**ORF library managing and organization**

Contact person for claimings:

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ORF library from Open Biosciences has a high error rate of cloning, so we ask our researchers to sequence each clone and that they give us the resulting information. With that information you must update the ORF database in the Note column:

* If the clone is correct, add “Seq OK”
* If the clone is not correct, add “Claming ongoing” and start a claiming. Change letter color to red for the complete row

To put a claiming just send an e-mail to Jesús from Cultek and ask for a replacement of the corresponding clone(s). Once the replacement is arrived the user can grow and sequence again the clone. In this case, two new options can occur:

* If the replacement clone is correct, add “Replacement OK” in green
* If it is not correct, add “NO replacement” in red.

Replacement clones are kept in a criobox in a -80º freezer (Big Sanyo freezer). Once you have 90 clones or so, you must do a new ORF library plate and add the information of each clone to the new plate coordinate.