

Identifying critical biodiversity areas through phylogenetic and species diversity

S. von der Heyden, Stellenbosch University

The main outputs will be the generation of 370 barcode sequences for nudibranchs, chitons and limpets; the bulk of samples comes from the Nudibranchia. For each taxonomic group, there are few South African representatives in the BOLD data base. The list detailing species is attached to this application.