the dataset DOI. If all the supporting data are included as additional files the section should state this. If your manuscript has any supporting sequence data, microarray data, or proteomic data this must be deposited in the appropriate repository and a link to the dataset should be included in the Availability of supporting data section.

We have done accordingly

3.) **Ethics:**
Please revise the Methods section of your manuscript to include the names of the ethics committees that approved your study. We recommend the following format: “This study was approved by the ethics committee of [xxx] Hospital [or University]”. If more than five ethics committees approved your study, you may wish to list the names of the committees in an additional file.

We have done accordingly

Best personal regards

Lorenzo Leoncini