**Conference Abstract** 

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## SYNTHESYS+ Molecular Collections in the Age of Genomics: Standards and processes

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## Abstract

The aim of the Global Genome Biodiversity Network (GGBN, <u>http://www.ggbn.org</u>) is to foster collaboration among biodiversity biobanks on a global scale in order to further compliance with standards, best practices, and to secure interoperability and exchange of material in accordance with national and international legislation and conventions. Thus, key aspects of GGBN's mission are to develop a network of trusted collections, establishing standards, and identifying best practices by reaching out to other communities. This is especially critical in the light of new international legislation such as the recent <u>Nagoya Protocol</u> on Access and Benefit Sharing (ABS).

Biological repositories such as but not limited to natural history collections, botanic gardens, culture collections and zoos are facing a series of challenges triggered by the rapid acceleration in sequencing technology that has put added pressure on the use of samples, which just a few years ago were considered inaccessible for sequencing.

ABS legislation applies to nearly all collection types, and with biodiversity biobanks increasing in number worldwide, there is an urgent need to streamline procedures and to ensure legislative compliance. Within Europe it is necessary to 1) reach common

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standards for biodiversity and environmental biobanks; 2) define best practices for the use of molecular collections; and 3) try to ease exchange of samples and related information, while staying compliant with legislation and conventions.

Within the EU funded SYNTHESYS+ project (<u>http://www.synthesys.info</u>), GGBN is leading Network Activity 3 (NA3). An overview of planned activities and tasks will be given here with special emphasis on linkages within and beyond SYNTHESYS+.

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