

Questionnaire

Related to the **CING Bioinformatics Workshop**
organized by the CING Bioinformatics ERA Chair (BIORISE) on April 3, 2017, titled
Next Generation Sequencing and Human Variation Data Analysis
at the Cyprus Institute of Neurology and Genetics



(1) How do you find the context of the Workshop?

Adequate	Good	Very Good	Excellent

(2) How do you find the clarity of the talks?

Adequate	Good	Very Good	Excellent

(3) Please rate the usefulness of the knowledge obtained from the Workshop with respect to your research/studies.

Adequate	Good	Very Good	Excellent

(4) Please add any comments/suggestions for future Workshops.

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Thank you !

