

APPLICATION FORM

NGS for evolutionary biologists: from basic scripting to variant calling

Rome, CINECA, 23-27 November 2015

PERSONAL DATA

Last Surname:

First and other names:

Date of birth:

Gender:

PROFESSIONAL DATA

Institutional affiliation (main):

PI or Project Name:

Institutional affiliation (Department/Institute):

Institutional address:

Business Telephone Number:

E-mail address:

Applicant questionnaire:

1) Are you familiar with Unix/Linux? (yes/no)

2) Are you familiar with R and BioConductor? (yes/no)

If yes, at what level? Choice of:

☐ *beginner (I can run scripts but not write them myself)*

☐ *intermediate (I can write my own scripts)*

☐ *advanced (use R routinely & help people with their analysis)*

3) Are you currently using NGS technologies? (yes/no)

If Yes, please briefly explain:

If No, how do you plan to use the things you will learn on this course?

4) Which tools have you used so far to analyse NGS data?

5) What do you expect to learn from this course?

6) How would this course benefit your research and/or the work from your group/department?

7) Given the choice which of the three main topics would you choose for the group project (1. population structure and admixture, 2. demographic changes and 3. natural selection)?