

Cambridge, 10th November 2020

To whom it may concern,

Re: University of Cambridge, Bioinformatics Training Facility

It is my pleasure to confirm that Marco Chiapello (Raven ID mc983) attended the following training courses conducted at the University of Cambridge, Bioinformatics Training Facility loacted in Cambridge, United Kingdom.

- An Introduction to Solving Biological Problems with PERL (10<sup>th</sup> – 11<sup>th</sup> March 2016)
- Using the Ensembl Genome Browser (14<sup>th</sup> March 2016)
- Molecular Phylogenetics  $(20^{th} - 22^{nd} \text{ April } 2016)$
- Analysis of single cell RNA-seq data (22<sup>nd</sup> June 2016)
- Beginners guide to version control with git (22<sup>nd</sup> July 2016)
- Software Carpentry: Instructor training (19<sup>th</sup> – 20<sup>th</sup> September 2016)
- R object orientated programming and package development (30<sup>th</sup> September 2016)
- EMBL-EBI: Exploring Protein Sequence and Functional Information with UniProt (24th October 2016)

Best regards,

Dr Matt Castle

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