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Author: Francesca Rosa

**Folder results/slost/**

This folder contains multiple sub-folders with the results of the application of the biodiversity model (the scripts of the biodiversity model are in the folder *scripts\_model*).

The list of code words used to identify the setting implemented for each sub-folder are listed here.

Almost all folders are labelled either with “bs” or “nobs”:

* bs: bootstrapping – in the biodiversity model, the bootstrapping procedure has been applied to both the response ratios and the z values to quantify the response ratio to quantify the confidence intervals.
* nobs: the biodiversity model has calculated the static results (no bootstrapping applied). They correspond to in Rosa et al. (2022) (S8) and are to the results to which the confidence intervals are applied.

Warning: there is only one folder which does not have this specification. This folder contains the files where the static results (nobs) are together with the confidence intervals (bs) according to equation S8.1 of Rosa et al. (2022).

* cutoff: in the biodiversity model, all response ratios above 1 have been set to 1 (default setting).
* timber: in the GLOBIOM model, part of EU clear-cut management has been replaced with timber plantations.
* nocutoff: no cutoff was applied.

The acronym “mg” which is the title of all .csv files stands for “marginal”. “Marginal” is an option in the GLOBIOM model and means that only the most productive forest area is assumed to be imported to the EU28 (Clear-cut area in Boreal/Temperate and Plantations in the Tropics) or exported from EU28 (Clear-cut area), given economical advantage of intensive management. This option was selected because according to IIASA’s experts it appeared the most likely.