

Georg-August-Universität Göttingen Module M.iPAB.0003: Statistical genetics, breeding informatics and experimental design	6 C 4 WLH
Learning outcome, core skills: Novel biotechnological methods allow the production of very large data sets (gene sequences, genotypes, transcriptomes) at decreasing costs. Students learn about statistical and computational methods to use these records for breeding issues. Furthermore, the main experimental designs to plan, implement, and evaluate targeted and efficient experiments for data generation will be treated.	Workload: Attendance time: 56 h Self-study time: 124 h
Course: Statistical genetics, breeding informatics and experimental design (Lecture, Exercise) <i>Contents:</i> <ul style="list-style-type: none"> • Gene Expression Analysis • Genome-wide association analysis • QTL mapping • Statistical hypothesis testing • Regression methods • Analysis of variance • Multiple testing • Experimental designs (block designs, randomized designs, Latin squares) • Sample size estimation • Introduction to programming • Fundamentals of databases Literature: Andrea Foulkes: Applied Statistical Genetics with R	4 WLH
Examination: Written examination (60 minutes) Examination requirements: Profound knowledge of statistics and informatics methods to use them for breeding issues.	6 C
Admission requirements: none	Recommended previous knowledge: Basics in statistics and genetics
Language: English	Person responsible for module: Prof. Dr. Armin Schmitt
Course frequency: each winter semester	Duration: 1 semester[s]
Number of repeat examinations permitted: twice	Recommended semester:
Maximum number of students: 20	