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    "from sklearn.neighbors import KNeighborsClassifier\n",
    "from sklearn.svm import SVC\n",
    "from sklearn import model_selection\n",
    "from sklearn.metrics import classification_report\n",
    "from sklearn.metrics import accuracy_score\n",
    "from pandas.plotting import scatter_matrix\n",
    "import matplotlib.pyplot as plt\n",
    "import pandas as pd\n",
    "from sklearn.model_selection import train_test_split"
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    "url=\"https://archive.ics.uci.edu/ml/machine-learning-databases/breast-cancer-wisconsin/breast-cancer-wisconsin.data\"\n",
    "names= ['id', 'clump_thickness', 'uniform_cell_size', 'uniform_cell_shape',\n",
    "         'marginal_adhesion', 'single_epithelial_size', 'bare_nuclei',\n",
    "         'bland_chromatin', 'normal_nucleoli', 'mitoses', 'class']\n",

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"dataset=pd.read_csv(url,names=names)\n",
"dataset.head()"
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"3 1016277          6          8          8 \n",
"4 1017023          4          1          1 \n",
"\n",
"      marginal_adhesion single_epithelial_size bare_nuclei bland_chromatin \\n",
"0          1          2          1          3 \n",
"1          5          7         10          3 \n",
"2          1          2          2          3 \n",
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"\n",
"      normal_nucleoli mitoses class \n",
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" vertical-align: middle;\n",
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"</div>\n",
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a925-4bcf-8ccd-9812cd2176f4')\">\n",
"    title=\"Convert this dataframe to an interactive table.\"\n",
"    style=\"display:none;\">\n",
"  \n",
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"    width=\"24px\">\n",
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"    <path d=\"M18.5 5.441 9.4 2.06 9.4-2.06 2.06-9.4 2.06-.94 2.06-.94
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```

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2.06.94-2.06 2.06-.94-2.06-.94-.94-2.06-.94 2.06-2.06.94z\"/><path d=\"M17.41 7.96l-1.37-
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2.83L4 21.41c.39.39.95.59 1.41.59.51 0 1.02-.2 1.41-.59l7.78-7.78 2.81-2.81c.8-.78.8-2.07 0-
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  "   display: none;\n",
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  "   width: 32px;\n",
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  " .colab-df-convert:hover {\n",
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0.15);\n",
  "   fill: #174EA6;\n",
  " } \n",
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  "   fill: #D2E3FC;\n",
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  "\n",
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  "   background-color: #434B5C;\n",
  "   box-shadow: 0px 1px 3px 1px rgba(0, 0, 0, 0.15);\n",
  "   filter: drop-shadow(0px 1px 2px rgba(0, 0, 0, 0.3));\n",
  "   fill: #FFFFFF;\n",
  " } \n",
  " </style>\n",
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  " <script>\n",
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button.colab-df-convert');\n",

```

```

        buttonEl.style.display =\n",
        google.colab.kernel.accessAllowed ? 'block' : 'none';\n",
        "\n",
        async function convertToInteractive(key) {\n",
        const element = document.querySelector('#df-34551bb8-a925-4bcf-8ccd-
9812cd2176f4');\n",
        const dataTable =\n",
        await google.colab.kernel.invokeFunction('convertToInteractive',\n",
        [key], {});\n",
        if (!dataTable) return;\n",
        "\n",
        const docLinkHtml = 'Like what you see? Visit the ' +\n",
        '<a target="_blank"
href=https://colab.research.google.com/notebooks/data_table.ipynb>data table
notebook</a>\n",
        + ' to learn more about interactive tables.';\n",
        element.innerHTML = ";\n",
        dataTable['output_type'] = 'display_data';\n",
        await google.colab.output.renderOutput(dataTable, element);\n",
        const docLink = document.createElement('div');\n",
        docLink.innerHTML = docLinkHtml;\n",
        element.appendChild(docLink);\n",
        }\n",
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[illegible]

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      "single_epithelial_size 1\n",
      "bare_nuclei           1\n",
      "bland_chromatin       3\n",
      "normal_nucleoli       1\n",
      "mitoses               1\n",
      "class                 2\n",
      "Name: 10, dtype: object\n",
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        "std         2.815741           3.051459           2.971913  \n",
        "min          1.000000           1.000000           1.000000  \n",
        "25%          2.000000           1.000000           1.000000  \n",
        "50%          4.000000           1.000000           1.000000  \n",
        "75%          6.000000           5.000000           5.000000  \n",
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```



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"25%        1.000000        2.000000        2.000000 \n",
"50%        1.000000        2.000000        3.000000 \n",
"75%        4.000000        4.000000        5.000000 \n",
"max        10.000000       10.000000       10.000000 \n",
"\n",
"    normal_nucleoli  mitoses    class \n",
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"std        3.053634   1.715078   0.951273 \n",
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"max        10.000000  10.000000   4.000000 \n"
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