# classify\_taxonomic\_resolution.R

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### **Description**

Classifies taxa in a data frame to a specified taxonomic resolution (family, genus, or species) using a classification table, and aggregates pollen proportions accordingly.

#### Usage

```
classify_taxonomic_resolution(
  data,
  data_classification_table,
  taxonomic_resolution
)
```

#### **Arguments**

data  $A \ data \ frame \ containing \ taxon \ data \ with \ columns \ including \ 'taxon', \ 'dataset\_name', \ 'age', \ and \ 'pollen\_prop'.$   $data\_classification\_table$ 

A data frame mapping 'sel\_name' to taxonomic levels (e.g., family, genus, species). taxonomic\_resolution

A character string specifying the taxonomic resolution to classify to. Must be one of 'family', 'genus', or 'species'.

#### **Details**

Performs a left join to map taxa to the desired resolution, aggregates pollen proportions, and ensures all dataset\_name-age-taxon combinations are present in the output.

# Value

A data frame with taxa classified to the specified resolution and pollen proportions aggregated accordingly. The output preserves all dataset\_name and age combinations for true negatives.

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