

```

{
  "cells": [
    {
      "cell_type": "markdown",
      "metadata": {
        "id": "K6aUpDf7KC_p"
      },
      "source": [
        "# Comprensión de los Datos"
      ]
    },
    {
      "cell_type": "code",
      "execution_count": 2,
      "metadata": {
        "id": "Xhds4njPKC_u"
      },
      "outputs": [],
      "source": [
        "#importa librerías\n",
        "import pandas as pd"
      ]
    },
    {
      "cell_type": "markdown",
      "metadata": {
        "id": "JdYf-8XyKC_x"
      },
      "source": [
        "# Descripción de Variables"
      ]
    },
    {
      "cell_type": "markdown",
      "metadata": {
        "id": "Y7EVetS8KC_y"
      },
      "source": [
        "\n",
        "Pclass Passenger Class (1 = 1st; 2 = 2nd; 3 = 3rd): Categórica Nominal \n",
        "survival Survival (0 = No; 1 = Yes) \n",
        "name Name \n",
        "sex Sex \n",
        "age Age \n",
        "sibsp Number of Siblings/Spouses Aboard \n",
        "parch Number of Parents/Children Aboard \n",
        "ticket Ticket Number \n",
        "fare Passenger Fare (British pound) \n",
        "cabin Cabin \n",
        "embarked Port of Embarkation (C = Cherbourg; Q = Queenstown; S = Southampton) \n",
        "boat Lifeboat \n",
        "body Body Identification Number \n",
        "home.dest Home/Destination"
      ]
    },
    {
      "cell_type": "markdown",
      "metadata": {
        "id": "B68geAa3KC_z"
      },
      "source": [
        "***Ejemplo:** Crear un objeto DataFrame con base en un archivo .csv (poner \"df = \")"
      ]
    }
  ]
}

```

```

{
  "cell_type": "code",
  "execution_count": 3,
  "metadata": {
    "id": "dLr3UCdTKC_1"
  },
  "outputs": [],
  "source": [
    "#lee archivo csv\n",
    "df = pd.read_csv(\"diabetes.csv\")"
  ]
},
{
  "cell_type": "code",
  "execution_count": 5,
  "metadata": {},
  "outputs": [
    {
      "data": {
        "text/plain": [
          "(768, 9)"
        ]
      },
      "execution_count": 5,
      "metadata": {},
      "output_type": "execute_result"
    }
  ],
  "source": [
    "#Usa función shape para revisar el total de renglones y columnas\n",
    "df.shape"
  ]
},
{
  "cell_type": "code",
  "execution_count": 6,
  "metadata": {
    "id": "NLa_otfWKC_1",
    "outputId": "8ce7dd9a-5f3d-4fe7-d34f-ee8a02afda94",
    "scrolled": true
  },
  "outputs": [
    {
      "data": {
        "text/html": [
          "<div>\n",
          "<style scoped>\n",
          "  .dataframe tbody tr th:only-of-type {\n",
          "    vertical-align: middle;\n",
          "  }\n",
          "\n",
          "  .dataframe tbody tr th {\n",
          "    vertical-align: top;\n",
          "  }\n",
          "\n",
          "  .dataframe thead th {\n",
          "    text-align: right;\n",
          "  }\n",
          "</style>\n",
          "<table border=\"1\" class=\"dataframe\">\n",
          "  <thead>\n",
          "    <tr style=\"text-align: right;\">\n",
          "      <th></th>\n",
          "      <th>Pregnancies</th>\n",
          "      <th>Glucose</th>\n",

```

```

"      <th>BloodPressure</th>\n",
"      <th>SkinThickness</th>\n",
"      <th>Insulin</th>\n",
"      <th>BMI</th>\n",
"      <th>DiabetesPedigreeFunction</th>\n",
"      <th>Age</th>\n",
"      <th>Outcome</th>\n",
"    </tr>\n",
"  </thead>\n",
"  <tbody>\n",
"    <tr>\n",
"      <th>0</th>\n",
"      <td>6</td>\n",
"      <td>148</td>\n",
"      <td>72</td>\n",
"      <td>35</td>\n",
"      <td>0</td>\n",
"      <td>33.6</td>\n",
"      <td>0.627</td>\n",
"      <td>50</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>1</th>\n",
"      <td>1</td>\n",
"      <td>85</td>\n",
"      <td>66</td>\n",
"      <td>29</td>\n",
"      <td>0</td>\n",
"      <td>26.6</td>\n",
"      <td>0.351</td>\n",
"      <td>31</td>\n",
"      <td>0</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>2</th>\n",
"      <td>8</td>\n",
"      <td>183</td>\n",
"      <td>64</td>\n",
"      <td>0</td>\n",
"      <td>0</td>\n",
"      <td>23.3</td>\n",
"      <td>0.672</td>\n",
"      <td>32</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>3</th>\n",
"      <td>1</td>\n",
"      <td>89</td>\n",
"      <td>66</td>\n",
"      <td>23</td>\n",
"      <td>94</td>\n",
"      <td>28.1</td>\n",
"      <td>0.167</td>\n",
"      <td>21</td>\n",
"      <td>0</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>4</th>\n",
"      <td>0</td>\n",
"      <td>137</td>\n",
"      <td>40</td>\n",
"      <td>35</td>\n",
"      <td>168</td>\n",

```

```

"      <td>43.1</td>\n",
"      <td>2.288</td>\n",
"      <td>33</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"  </tbody>\n",
"</table>\n",
"</div>"
],
"text/plain": [
"  Pregnancies  Glucose  BloodPressure  SkinThickness  Insulin   BMI   \\\n",
"0            6     148             72           35         0  33.6   \n",
"1            1      85             66           29         0  26.6   \n",
"2            8     183             64            0         0  23.3   \n",
"3            1      89             66           23        94  28.1   \n",
"4            0     137             40           35       168  43.1   \n",
"\n",
"  DiabetesPedigreeFunction  Age  Outcome  \n",
"0                0.627    50         1  \n",
"1                0.351    31         0  \n",
"2                0.672    32         1  \n",
"3                0.167    21         0  \n",
"4                2.288    33         1  "
]
},
"execution_count": 6,
"metadata": {},
"output_type": "execute_result"
}
],
"source": [
"#Revisa los primeros 5 renglones del dataset usando la función head()\n",
"df.head()"
]
},
{
"cell_type": "code",
"execution_count": 7,
"metadata": {},
"outputs": [
{
"data": {
"text/html": [
"<div>\n",
"<style scoped>\n",
"  .dataframe tbody tr th:only-of-type {\n",
"    vertical-align: middle;\n",
"  }\n",
"\n",
"  .dataframe tbody tr th {\n",
"    vertical-align: top;\n",
"  }\n",
"\n",
"  .dataframe thead th {\n",
"    text-align: right;\n",
"  }\n",
"</style>\n",
"<table border=\"1\" class=\"dataframe\">\n",
"  <thead>\n",
"    <tr style=\"text-align: right;\">\n",
"      <th></th>\n",
"      <th>Pregnancies</th>\n",
"      <th>Glucose</th>\n",
"      <th>BloodPressure</th>\n",
"      <th>SkinThickness</th>\n",

```

```

"      <th>Insulin</th>\n",
"      <th>BMI</th>\n",
"      <th>DiabetesPedigreeFunction</th>\n",
"      <th>Age</th>\n",
"      <th>Outcome</th>\n",
"    </tr>\n",
"  </thead>\n",
"  <tbody>\n",
"    <tr>\n",
"      <th>763</th>\n",
"      <td>10</td>\n",
"      <td>101</td>\n",
"      <td>76</td>\n",
"      <td>48</td>\n",
"      <td>180</td>\n",
"      <td>32.9</td>\n",
"      <td>0.171</td>\n",
"      <td>63</td>\n",
"      <td>0</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>764</th>\n",
"      <td>2</td>\n",
"      <td>122</td>\n",
"      <td>70</td>\n",
"      <td>27</td>\n",
"      <td>0</td>\n",
"      <td>36.8</td>\n",
"      <td>0.340</td>\n",
"      <td>27</td>\n",
"      <td>0</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>765</th>\n",
"      <td>5</td>\n",
"      <td>121</td>\n",
"      <td>72</td>\n",
"      <td>23</td>\n",
"      <td>112</td>\n",
"      <td>26.2</td>\n",
"      <td>0.245</td>\n",
"      <td>30</td>\n",
"      <td>0</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>766</th>\n",
"      <td>1</td>\n",
"      <td>126</td>\n",
"      <td>60</td>\n",
"      <td>0</td>\n",
"      <td>0</td>\n",
"      <td>30.1</td>\n",
"      <td>0.349</td>\n",
"      <td>47</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>767</th>\n",
"      <td>1</td>\n",
"      <td>93</td>\n",
"      <td>70</td>\n",
"      <td>31</td>\n",
"      <td>0</td>\n",
"      <td>30.4</td>\n",
"      <td>0.315</td>\n",

```

```

"      <td>23</td>\n",
"      <td>0</td>\n",
"    </tr>\n",
"  </tbody>\n",
"</table>\n",
"</div>"
],
"text/plain": [
"      Pregnancies  Glucose  BloodPressure  SkinThickness  Insulin   BMI   \\\n",
"763             10      101             76           48        180  32.9   \n",
"764              2      122             70           27          0  36.8   \n",
"765              5      121             72           23        112  26.2   \n",
"766              1      126             60            0          0  30.1   \n",
"767              1       93             70           31          0  30.4   \n",
"\n",
"      DiabetesPedigreeFunction  Age  Outcome  \n",
"763                0.171      63          0  \n",
"764                0.340      27          0  \n",
"765                0.245      30          0  \n",
"766                0.349      47          1  \n",
"767                0.315      23          0  "
]
},
"execution_count": 7,
"metadata": {},
"output_type": "execute_result"
}
],
"source": [
"#Revisa los últimos 5 renglones del dataset usando la función tail()\n",
"df.tail()"
]
},
{
"cell_type": "code",
"execution_count": 8,
"metadata": {
"id": "YcQfUpcLKC_3",
"outputId": "4206bddf-7216-4cdc-f9d9-35500b1591bb"
},
"outputs": [
{
"name": "stdout",
"output_type": "stream",
"text": [
"<class 'pandas.core.frame.DataFrame'>\n",
"RangeIndex: 768 entries, 0 to 767\n",
"Data columns (total 9 columns):\n",
" #   Column                Non-Null Count  Dtype  \n",
"---  ---                ---
" 0   Pregnancies            768 non-null   int64  \n",
" 1   Glucose                768 non-null   int64  \n",
" 2   BloodPressure          768 non-null   int64  \n",
" 3   SkinThickness          768 non-null   int64  \n",
" 4   Insulin                768 non-null   int64  \n",
" 5   BMI                   768 non-null   float64\n",
" 6   DiabetesPedigreeFunction 768 non-null   float64\n",
" 7   Age                   768 non-null   int64  \n",
" 8   Outcome                768 non-null   int64  \n",
"dtypes: float64(2), int64(7)\n",
"memory usage: 54.1 KB\n"
]
}
]
},
"source": [

```

```

    "#Revisa la información mas completa del conjunto de datos usando la función info()\n",
    "#Muestra el total de datos, las columnas y su tipo correspondiente, dice si contiene nulos o
no\n",
    "df.info()"
  ]
},
{
  "cell_type": "code",
  "execution_count": 9,
  "metadata": {
    "id": "d70yk1VbKC_4",
    "outputId": "e7d340fd-be4a-4b6a-a205-18e7833aef88"
  },
  "outputs": [
    {
      "data": {
        "text/plain": [
          "Pregnancies      17\n",
          "Glucose          136\n",
          "BloodPressure    47\n",
          "SkinThickness    51\n",
          "Insulin          186\n",
          "BMI              248\n",
          "DiabetesPedigreeFunction  517\n",
          "Age              52\n",
          "Outcome          2\n",
          "dtype: int64"
        ]
      },
      "execution_count": 9,
      "metadata": {},
      "output_type": "execute_result"
    }
  ],
  "source": [
    "#revisa cuántos valores únicos tiene cada atributo del archivo usando la función nunique()\n",
    "df.nunique()"
  ]
},
{
  "cell_type": "markdown",
  "metadata": {
    "id": "8nTyC2dzKC_5"
  },
  "source": [
    "## Exploración de Datos"
  ]
},
{
  "cell_type": "code",
  "execution_count": 10,
  "metadata": {
    "id": "0CsuORYXKC_5",
    "outputId": "8425ef66-1f29-492c-81cd-f72230bad9d2"
  },
  "outputs": [
    {
      "data": {
        "text/html": [
          "<div>\n",
          "<style scoped>\n",
          "  .dataframe tbody tr th:only-of-type {\n",
          "    vertical-align: middle;\n",
          "  }\n",
          "\n"
        ]
      }
    }
  ]
}

```

```

"    .dataframe tbody tr th {\n",
"        vertical-align: top;\n",
"    }\n",
"\n",
"    .dataframe thead th {\n",
"        text-align: right;\n",
"    }\n",
"</style>\n",
"<table border=\"1\" class=\"dataframe\">\n",
"  <thead>\n",
"    <tr style=\"text-align: right;\">\n",
"      <th></th>\n",
"      <th>Pregnancies</th>\n",
"      <th>Glucose</th>\n",
"      <th>BloodPressure</th>\n",
"      <th>SkinThickness</th>\n",
"      <th>Insulin</th>\n",
"      <th>BMI</th>\n",
"      <th>DiabetesPedigreeFunction</th>\n",
"      <th>Age</th>\n",
"      <th>Outcome</th>\n",
"    </tr>\n",
"  </thead>\n",
"  <tbody>\n",
"    <tr>\n",
"      <th>count</th>\n",
"      <td>768.000000</td>\n",
"      <td>768.000000</td>\n",
"      <td>768.000000</td>\n",
"      <td>768.000000</td>\n",
"      <td>768.000000</td>\n",
"      <td>768.000000</td>\n",
"      <td>768.000000</td>\n",
"      <td>768.000000</td>\n",
"      <td>768.000000</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>mean</th>\n",
"      <td>3.845052</td>\n",
"      <td>120.894531</td>\n",
"      <td>69.105469</td>\n",
"      <td>20.536458</td>\n",
"      <td>79.799479</td>\n",
"      <td>31.992578</td>\n",
"      <td>0.471876</td>\n",
"      <td>33.240885</td>\n",
"      <td>0.348958</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>std</th>\n",
"      <td>3.369578</td>\n",
"      <td>31.972618</td>\n",
"      <td>19.355807</td>\n",
"      <td>15.952218</td>\n",
"      <td>115.244002</td>\n",
"      <td>7.884160</td>\n",
"      <td>0.331329</td>\n",
"      <td>11.760232</td>\n",
"      <td>0.476951</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>min</th>\n",
"      <td>0.000000</td>\n",
"      <td>0.000000</td>\n",
"      <td>0.000000</td>\n",

```



```

"      <td>0.000000</td>\n",
"      <td>0.000000</td>\n",
"      <td>0.000000</td>\n",
"      <td>0.078000</td>\n",
"      <td>21.000000</td>\n",
"      <td>0.000000</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>25%</th>\n",
"      <td>1.000000</td>\n",
"      <td>99.000000</td>\n",
"      <td>62.000000</td>\n",
"      <td>0.000000</td>\n",
"      <td>0.000000</td>\n",
"      <td>27.300000</td>\n",
"      <td>0.243750</td>\n",
"      <td>24.000000</td>\n",
"      <td>0.000000</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>50%</th>\n",
"      <td>3.000000</td>\n",
"      <td>117.000000</td>\n",
"      <td>72.000000</td>\n",
"      <td>23.000000</td>\n",
"      <td>30.500000</td>\n",
"      <td>32.000000</td>\n",
"      <td>0.372500</td>\n",
"      <td>29.000000</td>\n",
"      <td>0.000000</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>75%</th>\n",
"      <td>6.000000</td>\n",
"      <td>140.250000</td>\n",
"      <td>80.000000</td>\n",
"      <td>32.000000</td>\n",
"      <td>127.250000</td>\n",
"      <td>36.600000</td>\n",
"      <td>0.626250</td>\n",
"      <td>41.000000</td>\n",
"      <td>1.000000</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>max</th>\n",
"      <td>17.000000</td>\n",
"      <td>199.000000</td>\n",
"      <td>122.000000</td>\n",
"      <td>99.000000</td>\n",
"      <td>846.000000</td>\n",
"      <td>67.100000</td>\n",
"      <td>2.420000</td>\n",
"      <td>81.000000</td>\n",
"      <td>1.000000</td>\n",
"    </tr>\n",
"  </tbody>\n",
"</table>\n",
"</div>"
],
"text/plain": [
"      Pregnancies      Glucose      BloodPressure      SkinThickness      Insulin  \\n",
"count      768.000000      768.000000      768.000000      768.000000      768.000000  \n",
"mean        3.845052     120.894531        69.105469        20.536458        79.799479  \n",
"std         3.369578      31.972618        19.355807        15.952218       115.244002  \n",
"min         0.000000      0.000000         0.000000         0.000000         0.000000  \n",

```

```

"25%      1.000000    99.000000    62.000000    0.000000    0.000000    \n",
"50%      3.000000   117.000000    72.000000    23.000000    30.500000    \n",
"75%      6.000000   140.250000    80.000000    32.000000   127.250000    \n",
"max      17.000000  199.000000   122.000000   99.000000   846.000000    \n",
"\n",
"          BMI  DiabetesPedigreeFunction      Age      Outcome  \n",
"count  768.000000      768.000000  768.000000  768.000000  \n",
"mean    31.992578          0.471876   33.240885    0.348958  \n",
"std      7.884160          0.331329   11.760232    0.476951  \n",
"min      0.000000          0.078000   21.000000    0.000000  \n",
"25%     27.300000          0.243750   24.000000    0.000000  \n",
"50%     32.000000          0.372500   29.000000    0.000000  \n",
"75%     36.600000          0.626250   41.000000    1.000000  \n",
"max     67.100000          2.420000   81.000000    1.000000  "
]
},
"execution_count": 10,
"metadata": {},
"output_type": "execute_result"
}
],
"source": [
"#utiliza la función describe() para obtener estadística básica. se puede incluir -0\n",
"df.describe()"
]
},
{
"cell_type": "code",
"execution_count": 14,
"metadata": {
"id": "6FZFAnZjKC_6",
"outputId": "753497de-4abb-4bcb-d1b7-34275424396f"
},
"outputs": [
{
"data": {
"text/plain": [
"Pregnancies      0\n",
"Glucose           0\n",
"BloodPressure     0\n",
"SkinThickness     0\n",
"Insulin           0\n",
"BMI              0\n",
"DiabetesPedigreeFunction  0\n",
"Age              0\n",
"Outcome          0\n",
"dtype: int64"
]
},
"execution_count": 14,
"metadata": {},
"output_type": "execute_result"
}
],
"source": [
"#Revisa Valores nulos con funcion isnull().sum()\n",
"df.isnull().sum()"
]
},
{
"cell_type": "code",
"execution_count": 16,
"metadata": {},
"outputs": [

```

```

"data": {
  "text/plain": [
    "array([ 6,  1,  8,  0,  5,  3, 10,  2,  4,  7,  9, 11, 13, 15, 17, 12, 14])"
  ]
},
"execution_count": 16,
"metadata": {},
"output_type": "execute_result"
},
"source": [
  "#Revisar valores únicos por columna usando función unique(): nombre-columna.unique()\n",
  "df.Pregnancies.unique()"
]
},
{
  "cell_type": "code",
  "execution_count": 17,
  "metadata": {},
  "outputs": [
    {
      "data": {
        "text/plain": [
          "array([148,  85, 183,  89, 137, 116,  78, 115, 197, 125, 110, 168, 139,\n",
          "      189, 166, 100, 118, 107, 103, 126,  99, 196, 119, 143, 147,  97,\n",
          "      145, 117, 109, 158,  88,  92, 122, 138, 102,  90, 111, 180, 133,\n",
          "      106, 171, 159, 146,  71, 105, 101, 176, 150,  73, 187,  84,  44,\n",
          "      141, 114,  95, 129,  79,  62, 131, 112, 113,  74,  83, 136,\n",
          "      80, 123,  81, 134, 142, 144,  93, 163, 151,  96, 155,  76, 160,\n",
          "      124, 162, 132, 120, 173, 170, 128, 108, 154,  57, 156, 153, 188,\n",
          "      152, 104,  87,  75, 179, 130, 194, 181, 135, 184, 140, 177, 164,\n",
          "      91, 165,  86, 193, 191, 161, 167,  77, 182, 157, 178,  61,  98,\n",
          "      127,  82,  72, 172,  94, 175, 195,  68, 186, 198, 121,  67, 174,\n",
          "      199,  56, 169, 149,  65, 190])"
        ]
      },
      "execution_count": 17,
      "metadata": {},
      "output_type": "execute_result"
    }
  ],
  "source": [
    "#Revisar valores únicos por columna usando función unique(): nombre-columna.unique()\n",
    "df.Glucose.unique()"
]
},
{
  "cell_type": "code",
  "execution_count": 18,
  "metadata": {},
  "outputs": [
    {
      "data": {
        "text/plain": [
          "array([1, 0])"
        ]
      },
      "execution_count": 18,
      "metadata": {},
      "output_type": "execute_result"
    }
  ],
  "source": [
    "#Revisar valores únicos por columna usando función unique(): nombre-columna.unique()\n",
    "df.Outcome.unique()"
]

```

```

]
},
{
  "cell_type": "markdown",
  "metadata": {
    "id": "ae13SbMmKC_7"
  },
  "source": [
    "## Variables Cuantitativas\n",
    "\n",
    "### Medidas de tendencia central\n"
  ]
},
{
  "cell_type": "code",
  "execution_count": 20,
  "metadata": {
    "id": "vDNofiEZKC_7",
    "outputId": "63524fd6-ca06-484e-fa9e-0dbbf5be2d6d"
  },
  "outputs": [
    {
      "name": "stdout",
      "output_type": "stream",
      "text": [
        "Mean_pregnancies: 3.8450520833333335\n",
        "Median_pregnancies: 3.0\n",
        "Mode_pregnancies: 0    1\n",
        "Name: Pregnancies, dtype: int64\n"
      ]
    }
  ],
  "source": [
    "#Pregnancies\n",
    "#Se puede obtener la media, mediana y moda para\n",
    "mean_pregnancies = df['Pregnancies'].mean()\n",
    "median_pregnancies = df['Pregnancies'].median()\n",
    "mode_pregnancies = df['Pregnancies'].mode()\n",
    "print(\"Mean_pregnancies:\",mean_pregnancies)\n",
    "print(\"Median_pregnancies:\",median_pregnancies)\n",
    "print(\"Mode_pregnancies:\",mode_pregnancies)"
  ]
},
{
  "cell_type": "markdown",
  "metadata": {
    "id": "Bx0aUF1lKC_8"
  },
  "source": [
    "Conclusiones:\n",
    "El promedio de embarazos fue de 3 \n",
    "La cantidad de embarazos al centro es 3 \n",
    "La cantidad de embarazos más repetida fue de 0 y 1"
  ]
},
{
  "cell_type": "code",
  "execution_count": 22,
  "metadata": {
    "id": "vDNofiEZKC_7",
    "outputId": "63524fd6-ca06-484e-fa9e-0dbbf5be2d6d"
  },
  "outputs": [
    {
      "name": "stdout",

```

```

    "output_type": "stream",
    "text": [
        "Mean_glucose: 120.89453125\n",
        "Median_glucose: 117.0\n",
        "Mode_glucose: 0      99\n",
        "1      100\n",
        "Name: Glucose, dtype: int64\n"
    ]
},
{
    "source": [
        "#Glucose\n",
        "#Se puede obtener la media, mediana y moda para\n",
        "mean_glucose = df['Glucose'].mean()\n",
        "median_glucose = df['Glucose'].median()\n",
        "mode_glucose = df['Glucose'].mode()\n",
        "print(\"Mean_glucose:\",mean_glucose)\n",
        "print(\"Median_glucose:\",median_glucose)\n",
        "print(\"Mode_glucose:\",mode_glucose)"
    ]
},
{
    "cell_type": "markdown",
    "metadata": {
        "id": "Bx0aUF1lKC_8"
    },
    "source": [
        "Conclusiones: \n",
        "El promedio de glucosa fue de 120 \n",
        "La glucosa al centro es 117 \n",
        "La glucosa más repetida fue de 0, 1, 99 y 100"
    ]
},
{
    "cell_type": "code",
    "execution_count": 24,
    "metadata": {
        "id": "vDNofIEZKC_7",
        "outputId": "63524fd6-ca06-484e-fa9e-0dbbf5be2d6d"
    },
    "outputs": [
        {
            "name": "stdout",
            "output_type": "stream",
            "text": [
                "Mean_outcome: 0.3489583333333333\n",
                "Median_outcome: 0.0\n",
                "Mode_outcome: 0      0\n",
                "Name: Outcome, dtype: int64\n"
            ]
        }
    ],
    "source": [
        "#Outcome\n",
        "#Se puede obtener la media, mediana y moda para\n",
        "mean_outcome = df['Outcome'].mean()\n",
        "median_outcome = df['Outcome'].median()\n",
        "mode_outcome = df['Outcome'].mode()\n",
        "print(\"Mean_outcome:\",mean_outcome)\n",
        "print(\"Median_outcome:\",median_outcome)\n",
        "print(\"Mode_outcome:\",mode_outcome)"
    ]
},
{
    "cell_type": "markdown",

```

```

"metadata": {
  "id": "Bx0aUF1lKC_8"
},
"source": [
  "Conclusiones:\n",
  "El promedio de resultado fue de 0.3 \n",
  "El resultado al centro es 0 \n",
  "El resultado más repetido fue de 0"
]
},
{
  "cell_type": "markdown",
  "metadata": {},
  "source": [
    "Cabe mencionar que, en las discusiones en Kaggle, se menciona que el dato \"0\" significa que el paciente es sano y el dato \"1\" significa que el paciente tiene diabetes. Por lo tanto, la mayoría de los datos indican un paciente sano."
  ]
},
{
  "cell_type": "markdown",
  "metadata": {
    "id": "ru1HJsXHKC_9"
  },
  "source": [
    "# Variables Categóricas"
  ]
},
{
  "cell_type": "code",
  "execution_count": 21,
  "metadata": {
    "id": "hu2J0Q7NKC_9",
    "outputId": "b0509fc7-7e9d-4b5b-8fbf-0a9563c10d55"
  },
  "outputs": [
    {
      "data": {
        "text/plain": [
          "Pregnancies\n",
          "1      135\n",
          "0      111\n",
          "2      103\n",
          "3       75\n",
          "4       68\n",
          "5       57\n",
          "6       50\n",
          "7       45\n",
          "8       38\n",
          "9       28\n",
          "10      24\n",
          "11      11\n",
          "13      10\n",
          "12       9\n",
          "14       2\n",
          "15       1\n",
          "17       1\n",
          "Name: count, dtype: int64"
        ]
      },
      "execution_count": 21,
      "metadata": {},
      "output_type": "execute_result"
    }
  ]
},

```

```

"source": [
  "#Para conteo de cada valor en una columna, en orden descendente usar función value_counts():
\n",
  "# nombreDataframe.columna.value_counts()\n",
  "# nombreDataframe['columna'].value_counts()\n",
  "df.Pregnancies.value_counts()"
]
},
{
  "cell_type": "code",
  "execution_count": 25,
  "metadata": {
    "id": "hu2J0Q7NKC_9",
    "outputId": "b0509fc7-7e9d-4b5b-8fbf-0a9563c10d55"
  },
  "outputs": [
    {
      "data": {
        "text/plain": [
          "Glucose\n",
          "99      17\n",
          "100     17\n",
          "111     14\n",
          "129     14\n",
          "125     14\n",
          "      ..\n",
          "191      1\n",
          "177      1\n",
          "44       1\n",
          "62       1\n",
          "190      1\n",
          "Name: count, Length: 136, dtype: int64"
        ]
      },
      "execution_count": 25,
      "metadata": {},
      "output_type": "execute_result"
    }
  ],
  "source": [
    "#Para conteo de cada valor en una columna, en orden descendente usar función value_counts():
\n",
    "# nombreDataframe.columna.value_counts()\n",
    "# nombreDataframe['columna'].value_counts()\n",
    "df.Glucose.value_counts()"
  ]
},
{
  "cell_type": "code",
  "execution_count": 26,
  "metadata": {
    "id": "hu2J0Q7NKC_9",
    "outputId": "b0509fc7-7e9d-4b5b-8fbf-0a9563c10d55"
  },
  "outputs": [
    {
      "data": {
        "text/plain": [
          "Outcome\n",
          "0      500\n",
          "1      268\n",
          "Name: count, dtype: int64"
        ]
      },
      "execution_count": 26,

```

```

    "metadata": {},
    "output_type": "execute_result"
  }
],
"source": [
  "#Para conteo de cada valor en una columna, en orden descendente usar función value_counts():
\n",
  "# nombreDataframe.columna.value_counts()\n",
  "# nombreDataframe['columna'].value_counts()\n",
  "df.Outcome.value_counts()"
]
},
{
  "cell_type": "code",
  "execution_count": 27,
  "metadata": {
    "id": "hu2J0Q7NKC_9",
    "outputId": "b0509fc7-7e9d-4b5b-8fbf-0a9563c10d55"
  },
  "outputs": [
    {
      "data": {
        "text/plain": [
          "Pregnancies\n",
          "1      135\n",
          "0      111\n",
          "2      103\n",
          "3       75\n",
          "4       68\n",
          "5       57\n",
          "6       50\n",
          "7       45\n",
          "8       38\n",
          "9       28\n",
          "10      24\n",
          "11      11\n",
          "13      10\n",
          "12       9\n",
          "14       2\n",
          "15       1\n",
          "17       1\n",
          "Name: count, dtype: int64"
        ]
      },
      "execution_count": 27,
      "metadata": {},
      "output_type": "execute_result"
    }
  ],
  "source": [
    "#Para conteo de cada valor en una columna, en orden descendente usar función value_counts():
\n",
    "# nombreDataframe.columna.value_counts()\n",
    "# nombreDataframe['columna'].value_counts()\n",
    "df[\"Pregnancies\"].value_counts()"
  ]
},
{
  "cell_type": "code",
  "execution_count": 28,
  "metadata": {
    "id": "hu2J0Q7NKC_9",
    "outputId": "b0509fc7-7e9d-4b5b-8fbf-0a9563c10d55"
  },
  "outputs": [

```



```

{
  "data": {
    "text/plain": [
      "Glucose\n",
      "99      17\n",
      "100     17\n",
      "111     14\n",
      "129     14\n",
      "125     14\n",
      "      ..\n",
      "191      1\n",
      "177      1\n",
      "44       1\n",
      "62       1\n",
      "190      1\n",
      "Name: count, Length: 136, dtype: int64"
    ]
  },
  "execution_count": 28,
  "metadata": {},
  "output_type": "execute_result"
},
{
  "source": [
    "#Para conteo de cada valor en una columna, en orden descendente usar función value_counts():\n",
    "# nombreDataframe.columna.value_counts()\n",
    "# nombreDataframe['columna'].value_counts()\n",
    "df[\"Glucose\"].value_counts()"
  ],
  "cell_type": "code",
  "execution_count": 29,
  "metadata": {
    "id": "hu2J0Q7NKC_9",
    "outputId": "b0509fc7-7e9d-4b5b-8fbf-0a9563c10d55"
  },
  "outputs": [
    {
      "data": {
        "text/plain": [
          "Outcome\n",
          "0      500\n",
          "1      268\n",
          "Name: count, dtype: int64"
        ]
      },
      "execution_count": 29,
      "metadata": {},
      "output_type": "execute_result"
    }
  ],
  "source": [
    "#Para conteo de cada valor en una columna, en orden descendente usar función value_counts():\n",
    "# nombreDataframe.columna.value_counts()\n",
    "# nombreDataframe['columna'].value_counts()\n",
    "df[\"Outcome\"].value_counts()"
  ],
  "cell_type": "code",
  "execution_count": 10,
  "metadata": {}
}

```

```

    "id": "r3T8XkonKC_-",
    "outputId": "cd7251fd-e296-48ac-eb48-d883e5af0f35"
  },
  "outputs": [],
  "source": [
    "#Revisa conteo de varias columnas"
  ]
},
{
  "cell_type": "code",
  "execution_count": 42,
  "metadata": {
    "id": "Q7VSqNEtKC__",
    "outputId": "567aec70-2bd2-4f2f-8128-5a3ce5459baa"
  },
  "outputs": [],
  "source": [
    "# Crear variable totalPregDiabetic que incluya la suma de las columnas Pregnancies y Outcome con
    valor \"1\\n\",
    "# Mostrar el total por cada tamaño de familia\\n\",
    "conteo_preg = df[\"Pregnancies\"].count()\\n\",
    "conteo_outcome = (df[\"Outcome\"] == 1).sum()\\n\",
    "df[\"totalPregDiabetic\"] = df[\"Pregnancies\"] + (df[\"Outcome\"] == 1).astype(int)"
  ]
},
{
  "cell_type": "code",
  "execution_count": 43,
  "metadata": {
    "id": "WLB1AfB0KDAA"
  },
  "outputs": [
    {
      "data": {
        "text/plain": [
          "<bound method NDFrame.head of
Insulin  BMI  \\n\",
    \"0      6    148      72      35      0  33.6  \\n\",
    \"1      1     85      66      29      0  26.6  \\n\",
    \"2      8    183      64       0      0  23.3  \\n\",
    \"3      1     89      66      23     94  28.1  \\n\",
    \"4      0    137      40      35    168  43.1  \\n\",
    \"..    ...    ...    ...    ...    ...    ...  \\n\",
    \"763    10    101      76      48    180  32.9  \\n\",
    \"764     2    122      70      27      0  36.8  \\n\",
    \"765     5    121      72      23    112  26.2  \\n\",
    \"766     1    126      60       0      0  30.1  \\n\",
    \"767     1     93      70      31      0  30.4  \\n\",
    \"\\n\",
    \"      DiabetesPedigreeFunction  Age  Outcome  totalPregDiabetic  \\n\",
    \"0                0.627    50      1          7  \\n\",
    \"1                0.351    31      0          1  \\n\",
    \"2                0.672    32      1          9  \\n\",
    \"3                0.167    21      0          1  \\n\",
    \"4                2.288    33      1          1  \\n\",
    \"..                ...    ...    ...    ...    ...  \\n\",
    \"763              0.171    63      0         10  \\n\",
    \"764              0.340    27      0          2  \\n\",
    \"765              0.245    30      0          5  \\n\",
    \"766              0.349    47      1          2  \\n\",
    \"767              0.315    23      0          1  \\n\",
    \"\\n\",
    \"[768 rows x 10 columns]>\"
        ]
      },
      "text/plain": [
        "<bound method NDFrame.head of
Insulin  BMI  \\n\",
    \"0      6    148      72      35      0  33.6  \\n\",
    \"1      1     85      66      29      0  26.6  \\n\",
    \"2      8    183      64       0      0  23.3  \\n\",
    \"3      1     89      66      23     94  28.1  \\n\",
    \"4      0    137      40      35    168  43.1  \\n\",
    \"..    ...    ...    ...    ...    ...    ...  \\n\",
    \"763    10    101      76      48    180  32.9  \\n\",
    \"764     2    122      70      27      0  36.8  \\n\",
    \"765     5    121      72      23    112  26.2  \\n\",
    \"766     1    126      60       0      0  30.1  \\n\",
    \"767     1     93      70      31      0  30.4  \\n\",
    \"\\n\",
    \"      DiabetesPedigreeFunction  Age  Outcome  totalPregDiabetic  \\n\",
    \"0                0.627    50      1          7  \\n\",
    \"1                0.351    31      0          1  \\n\",
    \"2                0.672    32      1          9  \\n\",
    \"3                0.167    21      0          1  \\n\",
    \"4                2.288    33      1          1  \\n\",
    \"..                ...    ...    ...    ...    ...  \\n\",
    \"763              0.171    63      0         10  \\n\",
    \"764              0.340    27      0          2  \\n\",
    \"765              0.245    30      0          5  \\n\",
    \"766              0.349    47      1          2  \\n\",
    \"767              0.315    23      0          1  \\n\",
    \"\\n\",
    \"[768 rows x 10 columns]>\"
        ]
      }
    }
  ]
},

```

```

    "execution_count": 43,
    "metadata": {},
    "output_type": "execute_result"
  }
],
"source": [
  "df.head"
]
},
{
  "cell_type": "markdown",
  "metadata": {
    "id": "G13IyhcdKDAT"
  },
  "source": [
    "## Consulta"
  ]
},
{
  "cell_type": "code",
  "execution_count": 44,
  "metadata": {},
  "outputs": [
    {
      "data": {
        "text/plain": [
          "Pregnancies      6.000\n",
          "Glucose          148.000\n",
          "BloodPressure    72.000\n",
          "SkinThickness    35.000\n",
          "Insulin          0.000\n",
          "BMI              33.600\n",
          "DiabetesPedigreeFunction 0.627\n",
          "Age              50.000\n",
          "Outcome          1.000\n",
          "totalPregDiabetic 7.000\n",
          "Name: 0, dtype: float64"
        ]
      },
      "execution_count": 44,
      "metadata": {},
      "output_type": "execute_result"
    }
  ],
  "source": [
    "# df.iloc[i]: Accede a la fila en la posición i.\n",
    "# Acceder a la primera fila\n",
    "df.iloc[0]"
  ]
},
{
  "cell_type": "code",
  "execution_count": 45,
  "metadata": {},
  "outputs": [
    {
      "data": {
        "text/html": [
          "<div>\n",
          "<style scoped>\n",
          "  .dataframe tbody tr th:only-of-type {\n",
          "    vertical-align: middle;\n",
          "  }\n",
          "\n",
          "  .dataframe tbody tr th {\n",

```

```

"         vertical-align: top;\n",
"     }\n",
"\n",
"     .dataframe thead th {\n",
"         text-align: right;\n",
"     }\n",
"</style>\n",
"<table border=\"1\" class=\"dataframe\">\n",
"  <thead>\n",
"    <tr style=\"text-align: right;\">\n",
"      <th></th>\n",
"      <th>Pregnancies</th>\n",
"      <th>Glucose</th>\n",
"      <th>BloodPressure</th>\n",
"      <th>SkinThickness</th>\n",
"      <th>Insulin</th>\n",
"      <th>BMI</th>\n",
"      <th>DiabetesPedigreeFunction</th>\n",
"      <th>Age</th>\n",
"      <th>Outcome</th>\n",
"      <th>totalPregDiabetic</th>\n",
"    </tr>\n",
"  </thead>\n",
"  <tbody>\n",
"    <tr>\n",
"      <th>0</th>\n",
"      <td>6</td>\n",
"      <td>148</td>\n",
"      <td>72</td>\n",
"      <td>35</td>\n",
"      <td>0</td>\n",
"      <td>33.6</td>\n",
"      <td>0.627</td>\n",
"      <td>50</td>\n",
"      <td>1</td>\n",
"      <td>7</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>1</th>\n",
"      <td>1</td>\n",
"      <td>85</td>\n",
"      <td>66</td>\n",
"      <td>29</td>\n",
"      <td>0</td>\n",
"      <td>26.6</td>\n",
"      <td>0.351</td>\n",
"      <td>31</td>\n",
"      <td>0</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>2</th>\n",
"      <td>8</td>\n",
"      <td>183</td>\n",
"      <td>64</td>\n",
"      <td>0</td>\n",
"      <td>0</td>\n",
"      <td>23.3</td>\n",
"      <td>0.672</td>\n",
"      <td>32</td>\n",
"      <td>1</td>\n",
"      <td>9</td>\n",
"    </tr>\n",
"  </tbody>\n",
"</table>\n",

```

```

"</div>"
],
"text/plain": [
  "   Pregnancies   Glucose   BloodPressure   SkinThickness   Insulin   BMI   \\n",
  "0             6       148             72           35         0   33.6   \n",
  "1             1        85             66           29         0   26.6   \n",
  "2             8       183             64           0         0   23.3   \n",
  "\\n",
  "   DiabetesPedigreeFunction   Age   Outcome   totalPregDiabetic   \n",
  "0                        0.627   50         1             7   \n",
  "1                        0.351   31         0             1   \n",
  "2                        0.672   32         1             9   "
]
},
"execution_count": 45,
"metadata": {},
"output_type": "execute_result"
}
],
"source": [
  "# Acceder a las dos primeras filas\\n",
  "df.iloc[:3]"
]
},
{
  "cell_type": "code",
  "execution_count": 46,
  "metadata": {},
  "outputs": [
    {
      "data": {
        "text/html": [
          "<div>\\n",
          "<style scoped>\\n",
          "  .dataframe tbody tr th:only-of-type {\\n",
          "    vertical-align: middle;\\n",
          "  }\\n",
          "\\n",
          "  .dataframe tbody tr th {\\n",
          "    vertical-align: top;\\n",
          "  }\\n",
          "\\n",
          "  .dataframe thead th {\\n",
          "    text-align: right;\\n",
          "  }\\n",
          "</style>\\n",
          "<table border='1' class='dataframe'>\\n",
          "  <thead>\\n",
          "    <tr style='text-align: right;'>\\n",
          "      <th></th>\\n",
          "      <th>Pregnancies</th>\\n",
          "      <th>Glucose</th>\\n",
          "    </tr>\\n",
          "  </thead>\\n",
          "  <tbody>\\n",
          "    <tr>\\n",
          "      <th>0</th>\\n",
          "      <td>6</td>\\n",
          "      <td>148</td>\\n",
          "    </tr>\\n",
          "    <tr>\\n",
          "      <th>1</th>\\n",
          "      <td>1</td>\\n",
          "      <td>85</td>\\n",
          "    </tr>\\n",

```

```

"      <tr>\n",
"          <th>2</th>\n",
"          <td>8</td>\n",
"          <td>183</td>\n",
"      </tr>\n",
"      <tr>\n",
"          <th>3</th>\n",
"          <td>1</td>\n",
"          <td>89</td>\n",
"      </tr>\n",
"      <tr>\n",
"          <th>4</th>\n",
"          <td>0</td>\n",
"          <td>137</td>\n",
"      </tr>\n",
"      <tr>\n",
"          <th>...</th>\n",
"          <td>...</td>\n",
"          <td>...</td>\n",
"      </tr>\n",
"      <tr>\n",
"          <th>763</th>\n",
"          <td>10</td>\n",
"          <td>101</td>\n",
"      </tr>\n",
"      <tr>\n",
"          <th>764</th>\n",
"          <td>2</td>\n",
"          <td>122</td>\n",
"      </tr>\n",
"      <tr>\n",
"          <th>765</th>\n",
"          <td>5</td>\n",
"          <td>121</td>\n",
"      </tr>\n",
"      <tr>\n",
"          <th>766</th>\n",
"          <td>1</td>\n",
"          <td>126</td>\n",
"      </tr>\n",
"      <tr>\n",
"          <th>767</th>\n",
"          <td>1</td>\n",
"          <td>93</td>\n",
"      </tr>\n",
"  </tbody>\n",
"</table>\n",
"<p>768 rows x 2 columns</p>\n",
"</div>"
],
"text/plain": [
"      Pregnancies  Glucose\n",
"0              6      148\n",
"1              1       85\n",
"2              8      183\n",
"3              1       89\n",
"4              0      137\n",
"..           ...      ... \n",
"763           10      101\n",
"764           2       122\n",
"765           5       121\n",
"766           1       126\n",
"767           1        93\n",
"\n",
"[768 rows x 2 columns]"

```

```

    ]
  },
  "execution_count": 46,
  "metadata": {},
  "output_type": "execute_result"
}
],
"source": [
  "#Seleccionar columnas, indicando entre corchetes [nombreColumna, nombreColumna]\n",
  "df[\"Pregnancies\", \"Glucose\"]"
]
},
{
  "cell_type": "code",
  "execution_count": 50,
  "metadata": {},
  "outputs": [],
  "source": [
    "#Selección de filas [indicar dataframe[columna] operador valor]\n",
    "embarazos = df[df[\"Pregnancies\"] == 0]"
  ]
},
{
  "cell_type": "code",
  "execution_count": 51,
  "metadata": {},
  "outputs": [
    {
      "data": {
        "text/html": [
          "<div>\n",
          "<style scoped>\n",
          "  .dataframe tbody tr th:only-of-type {\n",
          "    vertical-align: middle;\n",
          "  }\n",
          "\n",
          "  .dataframe tbody tr th {\n",
          "    vertical-align: top;\n",
          "  }\n",
          "\n",
          "  .dataframe thead th {\n",
          "    text-align: right;\n",
          "  }\n",
          "</style>\n",
          "<table border='1' class='dataframe'>\n",
          "  <thead>\n",
          "    <tr style='text-align: right;'>\n",
          "      <th></th>\n",
          "      <th>Pregnancies</th>\n",
          "      <th>Glucose</th>\n",
          "      <th>BloodPressure</th>\n",
          "      <th>SkinThickness</th>\n",
          "      <th>Insulin</th>\n",
          "      <th>BMI</th>\n",
          "      <th>DiabetesPedigreeFunction</th>\n",
          "      <th>Age</th>\n",
          "      <th>Outcome</th>\n",
          "      <th>totalPregDiabetic</th>\n",
          "    </tr>\n",
          "  </thead>\n",
          "  <tbody>\n",
          "    <tr>\n",
          "      <th>4</th>\n",
          "      <td>0</td>\n",
          "      <td>137</td>\n",

```

```

"      <td>40</td>\n",
"      <td>35</td>\n",
"      <td>168</td>\n",
"      <td>43.1</td>\n",
"      <td>2.288</td>\n",
"      <td>33</td>\n",
"      <td>1</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"  <tr>\n",
"    <th>528</th>\n",
"    <td>0</td>\n",
"    <td>117</td>\n",
"    <td>66</td>\n",
"    <td>31</td>\n",
"    <td>188</td>\n",
"    <td>30.8</td>\n",
"    <td>0.493</td>\n",
"    <td>22</td>\n",
"    <td>0</td>\n",
"    <td>0</td>\n",
"  </tr>\n",
"  <tr>\n",
"    <th>511</th>\n",
"    <td>0</td>\n",
"    <td>139</td>\n",
"    <td>62</td>\n",
"    <td>17</td>\n",
"    <td>210</td>\n",
"    <td>22.1</td>\n",
"    <td>0.207</td>\n",
"    <td>21</td>\n",
"    <td>0</td>\n",
"    <td>0</td>\n",
"  </tr>\n",
"  <tr>\n",
"    <th>506</th>\n",
"    <td>0</td>\n",
"    <td>180</td>\n",
"    <td>90</td>\n",
"    <td>26</td>\n",
"    <td>90</td>\n",
"    <td>36.5</td>\n",
"    <td>0.314</td>\n",
"    <td>35</td>\n",
"    <td>1</td>\n",
"    <td>1</td>\n",
"  </tr>\n",
"  <tr>\n",
"    <th>487</th>\n",
"    <td>0</td>\n",
"    <td>173</td>\n",
"    <td>78</td>\n",
"    <td>32</td>\n",
"    <td>265</td>\n",
"    <td>46.5</td>\n",
"    <td>1.159</td>\n",
"    <td>58</td>\n",
"    <td>0</td>\n",
"    <td>0</td>\n",
"  </tr>\n",
"  <tr>\n",
"    <th>...</th>\n",
"    <td>...</td>\n",
"    <td>...</td>\n",

```



```

"      <td>...</td>\n",
"      <td>...</td>\n",
"      <td>...</td>\n",
"      <td>...</td>\n",
"      <td>...</td>\n",
"      <td>...</td>\n",
"      <td>...</td>\n",
"      <td>...</td>\n",
"    </tr>\n",
"  <tr>\n",
"    <th>247</th>\n",
"    <td>0</td>\n",
"    <td>165</td>\n",
"    <td>90</td>\n",
"    <td>33</td>\n",
"    <td>680</td>\n",
"    <td>52.3</td>\n",
"    <td>0.427</td>\n",
"    <td>23</td>\n",
"    <td>0</td>\n",
"    <td>0</td>\n",
"  </tr>\n",
"  <tr>\n",
"    <th>239</th>\n",
"    <td>0</td>\n",
"    <td>104</td>\n",
"    <td>76</td>\n",
"    <td>0</td>\n",
"    <td>0</td>\n",
"    <td>18.4</td>\n",
"    <td>0.582</td>\n",
"    <td>27</td>\n",
"    <td>0</td>\n",
"    <td>0</td>\n",
"  </tr>\n",
"  <tr>\n",
"    <th>237</th>\n",
"    <td>0</td>\n",
"    <td>179</td>\n",
"    <td>90</td>\n",
"    <td>27</td>\n",
"    <td>0</td>\n",
"    <td>44.1</td>\n",
"    <td>0.686</td>\n",
"    <td>23</td>\n",
"    <td>1</td>\n",
"    <td>1</td>\n",
"  </tr>\n",
"  <tr>\n",
"    <th>297</th>\n",
"    <td>0</td>\n",
"    <td>126</td>\n",
"    <td>84</td>\n",
"    <td>29</td>\n",
"    <td>215</td>\n",
"    <td>30.7</td>\n",
"    <td>0.520</td>\n",
"    <td>24</td>\n",
"    <td>0</td>\n",
"    <td>0</td>\n",
"  </tr>\n",
"  <tr>\n",
"    <th>757</th>\n",
"    <td>0</td>\n",
"    <td>123</td>\n",

```

```

"      <td>72</td>\n",
"      <td>0</td>\n",
"      <td>0</td>\n",
"      <td>36.3</td>\n",
"      <td>0.258</td>\n",
"      <td>52</td>\n",
"      <td>1</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"  </tbody>\n",
"</table>\n",
"<p>111 rows x 10 columns</p>\n",
"</div>"
],
"text/plain": [
"      Pregnancies  Glucose  BloodPressure  SkinThickness  Insulin   BMI   \\\n",
"4                0      137             40             35      168   43.1   \n",
"528              0      117             66             31      188   30.8   \n",
"511              0      139             62             17      210   22.1   \n",
"506              0      180             90             26       90   36.5   \n",
"487              0      173             78             32      265   46.5   \n",
"...\n",
"247              0      165             90             33      680   52.3   \n",
"239              0      104             76              0       0   18.4   \n",
"237              0      179             90             27       0   44.1   \n",
"297              0      126             84             29      215   30.7   \n",
"757              0      123             72              0       0   36.3   \n",
"\n",
"      DiabetesPedigreeFunction  Age  Outcome  totalPregDiabetic  \n",
"4                2.288      33         1             1  \n",
"528              0.493      22         0             0  \n",
"511              0.207      21         0             0  \n",
"506              0.314      35         1             1  \n",
"487              1.159      58         0             0  \n",
"...\n",
"247              0.427      23         0             0  \n",
"239              0.582      27         0             0  \n",
"237              0.686      23         1             1  \n",
"297              0.520      24         0             0  \n",
"757              0.258      52         1             1  \n",
"\n",
"[111 rows x 10 columns]"
]
},
"execution_count": 51,
"metadata": {},
"output_type": "execute_result"
}
],
"source": [
"#ordenar usando funcion sort_values(by=atributo, ascending=True/false\n",
"embarazos.sort_values(by=\"Pregnancies\", ascending=True)"
]
},
{
"cell_type": "code",
"execution_count": 52,
"metadata": {},
"outputs": [
{
"data": {
"text/plain": [
"Glucose\n",
"57      0.0\n",
"67      0.0\n",

```

```

"73      0.0\n",
"74      0.0\n",
"78      0.0\n",
"      ... \n",
"180     1.0\n",
"181     1.0\n",
"188     1.0\n",
"189     1.0\n",
"198     1.0\n",
"Name: Outcome, Length: 64, dtype: float64"
]
},
"execution_count": 52,
"metadata": {},
"output_type": "execute_result"
}
],
"source": [
"#Agrupar por un atributo y calcular función de agregación utilizando groupby(atributo)
['atributoAgregacion'].funcion() (ej. mean() para promedio)\n",
"embarazos.groupby(\"Glucose\")[\"Outcome\"].mean()"
]
},
{
"cell_type": "markdown",
"metadata": {
"id": "PnlaaWEKDAU"
},
"source": [
"Crea un subconjunto de glucose para el valor mayor a 180"
]
},
{
"cell_type": "code",
"execution_count": 56,
"metadata": {
"id": "o7-EK3csKDAU",
"outputId": "498702ea-89b2-480c-dec1-d5a7704b6104"
},
"outputs": [
{
"data": {
"text/html": [
"<div>\n",
"<style scoped>\n",
"      .dataframe tbody tr th:only-of-type {\n",
"        vertical-align: middle;\n",
"      }\n",
"\n",
"      .dataframe tbody tr th {\n",
"        vertical-align: top;\n",
"      }\n",
"\n",
"      .dataframe thead th {\n",
"        text-align: right;\n",
"      }\n",
"</style>\n",
"<table border=\"1\" class=\"dataframe\">\n",
"  <thead>\n",
"    <tr style=\"text-align: right;\">\n",
"      <th></th>\n",
"      <th>Pregnancies</th>\n",
"      <th>Glucose</th>\n",
"      <th>BloodPressure</th>\n",
"      <th>SkinThickness</th>\n",

```

```

"      <th>Insulin</th>\n",
"      <th>BMI</th>\n",
"      <th>DiabetesPedigreeFunction</th>\n",
"      <th>Age</th>\n",
"      <th>Outcome</th>\n",
"      <th>totalPregDiabetic</th>\n",
"    </tr>\n",
"  </thead>\n",
"  <tbody>\n",
"    <tr>\n",
"      <th>440</th>\n",
"      <td>0</td>\n",
"      <td>189</td>\n",
"      <td>104</td>\n",
"      <td>25</td>\n",
"      <td>0</td>\n",
"      <td>34.3</td>\n",
"      <td>0.435</td>\n",
"      <td>41</td>\n",
"      <td>1</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>561</th>\n",
"      <td>0</td>\n",
"      <td>198</td>\n",
"      <td>66</td>\n",
"      <td>32</td>\n",
"      <td>274</td>\n",
"      <td>41.3</td>\n",
"      <td>0.502</td>\n",
"      <td>28</td>\n",
"      <td>1</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>595</th>\n",
"      <td>0</td>\n",
"      <td>188</td>\n",
"      <td>82</td>\n",
"      <td>14</td>\n",
"      <td>185</td>\n",
"      <td>32.0</td>\n",
"      <td>0.682</td>\n",
"      <td>22</td>\n",
"      <td>1</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"    <tr>\n",
"      <th>753</th>\n",
"      <td>0</td>\n",
"      <td>181</td>\n",
"      <td>88</td>\n",
"      <td>44</td>\n",
"      <td>510</td>\n",
"      <td>43.3</td>\n",
"      <td>0.222</td>\n",
"      <td>26</td>\n",
"      <td>1</td>\n",
"      <td>1</td>\n",
"    </tr>\n",
"  </tbody>\n",
"</table>\n",
"</div>"
],

```

```

"text/plain": [
  "      Pregnancies  Glucose  BloodPressure  SkinThickness  Insulin   BMI   \\\n",
  "440              0      189            104            25         0  34.3   \n",
  "561              0      198             66            32        274  41.3   \n",
  "595              0      188             82            14        185  32.0   \n",
  "753              0      181             88            44        510  43.3   \n",
  "\n",
  "      DiabetesPedigreeFunction  Age  Outcome  totalPregDiabetic  \n",
  "440                        0.435  41         1                 1  \n",
  "561                        0.502  28         1                 1  \n",
  "595                        0.682  22         1                 1  \n",
  "753                        0.222  26         1                 1  "
]
},
"execution_count": 56,
"metadata": {},
"output_type": "execute_result"
}
],
"source": [
"# usa el criterio para extraer solo los boletos caros con fare > 50\n",
"embarazos[embarazos[\"Glucose\"] > 180]"
]
},
{
"cell_type": "code",
"execution_count": null,
"metadata": {},
"outputs": [],
"source": []
}
],
"metadata": {
"colab": {
"provenance": []
},
"kernel_spec": {
"display_name": "Python 3 (ipykernel)",
"language": "python",
"name": "python3"
},
"language_info": {
"codemirror_mode": {
"name": "ipython",
"version": 3
},
"file_extension": ".py",
"mimetype": "text/x-python",
"name": "python",
"nbconvert_exporter": "python",
"pygments_lexer": "ipython3",
"version": "3.8.10"
}
},
"nbformat": 4,
"nbformat_minor": 4
}

```