Eleven species are distinguished in the genus Oxythyrea Mulsant, 1842 (Coleoptera: Scarabaeidae) nowadays. They are not divided into subspecies. Diversity of the genus is concentrated in the Mediterranean and Oxythyrea funesta (Poda, 1761) inhabit a wide area in the western Palearctic Region. It was observed in last decades, that O. funesta retreated from central Europe to south and then recolonized it back including new areas in northern regions.

Master thesis is focused on resolving population structure of O. funesta and partial phylogeny of the genus Oxythyrea using molecular genetic methods. 145 individuals of O. funesta and 15 individuals of five other species of the genus Oxythyrea appear in analysis. We acquired sequences of mitochondrial genes cytochrome oxidase I (807 bp), cytochrome b (381 bp) and nuclear gene internal transcribed spacer I (946 bp) from these specimens.

The results of phylogenetic analysis confirmed so far the only one existing interpretation of relationships within the genus Oxythyrea based on morphological data. We also confirmed complicated relationships between O. funesta and O. pantherina, which also appear in the historical development of their taxonomy. We detected different genetic lineage in Sicily, southern Italy and Tunisia using phylogenetic trees and haplotype networks. Haplotype networks and Tajima D test point to recent demographic expansion of the remaining individuals of O. funesta.