

get_species_association.R

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get_species_association	
<i>Compute Species Associations</i>	

Description

Computes species associations from a fitted Hmsc model object.

Usage

```
get_species_association(data_source)
```

Arguments

data_source	A list containing a fitted Hmsc model under the 'mod' element. Generally, this is the output of the function <code>add_model_evaluation()</code> or <code># get_better_model_based_on_fit()</code>
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Value

A matrix of species associations.

See Also

[`add_model_evaluation()`, `get_better_model_based_on_fit()`]

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