Folder 3D Data

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Custom Folder Dataset.

This script creates a custom dataset from a folder.

```
class anomalib.data.depth.folder_3d.Folder3D(name, normal_dir, root, abnormal_dir=None, normal_test_dir=None, mask_dir=None, normal_depth_dir=None, abnormal_depth_dir=None, normal_test_depth_dir=None, extensions=None, train_batch_size=32, eval_batch_size=32, num_workers=8, task=TaskType.SEGMENTATION, image_size=None, transform=None, train_transform=None, eval_transform=None, test_split_mode=TestSplitMode.FROM_DIR, test_split_ratio=0.2, val_split_mode=ValSplitMode.FROM_TEST, val_split_ratio=0.5, seed=None)
```

Bases: AnomalibDataModule

Folder DataModule.

Parameters:

- **name** (*str*) Name of the dataset. This is used to name the datamodule, especially when logging/saving.
- **normal_dir** (*str* | *Path*) Name of the directory containing normal images.
- **root** (*str* | *Path* | *None*) Path to the root folder containing normal and abnormal dirs. Defaults to None.
- **abnormal_dir** (*str* | *Path* | *None*) Name of the directory containing abnormal images. Defaults to abnormal.
- **normal_test_dir** (*str* | *Path* | *None, optional*) Path to the directory containing normal images for the test dataset. Defaults to None.
- mask_dir (str | Path | None, optional) Path to the directory containing the mask annotations. Defaults to None.
- **normal_depth_dir** (*str* | *Path* | *None, optional*) Path to the directory containing normal depth images for the test dataset. Normal test depth images will be a split of *normal_dir*
- **abnormal_depth_dir** (*str* | *Path* | *None, optional*) Path to the directory containing abnormal depth images for the test dataset.
- **normal_test_depth_dir** (*str* | *Path* | *None, optional*) Path to the directory containing normal depth images for the test dataset. Normal test images will be a split of *normal_dir* if *None*. Defaults to None.
- **normal_split_ratio** (*float, optional*) Ratio to split normal training images and add to the test set in case test set doesn't contain any normal images. Defaults to 0.2.
- **extensions** (*tuple[str, ...]* | *None, optional*) Type of the image extensions to read from the directory. Defaults to None.
- train_batch_size (int, optional) Training batch size. Defaults to 32.
- eval_batch_size (int, optional) Test batch size. Defaults to 32.
- **num_workers** (*int, optional*) Number of workers. Defaults to [8].
- **task** (*TaskType*, *optional*) Task type. Could be classification, detection or segmentation. Defaults to TaskType.SEGMENTATION.
- **image_size** (*tuple[int, int], optional*) Size to which input images should be resized. Defaults to None.
- **transform** (*Transform, optional*) Transforms that should be applied to the input images. Defaults to None.
- train transform (Transform ontional) Transforms that should be applied to

- the input images during training. Defaults to None.
- **eval_transform** (*Transform, optional*) Transforms that should be applied to the input images during evaluation. Defaults to None.
- **test_split_mode** (<u>TestSplitMode</u>) Setting that determines how the testing subset is obtained. Defaults to <u>TestSplitMode.FROM_DIR</u>.
- **test_split_ratio** (*float*) Fraction of images from the train set that will be reserved for testing. Defaults to 0.2.
- val_split_mode (<u>ValSplitMode</u>) Setting that determines how the validation subset is obtained. Defaults to <u>ValSplitMode.FROM_TEST</u>.
- val_split_ratio (*float*) Fraction of train or test images that will be reserved for validation. Defaults to 0.5.
- **seed** (*int* | *None*, *optional*) Seed used during random subset splitting. Defaults to None.

property name: str

Name of the datamodule.

Folder3D datamodule overrides the name property to provide a custom name.

class anomalib.data.depth.folder_3d.Folder3DDataset(name, task,
normal_dir, root=None, abnormal_dir=None, normal_test_dir=None,
mask_dir=None, normal_depth_dir=None, abnormal_depth_dir=None,
normal_test_depth_dir=None, transform=None, split=None, extensions=None)

Bases: AnomalibDepthDataset

Folder dataset.

Parameters:

- **name** (*str*) Name of the dataset.
- task (*TaskType*) Task type. (classification, detection or segmentation).
- **transform** (*Transform, optional*) Transforms that should be applied to the input images.
- **normal_dir** (*str* | *Path*) Path to the directory containing normal images.
- **root** (*str* | *Path* | *None*) Root folder of the dataset. Defaults to [None].
- **abnormal_dir** (*str* | *Path* | *None, optional*) Path to the directory containing abnormal images. Defaults to None.
- **normal_test_dir** (*str* | *Path* | *None, optional*) Path to the directory containing normal images for the test dataset. Defaults to None.
- mask_dir (str | Path | None, optional) Path to the directory containing the mask annotations. Defaults to None.
- **normal_depth_dir** (*str* | *Path* | *None*, *optional*) Path to the directory containing normal depth images for the test dataset. Normal test depth images will be a split of *normal_dir* Defaults to None.
- **abnormal_depth_dir** (*str* | *Path* | *None, optional*) Path to the directory containing abnormal depth images for the test dataset. Defaults to None.
- **normal_test_depth_dir** (*str* | *Path* | *None*, *optional*) Path to the directory containing normal depth images for the test dataset. Normal test images will be a split of *normal_dir* if *None*. Defaults to None.
- **transform** Transforms that should be applied to the input images. Defaults to None.
- **split** (*str* | *Split* | *None*) Fixed subset split that follows from folder structure on file system. Choose from [Split.FULL, Split.TRAIN, Split.TEST] Defaults to None.
- **extensions** (*tuple[str, ...]* | *None, optional*) Type of the image extensions to read from the directory. Defaults to None.

Raises:

ValueError – When task is set to classification and *mask_dir* is provided. When *mask_dir* is provided, *task* should be set to *segmentation*.

property name: str

Name of the dataset.

Folder3D dataset overrides the name property to provide a custom name.

```
anomalib.data.depth.folder_3d.make_folder3d_dataset(normal_dir,
root=None, abnormal_dir=None, normal_test_dir=None, mask_dir=None,
normal_depth_dir=None, abnormal_depth_dir=None,
normal_test_depth_dir=None, split=None, extensions=None)
```

Make Folder Dataset.

Parameters:

- **normal_dir** (*str* | *Path*) Path to the directory containing normal images.
- **root** (*str* | *Path* | *None*) Path to the root directory of the dataset. Defaults to None.
- **abnormal_dir** (*str* | *Path* | *None*, *optional*) Path to the directory containing abnormal images. Defaults to None.
- **normal_test_dir** (*str* | *Path* | *None, optional*) Path to the directory containing normal images for the test
- **None.** (dataset. Normal test images will be a split of normal_dir if) Defaults to None.
- mask_dir (str | Path | None, optional) Path to the directory containing the mask annotations. Defaults to None.
- **normal_depth_dir** (*str* | *Path* | *None, optional*) Path to the directory containing normal depth images for the test dataset. Normal test depth images will be a split of *normal_dir* Defaults to None.
- **abnormal_depth_dir** (*str* | *Path* | *None, optional*) Path to the directory containing abnormal depth images for the test dataset. Defaults to None.
- **normal_test_depth_dir** (*str* | *Path* | *None*, *optional*) Path to the directory containing normal depth images for the test dataset. Normal test images will be a split of *normal_dir* if *None*. Defaults to None.
- **split** (*str* | *Split* | *None*, *optional*) Dataset split (ie., Split.FULL, Split.TRAIN or Split.TEST). Defaults to None.
- **extensions** (*tuple[str, ...]* | *None, optional*) Type of the image extensions to read from the directory. Defaults to None.

Returns:

an output dataframe containing samples for the requested split (ie., train or test)

Return type:

DataFrame

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