

# Pocket-sized genomics and transcriptomics analyses: a look at the newborn BioVRPi project



Treccani M<sup>1</sup>, Veschetti L<sup>1</sup>, Locatelli E<sup>1</sup>, De Tomi E<sup>1</sup>, Leoni B<sup>1</sup>, Gallinaro M<sup>1</sup>, Dagnogo D<sup>1</sup>, Moron Dalla Tor L<sup>1</sup>, Patuzzo C<sup>1</sup>, Malerba G<sup>1</sup>

Section of Biology and Genetics, Department of Neurosciences, Biomedicine and Movement Sciences, University of Verona, Italy<sup>1</sup>

## INTRODUCTION

### BioVRPi

Our project has started in January 2021 and focuses on RPi usage in bioinformatics

### Raspberry Pi (RPi)

RPi is a small single board low-cost computer developed by the Raspberry Pi Foundation

### Goals

BioVRPi aims to offer a low cost, stable and tested bioinformatic environment to students and researchers involved in genomics and transcriptomics

## RESULTS

### Genomics

GWAS on *Homo sapiens*  
395 samples - 1047171 markers

### Devices (64-bit)

RPi model 4 - 8 GB RAM - 1.5 GHz  
Tower PC - 32 GB RAM - 3.0 GHz

Phases	RPi	PC
Pre-processing	1 min	3 sec
QC	2 min	4 sec
Association	5 min	1 min

### Transcriptomics

RNA-seq on *Strongyloides stercoralis*  
8 samples - 38.9 M reads/sample on avg

### Devices (64-bit)

RPi model 4 - 2 GB RAM - 1.5 GHz  
Laptop - 16 GB RAM - 3.2 GHz

Phases	RPi	Laptop
Alignment	60 min	30 min
Reconstruct	25 min	28 min
Expression	41 min	7 min

## CONCLUSION

RPis are effective devices for reproducible GWAS and RNA-seq analyses  
Availability of state-of-art programs with ARM-based customization  
Possibility to develop an having-fun environment, to learn and explore new strategies