

code review
[Preprocessing]

git 구성

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YL-428 train code merge project commit

b6bcd8 20 minutes ago

History

..



2.DataPrepare

train code merge project commit

20 minutes ago



3.DetectModel

필요없는 파일 정리

21 days ago



4.ClassifyModel/classification_EfficientNet

train code merge project commit

20 minutes ago



json_test

first

27 days ago



yolo_eff_train

train code merge project commit

20 minutes ago

Pre-Processing protocol

<2.DataPrepare> - Data Preprocessing

- **make_json_real.py** (dicom metadata와 data제공 csv를 각각 json으로 변환하여 하나의 json으로 병합하여 저장)
- **make_txt_c18.py** (json to txt convert for detection-YOLOv5)
- **crop_make_folder.py** (ROI crop & foldering image by class for classification-EfficientNetB4)

Dataset: SIIM COVID-19 Detection



- 약 7000(train-6000 / test-1000)장의 dicom 폐 이미지
- class 및 bounding box 정보를 담은 csv
- 4개의 class
(Negative for Pneumonia / Typical Appearance / Indeterminate Appearance / Atypical Appearance)

Pre-Processing; make_json_real.py

- dicom metadata와 data제공 csv를 각각 json으로 변환하여 하나의 json으로 병합하여 저장

- dataset 제공 csv

	A	B	C	D
1	StudyInstanceUID	imageID	boxes	label
6023	f26c50ef1bd0	9745a7b0789a_image	[[{'x': 663.25938, 'y': 995.39205, 'width': 886.33514, 'height': 1227.3858}], opacity 1	663.25938 995.39205 1549.59452 2222.7
6024	f29d4d4eb110	615950e322f0_image	[[{'x': 1812.06709, 'y': 555.68, 'width': 781.17333, 'height': 1457.65333}], opacity 1	1812.06709 555.68 2593.24042 2013.333
6025	f2a899251da3	e1b500e2ea12_image	[[{'x': 482.32795, 'y': 393.82, 'width': 917.56, 'height': 1485.96}], opacity 1	482.32795 393.82 1399.8879499999998
6026	f2a9e6e67eda	8438a6e0c60b_image	[[{'x': 2569.04277, 'y': 1221.87374, 'width': 949.89819, 'height': 790.14246}], opacity 1	2569.04277 1221.87374 3518.94096 201
6027	f2b77c3c70c5	1c1069e57757_image	none	1 0 0 1 1
6028	f2b77c3c70c5	bdd3115879aa_image	none	1 0 0 1 1
6029	f2bf2f65fe68	57c6e5c6f17a_image	[[{'x': 1018.49287, 'y': 1878.35439, 'width': 775.78815, 'height': 801.79236}], opacity 1	1018.49287 1878.35439 1794.28102 268
6030	f2d30ac3777b	8b69eac9fa7b_image	[[{'x': 1732.04784, 'y': 1367.93613, 'width': 845.30566, 'height': 524.36072}], opacity 1	1732.04784 1367.93613 2577.3535 1892
6031	f2d5dd65d7d2	fab0ab9d7d17_image	[[{'x': 175.4401, 'y': 879.37257, 'width': 527.23109, 'height': 667.64948}], opacity 1	175.4401 879.37257 702.67119 1547.022
6032	f2d890d8bea3	9b3d574321bf_image	[[{'x': 692.78252, 'y': 1550.99621, 'width': 339.9444, 'height': 269.12268}], opacity 1	692.78252 1550.99621 1032.72692 1820
6033	f2e430c3888c	4e1ed0596509_image	none	1 0 0 1 1
6034	f2e5da098e85	5a0ad476493b_image	[[{'x': 477.92002, 'y': 317.06667, 'width': 680.53333, 'height': 1407.46667}], opacity 1	477.92002 317.06667 1158.45335 1724.5

	E	F	G	H	I	J
StudyInstanceUID	studyID	Negative for Pneumonia	Typical Appearance	Indeterminate Appearance	Atypical Appearance	
f26c50ef1bd0	f26c50ef1bd0_study	0	0	0	1	
f29d4d4eb110	f29d4d4eb110_study	0	1	0	0	
f2a899251da3	f2a899251da3_study	0	1	0	0	
f2a9e6e67eda	f2a9e6e67eda_study	0	1	0	0	
f2b77c3c70c5	f2b77c3c70c5_study	1	0	0	0	
f2b77c3c70c5						
f2bf2f65fe68	f2bf2f65fe68_study	0	1	0	0	
f2d30ac3777b	f2d30ac3777b_study	0	0	0	1	
f2d5dd65d7d2	f2d5dd65d7d2_study	0	0	1	0	
f2d890d8bea3	f2d890d8bea3_study	0	0	1	0	
f2e430c3888c	f2e430c3888c_study	1	0	0	0	
f2e5da098e85	f2e5da098e85_study	0	1	0	0	
f2e15af6cad	f2e15af6cad_study	0	1	0	0	
f317383c5831	f317383c5831_study	0	0	1	0	
f32c2b029aca	f32c2b029aca_study	0	1	0	0	
f32dd337bd1a	f32dd337bd1a_study	1	0	0	0	

- dicom 메타데이터

```
D:\Anaconda\envs\colorectal\python.exe D:/final_colorectal_gastric_train/Train_Model/2.DataPrepare/hadling_json.py
Dataset.file_meta -----
(0002, 0002) Media Storage SOP Class UID          UI: Digital X-Ray Image Storage - For Presentation
(0002, 0003) Media Storage SOP Instance UID       UI: 1.2.840.113654.2.70.1.9822815699402501441770223933240180074
(0002, 0010) Transfer Syntax UID                  UI: Explicit VR Little Endian
-----
(0008, 0005) Specific Character Set                CS: 'ISO_IR 100'
(0008, 0008) Image Type                           CS: ['DERIVED', 'PRIMARY', '']
(0008, 001a) SOP Class UID                         UI: 71228e4340de
(0008, 0018) SOP Instance UID                      UI: 51759b55779bc
(0008, 0020) Study Date                           DA: '96fc21dd2b1f'
(0008, 0030) Study Time                           TM: '13e700cac7f0'
(0008, 0050) Accession Number                     SH: '0dc10cf540cf'
(0008, 0060) Modality                             CS: 'DX'
(0009, 0010) Private Creator                      LO: 'GEIIS'
(0010, 0010) Patient's Name                       PN: '3a0c965d2001'
(0010, 0020) Patient ID                           LO: '2c00dc1ead80'
(0010, 0040) Patient's Sex                         CS: 'M'
(0012, 0063) De-identification Method              LO: 'CTP Default: based on DICOM PS3.15 AnnexE. Details in 0012,0064'
(0012, 0064) De-identification Method Code Sequence & item(s) ----
(0008, 0100) Code Value                           SH: '113100'
(0008, 0102) Coding Scheme Designator              SH: 'DCM'
(0008, 0104) Code Meaning                           LO: 'Basic Application Confidentiality Profile'
-----
(0008, 0100) Code Value                           SH: '113105'
(0008, 0102) Coding Scheme Designator              SH: 'DCM'
(0008, 0104) Code Meaning                           LO: 'Clean Descriptors Option'
-----
(0008, 0100) Code Value                           SH: '113107'
(0008, 0102) Coding Scheme Designator              SH: 'DCM'
(0008, 0104) Code Meaning                           LO: 'Retain Longitudinal Temporal Information Modified Dates Option'
-----
(0008, 0100) Code Value                           SH: '113108'
(0008, 0102) Coding Scheme Designator              SH: 'DCM'
(0008, 0104) Code Meaning                           LO: 'Retain Patient Characteristics Option'
```




Pre-Processing; make_json_real.py

- dicom 메타데이터 / csv 정보 ⇒ 각각의 json으로 저장

```
def meta2json(dcm):  
    print("====start converting meta data to json====")  
    print("\n")  
    meta2json_list = list()  
    ds = pydicom.dcmread(dcm) # dicom파일 불러오기  
    #print("type(ds):", type(ds)) # metaData 출력  
    ds_json = ds.to_json_dict()  
    meta2json = dcm[:-4] + '_meta.json'  
    with open(meta2json, "w") as f: # json파일 저장위치  
        json.dump(ds_json, f)  
    #print("meta2json_list: ", meta2json_list, "\n")  
  
    return meta2json
```

```
def csv2json(csv_path, dcm_list):  
    print("====start converting csv data to json====")  
    print("\n")  
    train_df = pd.read_csv(csv_path + 'merged_csv.csv')  
    csv2json_list = list()  
    for i in range(train_df.shape[0]):  
        subj_series = train_df.loc[i] # (subj_series)type; pandas.core.series.Series  
        # print("=====subject", i, "=====")  
        # print("subj_series.index: ", subj_series.index) #['StudyInstanceUID', 'imageID', 'boxe'  
        # print("subj_series.values: ", subj_series.values, "\n")  
        # print(subj_series)  
        # print("=====\n")  
  
        subj_dict = subj_series.to_dict() # type; dict  
  
        # subj_json = json.dumps(subj_dict) #convert dictionary to json  
  
        cls_yn = subj_dict.get('Negative for Pneumonia')  
  
        if not pd.isna(cls_yn):  
            for k in range(len(dcm_list)):  
                if subj_dict.get('imageID')[:-6] in dcm_list[k]:  
                    path_split = dcm_list[k].split('\\')  
                    # print("path_split: ", path_split, "\n")  
                    save_path = path_split[:-1]  
                    # print("save_path: ", save_path, "\n")  
                    path_join = '/'.join(save_path)  
                    # print("path_join: ", path_join, "\n")  
                    csv2json_dir = path_join + '/' + subj_dict.get('imageID')[:-6] + '_csv.json'  
                    # print("csv2json_dir: ", csv2json_dir, "\n")  
  
                    with open(csv2json_dir, "w") as f: # json파일 저장위치  
                        json.dump(subj_dict, f)  
  
                    csv2json_list.append(csv2json_dir)  
  
    #print("csv2json_list: ", csv2json_list, "\n")  
    return csv2json_list
```

dataset > train > 00c241c3fc0d > eac6d8583ff9

이름	수정한 날짜	유형
 bcd2179fa24e	2021-05-13 오후 1:45	MicroDicom D
 bcd2179fa24e_csv	2021-11-10 오후 6:45	JSON 파일
 bcd2179fa24e_meta	2021-11-10 오후 5:58	JSON 파일

Pre-Processing; make_json_real.py

- 각 json에서 필요한 정보만 추출

```
def get_meta2json(meta2json):
```

```
    print("====start getting meta2json info====")
    with open(meta2json, 'r', encoding='UTF8') as f:
        meta_json = json.load(f, strict=False) # type; dictionary
        # print(type(meta_json))
```

```
    Modality = meta_json.get("00080060") # type; dictionary
```

```
    Modality = Modality.get("Value")
```

```
    Modality = Modality[0]
```

```
    Patient_ID = meta_json.get("00100020")
```

```
    Patient_ID = Patient_ID.get("Value")
```

```
    Patient_ID = Patient_ID[0]
```

```
    Sex = meta_json.get("00100040")
```

```
    Sex = Sex.get("Value")
```

```
    Sex = Sex[0]
```

```
    Study_Name = meta_json.get("00200000")
```

```
    Study_Name = Study_Name.get("Value")
```

```
    Study_Name = Study_Name[0]
```

```
    Series = meta_json.get("0020000E")
```

```
    Series = Series.get("Value")
```

```
    Series = Series[0]
```

```
    Dicom = meta_json.get("00080018")
```

```
    Dicom = Dicom.get("Value")
```

```
    Dicom = Dicom[0]
```

```
metajson = {'Modality': Modality, 'Patient_ID': Patient_ID, 'Sex': Sex, 'Study_Name': Study_Name,
            'Series': Series, 'Dicom': Dicom}
```

```
def get_csv2json(csv2json_list):
```

```
    print("====start getting csv2json info====")
```

```
    csvjson_list = list()
```

```
    for i in range(len(csv2json_list)):
```

```
        with open(csv2json_list[i], 'r', encoding='UTF8') as f:
```

```
            csv_json = json.load(f, strict=False) # type; dictionary
```

```
    Dicom_csv = csv_json.get("imageID")
```

```
    bbox = csv_json.get("boxes")
```

```
    label = csv_json.get("label")
```

```
    cls0 = csv_json.get("Negative for Pneumonia")
```

```
    if type(cls0) is str:
```

```
        cls0 = int(cls0)
```

```
    cls1 = csv_json.get("Typical Appearance")
```

```
    if type(cls1) is str:
```

```
        cls1 = int(cls1)
```

```
    cls2 = csv_json.get("Indeterminate Appearance")
```

```
    if type(cls2) is str:
```

```
        cls2 = int(cls2)
```

```
    cls3 = csv_json.get("Atypical Appearance")
```

```
    if type(cls3) is str:
```

```
        cls3 = int(cls3)
```

```
    csvjson = {'Dicom': Dicom_csv[:-6], 'boxes': bbox, 'label': label, 'Negative_for_Pneumonia': cls0,
              'Typical_Appearance': cls1, 'Indeterminate_Appearance': cls2, 'Atypical_Appearance': cls3}
```

```
    # print("csvjson: ", csvjson)
```

```
    csvjson_list.append(csvjson)
```


Pre-Processing; make_json_real.py

- class 분류 추가 및 최종 병합된 json 저장

```
def write_cls(csv2json_list):
    print("====start writhing class====")
    cls_list = list()
    for i in csv2json_list:
        cls0 = i.get('Negative_for_Pneumonia')
        cls0 = int(cls0)
        cls1 = i.get('Typical_Appearance')
        cls1 = int(cls1)
        cls2 = i.get('Indeterminate_Appearance')
        cls2 = int(cls2)
        cls3 = i.get('Atypical_Appearance')
        cls3 = int(cls3)

        if cls0 == 1:
            cls0 = 0
            cls = {'class': cls0}
        elif cls1 == 1:
            cls1 = 1
            cls = {'class': cls1}
        elif cls2 == 1:
            cls2 = 2
            cls = {'class': cls2}
        else:
            cls3 = 3
            cls = {'class': cls3}

    cls_list.append(cls)
```

ad8d4a5ba8f0_merge

파일(F) 편집(E) 서식(O) 보기(V) 도움말(H)

```
{"images": [{"Modality": "CR", "Patient_ID": "4df77458d988", "Sex": "F", "Study_Name": "00a76543ed93", "Series": "4a223cccbe04", "Dicom": "ad8d4a5ba8f0",
"boxes": "[['x': 639.68044, 'y': 0, 'width': 902.77729, 'height': 1840.64232]]", "label": "opacity 1 639.68044 0 1542.45773 1840.64232",
"Negative_for_Pneumonia": 0.0, "Typical_Appearance": 0.0, "Indeterminate_Appearance": 1.0, "Atypical_Appearance": 0.0, "class": 2}]}
```

```
def merge_json(csv2json_list, metajson, csvjson_list, cls_list):
    print("====start merging jsons====")
    for i in range(len(csvjson_list)):
        if metajson.get("Dicom") == csvjson_list[i].get("Dicom"):
            aa = metajson
            bb = csvjson_list[i]
            cc = cls_list[i]

            # match = csvjson_list[i].get("Dicom_csv")
            # print(metajson_list[i].get("Dicom"))
            # print(csvjson_list[i].get("Dicom"))

            dd = dict(bb, **cc)
            merge = dict(aa, **dd)
            merge = {"images": [merge]}
            # img = json.get('images')

            dir = csv2json_list[i][:-8] + 'merge.json'
            #print("dir:", dir)
            #print("\n")
            with open(dir, "w") as f: # json파일 저장위치
                json.dump(merge, f)
            print("final_json:", merge)
            print(type(merge))
            print("\n")
```


Pre-Processing; make_txt_c18.py

- json to txt convert for detection-YOLOv5

- json에서 class와 bbox 정보를 추출하여 txt로 저장

```
def make_annot_list(json_data, point_list):
    annot_list = list()
    if len(point_list) == 0:
        point = get_stage(json_data) + " " + str('none')
        annot_list.append(point)
    else:
        for i in range(len(point_list)):
            point = get_stage(json_data) + " " + point_list[i]
            #print("point:" + point)
            annot_list.append(point)

    return annot_list

def save_txt(save_path, filename, annot_list):
    print("Start creating text files...")
    #with open(folder_name, 'w') as f:
    with open(save_path + "\\\" + filename, 'w') as f:
        f.write('\n'.join(annot_list))
    f.close()
```

09cf9767a7bf - Windows 메모장

파일(F) 편집(E) 서식(O) 보기(V) 도움말(H)

```
1 0.7384837093321919 0.30826202045133994 0.21402918236301371 0.31075698871650215
1 0.3424693386130137 0.14939746121297603 0.20072788099315067 0.1862549717912553
1 0.30800700770547945 0.4108516502115656 0.19468190068493152 0.13545816643159378
```

- dicom to png 이미지 형식 변환

```
import mritopng
mritopng.convert_file(dcm_path, png_path)
```

mritopng 라이브러리 __init__.py

```
def convert_file(mri_file_path, png_file_path, auto_contrast=False):
    """ Function to convert an MRI binary file to a
    PNG image file.

    @param mri_file_path: Full path to the mri file
    @param png_file_path: Full path to the png file
    """

    # Making sure that the mri file exists
    if not os.path.exists(mri_file_path):
        raise Exception('Source file "%s" does not exists' % mri_file_path)

    # Making sure the png file does not exist
    if os.path.exists(png_file_path):
        print('Removing existing output file %s' %
              png_file_path)
        os.remove(png_file_path)

    mri_file = open(mri_file_path, 'rb')
    png_file = open(png_file_path, 'wb')

    mri_to_png(mri_file, png_file, auto_contrast)

    png_file.close()

def mri_to_png(mri_file, png_file, do_auto_contrast=False):
    """ Function to convert from a DICOM image to png

    @param mri_file: An opened file like object to read the dicom data
    @param png_file: An opened file like object to write the png data
    """

    image_2d = extract_grayscale_image(mri_file)

    if do_auto_contrast:
        image_2d = auto_contrast(image_2d)

    # Writing the PNG file
    w = png.Writer(image_2d.width, image_2d.height, grayscale=True)
    w.write(png_file, image_2d.image)
```

Pre-Processing; crop_make_folder.py

- ROI crop & foldering image by class for classification-EfficientNetB4

- ROI crop & 검은 배경에 센터 맞춰 붙이기

```
def crop_merge_img(img, line):
    global back_resize

    back_dir = 'back.png'
    back = cv2.imread(back_dir)
    img_size = img.shape

    if not line[1] == 'none':
        center_x = float(line[1])
        center_y = float(line[2])
        width = float(line[3])
        height = float(line[4])
        #print(center_x, center_y, width, height)
        #print("\n")

        #crop이미지
        x1 = int((center_x) * img_size[0])
        x2 = int((center_x + width) * img_size[0])
        w = x2 - x1
        print("x1", x1, "x2", x2)
        print("w", w)

        y1 = int((center_y) * img_size[1])
        y2 = int((center_y + height) * img_size[1])
        h = y2 - y1
        print("y1", y1, "y2", y2)
        print("h", h)

        # merge_img
        roi = img[y1:y2, x1:x2]
        print("roi.shape", roi.shape)

        if h > w:
            back_resize = cv2.resize(back, dsize=(h, h), interpolation=cv2.INTER_AREA)
            back_resize[0:h, int(h / 2 - w / 2):int(h / 2 + w / 2)] = roi
        else:
            back_resize = cv2.resize(back, dsize=(w, w), interpolation=cv2.INTER_AREA)
            back_resize[int(w / 2 - h / 2):int(w / 2 + h / 2), 0:w] = roi

    else:
        back_resize = img

    return back_resize
```

- class에 맞춰 foldering

```
for c in range(0, num_of_class):
    d = ""+str(c)
    path = os.path.join(dst_folder, d)
    #print(path)
    #os.mkdir(path)
    try:
        os.mkdir(path)
    except FileExistsError:
        pass

    image_list = os.listdir(image_path2)
    #print("image_list ", image_list)
    label_list = os.listdir(label_path2)
    #print("label_list ", label_list)

    for i in range(len(label_list)):
        image_path3 = image_path2 + label_list[i][:-4] + ".png"
        print("image_path3 ", image_path3)
        label_path3 = label_path2 + label_list[i]
        print("label_path3 ", label_path3)

        img = cv2.imread(image_path3)
        lines = open(label_path3).readlines()
        print("lines ", lines)

        uniq = 0
        for line in lines:
            line = line.split()
            # print("line ", line)
            for j in range(0, len(line)):
                # cls = coord[0]
                # print("cls ", cls)

                final = crop_merge_img(img, line)
                img_name = label_list[i][:-4]

                image_dd = dst_folder + "/" + str(int(line[0])) + "/" + img_name + ".png"

            if os.path.isfile(image_dd):
                uniq += 1
                img_name = img_name + '(' + str(uniq) + ')'
                image_dd = dst_folder + "/" + str(int(line[0])) + "/" + img_name + ".png"
```

Pre-Processing; crop_make_folder.py

- 이미지 처리 결과

