# get\_community\_taxa.R

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# Description

Extracts a vector of unique taxa from a community data frame.

#### Usage

```
get_community_taxa(data)
```

#### Arguments

data

A data frame containing a 'taxon' column.

# **Details**

Uses dplyr to select distinct values from the 'taxon' column and returns them as a vector.

#### Value

A character vector of unique taxon names present in the data.

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