### PERSONAL INFORMATION

#### **Chiara Reale**

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Fondazione IRCCS Istituto Neurologico Carlo Besta Unit of Medical Genetics and Neurogenetics Via Libero Temolo 4, 20126 Milan

Gender: F | Nationality: Italian

ORCID: 0000-0002-7493-758X

#### **WORK EXPERIENCE**

From 15-06-2020 to date

## Italian NHS Researcher

Health Researcher in Fondazione IRCCS Istituto Neurologico Carlo Besta, Unit of Medical Genetics and Neurogenetics

 Application of Next-generation sequencing (NGS) technology, specifically targeted resequencing, in movement disorders study

#### From 01-11-2014 to14-06-2020

# **Contract Researcher**

Molecular biologist in Fondazione IRCCS Istituto Neurologico Carlo Besta, Unit of Medical Genetics and Neurogenetics

 Application of Next-generation sequencing (NGS) technology, specifically targeted resequencing, in movement disorders and autism spectrum disorders study.

#### From 01-06-2009 to 30-09-2014

## **Fellowship**

Molecular biologist in Fondazione IRCCS Istituto Neurologico Carlo Besta, Unit of Medical Genetics and Neurogenetics

• Use of principal biomolecular techniques for the study of genetic basis of movement disorders

#### From 17-10-2006 to 31-05-2009

# Internship

Graduate student in Fondazione IRCCS Istituto Neurologico Carlo Besta, Unit of Medical Genetics and Neurogenetics

· Learning of principal biomolecular techniques for the study of genetic basis of movement disorders

#### **EDUCATION AND TRAINING**

25/09/2014

# Registration in the Italian National Order of Biologist

Section A, number AA\_071666

#### 07/07/2014

# Qualifying exam to exercise the biological profession

University of Insubria-Varese

#### 27-04-2009

# Master's Degree in Biomolecular Biology,

University of Milano-Bicocca

## **TECHNICAL SKILLS**

DNA and RNA extraction and quantification, PCR (Polymerase Chain Reaction), Real-time PCR, MLPA (Multiplex Ligation-Dependent Probe Amplification), horizontal and capillary elettrophoresis, RFLP, knowledge of bioinfarmatic tools for Sanger sequencing, variants pathogenicity prediction and Next-generation sequencing (NGS), preparation library for Next Generation Sequencing (NGS) (MiSeq, Illumina).