get_significant_associations.R

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Contents

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get_significant_associations

Get Significant Species Associations

Description

Identifies significant species associations based on support and mean values.

Usage

```
get_significant_associations(data_source, alpha = 0.05)
```

Arguments

data_source	A list containing association matrices from a fitted Hmsc model. Generally, this is the output of the function get_species_association().
alpha	Significance level for support threshold (default: 0.05).

Value

A vector of significant association values.

See Also

[get_species_association()]

Index