

INTEGRATED APPLIED GENETICS TRAINING - AppGENEdu

A Project Financed through the EEA Grants 2014-2021, Education, Scholarships, Apprenticeships and Youth Entrepreneurship Programme in Romania, Cooperation projects in The Higher Education Area

Project Curricula

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| Genetics perspective Epidemiology perspective |
| Genetics Fundamentals |
| Statistics and Bioinformatics <ul style="list-style-type: none"> • Fundamental elements of biostatistics • Fundamental elements of statistical genetics • Elements of statistical study design |
| Workshop <ul style="list-style-type: none"> • Statistical tools; R • Statistical basic processing |
| Workshop <ul style="list-style-type: none"> • Statistical basic processing • Basic data visualization |
| Fundamentals of Genetic Epidemiology <ul style="list-style-type: none"> • Quantitative Statistical Tests, linear regression (smoking) • Phenotype - genotype correlations |
| Workshop <ul style="list-style-type: none"> • Phenotype - genotype correlation • Example of epidemiology study design based on provided "use-cases" |
| Elements of genetic risk analysis, <ul style="list-style-type: none"> • Genetics: LD frequencies, HWE • Relative risk |
| Workshop biostatistical tools - plink <ul style="list-style-type: none"> • Missingness by phenotype • Missingness by genotype • Hardy-Weinberg • Allele frequencies • LD-based SNP pruning • Mendel errors |
| Elements of genetic risk analysis <ul style="list-style-type: none"> • Genetics: linkage disequilibrium, haplotypes • Example of population admixture • Bioinformatics methods & principles |

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| <ul style="list-style-type: none"> Genetics: LD frequencies, HWE |
| <p>Workshop</p> <ul style="list-style-type: none"> Case/control Fisher's exact Stratified analysis Quantitative trait Linear and logistic models Multiple-test correction |
| <p>Workshop</p> <ul style="list-style-type: none"> Missingness by phenotype Missingness by genotype Hardy-Weinberg Allele frequencies LD-based SNP pruning Mendel errors |
| <p>Elements of Data Science</p> <ul style="list-style-type: none"> Data load Data cleansing Data visualization Data processing Data comparison Reproducible research |
| <p>Workshop</p> <ul style="list-style-type: none"> Example of data processing (from problem to result) Bioinformatics exercises |
| <p>Bioinformatics</p> <ul style="list-style-type: none"> Biologic consequences of genetic variants/gene mutations Genetic variances in Genetic Epidemiology context (GWAS) |
| <p>Workshop</p> <ul style="list-style-type: none"> GWAS study Output analysis |
| <p>Advanced Applied Genetics</p> <ul style="list-style-type: none"> Replication studies Quantitative trait loci Genetic pathways & gene integration |
| <p>Workshop</p> <ul style="list-style-type: none"> Replication study Visual representation |
| <p>Bioethics principles</p> <ul style="list-style-type: none"> Informed consent GDPR elements Elements of ethics of scientific publication |
| <p>Elements of Research methodology</p> |
| <p>Specific steps and examples for genomics data science projects</p> |
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| <p>Prepare and writing a scientific article</p> |
| <p>Project bid and presentations</p> |
| <p>Visits to the clinics associated with the project topic</p> <ul style="list-style-type: none"> Phenotypic and epidemiological data collection Genetic sampling Clinical visit Clinical case report |
| <p>Project work</p> |
| <p>Project presentation and evaluation</p> |