



# DescribeML:

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## Language Reference Guide

(Version 0.0.5)

DescribeML is a VSCode language plugin to describe machine-learning datasets.

Full examples of the language can be found in the public open repository [here](#)

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**Metadata:**

- **Title:** **STRING** The public title of the dataset
- **Unique-identifier:** **ID** Machine-readable unique identifier of the dataset
- **Version:** **ID** The version of the dataset
- **Date:** The date of the dataset
  - **Created:** **DATE** The date where the dataset was initially created:
  - **Modified:** **DATE** The date where the dataset was last modified:
  - **Published:** **DATE** The publication date of the dataset:

**Example****Dates:**

Release **Date:** 10-08-20  
 Modified **Date:** 10-08-20  
 Published **Date:** 10-08-20

- **Citation:** The citation of the dataset, between chose between a raw citation and a structured format
  - **Raw Citation:** **STRING** Raw citation as text, or as Bibtex or equivalent format, of the dataset
  - **OR:**
    - **Title:** **STRING** The title of the dataset
    - **Authors:** **STRING** The authors of the dataset
    - **Year:** **DATE** The year of the dataset
    - **Journal/Conference:** **STRING** The publisher of the dataset
    - **Publisher:** **STRING** The publisher of the dataset:
    - **URL:** **URL** The URL of the dataset
    - **DOI:** **ID** The DOI of the dataset
    - **ISBN:** **ID** The ISBN of the dataset

**Example:****Citation:**

**Title:** "SIIM-ISIC 2020 Challenge Dataset. International Skin Imaging Collaboration"  
**Year:** 2020  
**Publisher:** "International Skin Imaging Collaboration"  
**DOI:** "doi.org/10.34970/2020-ds01"  
**Url:** "https://www.kaggle.com/c/siim-isic-melanoma-classification"

- **Description:** The description of the dataset
  - **Description:** **STRING** Textual description of the dataset **OR:**

- **Purposes** *STRING* For what purposes was the dataset created?
- **Tasks**: *TASKS ENUMERATE* List of ML tasks the dataset is intended for: *Autocomplete feature will guide you through the options*
- **Gaps**: *STRING* Which gaps does the dataset aims to fill
- **Areas**: *ID* Set a list of areas of the dataset
- **Tags**: *ID, ...* Set a list of Tags of the dataset

Example:

```
Description:
  Purposes:
    Purposes: "The 2020 SIIM-ISIC Melanoma"
    Tasks:    [classification]
    Gaps:      "As the leading healthcare organization for
informatics in medical imaging..."
    Areas:     HealthCare
    Tags:      Images Melanoma diagnosis SkinImage
```

- **Applications** Summerize the applications of the dataset
  - **Past Uses**: *STRING* Summerize the past uses of the dataset
  - **Recommended uses**: *STRING* Summerize the recommended uses of the dataset
  - **Non-recommended uses**: *STRING* Summerize the non-recommended uses of the dataset.
  - **Benchmarking**: Benchmarking of the dataset
    - **Task**: *TASKS ENUMERATE* Task to benchmark *Autocomplete feature will guide you through the options*
    - **Metric**: Metric to benchmark
      - **F1**: *NUMBER* F1 score
      - **Accuracy**: *NUMBER* Accuracy score
      - **Precision**: *NUMBER* Precision score
      - **Recall**: *NUMBER* Recall score
    - **Reference**: *STRING* Source of the benchmark

Example

```
Applications:
  Past Uses: "The 2020 SIIM-ISIC Melanoma Classification... "
  Recommended:
    "Identify melanoma in lesion images."
    "Predict incidence of melanoma in a population."
  Non-recommended: "Due to low population prevalence and
challenges with access."
  Benchmarking:
    Task: Language-model
    [
      Model: "ModelExample"
      Metrics:[
```

```

        F1: 81
        Accuracy: 81
        Precision: 81
        Recall: 81
    ]
    Reference: "https://www.kaggle.com/c/siim-isic-
melanoma-classification/leaderboard"
]
```

- **Distribution** Summarize the distribution of the dataset
  - **Is public?:** **BOOL** Indicate if the dataset is publicly available
  - **Licenses:** **LICENCES** **ENUMERATE** List of standard licenses, use others if not fit your case: **The Montreal data license** , **Creative Commons**, **CC0: Public Domain** ...
  - **Rights(stand-alone)** **ENUMERATE** Montreal data licence enumerate of stand-alone rights: **Access** | **Tagging** | **Distribute** | **Re-Represent**
  - **Rights(with models):** **ENUMERATE** Montreal data licence enumerate of model related rights: **Benchmark** | **Research** | **Publish** | **Internal Use** | **'Output Commercialization'** | **Model Commercialization**
  - **Credits/Attribution Notice:** **STRING** Who needs to be credited when using the dataset
  - **Designated Third Parties:** **STRING** Third parties in charge of licensing and distribution issues
  - **Additional Conditions:** **STRING** Other issues specified by the authors

#### Example

```

Distribution:
  Licences: CC BY 3.0 (Attribution 3.0 Unported)
  Rights(stand-alone): Access
  Rights(with models): Benchmark
  Additional Conditions "In addition to the CC-BY-NC license, the
dataset is governed by the ISIC Terms of Use ... "
```

- **Authoring** Authoring of the dataset
  - **Authors** Authors of the dataset
    - **Name:** **STRING** Name of the author
    - **Email:** **EMAIL** Email of the author
  - **Founders** Founders of the dataset
    - **Name:** **STRING** Name of the founder
    - **Type:** **ENUMERATE** Type of the founder **private** | **public** | **mixed**;
    - **Grantor** **STRING** Grantor of the dataset
    - **Grant ID:** **ID** Machine-readable name of the grant id
  - **Maintainers** Maintainers of the dataset
    - **Name:** **STRING** Name of the maintainer
    - **Email:** **EMAIL** Email of the maintainer
  - **Erratum?:** **STRING** Is there any erratum?

- **Data retention:** **STRING** Please indicate any data retention policy
- **Version lifecycle:** **STRING** Describe the planned version lifecycle
- **Contribution guidelines** **STRING** Is there any contribution guideline?

Example:

```

Authoring:
  Authors:
    Name Skin_Imaging_Collaboration_ISIC  email emailo@emailo.com
    [...]
  Funders:
    Name The_University_of_Queensland  type mixed
    grantor "National Health and Medical Research Council
(NHMRC) – Centre of Research Excellence Scheme"
    grantId: APP1099021
    [...]
  Erratum?: "There is no erratum known"
  Contribution guidelines: "No contribution guidelines provided"

```

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## Composition:

- **Rationale** **STRING** Provide a composition rationale
- **Total Size** **NUMBER** Total size of tuples of the dataset
- **Instances** A composition description of each instance of the dataset
  - **Instance:** **ID** Machine-readable name of the instance
  - **Size:** **NUMBER** Size of the instance
  - **Description:** **STRING** Description of the instance
  - **Type:** **ENUMERATE** Type of the instance **Record-Data** | **Time-Series** | **Ordered** | **Graph** | **Other**
  - **Attribute Number:** **NUMBER** Number of attributes
  - **Attributes:** Description of each attribute of the instance
    - **attribute:** **ID** Machine-readable name of the attribute
    - **Description:** **STRING** Description of the attribute
    - **Associated label:** **Labels** Reference to a declared label in a labeling process (first you should complete the provenance part)
    - **unique values:** **NUMBER** Type of the attribute
    - **ofType:** **ENUMERATE** Type of the attribute **Categorical** | **Nominal** **If** **ofType** is **Categorical**

- **Statistics:** Statistic of the attribute
  - **Unique:** **NUMBER** Unique tuples (without duplications)
  - **Unique Percentage:** **NUMBER** Percentage of unique tuples
  - **Missing Values:** **NUMBER** Number of missing values
  - **Completeness:** **NUMBER** Completeness of the attribute
  - **Mode:** **STRING** Mode of the attribute
  - **First Rows:** [**0**: **ROW1**, ...] Percentage of the mode
  - **Min-length:** **NUMBER** Min of the attribute
  - **Max-length:** **NUMBER** Max of the attribute
  - **Median-length:** **NUMBER** Median lengths of the attribute
  - **Length-histogram:** **STRING** Histogram of the attribute
  - **Chi-Squared:** Chi-Squared of the attribute
    - **statistic:** Statistic of the chi-square analysis
    - **p-value:** p-value of the chi-square analysis
- **Binary attribute:** **BOOL** Is a binary attribute?
  - **Symmetry:** **ENUMERATE** **Symmetryc** | **Asymmetryc**
  - **Attribute Sparsity:** **NUMBER** How sparse is the binary attribute?
- **Categoric Distribution:** [**"CATEGORY"**: **"NUMBER"**%, ...] Categoric distribution of the attribute

Example:

```
attribute: benignant_malignant
description: 'Type of the melanoma'
label: skinLabel
count: 33126
ofType: Categorical
Statistics:
  Missing Values: 0
  Completeness: 100
  Chi-Squared:
    p-value: 0
  Categoric Distribution:
    [
      "benignant": 80%,
      "malignant": 20%
    ]
```

Else **ofType** is **Nominal**

- **Statistics:** Statistics of the attribute
  - **Mean:** **NUMBER** Unique tuples (without duplications)
  - **Median:** **NUMBER** Percentage of unique tuples
  - **Mode:** **NUMBER** Mode of the attribute
  - **Minimmum:** **NUMBER** Min of the attribute
  - **Maximmum:** **NUMBER** Max of the attribute
  - **Quartiles:** **[Q1:NUMBER, ...]** Median lengths of the attribute
  - **IQR:** **NUMBER** Histogram of the attribute

#### Example

```
attribute: acidity
description: 'wine acidity mesure'
count: 33126
ofType: Numerical
Statistics:
  Mean: 4
  Median: 4.1
  Standard Desviation: 0.2
  Minimum: 5
  Maximum: 87
  Quartiles: Q1:17 Q2:27 Q3:30 Q4:30
  IQR: 1.2
```

- **Statistics:** (instance) Statistic of the instance
  - **Correlations:** Correlation of the instance, choose one calculation type
    - **Pearson:** **[INDEX:"NUMBER", ...]** Pearson correlation of the instance
    - **Spearman:** **[INDEX:"NUMBER", ...]** Spearman correlation of the instance
    - **Kendall:** **[INDEX:"NUMBER", ...]** Kendall correlation of the instance
    - **Cramers:** **[INDEX:"NUMBER", ...]** Cramers correlation of the instance
    - **Phi-k** **[INDEX:"NUMBER", ...]** Phi-k correlation of the instance
  - **Pair Correlation** **Between [ATTRIBUTE], and [ATTRIBUTE]** Points the relevant pair-correlation between two instances of declared attributes.
  - **Quality Metrics:** General quality metrics of the instance
    - **Sparsity:** **NUMBER** Sparsity of the instance
    - **Completeness:** **NUMBER** Completeness of the instance
    - **Class balance:** **STRING** Class balance of the instance
    - **Noisy labels:** **STRING** Noisy labels of the instance

#### Example:

```
Statistics:
  Correlations: Spearman: ['1': 0.2, '2':0.3, '3':0.4, '4':0.5,
    '5':0.6, '6':0.7, '7':0.8, '8':0.9]
```

```

Pair Correlation:
  between ImageId and diagnosis
  between age and external source
  From: "National statistical office"
  Rationale: "The age average is similar to the Nevada
state age average due to
              national statistical office average of
2022 of Nevada"
Quality Metrics:
  Completeness: 100

```

- **Consistency Rules:** Set the consistency rules of your dataset

- **Rule:** **OCLExpression** OCL expression of the rule

Example:

```

Consistency rules:
inv: skinImages : (age >= 0)

```

- **Dependencies:** Dependencies of the rule
  - **Description:** **STRING** Description of the dependencies
  - **Links:** **URL** Link to the dependency artifact
- **Instances relation:** **Relation: ID attribute: [ATTRIBUTE] is related to [INSTANCE]** Relation between instances

## Provenance:

- **Curation Rationale** **STRING** Provide a provenance rationale
- **Gathering Processes:**
  - **Process:** **ID** Machine-readable name of the process
  - **Description:** **STRING** Description of the process
  - **When data was collected:** **STRING** Date where data the process was performed
  - **How data was collected** **STRING** How data was collected
  - **Is language data:** Set the speech situation
    - **Language:** **STRING** Language of the data
    - **Time and place:** **STRING**
    - **Modality:** **ENUMERATE** Modality of the speech **spoken/signed | written**
    - **Type:** **ENUMERATE** Type of the speech **scripted/edited | spontaneous**
    - **Syncrony:** **ENUMERATE** Syncrony of the speech **synchronous | asynchronous**
    - **Inteded Audience:** **STRING** Intended audience of the speech
  - **Social Issues:** **[SOCIAL ISSUES]** Relation of the gathering process with an already declared social issue instance
  - **Source:** Source of the data



- **Source:** `ID` machine-readable name of the source
- **Description:** `STRING` Description of the source
- **Noise:** `STRING` Description of the source's noise
- **Links:** `URL` Link to the source artifact
- **Process Demographics:**
  - **Age:** `NUMBER` Median age of the participants
  - **Gender:** `STRING` Gender relation of the participants
  - **Country/Region** `STRING` Country/Region of the participants
  - **Race/Ethnicity** `STRING` Race or ethnicity of the participants
  - **Native Language** `STRING` Native language of the participants
  - **Socioeconomic status** `STRING` Socioeconomic status
  - **Number of speakers represented:** `NUMBER` Number of participants
  - **Precense of disorders in speech:** `STRING` Number of speakers
  - **Training in linguistics/other relevant disciplines** `STRING` Explain the training of the participants
- **Gathering Team** Team in charge of gathering the data
  - **Who collects the data:** `STRING` Who collects the data
  - **Type** `ENUMERATE` `Internal` | `External` | `Contractors` | `Crowdsourcing`
  - **Demographics:** Demographics of the gathering team
    - **Age:** `NUMBER` Median age of the participants
    - **Gender:** `STRING` Gender relation of the participants
    - **Country/Region** `STRING` Country/Region of the participants
    - **Race/Ethnicity** `STRING` Race or ethnicity of the participants
    - **Native Language** `STRING` Native language of the participants
    - **Socioeconomic status** `STRING` Socioeconomic status
    - **Training in linguistics/other relevant disciplines** `STRING` Explain the training of the participants
- **Gathering Requirements:** `Requirement:` `STRING`, ...

Example:

```
Data Provenance:
  Curation Rationale: "The curation process have been conducted by
several health institutions... "
  Gathering Processes:
    Process: GatheringProcess1
    Description:
      "The sources are: the Melanoma Institute Australia and
the ..."
```

```
    Source: GeneralHospital1
    Description: 'Source Description'
    Noise:
      "Inconsistent lighting in images may alter skin
type"

      "Duplicates:..."
    Related Instances: skinImages
    How data is collected: Manual Human Curator
    When data was collected:
```

```

Range: 1998 - 2019
Process Demographics:
  Country/Region: 'Australia'
  [...]
Gathering Team:
  Who collects the data: "A team of dermatologists and
  pathologists"
  Type Internal
Gather Requirements:
  Requirement: "We queried clinical imaging databases across
  the six centers to generate a ..."

```

- **LabelingProcesses:**

- **Labeling process:** **ID** Machine-readable name of the labeling process
- **Description:** **STRING** Description of the labeling process
- **Type:** **ENUMERATE** 'Bounding boxes' | 'Lines and splines' | 'Semantic Segmentation' | '3D cuboids' | 'Polygonal segmentation' | 'Landmark and key-point' | 'Image and video annotations' | 'Entity annotation' | 'Content and textual categorization'
- **Labels:** Labels of the labeling process
  - **Label:** **ID** Machine-readable name of the label
  - **Description:** **STRING** Description of the label
  - **Mapping:** [ATTRIBUTE,...] Relate a label with instances of attributes already declared in the documentation
- **Labeling Team:**
  - **Who collects the data:** **STRING** Who collects the data
  - **Type** **ENUMERATE** Internal, External, Contractors, Crowdsourcing
  - **Demographics:** Demographics of the gathering team
    - **Age:** **NUMBER** Median age of the participants
    - **Gender:** **STRING** Gender relation of the participants
    - **Country/Region** **STRING** Country/Region of the participants
    - **Race/Ethnicity** **STRING** Race or ethnicity of the participants
    - **Native Language** **STRING** Native language of the participants
    - **Socioeconomic status** **STRING** Socioeconomic status
    - **Number of speakers represented:** **NUMBER** Number of participants
    - **Precense of disorders in speech:** **STRING** Number of speakers
    - **Training in linguistics/other relevant disciplines** **STRING** Explain the training of the participants
- **Infrastructure:** Infrastructure used to annotate the data
  - **Tool:** **STRING** Tool used to annotate the data
  - **Platform:** **STRING** Platform where the tool works
  - **Version:** **STRING** Version of the tool and platform
  - **Language:** **STRING** Language of the tool
  - **Comments:** **STRING** Provide comments about the tool
- **Validation:** Validation methods to ensure annotation quality
  - **Validation Methods:** **STRING** Validation method used
  - **Validation Dates:** **STRING** Dates where the validation where done annotations

- **Golden Questions:** Golden Question pass to the annotators
  - **Question:** **STRING** Textual question
  - **Inter-annotation agreement:** **NUMBER** Inter-annotation agreement for each question. Low values mean low confidence in the annotation
- **Validation Requirements:** **Requirement:** **STRING**, ... Provide comments about the validation tool
- **Labeling Requirements:** **Requirement:** **STRING**, ...

Example:

```
LabelingProcesses:
  Labeling process: skinLabeling
  Description: "Medical staff looking at the data and images
and annotating the diagnosis"
  Type: Image and video annotations
  Labels:
    Label: skinLabel
    Description: "marked as beningnant or malignant"
    Mapping: beningnant_malignant
  Labeling Team:
    Who collects the data: "Internal Medical staff"
    Type Internal
    Country/Region: "Australia"
  Label Requirements:
    Requirement: "1) Images containing any potentially
identifying features, such as jewelry"
```

- **Preprocesses:** Data preprocesses done over the data
  - **Preprocess:** **ID** machine-readable name of the preprocess
  - **Type:** **ENUMERATE** Type of preprocess applied 'Missing Values' | 'Data Augmentation' | 'Outlier Filtering' | 'Remove Duplicates' | 'Data reduction' | 'Sampling' | 'Data Normalization' | 'Others'
  - **Description:** **STRING** Description of the preprocess
  - **Social Issues:** [**SOCIAL ISSUES**] Relation of the preprocess with an already declared social issue instance

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## Social Concerns

- **Social Concerns**
  - **Rationale:** **STRING** Rationale of the social concerns of the dataset
  - **Social Issues:** Social issues identified from the data
    - **Social Issue:** **ID** Machine-readable name of the social issue
    - **IssueType:** **ENUMERATE** Type of social concern 'Privacy' | 'Bias' | 'Sensitive Data' | 'Social Impact'
    - **Description:** **STRING** Description of the social issue
    - **Related Attributes** **attribute:** [**ATTRIBUTE**] Attributes related to the social issue
    - **Instace belong to people:**

- **Have sensitive attributes?** `[Attribute]`, ... List of sensitive attributes
- **Are there protected groups?** `ENUMERATE` (Yes, No, Unknown)
- **Might be offensive?** `STRING` Is there offensive content in the dataset

#### Examples

Social Concerns:

Rationale: 'Dataset may not be representative of the real world data, and the convenience sample is not representative of general incidence of melanoma'

Social Issue: raceRepresentative

IssueType: Bias

Description: "Dataset is not representative with respect to darker skin types"

Related Attributes:

attribute: ImageId

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For any related question, please contact the authors at: [jginermi@uoc.edu](mailto:jginermi@uoc.edu)