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    "import os\n",
    "import shutil\n",
    "import matplotlib.pyplot as plt"
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     "Copying files: 3660 files [01:27, 41.83 files/s] \n"
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    "#input folder=\"CT-KIDNEY-DATASET-Normal-Cyst-Tumor-Stone\"\n",
    "#splitfolders.ratio(input folder,output=\"CT-KIDNEY-DATASET-Normal-
Cyst-Tumor-Stone/train-test-
split'', seed=42, ratio=(0.7, 0.2, 0.1), group prefix=None)"
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```
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    "train path = r\"C:\\Users\\91995\\Major project\\CT-KIDNEY-DATASET-
Normal-Cyst-Tumor-Stone\\train-test-split\\train\\\n\\,
    "valid path = r\"C:\\Users\\91995\\Major project\\CT-KIDNEY-DATASET-
Normal-Cyst-Tumor-Stone\\train-test-split\\val\\"\n",
    "test path = r\"C:\\Users\\91995\\Major project\\CT-KIDNEY-DATASET-
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    "from keras.applications.densenet import preprocess input,
DenseNet121\n",
    "from keras.models import Model\n",
    "from keras.layers import Dense, MaxPool2D, Conv2D\n",
    "import keras"
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                                          zoom range= 0.2, \n",
    "
                                          horizontal flip= True, \n",
    "
                                          shear range= 0.2, \n",
    "
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    "train = train data gen.flow from directory(directory= train path,
\n",
                                                 target size=(224,224))"
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```
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target size=(224,224))"
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    "class type = {0:'Stone', 1 : 'Tumor'}"
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    "from keras.layers import Flatten , Dense, Dropout , MaxPool2D"
```

```
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include top will "
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   "for layer in res.layers:
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   "x = Flatten()(res.output)\n",
   "x = Dense(units=2 , activation='sigmoid', name = 'predictions'
)(x)\n",
   "\n",
   "# creating our model.\n",
   "model = Model(res.input, x)"
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Connected to
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\n",
\n",
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                                         (None, 112, 112, 64 0
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```
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```

```
" ization)
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```

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     "_ization)
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```
" ization)
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```
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     " n)
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\n",
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\n'',
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['conv3 block5 1 conv[0][0]'] \n",
      " ization)
\n",
```

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**
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     "<sup>-</sup>n)
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\n",
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\n",
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\n",
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                                                                36864
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    "from keras.callbacks import ModelCheckpoint\n",
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patience= 3, verbose=1) \n",
    "mc = ModelCheckpoint(filepath=\"bestmodel.h5\",
monitor=\"val_accuracy\", verbose=1, save_best_only= True)"
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to bestmodel.h5\n",
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6.4259 - accuracy: 0.5531 - val loss: 4.7980 - val accuracy: 0.6484\n",
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    accuracy: 0.7656\n",
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    accuracy: 0.8512\n",
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model to bestmodel.h5\n",
    1.1805 - accuracy: 0.8512 - val loss: 1.1290 - val accuracy: 0.8633\n",
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accuracy: 0.8824\n",
    "Epoch 4: val accuracy improved from 0.86328 to 0.90234, saving
model to bestmodel.h5\n",
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    accuracy: 0.9469\n",
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model to bestmodel.h5\n",
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model to bestmodel.h5\n",
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model to bestmodel.h5\n",
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0.1762 - accuracy: 0.9500 - val loss: 0.2382 - val accuracy: 0.9668\n",
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    accuracy: 0.9375\n",
    "Epoch 8: val accuracy improved from 0.96680 to 0.97070, saving
model to bestmodel.h5\n",
    0.3667 - accuracy: 0.9375 - val_loss: 0.1368 - val_accuracy: 0.9707\n",
    "Epoch 9/30\n",
    accuracy: 0.9312\n",
    "Epoch 9: val accuracy did not improve from 0.97070\n",
    0.3020 - accuracy: 0.9312 - val loss: 0.3666 - val accuracy: 0.9336\n",
    "Epoch 10/30\n",
    accuracy: 0.9594\n",
    "Epoch 10: val accuracy did not improve from 0.97070\n",
    0.1622 - accuracy: 0.9594 - val loss: 0.2072 - val accuracy: 0.9355\n",
    "Epoch 10: early stopping\n"
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validation data= valid , validation steps= 16, callbacks=[es,mc])"
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version. Please use `Model.evaluate`, which supports generators.\n",
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doing imports until\n"
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       Input : Takes in image path as input \n",
      Output: Gives out Pre-Processed image\n",
      \"\"\"\n",
      path = img path\n'',
      img = image.load img(path, target size=(224,224,3)) \n",
      img = image.img to array(img) \n",
      img = np.expand dims(img , axis= 0 )\n",
      \n",
    **
      return img"
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\{model.predict(img)[0][1]*100\}\ percent'")\n",
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    "\n",
    "# to display the image \n",
    "plt.imshow(img[0]/255, cmap = \"gray\")n",
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