PREDICTION OF MALADAPTATION FROM ECOLOGICAL AND GENOMIC DATA USING GENOMIC OFFSETS

FRANCISCO DE BORJA CAMPUZANO JIMÉNEZ, au726682

MASTER OF SCIENCE IN BIOINFORMATICS

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Supervisors: Thomas Bataillon and Genis Garcia Erill

AARHUS UNIVERSITY

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Bioinformatics Research Centre
Faculty of Natural Sciences
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ACRONYMS

SNP single-nucleotide polymorphism
QTL quantitative trait locus
RONA risk of non-adaptednes
RDA redundancy analysis
FDR false-discovery ratio
PCA principal component analysis
LFMM latent factor mixed model
GEA genotype $\times$ environment association