## 2\_0\_Lexical\_Complexity\_Binary\_Classification\_Prediction\_Data\_Preparate

### April 6, 2025

```
[1]: #@title Install Packages
[2]: !pip install -q transformers
     !pip install -q torchinfo
    !pip install -q datasets
     !pip install -q evaluate
    !pip install -q nltk
     !pip install -q contractions
                              491.2/491.2 kB
    24.1 MB/s eta 0:00:00
                              116.3/116.3 kB
    12.7 MB/s eta 0:00:00
                              183.9/183.9 kB
    17.9 MB/s eta 0:00:00
                              143.5/143.5 kB
    14.0 MB/s eta 0:00:00
                              194.8/194.8 kB
    17.6 MB/s eta 0:00:00
```

```
ERROR: pip's dependency resolver does not currently take into account
all the packages that are installed. This behaviour is the source of the
following dependency conflicts.
torch 2.6.0+cu124 requires nvidia-cublas-cu12==12.4.5.8; platform_system ==
"Linux" and platform machine == "x86 64", but you have nvidia-cublas-cu12
12.5.3.2 which is incompatible.
torch 2.6.0+cu124 requires nvidia-cuda-cupti-cu12==12.4.127; platform_system ==
"Linux" and platform machine == "x86_64", but you have nvidia-cuda-cupti-cu12
12.5.82 which is incompatible.
torch 2.6.0+cu124 requires nvidia-cuda-nvrtc-cu12==12.4.127; platform_system ==
"Linux" and platform_machine == "x86_64", but you have nvidia-cuda-nvrtc-cu12
12.5.82 which is incompatible.
torch 2.6.0+cu124 requires nvidia-cuda-runtime-cu12==12.4.127; platform_system
== "Linux" and platform_machine == "x86_64", but you have nvidia-cuda-runtime-
cu12 12.5.82 which is incompatible.
torch 2.6.0+cu124 requires nvidia-cudnn-cu12==9.1.0.70; platform_system ==
"Linux" and platform_machine == "x86_64", but you have nvidia-cudnn-cu12
9.3.0.75 which is incompatible.
torch 2.6.0+cu124 requires nvidia-cufft-cu12==11.2.1.3; platform_system ==
"Linux" and platform_machine == "x86_64", but you have nvidia-cufft-cu12
11.2.3.61 which is incompatible.
torch 2.6.0+cu124 requires nvidia-curand-cu12==10.3.5.147; platform system ==
"Linux" and platform_machine == "x86_64", but you have nvidia-curand-cu12
10.3.6.82 which is incompatible.
torch 2.6.0+cu124 requires nvidia-cusolver-cu12==11.6.1.9; platform system ==
"Linux" and platform_machine == "x86_64", but you have nvidia-cusolver-cu12
11.6.3.83 which is incompatible.
torch 2.6.0+cu124 requires nvidia-cusparse-cu12==12.3.1.170; platform system ==
"Linux" and platform_machine == "x86_64", but you have nvidia-cusparse-cu12
12.5.1.3 which is incompatible.
torch 2.6.0+cu124 requires nvidia-nvjitlink-cu12==12.4.127; platform_system ==
"Linux" and platform_machine == "x86_64", but you have nvidia-nvjitlink-cu12
12.5.82 which is incompatible.
```

gcsfs 2025.3.2 requires fsspec==2025.3.2, but you have fsspec 2024.12.0 which is

9/ 0/9/ 0 1-D

incompatible.

```
4.1 MB/s eta 0:00:00
                              289.9/289.9 kB
    23.5 MB/s eta 0:00:00
                              118.3/118.3 kB
    12.0 MB/s eta 0:00:00
[3]: sudo apt-get update
     ! sudo apt-get install tree
    Get:1 https://cloud.r-project.org/bin/linux/ubuntu jammy-cran40/ InRelease
    [3,632 B]
    Get:2 https://developer.download.nvidia.com/compute/cuda/repos/ubuntu2204/x86_64
    InRelease [1,581 B]
    Get:3 https://developer.download.nvidia.com/compute/cuda/repos/ubuntu2204/x86 64
    Packages [1,381 kB]
    Get:4 http://security.ubuntu.com/ubuntu jammy-security InRelease [129 kB]
    Hit:5 http://archive.ubuntu.com/ubuntu jammy InRelease
    Get:6 http://archive.ubuntu.com/ubuntu jammy-updates InRelease [128 kB]
    Hit:7 https://ppa.launchpadcontent.net/deadsnakes/ppa/ubuntu jammy InRelease
    Hit:8 https://ppa.launchpadcontent.net/graphics-drivers/ppa/ubuntu jammy
    InRelease
    Get:9 https://r2u.stat.illinois.edu/ubuntu jammy InRelease [6,555 B]
    Get:10 http://security.ubuntu.com/ubuntu jammy-security/restricted amd64
    Packages [3,978 kB]
    Get:11 http://archive.ubuntu.com/ubuntu jammy-backports InRelease [127 kB]
    Hit:12 https://ppa.launchpadcontent.net/ubuntugis/ppa/ubuntu jammy InRelease
    Get:13 http://archive.ubuntu.com/ubuntu jammy-updates/restricted amd64 Packages
    [4,148 kB]
    Get:14 https://r2u.stat.illinois.edu/ubuntu jammy/main all Packages [8,809 kB]
    Get:15 http://security.ubuntu.com/ubuntu jammy-security/universe amd64 Packages
    [1,241 kB]
    Get:16 http://security.ubuntu.com/ubuntu jammy-security/main amd64 Packages
    [2.775 \text{ kB}]
    Get:17 http://archive.ubuntu.com/ubuntu jammy-updates/universe amd64 Packages
    [1.540 kB]
    Get:18 http://archive.ubuntu.com/ubuntu jammy-updates/main amd64 Packages [3,092
    kBl
    Get:19 https://r2u.stat.illinois.edu/ubuntu jammy/main amd64 Packages [2,688 kB]
    Fetched 30.0 MB in 4s (6.824 \text{ kB/s})
    Reading package lists... Done
    W: Skipping acquire of configured file 'main/source/Sources' as repository
    'https://r2u.stat.illinois.edu/ubuntu jammy InRelease' does not seem to provide
    it (sources.list entry misspelt?)
    Reading package lists... Done
    Building dependency tree... Done
    Reading state information... Done
    The following NEW packages will be installed:
```

```
tree
    O upgraded, 1 newly installed, O to remove and 44 not upgraded.
    Need to get 47.9 kB of archives.
    After this operation, 116 kB of additional disk space will be used.
    Get:1 http://archive.ubuntu.com/ubuntu jammy/universe amd64 tree amd64 2.0.2-1
    [47.9 kB]
    Fetched 47.9 kB in 1s (70.5 kB/s)
    debconf: unable to initialize frontend: Dialog
    debconf: (No usable dialog-like program is installed, so the dialog based
    frontend cannot be used. at /usr/share/perl5/Debconf/FrontEnd/Dialog.pm line 78,
    <> line 1.)
    debconf: falling back to frontend: Readline
    debconf: unable to initialize frontend: Readline
    debconf: (This frontend requires a controlling tty.)
    debconf: falling back to frontend: Teletype
    dpkg-preconfigure: unable to re-open stdin:
    Selecting previously unselected package tree.
    (Reading database ... 126213 files and directories currently installed.)
    Preparing to unpack .../tree_2.0.2-1_amd64.deb ...
    Unpacking tree (2.0.2-1) ...
    Setting up tree (2.0.2-1) ...
    Processing triggers for man-db (2.10.2-1) ...
[4]: #@title Imports
     import nltk
     from nltk.tokenize import RegexpTokenizer
     import evaluate
     import transformers
     import contractions
     from torchinfo import summary
     from datasets import load_dataset
     from transformers import AutoTokenizer, AutoModel,
      → AutoModelForSequenceClassification
     from transformers import TrainingArguments, Trainer
     import os
     import pandas as pd
     import numpy as np
     import matplotlib.pyplot as plt
     import seaborn as sns
     import sklearn
```

```
import spacy
[5]: # @title Mount Google Drive
[6]: from google.colab import drive
     drive.mount('/content/drive')
    Mounted at /content/drive
[7]: dir root = '/content/drive/MyDrive/266-final/'
     # dir_data = '/content/drive/MyDrive/266-final/data/'
     # dir data = '/content/drive/MyDrive/266-final/data/se21-t1-comp-lex-master/'
     dir_data = '/content/drive/MyDrive/266-final/data/266-comp-lex-master'
     dir_models = '/content/drive/MyDrive/266-final/models/'
     dir_results = '/content/drive/MyDrive/266-final/results/'
[8]: | tree /content/drive/MyDrive/266-final/data/266-comp-lex-master/
    /content/drive/MyDrive/266-final/data/266-comp-lex-master/
       fe-test-labels
       fe-train
      fe-trial-val
       test-labels
          lcp_multi_test.tsv
          lcp_single_test.tsv
       train
          lcp_multi_train.tsv
          lcp_single_train.tsv
       trial
           lcp_multi_trial.tsv
          lcp_single_trial.tsv
    6 directories, 6 files
[9]: ||ls -R /content/drive/MyDrive/266-final/data/266-comp-lex-master/
    /content/drive/MyDrive/266-final/data/266-comp-lex-master/:
    fe-test-labels fe-train fe-trial-val test-labels train trial
    /content/drive/MyDrive/266-final/data/266-comp-lex-master/fe-test-labels:
    /content/drive/MyDrive/266-final/data/266-comp-lex-master/fe-train:
    /content/drive/MyDrive/266-final/data/266-comp-lex-master/fe-trial-val:
    /content/drive/MyDrive/266-final/data/266-comp-lex-master/test-labels:
    lcp_multi_test.tsv lcp_single_test.tsv
    /content/drive/MyDrive/266-final/data/266-comp-lex-master/train:
```

```
lcp_multi_train.tsv lcp_single_train.tsv
/content/drive/MyDrive/266-final/data/266-comp-lex-master/trial:
lcp_multi_trial.tsv lcp_single_trial.tsv
```

```
[10]: #@title Import Data
[11]: | # train single df = pd.read csv(os.path.join(dir data, "train", |
                          \hookrightarrow "lcp single train.tsv"), sep="\t")
                        # train_multi_df = pd.read_csv(os.path.join(dir_data, "train", "lcp_multi_train.
                            \hookrightarrow tsv''), sep='' \setminus t'')
                       # trail_val_single_df = pd.read_csv(os.path.join(dir_data, "trial", __
                           \hookrightarrow "lcp_single_trial.tsv"), sep="\t")
                        \# trail\_val\_multi\_df = pd.read\_csv(os.path.join(dir\_data, "trial", "trial
                          \hookrightarrow "lcp_multi_trial.tsv"), sep="\t")
                        \# test\_single\_df = pd.read\_csv(os.path.join(dir\_data, "test-labels", \sqcup test-labels", \sqcup test-labels", \sqcup test-labels = test-labe
                           \hookrightarrow "lcp_single_test.tsv"), sep="\t")
                        # test_multi_df = pd.read_csv(os.path.join(dir_data, "test-labels",_
                            \hookrightarrow "lcp_multi_test.tsv"), sep="\t")
[12]: # # Try to load the files containing unterminated strings
                        # try:
                                               # Approach 1: Try with the C engine but with error handling
                        #
                                               multi_test_df = pd.read_csv(
                        #
                                                               os.path.join(dir_data, "test", "lcp_multi_test.tsv"),
                        #
                                                               sep="\t",
                        #
                                                               on bad lines='skip' # Skip bad lines
                        #
                                               print("Loaded with skipping bad lines")
                        # except Exception as e:
                        #
                                               print(f"First approach failed: {e}")
                        #
                                               try:
                        #
                                                               # Approach 2: Try with the Python engine which might be more forgiving
                        #
                                                               multi_test_df = pd.read_csv(
                        #
                                                                               os.path.join(dir_data, "test", "lcp_multi_test.tsv"),
                        #
                                                                               sep="\t",
                        #
                                                                               engine="python",
                        #
                                                                               quoting=3 # QUOTE_NONE
                        #
                                                               print("Loaded with Python engine")
[13]: # Load train data into train_*_df
                       train_single_df = pd.read_csv(
                                       os.path.join(dir_data, "train", "lcp_single_train.tsv"),
                                       sep = "\t".
```

```
engine = "python",
    quoting = 3
train_multi_df = pd.read_csv(
    os.path.join(dir_data, "train", "lcp_multi_train.tsv"),
    sep = "\t",
    engine = "python",
    quoting = 3
)
# Load trial data into trial val * df
trial_val_single_df = pd.read_csv(
    os.path.join(dir_data, "trial", "lcp_single_trial.tsv"),
    sep = "\t",
    engine = "python",
    quoting = 3
trial_val_multi_df = pd.read_csv(
    os.path.join(dir_data, "trial", "lcp_multi_trial.tsv"),
    sep = "\t",
    engine = "python",
    quoting = 3
)
# Load test data (with labels) into test_*_df
test_single_df = pd.read_csv(
    os.path.join(dir_data, "test-labels", "lcp_single_test.tsv"),
    sep = "\t",
    engine = "python",
    quoting = 3
)
test_multi_df = pd.read_csv(
    os.path.join(dir_data, "test-labels", "lcp_multi_test.tsv"),
    sep = "\t",
    engine = "python",
    quoting = 3
)
print("Data successfully loaded into train, trial-val, and test variables")
```

Data successfully loaded into train, trial-val, and test variables

```
# Shape and Columns
   print(f"Shape: {df.shape}")
   print(f"Columns: {list(df.columns)}\n")
   # Data Types
   print("Data Types:")
   print(df.dtypes)
   print()
    # Missing Values
   print("Missing Values (by column):")
   print(df.isna().sum())
   print()
   # 'complexity' column stats
   desc = df['complexity'].describe() # count, mean, std, min, 25%, 50%, 75%, u
   print("'complexity' Column Stats (incl. quartiles and median):")
   print(desc)
   # Calculate frequency counts for each quartile range
   q1 = desc['25\%']
   q2 = desc['50\%'] # This is the median
   q3 = desc['75\%']
   q_max = desc['max']
    # Note: We'll define the ranges as:
    # <= Q1
    # > Q1 and <= Q2
    # > Q2 and <= Q3
    # > Q3
   freq_q1 = np.sum(df['complexity'] <= q1)</pre>
   freq_q2 = np.sum((df['complexity'] > q1) & (df['complexity'] <= q2))</pre>
   freq_q3 = np.sum((df['complexity'] > q2) & (df['complexity'] <= q3))</pre>
   freq_q4 = np.sum(df['complexity'] > q3)
   print()
   print("Quartile Frequency Counts (tab-separated next to each quartile):")
   print(f"25%: {q1}\tCount (<= Q1): {freq_q1}")</pre>
   print(f"50\% (Median): {q2}\tCount (Q1 < x <= Q2): {freq_q2}")
   print(f"75\%: {q3}\tCount (Q2 < x <= Q3): {freq_q3}")
   print(f"100\% (Max): {q_max}\tCount (Q3 < x <= Max): {freq_q4}")
   print("=======\n")
# Now we call this for each of our dataframes
```

```
print_dataframe_summary("train_single_df", train_single_df)
print_dataframe_summary("train_multi_df", train_multi_df)
print_dataframe_summary("trial_val_single_df", trial_val_single_df)
print_dataframe_summary("trial_val_multi_df", trial_val_multi_df)
print_dataframe_summary("test_single_df", test_single_df)
print_dataframe_summary("test_multi_df", test_multi_df)
====== train single df ======
Shape: (7662, 5)
Columns: ['id', 'corpus', 'sentence', 'token', 'complexity']
Data Types:
id
             object
corpus
             object
sentence
             object
token
             object
complexity
             float64
dtype: object
Missing Values (by column):
id
            0
corpus
sentence
            0
token
            7
complexity
dtype: int64
'complexity' Column Stats (incl. quartiles and median):
        7662.000000
count
           0.302288
mean
           0.132977
std
           0.000000
min
25%
           0.211538
50%
           0.279412
75%
           0.375000
           0.861111
max
Name: complexity, dtype: float64
Quartile Frequency Counts (tab-separated next to each quartile):
25%: 0.2115384615384615 Count (<= Q1): 1928
50% (Median): 0.2794117647058823
                                     Count (Q1 < x \le Q2): 1937
75%: 0.375
              Count (Q2 < x \le Q3): 1984
====== train_multi_df ======
Shape: (1517, 5)
Columns: ['id', 'corpus', 'sentence', 'token', 'complexity']
```

```
Data Types:
id
             object
             object
corpus
sentence
             object
             object
token
complexity
            float64
dtype: object
Missing Values (by column):
            0
id
            0
corpus
            0
sentence
token
complexity
dtype: int64
'complexity' Column Stats (incl. quartiles and median):
count
        1517.000000
mean
          0.418362
std
          0.155536
min
          0.027778
25%
         0.302632
50%
          0.409091
75%
         0.529412
          0.975000
max
Name: complexity, dtype: float64
Quartile Frequency Counts (tab-separated next to each quartile):
25%: 0.3026315789473685 Count (<= Q1): 382
75%: 0.5294117647058824 Count (Q2 < x <= Q3): 380
100% (Max): 0.975
                     Count (Q3 < x <= Max): 378
====== trial_val_single_df =======
Shape: (421, 5)
Columns: ['id', 'subcorpus', 'sentence', 'token', 'complexity']
Data Types:
id
             object
subcorpus
             object
sentence
             object
token
             object
complexity
            float64
dtype: object
Missing Values (by column):
```

```
id
              0
subcorpus
              0
sentence
              0
token
              0
complexity
dtype: int64
'complexity' Column Stats (incl. quartiles and median):
count
       421.000000
mean
           0.298631
           0.137619
std
          0.000000
min
25%
           0.214286
50%
           0.266667
75%
           0.359375
           0.875000
max
Name: complexity, dtype: float64
Quartile Frequency Counts (tab-separated next to each quartile):
25%: 0.2142857142857143 Count (<= Q1): 106
50% (Median): 0.266666666666667
                                        Count (Q1 < x \le Q2): 107
75%: 0.359375
              Count (Q2 < x <= Q3): 103
100% (Max): 0.875
                        Count (Q3 < x <= Max): 105
======= trial_val_multi_df =======
Shape: (99, 5)
Columns: ['id', 'subcorpus', 'sentence', 'token', 'complexity']
Data Types:
               object
id
subcorpus
               object
sentence
               object
token
               object
              float64
complexity
dtype: object
Missing Values (by column):
id
subcorpus
              0
sentence
              0
token
              0
complexity
dtype: int64
'complexity' Column Stats (incl. quartiles and median):
         99.000000
count
          0.417961
mean
```

```
std
         0.153752
         0.000000
min
25%
         0.309028
50%
         0.421875
75%
         0.513932
         0.825000
max
Name: complexity, dtype: float64
Quartile Frequency Counts (tab-separated next to each quartile):
25%: 0.309027777777778 Count (<= Q1): 25
50% (Median): 0.421875 Count (Q1 < x <= Q2): 25
75%: 0.5139318885448916 Count (Q2 < x <= Q3): 24
100% (Max): 0.825
                       Count (Q3 < x <= Max): 25
______
====== test_single_df ======
Shape: (917, 5)
Columns: ['id', 'corpus', 'sentence', 'token', 'complexity']
Data Types:
id
              object
corpus
              object
              object
sentence
token
              object
complexity
             float64
dtype: object
Missing Values (by column):
             0
id
corpus
             0
sentence
             0
token
             0
complexity
dtype: int64
'complexity' Column Stats (incl. quartiles and median):
count 917.000000
mean
          0.296362
std
          0.127290
min
          0.000000
25%
          0.214286
50%
          0.276316
75%
          0.357143
          0.777778
Name: complexity, dtype: float64
Quartile Frequency Counts (tab-separated next to each quartile):
25%: 0.2142857142857143 Count (<= Q1): 237
```

```
50% (Median): 0.2763157894736842
                                       Count (Q1 < x \le Q2): 224
    75%: 0.3571428571428571 Count (Q2 < x <= Q3): 229
    ======= test multi df =======
    Shape: (184, 5)
    Columns: ['id', 'corpus', 'sentence', 'token', 'complexity']
    Data Types:
    id
                 object
    corpus
                 object
    sentence
                 object
    token
                 object
    complexity
                 float64
    dtype: object
    Missing Values (by column):
    id
                 0
    corpus
                 0
    sentence
                 0
    token
    complexity
    dtype: int64
    'complexity' Column Stats (incl. quartiles and median):
            184.000000
    count
    mean
              0.422312
              0.155785
    std
    min
              0.000000
    25%
              0.316667
    50%
              0.428571
    75%
              0.527778
              0.800000
    max
    Name: complexity, dtype: float64
    Quartile Frequency Counts (tab-separated next to each quartile):
    25%: 0.316666666666666 Count (<= Q1): 47
    50% (Median): 0.4285714285714286
                                       Count (Q1 < x \le Q2): 46
    75%: 0.52777777777778 Count (Q2 < x <= Q3): 46
    100% (Max): 0.8 Count (Q3 < x <= Max): 45
    [16]: print(train_single_df.head())
                                id corpus \
```

0 3ZLW647WALVGE8EBR50EGUBPU4P32A bible

```
1 34ROBODSP1ZBN3DVY8J8XSIY551E5C bible
     2 3S1WOPCJFGTJU2SGNAN2Y213N6WJE3 bible
     3 3BFNCI9LYKQN09BHXHH9CLSX5KP738 bible
     4 3G5RUKN2EC3YIWSKUXZ8ZVH95R49N2 bible
                                                 sentence
                                                              token complexity
     O Behold, there came up out of the river seven c...
                                                                     0.00000
     1 I am a fellow bondservant with you and with yo... brothers
                                                                     0.000000
     2 The man, the lord of the land, said to us, 'By... brothers
                                                                     0.050000
     3 Shimei had sixteen sons and six daughters; but... brothers
                                                                     0.150000
     4
                     "He has put my brothers far from me. brothers
                                                                       0.263889
[17]: print(train multi df.head())
                                    id corpus \
     0 3S37Y8CWI80N8KVM53U4E6JKCDC4WE bible
     1 3WGCNLZJKF877FYC1Q6COKNWTDWD11
                                        bible
     2 3UOMW19E6D6WQ5TH2HDD74IVKTP5CB bible
     3 36JW4WBR06KF9AXMUL4N4760MF8FHD bible
     4 3HRWUH63QU2FH9Q8R7MRNFC7JX2N5A bible
                                                                     token \
                                                 sentence
     0 but the seventh day is a Sabbath to Yahweh you...
                                                             seventh day
     1 But let each man test his own work, and then h...
                                                                own work
     2 To him who by understanding made the heavens; ... loving kindness
     3 Remember to me, my God, this also, and spare m... loving kindness
     4 Because your loving kindness is better than li... loving kindness
        complexity
     0
          0.027778
          0.050000
     1
     2
          0.050000
     3
          0.050000
          0.075000
     4
[18]: #@title Data Engineering
[19]: # Assuming you have already loaded the DataFrames:
      # train_single_df, train_multi_df, trial_val_single_df, trial_val_multi_df,_
       ⇔test_single_df, test_multi_df
      def print_distinct_values(df, column_name):
          """Prints the distinct values of a specified column in a DataFrame."""
          distinct_values = df[column_name].unique()
          print(f"Distinct values in '{column_name}' column:")
          for value in distinct_values:
              print(value)
          print("-" * 30)  # Separator
```

```
# Print distinct values for each DataFrame
print_distinct_values(train_single_df, "corpus")
print_distinct_values(train_multi_df, "corpus")
print_distinct_values(trial_val_single_df, "subcorpus")
print_distinct_values(trial_val_multi_df, "subcorpus")
print_distinct_values(test_single_df, "corpus")
print_distinct_values(test_multi_df, "corpus")
Distinct values in 'corpus' column:
bible
biomed
europarl
Distinct values in 'corpus' column:
bible
biomed
europarl
______
Distinct values in 'subcorpus' column:
bible
biomed
europarl
Distinct values in 'subcorpus' column:
bible
biomed
europarl
Distinct values in 'corpus' column:
bible
biomed
europarl
_____
Distinct values in 'corpus' column:
bible
biomed
europarl
```

0.1 standardize column headers: convert trial\_val header from 'subcorpus' to 'corpus'

```
# Verify the change (optional)
      print(trial_val_single_df.columns)
      print(trial_val_multi_df.columns)
     Index(['id', 'corpus', 'sentence', 'token', 'complexity'], dtype='object')
     Index(['id', 'corpus', 'sentence', 'token', 'complexity'], dtype='object')
[21]: dataframes = [train_single_df, train_multi_df, trial_val_single_df,_
       otrial_val_multi_df, test_single_df, test_multi_df]
      # Get the headers (column names) of the first DataFrame as a reference
      reference_headers = list(dataframes[0].columns)
      # Loop through the remaining DataFrames and compare headers
      all_headers_match = True
      for df in dataframes[1:]:
          if list(df.columns) != reference_headers:
              all_headers_match = False
              print(f"Headers do not match for DataFrame: {df.head(0)}") # Print<sub>||</sub>
       →which DataFrame has different headers
              break # Exit the loop if a mismatch is found
      # Print the result
      if all_headers_match:
          print("All DataFrames have matching headers.")
      else:
          print("Headers do not match for all DataFrames.")
```

All DataFrames have matching headers.

#### 0.2 Interrogate Span Length by Corpus Value by Data Split

```
[22]: # Analyzing sentence spans by complexity quartile and corpus

tokenizer = RegexpTokenizer(r'\w+') # setup tokenizer

def analyze_sentence_spans_by_corpus_and_quartile(dfs_dict):
    """
    Analyze sentence spans (length metrics) grouped by corpus and complexity_
    quartile
    for multiple dataframes.
    """
    results = []

for df_name, df in dfs_dict.items():
    print(f"Processing {df_name}...")
```

```
# Calculate complexity quartiles for this dataframe
      q1 = df['complexity'].quantile(0.25)
      q2 = df['complexity'].quantile(0.50)
      q3 = df['complexity'].quantile(0.75)
       # Define quartile ranges for labeling
      def get_quartile(x):
           if x <= q1:</pre>
               return 'Q1'
           elif x \ll q2:
               return 'Q2'
           elif x \ll q3:
               return 'Q3'
           else:
               return 'Q4'
       # Add quartile column
      df = df.copy()
      df['quartile'] = df['complexity'].apply(get_quartile)
       # Compute sentence metrics using RegexpTokenizer instead of \Box
⇒word tokenize
      def compute_span_metrics(sentence):
           if pd.isna(sentence):
               return pd.Series({'word_count': 0, 'char_count': 0, |

¬'avg_word_len': 0})
           # Use our tokenizer that doesn't require punkt_tab
           words = tokenizer.tokenize(sentence)
           word count = len(words)
           char_count = len(sentence)
           avg_word_len = np.mean([len(word) for word in words]) if word_count_
→> 0 else 0
           return pd.Series({'word_count': word_count, 'char_count':__
⇔char_count, 'avg_word_len': avg_word_len})
       # Apply the function to each sentence
      span_metrics = df['sentence'].apply(compute_span_metrics)
      df = pd.concat([df, span_metrics], axis=1)
       # Get corpus column name (could be 'corpus' or 'subcorpus')
      corpus_col = 'corpus' if 'corpus' in df.columns else 'subcorpus'
       # Group by corpus and quartile
      for corpus_name, corpus_df in df.groupby(corpus_col):
           for quartile, quartile_df in corpus_df.groupby('quartile'):
               # Calculate statistics
```

```
complexity_range = f"{quartile_df['complexity'].min():.

¬3f}-{quartile_df['complexity'].max():.3f}"

                stats = {
                    'Dataframe': df name,
                    'Corpus': corpus_name,
                    'Quartile': quartile,
                    'Complexity Range': complexity_range,
                    'Count': len(quartile_df),
                    'Avg Words': quartile_df['word_count'].mean(),
                    'Median Words': quartile_df['word_count'].median(),
                    'Min Words': quartile_df['word_count'].min(),
                    'Max Words': quartile_df['word_count'].max(),
                    'Std Words': quartile_df['word_count'].std(),
                    'Avg Chars': quartile_df['char_count'].mean(),
                    'Avg Word Len': quartile_df['avg_word_len'].mean()
                }
                results.append(stats)
    # Convert to DataFrame and sort
    results_df = pd.DataFrame(results)
    results df = results df.sort values(['Dataframe', 'Corpus', 'Quartile'])
    return results_df
# Create dictionary of dataframes
dfs = {
    'train_single_df': train_single_df,
    'train_multi_df': train_multi_df,
    'trial_val_single_df': trial_val_single_df,
    'trial_val_multi_df': trial_val_multi_df,
    'test_single_df': test_single_df,
    'test_multi_df': test_multi_df
}
# Run analysis
span_analysis = analyze_sentence_spans_by_corpus_and_quartile(dfs)
# Display results
pd.set_option('display.max_rows', None)
pd.set_option('display.max_columns', None)
pd.set_option('display.width', 1000)
display(span_analysis)
# Save the analysis results
results_path = os.path.join(dir_results, 'sentence_span_analysis.csv')
span_analysis.to_csv(results_path, index=False)
print(f"Analysis saved to: {results path}")
```

Processing train\_single\_df...
Processing train\_multi\_df...
Processing trial\_val\_single\_df...
Processing trial\_val\_multi\_df...
Processing test\_single\_df...
Processing test\_multi\_df...

| Dataframe               |                | -          | Corpus Quartile Complexity Range |             | Count Avg Words u |   |
|-------------------------|----------------|------------|----------------------------------|-------------|-------------------|---|
| ⊶Median Words Min Words |                | s Max Word | s Std Word                       | s Avg Chars | Avg Word Len      |   |
| 60                      | test_multi_df  | bible      | Q1                               | 0.025-0.317 | 26 23.076923      | Ш |
| $\hookrightarrow$       | 22.0 4.0       | 48.0       | 11.831900                        | 118.653846  | 4.128898          |   |
| 61                      | test_multi_df  | bible      | Q2                               | 0.325-0.417 | 11 20.545455      | Ш |
| $\hookrightarrow$       | 17.0 7.0       | 47.0       | 12.917923                        | 109.545455  | 4.209752          |   |
| 62                      | test_multi_df  | bible      | Q3                               | 0.432-0.528 | 18 21.111111      | Ш |
| $\hookrightarrow$       | 21.5 4.0       | 43.0       | 10.889222                        | 112.777778  | 4.474206          |   |
| 63                      | test_multi_df  | bible      | Q4                               | 0.542-0.694 | 11 22.363636      | Ш |
| $\hookrightarrow$       | 20.0 7.0       | 51.0       | 11.935432                        | 126.181818  | 4.605062          |   |
| 64                      | test_multi_df  | biomed     | Q1                               | 0.000-0.312 | 11 29.818182      | Ш |
| $\hookrightarrow$       | 29.0 17.0      | 47.0       | 8.388304                         | 195.727273  | 5.491145          |   |
| 65                      | test_multi_df  | biomed     | Q2                               | 0.324-0.417 | 11 27.090909      | Ш |
| $\hookrightarrow$       | 24.0 9.0       | 47.0       | 11.449494                        | 171.818182  | 5.436237          |   |
| 66                      | test_multi_df  | biomed     | Q3                               | 0.456-0.528 | 10 26.900000      | Ш |
| $\hookrightarrow$       | 26.5 10.0      | 49.0       | 10.712921                        | 177.500000  | 5.497409          |   |
| 67                      | test_multi_df  | biomed     | Q4                               | 0.562-0.800 | 21 32.285714      | Ш |
| $\hookrightarrow$       | 34.0 14.0      | 56.0       | 13.598319                        | 209.285714  | 5.460101          |   |
| 68                      | test_multi_df  | europarl   | Q1                               | 0.214-0.303 | 10 24.700000      | Ш |
| $\hookrightarrow$       |                | 56.0       | 14.189589                        |             | 5.049688          |   |
| 69                      | test_multi_df  | europarl   | Q2                               | 0.321-0.429 | 24 27.833333      | Ш |
| $\hookrightarrow$       | 27.0 9.0       | 73.0       | 15.352855                        | 172.291667  | 5.269610          |   |
| 70                      | test_multi_df  | europarl   | Q3                               | 0.432-0.516 | 18 32.944444      | Ш |
| $\hookrightarrow$       | 32.0 6.0       | 68.0       | 19.129504                        | 209.888889  | 5.512245          |   |
| 71                      | test_multi_df  | europarl   | Q4                               | 0.531-0.562 | 13 39.000000      | Ш |
| $\hookrightarrow$       | 36.0 6.0       | 95.0       | 29.631065                        | 237.076923  | 5.100616          |   |
| 48                      | test_single_df | bible      | Q1                               | 0.000-0.214 | 79 22.835443      | Ш |
| $\hookrightarrow$       | 22.0 7.0       | 49.0       | 10.602891                        | 116.797468  | 4.031532          |   |
| 49                      | test_single_df | bible      | Q2                               | 0.217-0.276 | 68 24.176471      | Ш |
| $\hookrightarrow$       | 21.0 2.0       | 77.0       | 14.393138                        | 125.955882  | 4.167352          |   |
| 50                      | test_single_df | bible      | Q3                               | 0.278-0.353 | 67 22.388060      | Ш |
| $\hookrightarrow$       | 20.0 4.0       | 63.0       | 11.306950                        | 119.731343  | 4.254090          |   |
| 51                      | test_single_df | bible      | Q4                               | 0.359-0.732 | 69 20.579710      | Ш |
| $\hookrightarrow$       | 19.0 1.0       | 55.0       | 11.264736                        | 110.550725  | 4.337010          |   |
| 52                      | test_single_df | biomed     | Q1                               | 0.000-0.214 | 75 27.080000      | Ш |
| $\hookrightarrow$       | 25.0 10.0      | 84.0       | 12.025603                        | 172.893333  | 5.271985          |   |
| 53                      | test_single_df | biomed     | Q2                               | 0.217-0.275 | 58 30.275862      | Ш |
| $\hookrightarrow$       | 26.0 10.0      | 83.0       |                                  | 197.775862  | 5.434573          |   |
| 54                      | test_single_df | biomed     | QЗ                               | 0.278-0.357 | 66 29.833333      | Ш |
| $\hookrightarrow$       | 29.0 13.0      | 85.0       |                                  | 191.863636  |                   | _ |
|                         |                |            |                                  |             |                   |   |

| 55                |                          |          |           |             | 90 31.144444  | Ш |
|-------------------|--------------------------|----------|-----------|-------------|---------------|---|
| $\hookrightarrow$ | 30.0 14.0                | 83.0     | 12.089146 | 203.055556  | 5.393138      |   |
| 56                | test_single_df           | -        |           |             | 83 25.337349  | ш |
| $\hookrightarrow$ | 21.0 3.0                 |          |           | 151.891566  | 5.044222      |   |
| 57                | test_single_df           | _        |           | 0.217-0.276 |               | ш |
| $\hookrightarrow$ | 30.0 1.0                 | 97.0     | 18.707061 | 195.653061  | 5.062296      |   |
| 58                | test_single_df           | -        |           |             | 96 33.000000  | Ш |
| $\hookrightarrow$ | 30.0 3.0                 |          |           |             | 5.124551      |   |
| 59                | test_single_df           | -        |           |             | 68 33.235294  | ш |
| $\hookrightarrow$ | 29.0 1.0                 | 130.0    | 20.440023 | 206.514706  | 5.164123      |   |
| 12                |                          | bible    |           |             |               | ш |
| $\hookrightarrow$ | 22.0 3.0                 | 67.0     | 12.429421 | 124.834356  | 4.232989      |   |
| 13                | train_multi_df           |          | Q2        | 0.304-0.409 | 132 24.053030 | Ш |
| $\hookrightarrow$ | 22.0 6.0                 | 65.0     | 11.738444 | 129.575758  | 4.302615      |   |
| 14                | train_multi_df           |          |           |             | 131 23.770992 | ш |
| $\hookrightarrow$ | 23.0 4.0                 | 50.0     | 11.158691 | 127.389313  | 4.324088      |   |
| 15                |                          | bible    | •         |             | 79 25.481013  | Ш |
| $\hookrightarrow$ | 24.0 3.0                 | 81.0     | 13.490605 | 139.240506  | 4.486716      |   |
| 16                | train_multi_df           | biomed   | Q1        | 0.028-0.303 | 87 29.091954  | ш |
| $\hookrightarrow$ | 28.0 9.0                 | 77.0     | 11.882792 | 185.954023  | 5.276290      |   |
| 17                | ${\tt train\_multi\_df}$ | biomed   | Q2        | 0.304-0.408 | 74 30.716216  | ш |
| $\hookrightarrow$ | 28.0 11.0                | 85.0     | 13.521693 | 195.864865  | 5.370313      |   |
| 18                | train_multi_df           | biomed   | QЗ        | 0.411-0.529 | 111 29.783784 | ш |
| $\hookrightarrow$ | 29.0 8.0                 | 61.0     | 10.912383 | 193.855856  | 5.430133      |   |
| 19                | train_multi_df           | biomed   | Q4        | 0.531-0.975 | 242 29.595041 | ш |
| $\hookrightarrow$ | 28.0 10.0                | 75.0     | 12.040443 | 194.995868  | 5.534629      |   |
| 20                | train_multi_df           | europarl | Q1        | 0.118-0.303 | 132 29.363636 | ш |
| $\hookrightarrow$ | 27.0 3.0                 | 101.0    | 17.874146 | 176.553030  | 5.002618      |   |
| 21                | train_multi_df           | europarl | Q2        | 0.304-0.409 | 171 31.654971 | Ш |
| $\hookrightarrow$ | 28.0 3.0                 | 108.0    | 19.099221 | 195.152047  | 5.176834      |   |
| 22                | train_multi_df           | europarl | Q3        | 0.411-0.529 | 138 33.398551 | Ш |
| $\hookrightarrow$ | 30.0 7.0                 | 101.0    | 18.992715 | 208.304348  | 5.286607      |   |
| 23                | train_multi_df           | europarl | Q4        | 0.533-0.750 | 57 34.596491  | ш |
| $\hookrightarrow$ | 31.0 6.0                 | 96.0     | 20.318763 | 218.350877  | 5.345891      |   |
| 0                 | train_single_df          | bible    | Q1        | 0.000-0.212 | 701 23.275321 | ш |
| $\hookrightarrow$ | 22.0 4.0                 | 61.0     | 11.760701 | 121.607703  | 4.126789      |   |
| 1                 | train_single_df          | bible    | Q2        | 0.212-0.279 | 640 23.753125 | ш |
| $\hookrightarrow$ | 22.0 3.0                 | 60.0     | 11.577932 | 124.576562  | 4.148961      |   |
| 2                 | train_single_df          | bible    | QЗ        | 0.281-0.375 | 624 23.823718 | ш |
| $\hookrightarrow$ | 22.0 3.0                 | 70.0     | 11.958906 | 126.230769  | 4.208102      |   |
| 3                 | train_single_df          | bible    | Q4        | 0.380-0.861 | 609 23.577997 | ш |
| $\hookrightarrow$ | 21.0 3.0                 | 69.0     | 12.461688 |             | 4.295608      | _ |
| 4                 | train_single_df          | biomed   | Q1        | 0.000-0.212 | 586 28.534130 | Ш |
| $\hookrightarrow$ | 27.0 2.0                 | 85.0     | 12.115387 |             | 5.319754      | _ |
| 5                 | train_single_df          | biomed   | Q2        | 0.212-0.279 | 583 30.435678 | Ш |
| $\hookrightarrow$ | 29.0 7.0                 | 92.0     |           |             | 5.285758      | _ |
|                   |                          |          |           |             |               |   |

| 6                 | train_single_df<br>28.0 4.0 |          |                 | 0.281-0.375<br>191.050076 |                           | Ш |
|-------------------|-----------------------------|----------|-----------------|---------------------------|---------------------------|---|
|                   |                             |          |                 |                           |                           |   |
| 7 ↔               | train_single_df<br>28.0 3.0 | 85.0     | Q4<br>12.246613 |                           | 748 29.176471<br>5.298112 | Ш |
| 8                 | train_single_df             | europarl | Q1              | 0.025-0.212               | 641 26.761310             | ш |
| $\hookrightarrow$ | 24.0 2.0                    | _        | 15.230853       | 159.180967                | 4.942557                  | _ |
| 9                 | train_single_df             | europarl | Q2              | 0.212-0.279               | 714 30.420168             | Ш |
| $\hookrightarrow$ | 27.0 1.0                    | -        |                 | 183.093838                | 4.995672                  |   |
| 10                | train_single_df             | europarl | QЗ              | 0.281-0.375               | 701 30.523538             | Ш |
| $\hookrightarrow$ | 28.0 1.0                    | 122.0    | 18.163026       | 185.840228                | 5.114587                  |   |
| 11                | train_single_df             | europarl | Q4              | 0.381-0.775               | 456 33.528509             | Ш |
| $\hookrightarrow$ | 31.0 2.0                    | 235.0    | 21.704693       | 203.592105                | 5.054701                  |   |
| 36                | trial_val_multi_df          | bible    | Q1              | 0.000-0.292               | 11 26.272727              | Ш |
| $\hookrightarrow$ | 21.0 13.0                   | 64.0     | 13.950562       | 141.363636                | 4.282457                  |   |
| 37                | trial_val_multi_df          | bible    | Q2              | 0.333-0.400               | 7 20.571429               | Ш |
| $\hookrightarrow$ | 23.0 5.0                    | 28.0     | 7.412987        | 110.857143                | 4.279406                  |   |
| 38                | trial_val_multi_df          | bible    | Q3              | 0.425-0.500               | 5 19.600000               | Ш |
| $\hookrightarrow$ | 19.0 9.0                    | 32.0     | 8.905055        | 109.200000                | 4.431391                  |   |
| 39                | trial_val_multi_df          | bible    | Q4              | 0.525-0.661               | 6 22.333333               | Ш |
| $\hookrightarrow$ | 20.5 9.0                    | 44.0     | 12.242004       | 117.833333                | 4.178525                  |   |
| 40                | trial_val_multi_df          | biomed   | Q1              | 0.083-0.303               | 6 26.833333               | Ш |
| $\hookrightarrow$ | 25.0 15.0                   | 49.0     | 11.771434       | 159.166667                | 4.899969                  |   |
| 41                | trial_val_multi_df          | biomed   | Q2              | 0.317-0.422               | 7 25.428571               | Ш |
| $\hookrightarrow$ | 21.0 15.0                   | 48.0     | 11.588171       | 156.000000                | 5.194383                  |   |
| 42                | trial_val_multi_df          | biomed   | QЗ              | 0.438-0.513               | 6 37.833333               | Ш |
| $\hookrightarrow$ | 39.5 26.0                   | 44.0     | 6.675827        | 247.500000                | 5.438593                  |   |
| 43                | trial_val_multi_df          | biomed   | Q4              | 0.537-0.825               | 14 30.642857              | Ш |
| $\hookrightarrow$ | 29.5 17.0                   | 43.0     | 9.849695        | 211.428571                | 5.730623                  |   |
| 44                | trial_val_multi_df          | europarl | Q1              | 0.176-0.306               | 8 30.000000               | Ш |
| $\hookrightarrow$ | 25.5 4.0                    | 64.0     | 20.361027       | 186.750000                | 5.306837                  |   |
| 45                | trial_val_multi_df          | europarl | Q2              | 0.312-0.412               | 11 47.909091              | Ш |
| $\hookrightarrow$ | 46.0 24.0                   | 78.0     | 18.651834       | 296.909091                | 5.058375                  |   |
| 46                | trial_val_multi_df          | europarl | QЗ              | 0.432-0.500               | 13 26.307692              | Ш |
| $\hookrightarrow$ | 26.0 5.0                    | 66.0     | 18.167666       | 166.153846                | 5.263847                  |   |
| 47                | trial_val_multi_df          | europarl | Q4              | 0.515-0.714               | 5 26.400000               | ш |
| $\hookrightarrow$ | 15.0 6.0                    | 66.0     | 24.316661       | 164.600000                | 4.998182                  |   |
| 24                | trial_val_single_df         | bible    | Q1              | 0.000-0.214               | 52 26.750000              | ш |
| $\hookrightarrow$ | 26.0 5.0                    | 73.0     | 15.530962       | 137.230769                | 4.071006                  |   |
| 25                | trial_val_single_df         | bible    | Q2              | 0.217-0.266               | 38 24.868421              | Ш |
| $\hookrightarrow$ | 23.0 7.0                    | 50.0     | 10.768249       | 131.236842                | 4.195550                  |   |
| 26                | trial_val_single_df         | bible    | QЗ              | 0.268-0.355               | 26 22.884615              | Ш |
| $\hookrightarrow$ | 20.5 5.0                    | 44.0     | 9.961233        | 121.269231                | 4.312026                  |   |
| 27                | trial_val_single_df         | bible    | Q4              | 0.361-0.633               | 27 25.666667              | ш |
| $\hookrightarrow$ | 23.0 6.0                    | 49.0     | 12.554497       | 137.555556                | 4.212685                  |   |
| 28                | trial_val_single_df         | biomed   | Q1              | 0.028-0.214               | 21 25.571429              | Ш |
| $\hookrightarrow$ | 21.0 13.0                   | 65.0     | 11.543706       | 163.904762                | 5.305404                  |   |
|                   |                             |          |                 |                           |                           |   |

```
29 trial_val_single_df
                          biomed
                                       Q2
                                               0.217-0.267
                                                              28 30.571429
        27.5
                              57.0 12.099674 198.142857
                                                              5.315287
                   11.0
30 trial val single df
                                                              38 32.105263
                          biomed
                                       QЗ
                                               0.268-0.359
        29.0
                              61.0
                                   12.710476
                                              206.947368
                                                              5.364934
                   11.0
                                                              48 25.145833
31 trial_val_single_df
                          biomed
                                       Q4
                                               0.364-0.875
        25.5
                                   11.721937
                                                              5.439709
                    6.0
                              56.0
                                              163.979167
32 trial_val_single_df europarl
                                       Q1
                                               0.050 - 0.214
                                                              33 31.969697
        28.0
                    5.0
                             81.0
                                   20.356947
                                              185.969697
                                                              4.799024
33 trial_val_single_df
                                               0.217-0.267
                                                              41 28.463415
                        europarl
                                       Q2
                              71.0 15.386841
        28.0
                    4.0
                                              172.780488
                                                              4.997706
                                                              39 30.282051
34 trial_val_single_df europarl
                                       QЗ
                                               0.268-0.359
        28.0
                    3.0
                              99.0 20.040681
                                              184.358974
                                                              5.086945
                                               0.367-0.605
                                                              30 35.700000
35 trial_val_single_df
                        europarl
                                       Q4
        30.5
                    5.0
                             77.0
                                   20.142852 215.400000
                                                              4.910759
```

#### Analysis saved to:

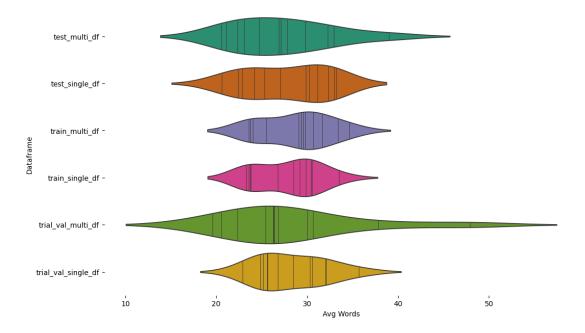
/content/drive/MyDrive/266-final/results/sentence\_span\_analysis.csv

```
[23]: from matplotlib import pyplot as plt
import seaborn as sns
figsize = (12, 1.2 * len(span_analysis['Dataframe'].unique()))
plt.figure(figsize=figsize)
sns.violinplot(span_analysis, x='Avg Words', y='Dataframe', inner='stick',
palette='Dark2')
sns.despine(top=True, right=True, bottom=True, left=True)
```

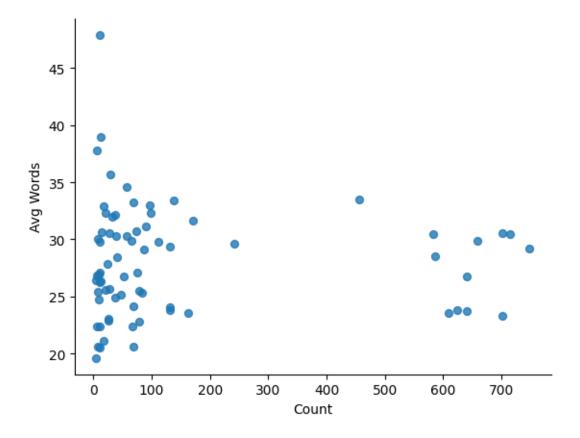
<ipython-input-23-00a8ad5642c1>:5: FutureWarning:

Passing `palette` without assigning `hue` is deprecated and will be removed in v0.14.0. Assign the `y` variable to `hue` and set `legend=False` for the same effect.

sns.violinplot(span\_analysis, x='Avg Words', y='Dataframe', inner='stick',
palette='Dark2')



[24]: from matplotlib import pyplot as plt span\_analysis.plot(kind='scatter', x='Count', y='Avg Words', s=32, alpha=.8) plt.gca().spines[['top', 'right',]].set\_visible(False)

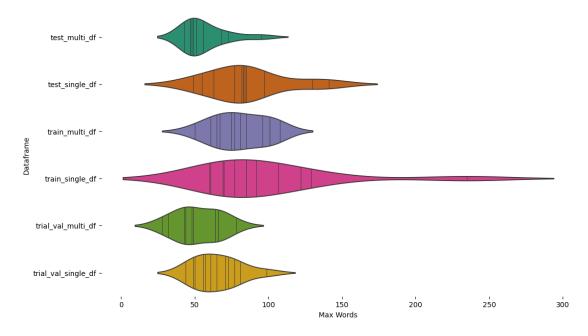


```
[25]: from matplotlib import pyplot as plt
import seaborn as sns
figsize = (12, 1.2 * len(span_analysis['Dataframe'].unique()))
plt.figure(figsize=figsize)
sns.violinplot(span_analysis, x='Max Words', y='Dataframe', inner='stick',
palette='Dark2')
sns.despine(top=True, right=True, bottom=True, left=True)
```

<ipython-input-25-01bf0c89d620>:5: FutureWarning:

Passing `palette` without assigning `hue` is deprecated and will be removed in v0.14.0. Assign the `y` variable to `hue` and set `legend=False` for the same effect.

sns.violinplot(span\_analysis, x='Max Words', y='Dataframe', inner='stick',
palette='Dark2')



```
# Remove spines for cleaner look
g.despine(top=True, right=True, bottom=True, left=True)

# Adjust layout and display the plot
plt.tight_layout()
plt.show()
```

/usr/local/lib/python3.11/dist-packages/seaborn/axisgrid.py:718: UserWarning: Using the violinplot function without specifying `order` is likely to produce an incorrect plot.

warnings.warn(warning)

/usr/local/lib/python3.11/dist-packages/seaborn/axisgrid.py:854: FutureWarning:

Passing `palette` without assigning `hue` is deprecated and will be removed in v0.14.0. Assign the `y` variable to `hue` and set `legend=False` for the same effect.

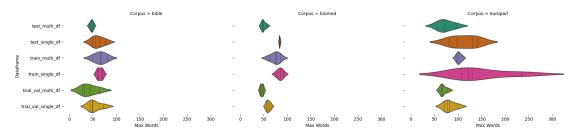
```
func(*plot_args, **plot_kwargs)
/usr/local/lib/python3.11/dist-packages/seaborn/axisgrid.py:854: FutureWarning:
```

Passing `palette` without assigning `hue` is deprecated and will be removed in v0.14.0. Assign the `y` variable to `hue` and set `legend=False` for the same effect.

```
func(*plot_args, **plot_kwargs)
/usr/local/lib/python3.11/dist-packages/seaborn/axisgrid.py:854: FutureWarning:
```

Passing `palette` without assigning `hue` is deprecated and will be removed in v0.14.0. Assign the `y` variable to `hue` and set `legend=False` for the same effect.

func(\*plot\_args, \*\*plot\_kwargs)



• decision: no modifications to sentence spans will be applied, except for Contraction standardization

### 0.3 Normalize / Eliminate Contractions

```
[27]: # --- STEP 1: CREATE A UTILITY FUNCTION ---
      def expand_contractions_in_df(df):
          1) Creates a new column 'sentence_no_contractions' by expanding any_
       \hookrightarrow contractions.
          2) Identifies rows where a contraction was actually expanded (the text_{\sqcup}
       \hookrightarrow changed).
          3) Returns the updated DataFrame and a grouped subset of rows for printing \Box
       \hookrightarrow examples.
          11 11 11
          df = df.copy()
          # Create the new column with expanded contractions
          df['sentence_no_contractions'] = df['sentence'].apply(
              lambda s: contractions.fix(s) if pd.notna(s) else s
          )
          # Check if anything changed after expansion:
          df['contraction_expanded'] = df.apply(
              lambda row: row['sentence'] != row['sentence_no_contractions'], axis=1
          )
          # Group by corpus
          results_by_corpus = {}
          for corpus_val, group in df.groupby('corpus'):
               # Filter to rows that actually had a contraction
              changed_rows = group[group['contraction_expanded']]
               # Take the first 3
              first_three = changed_rows.head(3)
              results_by_corpus[corpus_val] = first_three
          return df, results_by_corpus
      # --- STEP 2: APPLY TO ALL 6 DATAFRAMES ---
      dataframes info = [
          ("train_single_df", train_single_df),
          ("train_multi_df", train_multi_df),
          ("trial_val_single_df", trial_val_single_df),
          ("trial_val_multi_df", trial_val_multi_df),
          ("test_single_df", test_single_df),
          ("test_multi_df", test_multi_df),
      ]
```

```
for df_name, df in dataframes_info:
    # Expand contractions and collect examples
   updated_df, corpus_examples = expand_contractions_in_df(df)
    # Overwrite the old DataFrame variable if desired:
   # (So that we retain the new column for future tasks)
    # Otherwise, you can store it in a new variable.
   globals()[df_name] = updated_df
    # Print 3 "before" and 3 "after" examples per corpus
   print(f"\n{'='*60}")
   print(f"DataFrame: {df_name}")
   print(f"{'='*60}")
   for corpus_val in sorted(corpus_examples.keys()):
       subset = corpus_examples[corpus_val]
        # Only proceed if we found any changed rows in this corpus
        if len(subset) == 0:
            continue
       print(f"\n Corpus: {corpus_val}")
        # Print BEFORE lines
       print(" -- BEFORE --")
       for _, row in subset.iterrows():
           print(f" {row['sentence']}")
        # Print AFTER lines
       print(" -- AFTER --")
       for _, row in subset.iterrows():
           print(f"
                          {row['sentence_no_contractions']}")
```

\_\_\_\_\_

```
Corpus: bible -- BEFORE --
```

Shimei had sixteen sons and six daughters; but his brothers didn't have many children, neither did all their family multiply like the children of Judah.

When his speech is charming, don't believe him; for there are seven abominations in his heart.

Jesus said, "Father, forgive them, for they don't know what they are doing."  $% \frac{1}{2} \left( \frac{1}{2} \right) = \frac{1}{2} \left( \frac{1}{2} \right) \left$ 

-- AFTER --

Shimei had sixteen sons and six daughters; but his brothers did not have many children, neither did all their family multiply like the children of Judah.

When his speech is charming, do not believe him; for there are seven abominations in his heart.

Jesus said, "Father, forgive them, for they do not know what they are doing."

Corpus: biomed -- BEFORE --

Although missense mutation of ITPR1 had previously been ruled out [2] and the mode of inheritance was inconsistent with that seen in the Itpr1 $\Delta$ 18 and Itpr1opt mice, the phenotypic presence of ataxia in the mice led us to reexamine this candidate gene as a possible cause of SCA15.

Human germline mutations in APC cause FAP [4,5], which is characterized by hundreds of adenomatous colorectal polyps, with an almost inevitable progression to colorectal cancer in the third and fourth decades of life.

Null mutations in Bmpr1a cause early embryonic lethality, with defects in gastrulation similar to those seen in mice with mutations in Bmp4 (Mishina et al. 1995; Winnier et al. 1995).

-- AFTER --

Although missense mutation of ITPR1 had previously been ruled out [2] and the mode of inheritance was inconsistent with that seen in the Itpr1 $\Delta$ 18 and Itpr1opt mice, the phenotypic presence of ataxia in the mice led us to reexamine this candidate gene as a possible because of SCA15.

Human germline mutations in APC because FAP [4,5], which is characterized by hundreds of adenomatous colorectal polyps, with an almost inevitable progression to colorectal cancer in the third and fourth decades of life.

Null mutations in Bmpr1a because early embryonic lethality, with defects in gastrulation similar to those seen in mice with mutations in Bmp4 (Mishina et al. 1995; Winnier et al. 1995).

Corpus: europarl
-- BEFORE --

At the same time, you will also have an important role in winning over the general public of the Member States to the cause of enlargement, of enlargement based on conditionality.

the recommendation for second reading from the Committee on Transport and Tourism on the common position adopted by the Council with a view to the adoption of a Regulation of the European Parliament and of the Council establishing common rules concerning the conditions to be complied with to pursue the occupation of road transport operator and repealing Council Directive 96/26/EC (11783/1/2008 - C6-0015/2009 - (Rapporteur: Silvia-Adriana Ţicău), and

Yet, although credit rating agencies were not the main cause of the recent financial crisis, they did have a harmful influence.

-- AFTER --

At the same time, you will also have an important role in winning over the general public of the Member States to the because of enlargement, of enlargement based on conditionality. the recommendation for second reading from the Committee on Transport and Tourism on the common position adopted by the Council with a view to the adoption of a Regulation of the European Parliament and of the Council establishing common rules concerning the conditions to be complied with to pursue the occupation of road transport operator and repealing Council Directive 96/26/EC (11783/1/2008 - C6-0015/2009 - (Rapporteur: Silvia-Adriana Ţicāyou), and

Yet, although credit rating agencies were not the main because of the recent financial crisis, they did have a harmful influence.

\_\_\_\_\_\_

DataFrame: train\_multi\_df

\_\_\_\_\_

Corpus: bible -- BEFORE --

Jahath was the chief, and Zizah the second: but Jeush and Beriah didn't have many sons; therefore they became a fathers' house in one reckoning.

But Yahweh said to Samuel, "Don't look on his face, or on the height of his stature; because I have rejected him: for I see not as man sees; for man looks at the outward appearance, but Yahweh looks at the heart."

Because indeed a notable miracle has been done through them, as can be plainly seen by all who dwell in Jerusalem, and we can't deny it.

-- AFTER --

Jahath was the chief, and Zizah the second: but Jeush and Beriah did not have many sons; therefore they became a fathers' house in one reckoning.

But Yahweh said to Samuel, "Do not look on his face, or on the height of his stature; because I have rejected him: for I see not as man sees; for man looks at the outward appearance, but Yahweh looks at the heart."

Because indeed a notable miracle has been done through them, as can be plainly seen by all who dwell in Jerusalem, and we cannot deny it.

Corpus: biomed -- BEFORE --

The aim in the present study was to determine the location of pendrin and the cause of deafness in Slc26a4-/- mice.

These characteristics should make RMCE-ASAP a robust and general technology for analysis of mammalian genes under conditions that preserve normal control mechanisms in different tissues.

It was also demonstrated that mutations leading to abolishment of the enzymatic activity of CLN2 were the direct cause of a fatal inherited neurodegenerative disease, classical late-infantile neuronal ceroid lipofuscinosis [2].

-- AFTER --

The aim in the present study was to determine the location of pendrin and the because of deafness in Slc26a4-/- mice.

These characteristics should make RMCE-AS SOON AS POSSIBLE a robust and general technology for analysis of mammalian genes under conditions that

preserve normal control mechanisms in different tissues.

It was also demonstrated that mutations leading to abolishment of the enzymatic activity of CLN2 were the direct because of a fatal inherited neurodegenerative disease, classical late-infantile neuronal ceroid lipofuscinosis [2].

Corpus: europarl
-- BEFORE --

Account must also be taken of the costs to health, the environment and the climate of the fact that vehicles emit different types of particles and that, in burning fossil fuels, they cause increased pollution and thus more global warming.

However, this unequal trade relationship is not the only cause for concern; another is the case of unsafe products coming from China.

(IT) Madam President, ladies and gentlemen, the oral amendment that our Group is proposing involves replacing the words 'all forms of glorifying' by the word 'apology'.

-- AFTER --

Account must also be taken of the costs to health, the environment and the climate of the fact that vehicles emit different types of particles and that, in burning fossil fuels, they because increased pollution and thus more global warming.

However, this unequal trade relationship is not the only because for concern; another is the case of unsafe products coming from China.

(IT) Madam President, ladies and gentlemen, the oral amendment that our Group is proposing involves replacing the words forms of glorifying' by the word 'apology'.

\_\_\_\_\_

DataFrame: trial\_val\_single\_df

\_\_\_\_\_\_

Corpus: bible -- BEFORE --

Don't curse the king, no, not in your thoughts; and don't curse the rich in your bedroom: for a bird of the sky may carry your voice, and that which has wings may tell the matter.

The young man didn't wait to do this thing, because he had delight in Jacob's daughter, and he was honored above all the house of his father.

If the axe is blunt, and one doesn't sharpen the edge, then he must use more strength; but skill brings success.

-- AFTER --

Do not curse the king, no, not in your thoughts; and do not curse the rich in your bedroom: for a bird of the sky may carry your voice, and that which has wings may tell the matter.

The young man did not wait to do this thing, because he had delight in Jacob's daughter, and he was honored above all the house of his father.

If the axe is blunt, and one does not sharpen the edge, then he must use

more strength; but skill brings success.

Corpus: biomed -- BEFORE --

For example, the non-BC individual and BC individual groups are not perfectly matched with respect to age, gender or smoking history (Table 1) and each of these factors could contribute to the observed difference in correlation between groups.

EM and ER conducted transmission electron microscopy.

-- AFTER --

For example, the non-BECAUSE individual and BECAUSE individual groups are not perfectly matched with respect to age, gender or smoking history (Table 1) and each of these factors could contribute to the observed difference in correlation between groups.

THEM and ER conducted transmission electron microscopy.

Corpus: europarl
-- BEFORE --

With their help, John has sought to shed light on what has been a very murky area, and to bring clarity where uncertainty prevailed before, based consistently on the twin principles that the patient must always come first and that patient choice should be determined by needs and not by means.

-- AFTER --

With their help, John has sought to she would light on what has been a very murky area, and to bring clarity where uncertainty prevailed before, based consistently on the twin principles that the patient must always come first and that patient choice should be determined by needs and not by means.

Corpus: bible -- BEFORE --

the ten sons of Haman the son of Hammedatha, the Jew's enemy, but they didn't lay their hand on the plunder.

Hezekiah listened to them, and showed them all the house of his precious things, the silver, and the gold, and the spices, and the precious oil, and the house of his armor, and all that was found in his treasures: there was nothing in his house, nor in all his dominion, that Hezekiah didn't show them.

Of Manasseh also there fell away some to David, when he came with the Philistines against Saul to battle; but they didn't help them; for the lords of the Philistines sent him away after consultation, saying, "He will fall away to his master Saul to the jeopardy of our heads."

#### -- AFTER --

the ten sons of Haman the son of Hammedatha, the Jew's enemy, but they did not lay their hand on the plunder.

Hezekiah listened to them, and showed them all the house of his precious things, the silver, and the gold, and the spices, and the precious oil, and the house of his armor, and all that was found in his treasures: there was nothing in his house, nor in all his dominion, that Hezekiah did not show them.

Of Manasseh also there fell away some to David, when he came with the Philistines against Saul to battle; but they did not help them; for the lords of the Philistines sent him away after consultation, saying, "He will fall away to his master Saul to the jeopardy of our heads."

## Corpus: biomed -- BEFORE --

In that study, there was a tendency towards correlation in transcript abundance between several pairs of antioxidant or DNA repair genes in non-BC individuals, but not in BC individuals.

This, in turn, leads to increased representation among BC individuals of individuals with lack of correlation between CEBPG and each of the affected antioxidant and/or DNA repair genes.

The 'pregnancy rate' in mice is defined as successful pregnancies per detected vaginal plug, a phenotype associated with early pregnancy failure, which in turn possibly could have an inflammatory cause.

#### -- AFTER --

In that study, there was a tendency towards correlation in transcript abundance between several pairs of antioxidant or DNA repair genes in non-BECAUSE individuals, but not in BECAUSE individuals.

This, in turn, leads to increased representation among BECAUSE individuals of individuals with lack of correlation between CEBPG and each of the affected antioxidant and/or DNA repair genes.

The 'pregnancy rate' in mice is defined as successful pregnancies per detected vaginal plug, a phenotype associated with early pregnancy failure, which in turn possibly could have an inflammatory because.

# Corpus: europarl -- BEFORE --

The next item is the oral question to the Commission (B7-0240/2009) by Silvia-Adriana Ţicău, Brian Simpson, János Áder, Hannes Swoboda, Eva Lichtenberger, Michael Cramer, Saïd El Khadraoui, Mathieu Grosch, Iuliu Winkler, Victor Boştinaru, Ioan Mircea Paşcu, Marian-Jean Marinescu, Ivailo Kalfin, Norica Nicolai, Dirk Sterckx, Csaba Sándor Tabajdi, Michael Theurer, Ismail Ertug, Inés Ayala Sender, Jiří Havel, Edit Herczog, Stanimir Ilchev, Iliana Malinova Iotova, Jelko Kacin, Evgeni Kirilov, Ádám Kósa, Ioan Enciu, Eduard Kukan, Gesine Meissner, Alajos Mészáros, Nadezhda Neynsky, Katarína Neveďalová, Daciana Octavia Sârbu, Vilja Savisaar, Olga Sehnalová, Catherine Stihler, Peter van Dalen, Louis Grech, Corina Creţu, George Sabin Cutaş, Vasilica Viorica Dăncilă, Cătălin Sorin Ivan, Tanja Fajon, Kinga Göncz, Antonyia Parvanova, Adina-Ioana Vălean and Rovana Plumb, on the European Strategy for the Danube

Region.

#### -- AFTER --

The next item is the oral question to the Commission (B7-0240/2009) by Silvia-Adriana Ţicăyou, Brian Simpson, János Áder, Hannes Swoboda, Eva Lichtenberger, Michael Cramer, Saïd El Khadraoui, Mathieu Grosch, Iuliu Winkler, Victor Boştinaru, Ioan Mircea Paşcu, Marian-Jean Marinescu, Ivailo Kalfin, Norica Nicolai, Dirk Sterckx, Csaba Sándor Tabajdi, Michael Theurer, Ismail Ertug, Inés Ayala Sender, Jiří Havel, Edit Herczog, Stanimir Ilchev, Iliana Malinova Iotova, Jelko Kacin, Evgeni Kirilov, Ádám Kósa, Ioan Enciu, Eduard Kukan, Gesine Meissner, Alajos Mészáros, Nadezhda Neynsky, Katarína Neveďalová, Daciana Octavia Sârbu, Vilja Savisaar, Olga Sehnalová, Catherine Stihler, Peter van Dalen, Louis Grech, Corina Creţyou, George Sabin Cutaş, Vasilica Viorica Dăncilă, Cătălin Sorin Ivan, Tanja Fajon, Kinga Göncz, Antonyia Parvanova, Adina-Ioana Vălean and Rovana Plumb, on the European Strategy for the Danube Region.

\_\_\_\_\_\_

DataFrame: test\_multi\_df

\_\_\_\_\_

Corpus: bible -- BEFORE --

Yet he didn't leave himself without witness, in that he did good and gave you rains from the sky and fruitful seasons, filling our hearts with food and gladness."

When he has leveled its surface, doesn't he plant the dill, and scatter the cumin seed, and put in the wheat in rows, the barley in the appointed place, and the spelt in its place?

Don't count your handmaid for a wicked woman; for I have been speaking out of the abundance of my complaint and my provocation."

-- AFTER --

Yet he did not leave himself without witness, in that he did good and gave you rains from the sky and fruitful seasons, filling our hearts with food and gladness."

When he has leveled its surface, does not he plant the dill, and scatter the cumin seed, and put in the wheat in rows, the barley in the appointed place, and the spelt in its place?

Do not count your handmaid for a wicked woman; for I have been speaking out of the abundance of my complaint and my provocation."

False

```
False
     False
     False
     False
     False
[29]: dataframes = {
          "train_single_df": train_single_df,
          "train_multi_df": train_multi_df,
          "trial_val_single_df": trial_val_single_df,
          "trial val multi df": trial val multi df,
          "test_single_df": test_single_df,
          "test_multi_df": test_multi_df
      }
      total_true_counts = 0
      for df_name, df in dataframes.items():
          true_count = df['contraction_expanded'].sum()
          print(f"{df_name}: {true_count} True values in 'contraction_expanded'")
          total_true_counts += true_count
      print(f"\nTotal True values across all dataframes: {total_true_counts}")
     train_single_df: 254 True values in 'contraction_expanded'
     train_multi_df: 54 True values in 'contraction_expanded'
     trial_val_single_df: 16 True values in 'contraction_expanded'
     trial_val_multi_df: 0 True values in 'contraction_expanded'
     test_single_df: 31 True values in 'contraction_expanded'
     test_multi_df: 7 True values in 'contraction_expanded'
     Total True values across all dataframes: 362
[30]: # verify column headers
      dataframes = [train_single_df, train_multi_df, trial_val_single_df,_

¬trial_val_multi_df, test_single_df, test_multi_df]
      for df in dataframes:
        print(df.info())
     <class 'pandas.core.frame.DataFrame'>
     RangeIndex: 7662 entries, 0 to 7661
     Data columns (total 7 columns):
          Column
                                    Non-Null Count Dtype
         _____
      0
          id
                                    7662 non-null
                                                    object
                                    7662 non-null
         corpus
                                                    object
      2
          sentence
                                    7662 non-null
                                                    object
         token
                                    7655 non-null
                                                    object
          complexity
                                    7662 non-null
                                                    float64
```

```
sentence_no_contractions 7662 non-null
                                              object
 6
    contraction_expanded
                              7662 non-null
                                              bool
dtypes: bool(1), float64(1), object(5)
memory usage: 366.8+ KB
None
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1517 entries, 0 to 1516
Data columns (total 7 columns):
    Column
                              Non-Null Count Dtype
    -----
                              -----
 0
    id
                              1517 non-null
                                              object
 1
    corpus
                              1517 non-null
                                              object
 2
    sentence
                              1517 non-null
                                              object
 3
    token
                              1517 non-null
                                              object
 4
    complexity
                              1517 non-null
                                              float64
 5
    sentence_no_contractions 1517 non-null
                                              object
    contraction_expanded
                              1517 non-null
                                              bool
dtypes: bool(1), float64(1), object(5)
memory usage: 72.7+ KB
None
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 421 entries, 0 to 420
Data columns (total 7 columns):
    Column
                              Non-Null Count Dtype
    _____
                              -----
 0
    id
                              421 non-null
                                              object
 1
                              421 non-null
                                              object
    corpus
 2
    sentence
                              421 non-null
                                              object
 3
    token
                              421 non-null
                                              object
 4
    complexity
                              421 non-null
                                              float64
 5
    sentence_no_contractions 421 non-null
                                              object
    contraction_expanded
                              421 non-null
                                              bool
dtypes: bool(1), float64(1), object(5)
memory usage: 20.3+ KB
None
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 99 entries, 0 to 98
Data columns (total 7 columns):
```

| Data  | columns (total / columns) | •              |         |  |  |
|---|---------------------------|----------------|---------|--|--|
| #   | Column                    | Non-Null Count | Dtype   |  |  |
|   |                           |                |         |  |  |
| 0   | id                        | 99 non-null    | object  |  |  |
| 1   | corpus                    | 99 non-null    | object  |  |  |
| 2   | sentence                  | 99 non-null    | object  |  |  |
| 3   | token                     | 99 non-null    | object  |  |  |
| 4   | complexity                | 99 non-null    | float64 |  |  |
| 5   | sentence_no_contractions  | 99 non-null    | object  |  |  |
| 6   | contraction_expanded      | 99 non-null    | bool    |  |  |
| <pre>dtypes: bool(1), float64(1), object(5)</pre> |                           |                |         |  |  |

<class 'pandas.core.frame.DataFrame'> RangeIndex: 917 entries, 0 to 916 Data columns (total 7 columns): Column Non-Null Count Dtype 0 id 917 non-null object 1 corpus 917 non-null object 2 sentence 917 non-null object 3 token 917 non-null object 4 complexity 917 non-null float64 5 sentence\_no\_contractions 917 non-null object contraction\_expanded 917 non-null bool dtypes: bool(1), float64(1), object(5) memory usage: 44.0+ KB None <class 'pandas.core.frame.DataFrame'> RangeIndex: 184 entries, 0 to 183 Data columns (total 7 columns): Column Non-Null Count Dtype \_\_\_\_\_ \_\_\_\_\_ 0 id 184 non-null object 1 184 non-null corpus object 2 sentence 184 non-null object 3 token 184 non-null object 4 float64 complexity 184 non-null sentence\_no\_contractions 184 non-null object contraction\_expanded 184 non-null bool dtypes: bool(1), float64(1), object(5) memory usage: 8.9+ KB None [31]: # inspect each df dataframes = [train\_single\_df, train\_multi\_df, trial\_val\_single\_df,\_ strial\_val\_multi\_df, test\_single\_df, test\_multi\_df] for df in dataframes: print(df.head()) id corpus token complexity sentence sentence\_no\_contractions contraction\_expanded O 3ZLW647WALVGE8EBR50EGUBPU4P32A bible Behold, there came up out of the river 0.000000 Behold, there came up out of the river seven seven c... river False C... 1 34R0B0DSP1ZBN3DVY8J8XSIY551E5C bible I am a fellow bondservant with you and 0.000000 I am a fellow bondservant with you and with with yo... brothers

memory usage: 4.9+ KB

None

yo... False

2 3S1WOPCJFGTJU2SGNAN2Y213N6WJE3 bible The man, the lord of the land, said to us, 'By... brothers 0.050000 The man, the lord of the land, said to us, 'By... False

- 3 3BFNCI9LYKQN09BHXHH9CLSX5KP738 bible Shimei had sixteen sons and six daughters; but... brothers 0.150000 Shimei had sixteen sons and six daughters; but... True
- 4 3G5RUKN2EC3YIWSKUXZ8ZVH95R49N2 bible "He has put my brothers far from me. brothers 0.263889 "He has put my brothers far from me. False

id corpus

sentence token complexity

sentence\_no\_contractions contraction\_expanded

- 0 3S37Y8CWI80N8KVM53U4E6JKCDC4WE bible but the seventh day is a Sabbath to Yahweh you... seventh day 0.027778 but the seventh day is a Sabbath to Yahweh you... False
- 1 3WGCNLZJKF877FYC1Q6COKNWTDWD11 bible But let each man test his own work, and then h... own work 0.050000 But let each man test his own work, and then h... False
- 2 3UOMW19E6D6WQ5TH2HDD74IVKTP5CB bible To him who by understanding made the heavens; ... loving kindness 0.050000 To him who by understanding made the heavens; ... False
- 3 36JW4WBR06KF9AXMUL4N4760MF8FHD bible Remember to me, my God, this also, and spare m... loving kindness 0.050000 Remember to me, my God, this also, and spare m... False
- 4 3HRWUH63QU2FH9Q8R7MRNFC7JX2N5A bible Because your loving kindness is better than li... loving kindness 0.075000 Because your loving kindness is better than li... False

id corpus

sentence token complexity contraction\_expanded

 ${\tt sentence\_no\_contractions}$ 

- 0 3QI9WAYOGQB8GQIR4MDIEF0D2RLS67 bible They will not hurt nor destroy in all my holy  $\dots$  sea 0.000000 They will not hurt nor destroy in all my holy  $\dots$  False
- 1 3T8DUCXYON6WD9X4RTLK8UN1U929TF bible that sends ambassadors by the sea, even in ves... sea 0.102941 that sends ambassadors by the sea, even in ves... False
- 2 317KR83SNADXAQ7HXK7S7305BYB9KD bible and they entered into the boat, and were going... sea 0.109375 and they entered into the boat, and were going... False
- 3 3BO3NEOQMOHK9ERYPNOGQIWCPC4IAQ bible Joseph laid up grain as the sand of the sea, v... sea 0.160714 Joseph laid up grain as the sand of the sea, v... False
- 4  $\,$  3Y3CZJSZ9KT0W7I0KE38WZHHKSW5RH  $\,$  bible  $\,$  There will be a highway for the remnant that i...  $\,$  1  $\,$

id corpus

sentence token complexity

37

sentence\_no\_contractions contraction\_expanded

- 0 31HLTCK4BLVQ5B01AUR91TX9V9IVGH bible The name of one son was Gershom, for Moses sai... foreign land 0.000000 The name of one son was Gershom, for Moses sai... False
- 1 389A2A304OIXVY7G5B71Q9M43LEOCL bible unleavened bread, unleavened cakes mixed with ... wheat flour 0.157895 unleavened bread, unleavened cakes mixed with ... False
- 2 31N9JPQXIPIRX2A3S9NOCCFXO6TNHR bible However the high places were not taken away; t... burnt incense 0.200000 However the high places were not taken away; t... False
- 3 3JVP4ZJHDPSO81TGXL3N1CKZGQYOIN bible and he burnt incense of sweet spices on it, as... burnt incense 0.250000 and he burnt incense of sweet spices on it, as... False
- 4 3JAOYN9IHL25ZQAUV5EJZ4GHOKL33L bible The same day the king made the middle of the c... bronze altar 0.214286 The same day the king made the middle of the c... False

id corpus

sentence token complexity

 ${\tt sentence\_no\_contractions} \quad {\tt contraction\_expanded}$ 

- O 3K8CQCU3KE19US5SN890DFPK3SANWR bible But he, beckoning to them with his hand to be ... hand 0.000000 But he, beckoning to them with his hand to be ... False
- 1 3Q2T3FD00N86LCI41NJYV3PN0BW3MV bible If I forget you, Jerusalem, let my right hand ... hand 0.197368 If I forget you, Jerusalem, let my right hand ... False
- 2 3ULIZOH1VA5C32JJMKOTQ8Z4GUS51B bible the ten sons of Haman the son of Hammedatha, t... hand 0.200000 the ten sons of Haman the son of Hammedatha, t... True
- 3 3BFFODJK8XCEIOT30ZLBPPSRMZQTSD bible Let your hand be lifted up above your adversar... hand 0.267857 Let your hand be lifted up above your adversar... False
- 4 3QREJ3J433XSBS8QMHAICCROBQ1LKR bible Abimelech chased him, and he fled before him, ... entrance 0.000000 Abimelech chased him, and he fled before him, ... False

id corpus

sentence token complexity

sentence\_no\_contractions contraction\_expanded

- O 3UXQ63NLAAMRIP4WG4XPD98AOYOBLX bible for he had an only daughter, about twelve year... only daughter 0.025000 for he had an only daughter, about twelve year... False
- 1 3FJ2RVH25Z62TA3R8E1077EBUYU92W bible All these were cities fortified with high wall... high walls 0.100000 All these were cities fortified with high wall... False
- 2 3YO4AH2FPDK1PZHZAT8WAEBL70EQOF bible In the morning, 'It will be foul weather today... weather today 0.125000 In the morning, 'It will be foul weather today... False
- 3 3X52SWXEOX5Q3081YIOMX4V84QTCWZ bible Her young children also were dashed in pieces ... young children 0.160714 Her young children also were dashed in

```
[32]: from nltk.tokenize import RegexpTokenizer
      import pandas as pd
      import numpy as np
      tokenizer = RegexpTokenizer(r'\w+') # same tokenizer as before
      def analyze_sentence_spans_by_corpus_and_quartile_no_contracts(dfs_dict):
          Analyze sentence spans (length metrics) grouped by corpus and complexity \Box
       \hookrightarrow quartile
          for multiple dataframes, but this time using the 'sentence no contractions'
          instead of the original 'sentence'.
          results = []
          for df_name, df in dfs_dict.items():
              print(f"Processing {df_name} on 'sentence_no_contractions'...")
              # Make a copy to avoid altering the original
              df = df.copy()
              # Calculate complexity quartiles (unchanged logic)
              q1 = df['complexity'].quantile(0.25)
              q2 = df['complexity'].quantile(0.50)
              q3 = df['complexity'].quantile(0.75)
              # Function to map complexity to quartiles
              def get_quartile(x):
                  if x <= q1:</pre>
                      return 'Q1'
                  elif x \ll q2:
                      return 'Q2'
                  elif x \ll q3:
                      return 'Q3'
                  else:
                      return 'Q4'
              # Create a new column storing quartile label
              df['quartile'] = df['complexity'].apply(get_quartile)
              # Define a local function to compute sentence metrics
```

```
def compute_span_metrics_no_contracts(sentence):
           if pd.isna(sentence):
               return pd.Series({'word_count': 0, 'char_count': 0, |

¬'avg_word_len': 0})
           words = tokenizer.tokenize(sentence)
           word count = len(words)
           char count = len(sentence)
           avg_word_len = np.mean([len(w) for w in words]) if word_count > 0_{\sqcup}
⊶else 0
          return pd.Series({
               'word_count': word_count,
               'char_count': char_count,
               'avg_word_len': avg_word_len
           })
       # Apply our function to 'sentence_no_contractions'
      span_metrics_nc = df['sentence_no_contractions'].
apply(compute_span_metrics_no_contracts)
       # Merge the results back into df
      df = pd.concat([df, span_metrics_nc], axis=1)
       # Group by corpus + quartile, same as before
      corpus col = 'corpus' # or subcorpus if you have that column
      for corpus_name, corpus_df in df.groupby(corpus_col):
           for quartile, quartile_df in corpus_df.groupby('quartile'):
               complexity_range = f"{quartile_df['complexity'].min():.
→3f}-{quartile_df['complexity'].max():.3f}"
               stats = {
                   'Dataframe': df_name,
                   'Corpus': corpus_name,
                   'Quartile': quartile,
                   'Complexity Range': complexity_range,
                   'Count': len(quartile df),
                   'Avg Words': quartile_df['word_count'].mean(),
                   'Median Words': quartile_df['word_count'].median(),
                   'Min Words': quartile_df['word_count'].min(),
                   'Max Words': quartile_df['word_count'].max(),
                   'Std Words': quartile_df['word_count'].std(),
                   'Avg Chars': quartile_df['char_count'].mean(),
                   'Avg Word Len': quartile_df['avg_word_len'].mean()
               }
               results.append(stats)
  results_df = pd.DataFrame(results)
  results_df = results_df.sort_values(['Dataframe', 'Corpus', 'Quartile'])
```

```
return results_df
# Example usage:
dfs = {
    'train_single_df': train_single_df,
    'train_multi_df': train_multi_df,
    'trial_val_single_df': trial_val_single_df,
    'trial_val_multi_df': trial_val_multi_df,
    'test_single_df': test_single_df,
    'test_multi_df': test_multi_df
}
span_analysis_nc =__
 →analyze_sentence_spans_by_corpus_and_quartile_no_contracts(dfs)
# Show results
pd.set option('display.max rows', None)
pd.set_option('display.max_columns', None)
pd.set option('display.width', 1000)
display(span_analysis_nc)
# Save results
results_path_nc = os.path.join(dir_results,_
 ⇔'sentence_span_analysis_no_contractions.csv')
span analysis nc.to csv(results path nc, index=False)
print(f"Analysis (NO CONTRACTIONS) saved to: {results path nc}")
```

```
Processing train_single_df on 'sentence_no_contractions'...

Processing train_multi_df on 'sentence_no_contractions'...

Processing trial_val_single_df on 'sentence_no_contractions'...

Processing trial_val_multi_df on 'sentence_no_contractions'...

Processing test_single_df on 'sentence_no_contractions'...

Processing test_multi_df on 'sentence_no_contractions'...
```

|                   | Da       | ataframe  | Corpus Qu | artile Comp | plexity Range | Count Avg Words | ш |
|-------------------|----------|-----------|-----------|-------------|---------------|-----------------|---|
| ⊶Medi             | an Words | Min Words | Max Word  | s Std Word  | s Avg Chars   | Avg Word Len    |   |
| 60                | test_n   | nulti_df  | bible     | Q1          | 0.025-0.317   | 26 23.076923    | ш |
| $\hookrightarrow$ | 22.0     | 4.0       | 48.0      | 11.831900   | 118.730769    | 4.131249        |   |
| 61                | test_n   | nulti_df  | bible     | Q2          | 0.325-0.417   | 11 20.545455    | ш |
| $\hookrightarrow$ | 17.0     | 7.0       | 47.0      | 12.917923   | 109.636364    | 4.213539        |   |
| 62                | test_n   | nulti_df  | bible     | Q3          | 0.432-0.528   | 18 21.055556    | ш |
| $\hookrightarrow$ | 21.5     | 4.0       | 43.0      | 10.843660   | 113.166667    | 4.498610        |   |
| 63                | test_n   | nulti_df  | bible     | Q4          | 0.542-0.694   | 11 22.363636    | ш |
| $\hookrightarrow$ | 20.0     | 7.0       | 51.0      | 11.935432   | 126.181818    | 4.605062        |   |
| 64                | test_n   | nulti_df  | biomed    | Q1          | 0.000-0.312   | 11 29.818182    | ш |
| $\hookrightarrow$ | 29.0     | 17.0      | 47.0      | 8.388304    | 195.727273    | 5.491145        |   |

| 65                 |                            |        |                 |                           | 11 27.090909              | Ш |
|--------------------|----------------------------|--------|-----------------|---------------------------|---------------------------|---|
|                    | 24.0 9.0                   |        |                 |                           |                           |   |
| 66                 | test_multi_df              |        | Q3              |                           | 10 26.900000              | Ц |
| ⇔                  | 26.5 10.0                  | 49.0   |                 |                           | 5.497409                  |   |
| 67                 | test_multi_df              |        | •               | 0.562-0.800               |                           | П |
| <b>⇔</b>           |                            |        |                 | 209.285714                |                           |   |
| 68                 | 24.5 7.0                   | _      |                 | 146.900000                | 10 24.700000<br>5.049688  | П |
| <sup>↔</sup><br>69 | test_multi_df              |        |                 |                           |                           |   |
| <i>⇔</i>           | 27.0 9.0                   | 73.0   |                 |                           |                           | Ш |
| 70                 | test_multi_df              |        |                 |                           | 18 32.944444              | Ш |
| <b>→</b>           |                            | -      |                 | 209.888889                |                           | П |
| 71                 |                            |        |                 |                           | 13 39.000000              | Ш |
| · <b>-</b>         |                            | 95.0   |                 |                           |                           |   |
| 48                 | test_single_df             |        |                 |                           | 79 22.822785              | Ш |
|                    | 22.0 7.0                   |        |                 |                           |                           |   |
| 49                 | test_single_df             |        |                 |                           | 68 24.176471              | ш |
| $\hookrightarrow$  |                            |        |                 | 126.088235                |                           |   |
| 50                 | test_single_df             | bible  | QЗ              | 0.278-0.353               | 67 22.388060              | Ш |
| $\hookrightarrow$  | 20.0 4.0                   | 63.0   | 11.306950       |                           |                           |   |
| 51                 | test_single_df             | bible  | Q4              | 0.359-0.732               | 69 20.579710              | Ш |
| $\hookrightarrow$  | 19.0 1.0                   | 55.0   | 11.264736       | 110.637681                | 4.341070                  |   |
| 52                 | test_single_df             | biomed | Q1              | 0.000-0.214               | 75 27.080000              | Ш |
| $\hookrightarrow$  | 25.0 10.0                  | 84.0   | 12.025603       | 172.986667                | 5.277318                  |   |
| 53                 | test_single_df             | biomed | Q2              | 0.217-0.275               | 58 30.275862              | ш |
| $\hookrightarrow$  | 26.0 10.0                  | 83.0   | 15.856587       | 198.293103                | 5.446788                  |   |
| 54                 | test_single_df             | biomed | QЗ              | 0.278-0.357               | 66 29.833333              | ш |
| $\hookrightarrow$  | 29.0 13.0                  | 85.0   | 11.754650       | 191.863636                | 5.334048                  |   |
| 55                 | test_single_df             | biomed |                 |                           | 90 31.144444              | Ш |
| $\hookrightarrow$  | 30.0 14.0                  |        |                 | 203.077778                |                           |   |
| 56                 |                            | -      |                 |                           | 83 25.337349              | Ш |
| $\hookrightarrow$  | 21.0 3.0                   |        |                 |                           |                           |   |
| 57                 | _                          | -      |                 |                           | 98 32.326531              | П |
| $\hookrightarrow$  | 30.0 1.0                   |        |                 | 195.653061                |                           |   |
| 58                 |                            | -      |                 |                           | 96 33.000000              | Ш |
| →                  | 30.0 3.0                   |        |                 |                           | 5.124551                  |   |
| 59                 | test_single_df             | _      |                 | 0.361-0.583               |                           | Ц |
| <b>⇔</b>           |                            |        |                 | 206.573529                |                           |   |
| 12                 |                            | bible  |                 | 0.028-0.300               |                           | П |
| <b>↔</b>           | 22.0 3.0                   |        |                 | 124.871166                | 4.237932                  |   |
| 13                 | train_multi_df<br>22.0 6.0 | 65.0   | Q2<br>11.738444 | 0.304-0.409               |                           | П |
| ↔<br>14            | train_multi_df             |        |                 | 129.659091<br>0.411-0.529 | 4.305703<br>131 23.778626 |   |
| 14                 |                            | 50.0   |                 | 127.564885                |                           | П |
| 15                 |                            |        |                 |                           | 79 25.481013              |   |
| 5                  | 24.0 3.0                   |        |                 | 139.405063                | 4.491816                  | Ш |
| 7                  | 21.0                       | 01.0   | 10.10000        | 100.40000                 | 1.101010                  |   |

| 16                |                          |          |           |             | 87 29.091954  | ш |
|-------------------|--------------------------|----------|-----------|-------------|---------------|---|
| $\hookrightarrow$ | 28.0 9.0                 | 77.0     | 11.882792 | 185.977011  |               |   |
| 17                | train_multi_df           |          |           |             | 74 30.756757  | ш |
| $\hookrightarrow$ | 28.0 11.0                | 85.0     | 13.511853 | 196.067568  | 5.367302      |   |
| 18                | ${\tt train\_multi\_df}$ | biomed   | Q3        |             |               | ш |
| $\hookrightarrow$ | 29.0 8.0                 | 61.0     | 10.912383 | 193.873874  | 5.430754      |   |
| 19                | train_multi_df           |          | Q4        | 0.531-0.975 | 242 29.607438 | ш |
| $\hookrightarrow$ | 28.0 10.0                | 75.0     | 12.029995 |             | 5.535387      |   |
| 20                | train_multi_df           | -        |           |             | 132 29.363636 | ш |
| $\hookrightarrow$ | 27.0 3.0                 | 101.0    |           | 176.583333  | 5.003685      |   |
| 21                | train_multi_df           | -        |           |             |               | ш |
| $\hookrightarrow$ | 28.0 3.0                 | 108.0    | 19.112977 | 195.198830  | 5.176456      |   |
| 22                | train_multi_df           | -        |           |             | 138 33.398551 | ш |
| $\hookrightarrow$ | 30.0 7.0                 |          |           |             | 5.286607      |   |
| 23                | train_multi_df           | _        |           |             | 57 34.596491  | ш |
| $\hookrightarrow$ | 31.0 6.0                 | 96.0     | 20.318763 | 218.350877  | 5.345891      |   |
| 0                 | train_single_df          | bible    | Q1        |             |               | ш |
| $\hookrightarrow$ | 22.0 4.0                 |          |           | 121.714693  | 4.135685      |   |
| 1                 | train_single_df          |          |           |             | 640 23.750000 | ш |
| $\hookrightarrow$ | 22.0 3.0                 | 60.0     | 11.579622 |             | 4.153925      |   |
| 2                 | train_single_df          | bible    | Q3        | 0.281-0.375 | 624 23.825321 | ш |
| $\hookrightarrow$ | 22.0 3.0                 |          | 11.963291 | 126.338141  | 4.213931      |   |
| 3                 | train_single_df          | bible    | Q4        | 0.380-0.861 |               | ш |
| $\hookrightarrow$ | 21.0 3.0                 | 69.0     | 12.460182 | 126.602627  | 4.298065      |   |
| 4                 | train_single_df          |          |           | 0.000-0.212 | 586 28.534130 | ш |
| $\hookrightarrow$ | 27.0 2.0                 | 85.0     | 12.115387 |             | 5.322266      |   |
| 5                 | train_single_df          | biomed   | Q2        | 0.212-0.279 |               | ш |
| $\hookrightarrow$ | 29.0 7.0                 |          |           | 193.921098  | 5.289166      |   |
| 6                 | train_single_df          | biomed   |           | 0.281-0.375 |               | ш |
| $\hookrightarrow$ | 28.0 4.0                 | 77.0     |           | 191.098634  | 5.329940      |   |
| 7                 | train_single_df          |          | Q4        |             |               | ш |
| $\hookrightarrow$ | 28.0 3.0                 | 85.0     | 12.249267 | 186.978610  | 5.299963      |   |
| 8                 | train_single_df          | -        |           | 0.025-0.212 | 641 26.761310 | ш |
| $\hookrightarrow$ | 24.0 2.0                 | 107.0    | 15.230853 | 159.190328  | 4.942926      |   |
| 9                 | train_single_df          | _        |           | 0.212-0.279 | 714 30.420168 | ш |
| $\hookrightarrow$ | 27.0 1.0                 | 129.0    | 18.383783 | 183.105042  | 4.995897      |   |
| 10                | train_single_df          | _        |           |             | 701 30.523538 | ш |
| $\hookrightarrow$ | 28.0 1.0                 | 122.0    | 18.163026 | 185.843081  | 5.114626      |   |
| 11                | train_single_df          | europarl | Q4        | 0.381-0.775 | 456 33.543860 | ш |
| $\hookrightarrow$ | 31.0 2.0                 | 235.0    | 21.708515 | 203.664474  | 5.054387      |   |
| 36                | trial_val_multi_df       | bible    | Q1        | 0.000-0.292 | 11 26.272727  | ш |
| $\hookrightarrow$ | 21.0 13.0                | 64.0     | 13.950562 | 141.363636  | 4.282457      |   |
| 37                | trial_val_multi_df       | bible    | Q2        | 0.333-0.400 | 7 20.571429   | ш |
| $\hookrightarrow$ | 23.0 5.0                 | 28.0     | 7.412987  | 110.857143  | 4.279406      |   |
| 38                | trial_val_multi_df       | bible    | QЗ        | 0.425-0.500 | 5 19.600000   | ш |
| $\hookrightarrow$ | 19.0 9.0                 | 32.0     | 8.905055  | 109.200000  | 4.431391      |   |

| 39                | trial_val_multi_df  | bible    | Q4        | 0.525-0.661 | 6 22.333333  | Ш |
|-------------------|---------------------|----------|-----------|-------------|--------------|---|
| $\hookrightarrow$ | 20.5 9.0            | 44.0     | 12.242004 |             | 4.178525     | _ |
| 40                | trial_val_multi_df  | biomed   | Q1        | 0.083-0.303 | 6 26.833333  | ш |
| $\hookrightarrow$ | 25.0 15.0           | 49.0     | 11.771434 | 159.166667  | 4.899969     |   |
| 41                | trial_val_multi_df  | biomed   | Q2        | 0.317-0.422 | 7 25.428571  | ш |
| $\hookrightarrow$ | 21.0 15.0           | 48.0     | 11.588171 | 156.000000  | 5.194383     |   |
| 42                | trial_val_multi_df  | biomed   | QЗ        | 0.438-0.513 | 6 37.833333  | ш |
| $\hookrightarrow$ | 39.5 26.0           | 44.0     | 6.675827  | 247.500000  | 5.438593     |   |
| 43                | trial_val_multi_df  | biomed   | Q4        | 0.537-0.825 | 14 30.642857 | ш |
| $\hookrightarrow$ | 29.5 17.0           | 43.0     | 9.849695  | 211.428571  | 5.730623     |   |
| 44                | trial_val_multi_df  | europarl | Q1        | 0.176-0.306 | 8 30.000000  | ш |
| $\hookrightarrow$ | 25.5 4.0            | 64.0     | 20.361027 | 186.750000  | 5.306837     |   |
| 45                | trial_val_multi_df  | europarl | Q2        | 0.312-0.412 | 11 47.909091 | ш |
| $\hookrightarrow$ | 46.0 24.0           | 78.0     | 18.651834 | 296.909091  | 5.058375     |   |
| 46                | trial_val_multi_df  | europarl | QЗ        | 0.432-0.500 | 13 26.307692 | ш |
| $\hookrightarrow$ | 26.0 5.0            | 66.0     | 18.167666 | 166.153846  | 5.263847     |   |
| 47                | trial_val_multi_df  | europarl | Q4        | 0.515-0.714 | 5 26.400000  | ш |
| $\hookrightarrow$ | 15.0 6.0            | 66.0     | 24.316661 | 164.600000  | 4.998182     |   |
| 24                | trial_val_single_df | bible    | Q1        | 0.000-0.214 | 52 26.769231 | ш |
| $\hookrightarrow$ | 26.0 5.0            | 74.0     | 15.589860 | 137.423077  | 4.074456     |   |
| 25                | trial_val_single_df | bible    | Q2        | 0.217-0.266 | 38 24.868421 | ш |
| $\hookrightarrow$ | 23.0 7.0            | 50.0     | 10.768249 | 131.342105  | 4.200230     |   |
| 26                | trial_val_single_df | bible    | QЗ        | 0.268-0.355 | 26 22.884615 | ш |
| $\hookrightarrow$ | 20.5 5.0            | 44.0     | 9.961233  | 121.423077  | 4.316593     |   |
| 27                | trial_val_single_df | bible    | Q4        | 0.361-0.633 | 27 25.666667 | ш |
| $\hookrightarrow$ | 23.0 6.0            | 49.0     | 12.554497 | 137.592593  | 4.213842     |   |
| 28                | trial_val_single_df | biomed   | Q1        | 0.028-0.214 | 21 25.571429 | ш |
| $\hookrightarrow$ | 21.0 13.0           | 65.0     | 11.543706 | 164.380952  | 5.317614     |   |
| 29                | trial_val_single_df | biomed   | Q2        | 0.217-0.267 | 28 30.571429 | ш |
| $\hookrightarrow$ | 27.5 11.0           | 57.0     | 12.099674 | 198.142857  | 5.315287     |   |
| 30                | trial_val_single_df | biomed   | QЗ        | 0.268-0.359 | 38 32.105263 | ш |
| $\hookrightarrow$ | 29.0 11.0           | 61.0     | 12.710476 | 206.947368  | 5.364934     |   |
| 31                | trial_val_single_df | biomed   | Q4        | 0.364-0.875 | 48 25.145833 | ш |
| $\hookrightarrow$ | 25.5 6.0            | 56.0     | 11.721937 | 164.020833  | 5.445661     |   |
| 32                | trial_val_single_df | europarl | Q1        | 0.050-0.214 | 33 31.969697 | ш |
| $\hookrightarrow$ | 28.0 5.0            | 81.0     | 20.356947 | 185.969697  | 4.799024     |   |
| 33                | trial_val_single_df | europarl | Q2        | 0.217-0.267 | 41 28.487805 | ш |
| $\hookrightarrow$ | 28.0 4.0            | 71.0     | 15.424205 | 172.902439  | 4.997384     |   |
| 34                | trial_val_single_df | europarl | QЗ        | 0.268-0.359 | 39 30.282051 | ш |
| $\hookrightarrow$ | 28.0 3.0            | 99.0     | 20.040681 | 184.358974  | 5.086945     |   |
| 35                | trial_val_single_df | _        |           | 0.367-0.605 | 30 35.700000 | ш |
| $\hookrightarrow$ | 30.5 5.0            | 77.0     | 20.142852 | 215.400000  | 4.910759     |   |

Analysis (NO CONTRACTIONS) saved to: /content/drive/MyDrive/266-final/results/sentence\_span\_analysis\_no\_contractions.csv

• contraction processing successfuly, confirmed with Avg Word deltas between 'sentence' and

## 0.4 Enrich Datset with PoS Tags, Dependency Parsing, and Morphological Complexity

```
[33]: # !pip install -q spacy
      # !python -m spacy download en_core_web_trf
      !python -m spacy download en_core_web_lg
     Collecting en-core-web-lg==3.8.0
       Downloading https://github.com/explosion/spacy-
     models/releases/download/en_core_web_lg-3.8.0/en_core_web_lg-3.8.0-py3-none-
     any.whl (400.7 MB)
                                 400.7/400.7
     MB 2.6 MB/s eta 0:00:00
     Installing collected packages: en-core-web-lg
     Successfully installed en-core-web-lg-3.8.0
      Download and installation successful
     You can now load the package via spacy.load('en_core_web_lg')
      Restart to reload dependencies
     If you are in a Jupyter or Colab notebook, you may need to restart Python in
     order to load all the package's dependencies. You can do this by selecting the
     'Restart kernel' or 'Restart runtime' option.
[34]: nlp = spacy.load("en_core_web_lg")
[35]: # prompt: run a test with nlp, my spacy instance
      # Sample text for testing
      text = "This is a sample sentence for testing spaCy."
      # Process the text with spaCy
      doc = nlp(text)
      # Print the tokens and their part-of-speech tags
      for token in doc:
          print(f"Token: {token.text}, POS: {token.pos_}, Dependency: {token.dep_}")
     Token: This, POS: PRON, Dependency: nsubj
     Token: is, POS: AUX, Dependency: ROOT
     Token: a, POS: DET, Dependency: det
     Token: sample, POS: NOUN, Dependency: compound
     Token: sentence, POS: NOUN, Dependency: attr
     Token: for, POS: ADP, Dependency: prep
     Token: testing, POS: VERB, Dependency: pcomp
     Token: spaCy, POS: PROPN, Dependency: dobj
     Token: ., POS: PUNCT, Dependency: punct
```

```
[36]: def enrich_with_spacy(df, text_col='sentence_no_contractions'):
         Processes the 'text_col' with spaCy and appends:
           pos_sequence, dep_sequence, morph_sequence,
           and morph_complexity (float) per row.
         df = df.copy()
         pos_tags = []
         dep_tags = []
         morph tags = []
         morph_complexities = []
         for text in df[text_col]:
              if pd.isna(text) or not text.strip():
                  # If text is NaN or empty, store defaults
                  pos_tags.append([])
                 dep_tags.append([])
                 morph_tags.append([])
                 morph_complexities.append(0.0)
                  continue
             doc = nlp(text)
              # Build sequences
                                                       # e.g. ["DET",⊔
             pos_seq = [token.pos_ for token in doc]
       →"NOUN", "VERB"]
              dep_seq = [token.dep_ for token in doc]
                                                               # e.q. ["det",
       → "nsubj", "ROOT"]
             morph_seq = [token.morph for token in doc]
                                                                # e.g.,
       → [Number=Sing|Person=3, ...]
              # Count morphological features
              total_features = 0
              for token in doc:
                  # token.morph.to_dict() => a dict of morphological attributes
                  features_dict = token.morph.to_dict()
                  total_features += len(features_dict)
              # Average morphological features across tokens
              avg_morph = total_features / len(doc)
             pos_tags.append(pos_seq)
             dep_tags.append(dep_seq)
             morph_tags.append(morph_seq)
             morph_complexities.append(avg_morph)
```

```
# Add these lists as new columns
          df['pos_sequence'] = pos_tags
          df['dep_sequence'] = dep_tags
          df['morph_sequence'] = morph_tags
          df['morph_complexity'] = morph_complexities
          return df
[37]: dataframes info = [
          ("train_single_df", train_single_df),
          ("train multi df", train multi df),
          ("trial_val_single_df", trial_val_single_df),
          ("trial_val_multi_df", trial_val_multi_df),
          ("test_single_df", test_single_df),
          ("test_multi_df", test_multi_df),
      ]
      for df_name, df in dataframes_info:
          print(f"Enriching {df_name} with spaCy features...")
          enriched_df = enrich_with_spacy(df, text_col='sentence_no_contractions')
          # Overwrite the old DataFrame variable (or store it somewhere new)
          globals()[df_name] = enriched_df
          print(f"Done! Now '{df_name}' has columns: pos_sequence, dep_sequence,
       →morph_sequence, morph_complexity.\n")
     Enriching train_single_df with spaCy features...
     Done! Now 'train_single_df' has columns: pos_sequence, dep_sequence,
     morph_sequence, morph_complexity.
     Enriching train_multi_df with spaCy features...
     Done! Now 'train_multi_df' has columns: pos_sequence, dep_sequence,
     morph_sequence, morph_complexity.
     Enriching trial val single df with spaCy features...
     Done! Now 'trial_val_single_df' has columns: pos_sequence, dep_sequence,
     morph sequence, morph complexity.
     Enriching trial_val_multi_df with spaCy features...
     Done! Now 'trial_val_multi_df' has columns: pos_sequence, dep_sequence,
     morph_sequence, morph_complexity.
     Enriching test_single_df with spaCy features...
     Done! Now 'test_single_df' has columns: pos_sequence, dep_sequence,
     morph_sequence, morph_complexity.
     Enriching test_multi_df with spaCy features...
```

Done! Now 'test multi df' has columns: pos sequence, dep sequence,

morph\_sequence, morph\_complexity.

```
[38]: for df_name, df in dataframes_info:
          print(f"\n{'='*50}")
          print(f"DataFrame: {df name}")
          print(f"{'='*50}\n")
          sample_df = globals()[df_name].sample(3, random_state=42)
          display(sample_df[['sentence_no_contractions', 'pos_sequence',_

¬'dep_sequence', 'morph_sequence', 'morph_complexity']])

     DataFrame: train_single_df
     _____
                                    sentence_no_contractions
                    pos_sequence
                                                                        dep_sequence _
                                         morph_sequence morph_complexity
     5061 The transgenic approach that was used to creat... [DET, ADJ, NOUN, PRON,
      AUX, VERB, PART, VERB, ... [det, amod, nsubjpass, nsubjpass, auxpass, rel... _
      →[(Definite=Def, PronType=Art), (Degree=Pos), (...
     2471 When the report comes to Egypt, they will be i... [SCONJ, DET, NOUN, VERB,
      ADP, PROPN, PUNCT, PR... [advmod, det, nsubj, advcl, prep, pobj, punct,... [(),
      → (Definite=Def, PronType=Art), (Number=Sin...
                                                            1.166667
           Saul asked counsel of God, "Shall I go down af ... [PROPN, VERB, NOUN, ADP, __
      →PROPN, PUNCT, PUNCT, ... [nsubj, ROOT, dobj, prep, pobj, punct, punct, ... u
      →[(Number=Sing), (Tense=Past, VerbForm=Fin), (N...
                                                                 1.200000
     DataFrame: train_multi_df
                                    sentence no contractions
                    pos_sequence
                                                                        dep_sequence _
                                         morph_sequence morph_complexity
           BRCA2 may thus promote RAD51 assembly into rec... [PROPN, AUX, ADV, VERB,
     724
      PROPN, NOUN, ADP, ADJ,... [nsubj, aux, advmod, ROOT, compound, dobj, pre... []
      →[(Number=Sing), (VerbForm=Fin), (), (VerbForm=...
                                                                 1.222222
           Therefore, BMPR1A appears to maintain articula... [ADV, PUNCT, PROPN, VERB, ]
      →PART, VERB, ADJ, NOU... [advmod, punct, nsubj, ROOT, aux, xcomp, amod,... [(), __
      → (PunctType=Comm), (Number=Sing), (Number=...
                                                            1.000000
     1466 Continued support for the renewal and modernis... [VERB, NOUN, ADP, DET, __
      →NOUN, CCONJ, NOUN, ADP,... [amod, nsubj, prep, det, pobj, cc, conj, prep,... 🛭
      →[(Aspect=Perf, Tense=Past, VerbForm=Part), (Nu...
                                                                 1.205882
```

```
DataFrame: trial_val_single_df
_____
                            sentence_no_contractions
                                                              dep_sequence
             pos_sequence
                                 morph_sequence morph_complexity
145 However, this reduction in bone resorption occ... [ADV, PUNCT, DET, NOUN,
 ADP, NOUN, NOUN, VERB,... [advmod, punct, det, nsubj, prep, compound, po... [(),
 → (PunctType=Comm), (Number=Sing, PronType=...
                                                     1.0000
335 A word of thanks is also due to many non-gover... [DET, NOUN, ADP, NOUN,
 AUX, ADV, ADJ, ADP, ADJ... [det, nsubj, prep, pobj, ROOT, advmod, prep, p...
 →[(Definite=Ind, PronType=Art), (Number=Sing), ...
                                                          1.0625
175 To test the hypothesis that a temporal delay i... [PART, VERB, DET, NOUN,
 SCONJ, DET, ADJ, NOUN, ... [aux, advcl, det, dobj, mark, det, amod, nsubj... [(),
 → (VerbForm=Inf), (Definite=Def, PronType=A...
                                                     1.2000
DataFrame: trial_val_multi_df
                           sentence_no_contractions
                                                             dep_sequence
            pos_sequence
                                morph_sequence morph_complexity
62 by Mr Virrankoski, on behalf of the Committee ... [ADP, PROPN, PROPN, PUNCT,
 →ADP, NOUN, ADP, DET… [prep, compound, pobj, punct, prep, pobj, prep… [(), □
 →(Number=Sing), (Number=Sing), (PunctType=...
                                                   0.892857
40 Indeed, we recently showed that neural crest c... [ADV, PUNCT, PRON, ADV,
 →VERB, SCONJ, ADJ, PROP... [advmod, punct, nsubj, advmod, ROOT, mark, com... [(),
 → (PunctType=Comm), (Case=Nom, Number=Plur,...
95 It is not an easy task, particularly for the c... [PRON, AUX, PART, DET, ADJ,
 NOUN, PUNCT, ADV, ... [nsubj, ROOT, neg, det, amod, attr, punct, adv... u
 →[(Case=Nom, Gender=Neut, Number=Sing, Person=3...
                                                        1.180328
  ______
DataFrame: test_single_df
_____
                            sentence_no_contractions
                                                              dep sequence
             pos_sequence
                                 morph_sequence morph_complexity
668 It is therefore not a matter of indifference h... [PRON, AUX, ADV, PART,
 →DET, NOUN, ADP, NOUN, S... [nsubj, ROOT, advmod, neg, det, attr, prep, po... ⊔
 →[(Case=Nom, Gender=Neut, Number=Sing, Person=3...
                                                        1.200000
```

then shall he offer with the bull a meal offer... [ADV, AUX, PRON, VERB, \_\_ ADP, DET, NOUN, DET, NO... [advmod, aux, nsubj, ROOT, prep, det, pobj, de... \_ →[(PronType=Dem), (VerbType=Mod), (Case=Nom, Ge... 1.071429 377 While they do have their limitations (e.g. dev... [SCONJ, PRON, AUX, VERB, PRON, NOUN, PUNCT, AD... [mark, nsubj, aux, advcl, poss, dobj, punct, a... [(), →(Case=Nom, Number=Plur, Person=3, PronTyp... 1.157895 DataFrame: test\_multi\_df sentence\_no\_contractions pos\_sequence dep\_sequence morph\_sequence morph\_complexity God said, "Let the earth yield grass, herbs yi... [PROPN, VERB, PUNCT, \_ 19 →PUNCT, VERB, DET, NOUN, V... [nsubj, ROOT, punct, punct, xcomp, det, nsubj,... u →[(Number=Sing), (Tense=Past, VerbForm=Fin), (P... 1.564103 Moreover I will make a covenant of peace with ... [ADV, PRON, AUX, VERB, \_\_ →DET, NOUN, ADP, NOUN, A... [advmod, nsubj, aux, ccomp, det, dobj, prep, p... ⊔ →[(), (Case=Nom, Number=Sing, Person=1, PronTyp... 1.550000 156 Developing innovation policy is crucial to EU ... [VERB, NOUN, NOUN, AUX, ADJ, ADP, PROPN, NOUN,... [csubj, compound, dobj, ROOT, acomp, prep, com... \_ →[(Aspect=Prog, Tense=Pres, VerbForm=Part), (Nu... 1.333333 [39]: # verify column headers dataframes = [train\_single\_df, train\_multi\_df, trial\_val\_single\_df,\_ →trial\_val\_multi\_df, test\_single\_df, test\_multi\_df] for df in dataframes: print(df.info()) <class 'pandas.core.frame.DataFrame'> RangeIndex: 7662 entries, 0 to 7661 Data columns (total 11 columns): Column Non-Null Count Dtype 0 7662 non-null id object 1 corpus 7662 non-null object 2 sentence 7662 non-null object 3 token 7655 non-null object 4 complexity 7662 non-null float64 5 sentence\_no\_contractions 7662 non-null object

bool

object

object

object

float64

7662 non-null

7662 non-null

7662 non-null

7662 non-null

7662 non-null

6

7

8

contraction\_expanded

pos\_sequence

dep\_sequence

morph\_sequence

morph\_complexity

dtypes: bool(1), float64(2), object(8)

memory usage: 606.2+ KB

None

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1517 entries, 0 to 1516
Data columns (total 11 columns):

| #  | Column                   | Non-Null Count | Dtype   |
|----|--------------------------|----------------|---------|
|    |                          |                |         |
| 0  | id                       | 1517 non-null  | object  |
| 1  | corpus                   | 1517 non-null  | object  |
| 2  | sentence                 | 1517 non-null  | object  |
| 3  | token                    | 1517 non-null  | object  |
| 4  | complexity               | 1517 non-null  | float64 |
| 5  | sentence_no_contractions | 1517 non-null  | object  |
| 6  | contraction_expanded     | 1517 non-null  | bool    |
| 7  | pos_sequence             | 1517 non-null  | object  |
| 8  | dep_sequence             | 1517 non-null  | object  |
| 9  | morph_sequence           | 1517 non-null  | object  |
| 10 | morph_complexity         | 1517 non-null  | float64 |

dtypes: bool(1), float64(2), object(8)

memory usage: 120.1+ KB

None

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 421 entries, 0 to 420
Data columns (total 11 columns):

| #  | Column                        | Non-Null Count | Dtype   |
|----|-------------------------------|----------------|---------|
|    |                               |                |         |
| 0  | id                            | 421 non-null   | object  |
| 1  | corpus                        | 421 non-null   | object  |
| 2  | sentence                      | 421 non-null   | object  |
| 3  | token                         | 421 non-null   | object  |
| 4  | complexity                    | 421 non-null   | float64 |
| 5  | sentence_no_contractions      | 421 non-null   | object  |
| 6  | ${\tt contraction\_expanded}$ | 421 non-null   | bool    |
| 7  | pos_sequence                  | 421 non-null   | object  |
| 8  | dep_sequence                  | 421 non-null   | object  |
| 9  | morph_sequence                | 421 non-null   | object  |
| 10 | morph_complexity              | 421 non-null   | float64 |
|    |                               |                |         |

dtypes: bool(1), float64(2), object(8)

memory usage: 33.4+ KB

None

<class 'pandas.core.frame.DataFrame'>

RangeIndex: 99 entries, 0 to 98
Data columns (total 11 columns):

| # | Column | Non-Null Count | Dtype  |
|---|--------|----------------|--------|
|   |        |                |        |
| 0 | id     | 99 non-null    | object |
| 1 | corpus | 99 non-null    | object |

```
99 non-null
                                             object
 2
    sentence
 3
    token
                              99 non-null
                                             object
 4
                              99 non-null
                                             float64
    complexity
 5
    sentence_no_contractions 99 non-null
                                             object
 6
    contraction expanded
                              99 non-null
                                             bool
 7
    pos_sequence
                              99 non-null
                                             object
 8
    dep_sequence
                              99 non-null
                                             object
    morph_sequence
                              99 non-null
                                             object
 10 morph_complexity
                                             float64
                              99 non-null
dtypes: bool(1), float64(2), object(8)
memory usage: 8.0+ KB
None
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 917 entries, 0 to 916
Data columns (total 11 columns):
                              Non-Null Count Dtype
    Column
 #
    _____
                              _____
 0
    id
                              917 non-null
                                             object
 1
    corpus
                              917 non-null
                                             object
 2
    sentence
                              917 non-null
                                             object
 3
    token
                              917 non-null
                                             object
 4
    complexity
                              917 non-null
                                             float64
 5
    sentence_no_contractions 917 non-null
                                             object
 6
    contraction_expanded
                              917 non-null
                                             bool
 7
    pos_sequence
                              917 non-null
                                             object
 8
    dep_sequence
                              917 non-null
                                             object
    morph_sequence
                              917 non-null
                                             object
 10 morph_complexity
                              917 non-null
                                             float64
dtypes: bool(1), float64(2), object(8)
memory usage: 72.7+ KB
None
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 184 entries, 0 to 183
Data columns (total 11 columns):
    Column
                              Non-Null Count Dtype
                              _____
--- ----
 0
    id
                              184 non-null
                                             object
 1
    corpus
                              184 non-null
                                             object
 2
    sentence
                              184 non-null
                                             object
                              184 non-null
 3
    token
                                             object
 4
    complexity
                              184 non-null
                                             float64
 5
    sentence_no_contractions 184 non-null
                                              object
    contraction_expanded
                              184 non-null
                                             bool
 7
    pos_sequence
                              184 non-null
                                             object
    dep_sequence
                              184 non-null
                                             object
    morph_sequence
                              184 non-null
                                             object
```

dtypes: bool(1), float64(2), object(8)

10 morph\_complexity

float64

184 non-null

```
memory usage: 14.7+ KB None
```

0.5 Create Binarized Outcome Variable, based on train\_single\_df median and train\_multi\_df median, applied to trial-val and test

```
[40]: # Step 1: Compute the median for single-token train set
      train_single_median = train_single_df['complexity'].median()
      # Step 2: Define a function that applies the threshold
      def binarize_complexity(value, threshold):
          If value <= threshold, return 0, else return 1.
          if value <= threshold:</pre>
              return 0
          else:
              return 1
      # Step 3: Create 'binary_complexity' in the three single-token DataFrames
      train_single_df['binary_complexity'] = train_single_df['complexity'].apply(
          lambda x: binarize_complexity(x, train_single_median)
      trial_val_single_df['binary_complexity'] = trial_val_single_df['complexity'].
       →apply(
          lambda x: binarize_complexity(x, train_single_median)
      test_single_df['binary_complexity'] = test_single_df['complexity'].apply(
          lambda x: binarize_complexity(x, train_single_median)
      )
      # Step 4: Compute the median for multi-token train set
      train_multi_median = train_multi_df['complexity'].median()
      # Step 5: Create 'binary_complexity' in the three multi-token DataFrames
      train_multi_df['binary_complexity'] = train_multi_df['complexity'].apply(
          lambda x: binarize_complexity(x, train_multi_median)
      trial_val_multi_df['binary_complexity'] = trial_val_multi_df['complexity'].
       →apply(
          lambda x: binarize_complexity(x, train_multi_median)
      test_multi_df['binary_complexity'] = test_multi_df['complexity'].apply(
          lambda x: binarize_complexity(x, train_multi_median)
      )
      # Step 6: (Optional) Print out the medians and some basic info
```

```
print(f"Median complexity (single): {train_single_median}")
      print(f"Median complexity (multi): {train_multi_median}")
      print("\nSample rows from train_single_df:")
      print(train_single_df[['id', 'complexity', 'binary_complexity']].head())
      print("\nSample rows from train multi df:")
      print(train_multi_df[['id', 'complexity', 'binary_complexity']].head())
     Median complexity (single): 0.2794117647058823
     Median complexity (multi): 0.409090909090909
     Sample rows from train_single_df:
                                    id complexity
                                                    binary_complexity
     O 3ZLW647WALVGE8EBR50EGUBPU4P32A
                                          0.000000
     1 34ROBODSP1ZBN3DVY8J8XSIY551E5C
                                          0.000000
                                                                    0
                                                                    0
     2 3S1WOPCJFGTJU2SGNAN2Y213N6WJE3
                                          0.050000
     3 3BFNCI9LYKQN09BHXHH9CLSX5KP738
                                                                    0
                                          0.150000
     4 3G5RUKN2EC3YIWSKUXZ8ZVH95R49N2
                                          0.263889
                                                                    0
     Sample rows from train_multi_df:
                                    id complexity binary_complexity
     O 3S37Y8CWI8ON8KVM53U4E6JKCDC4WE
                                          0.027778
     1 3WGCNLZJKF877FYC1Q6COKNWTDWD11
                                          0.050000
                                                                    0
     2 3UOMW19E6D6WQ5TH2HDD74IVKTP5CB
                                          0.050000
                                                                    0
     3 36JW4WBR06KF9AXMUL4N4760MF8FHD
                                          0.050000
                                                                    0
     4 3HRWUH63QU2FH9Q8R7MRNFC7JX2N5A
                                          0.075000
                                                                    0
[44]: # verify column headers
      dataframes = [train_single_df, train_multi_df, trial_val_single_df,_

¬trial_val_multi_df, test_single_df, test_multi_df]
      for df in dataframes:
        print(df.info())
     <class 'pandas.core.frame.DataFrame'>
     RangeIndex: 7662 entries, 0 to 7661
     Data columns (total 12 columns):
          Column
                                    Non-Null Count Dtype
         _____
                                    _____
      0
                                    7662 non-null
                                                    object
          id
      1
          corpus
                                    7662 non-null
                                                    object
      2
          sentence
                                    7662 non-null
                                                    object
      3
          token
                                    7655 non-null
                                                    object
      4
          complexity
                                    7662 non-null
                                                    float64
          sentence_no_contractions 7662 non-null
                                                    object
      6
          contraction_expanded
                                    7662 non-null
                                                    bool
          pos_sequence
                                    7662 non-null
                                                    object
```

```
object
 8
    dep_sequence
                              7662 non-null
    morph_sequence
                              7662 non-null
                                              object
 10 morph_complexity
                              7662 non-null
                                              float64
 11 binary_complexity
                              7662 non-null
                                              int64
dtypes: bool(1), float64(2), int64(1), object(8)
memory usage: 666.1+ KB
None
```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1517 entries, 0 to 1516
Data columns (total 12 columns):

| #  | Column                   | Non-Null Count | Dtype      |
|----|--------------------------|----------------|------------|
|    |                          |                |            |
| 0  | id                       | 1517 non-null  | object     |
| 1  | corpus                   | 1517 non-null  | object     |
| 2  | sentence                 | 1517 non-null  | object     |
| 3  | token                    | 1517 non-null  | object     |
| 4  | complexity               | 1517 non-null  | float64    |
| 5  | sentence_no_contractions | 1517 non-null  | object     |
| 6  | contraction_expanded     | 1517 non-null  | bool       |
| 7  | pos_sequence             | 1517 non-null  | object     |
| 8  | dep_sequence             | 1517 non-null  | object     |
| 9  | morph_sequence           | 1517 non-null  | object     |
| 10 | morph_complexity         | 1517 non-null  | float64    |
| 11 | binary_complexity        | 1517 non-null  | int64      |
|    | 1 7 (4) 47 . 44 (6)      |                | <b>~</b> ` |

dtypes: bool(1), float64(2), int64(1), object(8)

memory usage: 132.0+ KB

None

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 421 entries, 0 to 420
Data columns (total 12 columns):

|    | 00_4444   | , ·             |         |
|----|---|-----------------|---------|
| #  | Column  | Non-Null Count  | Dtype   |
|    |   |                 |         |
| 0  | id  | 421 non-null    | object  |
| 1  | corpus  | 421 non-null    | object  |
| 2  | sentence  | 421 non-null    | object  |
| 3  | token   | 421 non-null    | object  |
| 4  | complexity  | 421 non-null    | float64 |
| 5  | sentence_no_contractions  | 421 non-null    | object  |
| 6  | contraction_expanded  | 421 non-null    | bool    |
| 7  | pos_sequence  | 421 non-null    | object  |
| 8  | dep_sequence  | 421 non-null    | object  |
| 9  | morph_sequence  | 421 non-null    | object  |
| 10 | morph_complexity  | 421 non-null    | float64 |
| 11 | binary_complexity   | 421 non-null    | int64   |
| 4+ | $a_{0}$ , $b_{0}$ $a_{0}$ $a_{0}$ $a_{0}$ $a_{0}$ $a_{0}$ $a_{0}$ $a_{0}$ | n+61(1) abias+( | 0)      |

dtypes: bool(1), float64(2), int64(1), object(8)

memory usage: 36.7+ KB

None

<class 'pandas.core.frame.DataFrame'>

RangeIndex: 99 entries, 0 to 98
Data columns (total 12 columns):

| #  | Column                   | Non-Null Count | Dtype   |
|----|--------------------------|----------------|---------|
|    |                          |                |         |
| 0  | id                       | 99 non-null    | object  |
| 1  | corpus                   | 99 non-null    | object  |
| 2  | sentence                 | 99 non-null    | object  |
| 3  | token                    | 99 non-null    | object  |
| 4  | complexity               | 99 non-null    | float64 |
| 5  | sentence_no_contractions | 99 non-null    | object  |
| 6  | contraction_expanded     | 99 non-null    | bool    |
| 7  | pos_sequence             | 99 non-null    | object  |
| 8  | dep_sequence             | 99 non-null    | object  |
| 9  | morph_sequence           | 99 non-null    | object  |
| 10 | morph_complexity         | 99 non-null    | float64 |
| 11 | binary_complexity        | 99 non-null    | int64   |
|    | 1 7 (4) 07 .04 (0) 4     |                | ~ `     |

dtypes: bool(1), float64(2), int64(1), object(8)

memory usage: 8.7+ KB

None

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 917 entries, 0 to 916
Data columns (total 12 columns):

| #  | Column                   | Non-Null Count | Dtype   |
|----|--------------------------|----------------|---------|
|    |                          |                |         |
| 0  | id                       | 917 non-null   | object  |
| 1  | corpus                   | 917 non-null   | object  |
| 2  | sentence                 | 917 non-null   | object  |
| 3  | token                    | 917 non-null   | object  |
| 4  | complexity               | 917 non-null   | float64 |
| 5  | sentence_no_contractions | 917 non-null   | object  |
| 6  | contraction_expanded     | 917 non-null   | bool    |
| 7  | pos_sequence             | 917 non-null   | object  |
| 8  | dep_sequence             | 917 non-null   | object  |
| 9  | morph_sequence           | 917 non-null   | object  |
| 10 | morph_complexity         | 917 non-null   | float64 |
| 11 | binary_complexity        | 917 non-null   | int64   |

dtypes: bool(1), float64(2), int64(1), object(8)

memory usage: 79.8+ KB

None

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 184 entries, 0 to 183
Data columns (total 12 columns):

| # | Column   | Non-Null Count | Dtype  |
|---|----------|----------------|--------|
|   |          |                |        |
| 0 | id       | 184 non-null   | object |
| 1 | corpus   | 184 non-null   | object |
| 2 | sentence | 184 non-null   | object |
| 3 | token    | 184 non-null   | object |

```
complexity
                               184 non-null
                                               float64
 4
 5
    sentence_no_contractions 184 non-null
                                               object
 6
    contraction_expanded
                               184 non-null
                                               bool
 7
    pos_sequence
                               184 non-null
                                               object
    dep sequence
 8
                               184 non-null
                                               object
    morph sequence
                               184 non-null
                                               object
 10 morph complexity
                               184 non-null
                                               float64
 11 binary_complexity
                               184 non-null
                                               int64
dtypes: bool(1), float64(2), int64(1), object(8)
memory usage: 16.1+ KB
None
```

## [45]: # inspect each df

dataframes = [train single\_df, train multi\_df, trial\_val\_single\_df,\_ ¬trial\_val\_multi\_df, test\_single\_df, test\_multi\_df] for df in dataframes: print(df.head())

id corpus

```
sentence
             token complexity
sentence_no_contractions contraction_expanded
pos_sequence
                                                   dep_sequence
morph_sequence morph_complexity binary_complexity
O 3ZLW647WALVGE8EBR50EGUBPU4P32A bible Behold, there came up out of the river
                      0.000000 Behold, there came up out of the river seven
                   False [ADV, PUNCT, PRON, VERB, ADP, ADP, ADP, DET, N...
C...
[advmod, punct, expl, ROOT, prt, prep, prep, d... [(), (PunctType=Comm), (),
(Tense=Past, VerbFo...
                              1.041667
1 34ROBODSP1ZBN3DVY8J8XSIY551E5C bible I am a fellow bondservant with you and
with yo... brothers
                      0.000000 I am a fellow bondservant with you and with
                    False [PRON, AUX, DET, ADJ, NOUN, ADP, PRON, CCONJ, ...
[nsubj, ROOT, det, amod, attr, prep, pobj, cc,... [(Case=Nom, Number=Sing,
                                1.461538
Person=1, PronType=Pr...
2 3S1WOPCJFGTJU2SGNAN2Y213N6WJE3 bible The man, the lord of the land, said to
us, 'By... brothers
                      0.050000 The man, the lord of the land, said to us,
                     False [DET, NOUN, PUNCT, DET, PROPN, ADP, DET, NOUN, ...
[det, nsubj, punct, det, appos, prep, det, pob... [(Definite=Def,
PronType=Art), (Number=Sing), ...
                                         1.354167
3 3BFNCI9LYKQN09BHXHH9CLSX5KP738 bible Shimei had sixteen sons and six
daughters; but... brothers
                             0.150000 Shimei had sixteen sons and six
                                 True [PROPN, VERB, NUM, NOUN, CCONJ, NUM,
daughters; but...
NOUN, PUN... [nsubj, ROOT, nummod, dobj, cc, nummod, conj, ... [(Number=Sing),
(Tense=Past, VerbForm=Fin), (N...
                                         1.275862
                                                                   0
4 3G5RUKN2EC3YIWSKUXZ8ZVH95R49N2 bible
                                                       "He has put my brothers
far from me. brothers
                                                 "He has put my brothers far
                          0.263889
                         False [PUNCT, PRON, AUX, VERB, PRON, NOUN, ADV,
from me.
ADP,... [punct, nsubj, aux, ROOT, poss, dobj, advmod, ... [(PunctSide=Ini,
```

```
PunctType=Quot), (Case=Nom, G...
                                        2.500000
                                                                   0
                               id corpus
                    token complexity
sentence
sentence_no_contractions contraction_expanded
pos sequence
                                                   dep sequence
morph_sequence morph_complexity binary_complexity
O 3S37Y8CWI8ON8KVM53U4E6JKCDC4WE bible but the seventh day is a Sabbath to
Yahweh you...
                 seventh day
                                0.027778 but the seventh day is a Sabbath to
                            False [CCONJ, DET, ADJ, NOUN, AUX, DET, PROPN,
Yahweh you...
ADP, ... [cc, det, amod, nsubj, ccomp, det, attr, prep,... [(ConjType=Cmp),
(Definite=Def, PronType=Art),...
                                        1.341772
1 3WGCNLZJKF877FYC1Q6COKNWTDWD11 bible But let each man test his own work,
                                0.050000 But let each man test his own work,
and then h...
                    own work
                            False [CCONJ, VERB, DET, NOUN, VERB, PRON, ADJ,
and then h...
NOUN... [cc, ROOT, det, nsubj, ccomp, poss, amod, dobj... [(ConjType=Cmp),
(VerbForm=Inf), (), (Number=S...
                                        1.608696
2 3UOMW19E6D6WQ5TH2HDD74IVKTP5CB bible To him who by understanding made the
                               0.050000 To him who by understanding made the
heavens; ... loving kindness
                           False [ADP, PRON, PRON, ADP, VERB, VERB, DET,
heavens; ...
NOUN, ... [prep, pobj, nsubj, prep, pcomp, advcl, det, d... [(), (Case=Acc,
Gender=Masc, Number=Sing, Pers...
                                         1.562500
3 36JW4WBR06KF9AXMUL4N4760MF8FHD bible Remember to me, my God, this also, and
spare m... loving kindness
                             0.050000 Remember to me, my God, this also, and
                         False [VERB, ADP, PRON, PUNCT, PRON, PROPN, PUNCT,
spare m...
P... [ROOT, prep, pobj, punct, poss, npadvmod, punc... [(VerbForm=Inf), (),
                                    1.590909
(Case=Acc, Number=Sing, P...
4 3HRWUH63QU2FH9Q8R7MRNFC7JX2N5A bible Because your loving kindness is better
than li... loving kindness
                             0.075000 Because your loving kindness is better
                         False [SCONJ, PRON, ADJ, NOUN, AUX, ADJ, ADP, NOUN,
... [mark, poss, amod, nsubj, advcl, acomp, prep, ... [(), (Person=2,
Poss=Yes, PronType=Prs), (Degr...
                                         1.600000
                               id corpus
sentence token complexity
                                                     sentence_no_contractions
contraction_expanded
                                                           pos_sequence
dep sequence
                                                 morph sequence
morph complexity binary complexity
O 3QI9WAYOGQB8GQIR4MDIEFOD2RLS67 bible They will not hurt nor destroy in all
                   0.000000 They will not hurt nor destroy in all my holy ...
my holy ...
False [PRON, AUX, PART, VERB, CCONJ, VERB, ADP, PRON... [nsubj, aux, neg,
ccomp, cc, conj, prep, prede... [(Case=Nom, Number=Plur, Person=3,
                      1.129032
PronType=Pr...
1 3T8DUCXY0N6WD9X4RTLK8UN1U929TF bible that sends ambassadors by the sea,
                      0.102941 that sends ambassadors by the sea, even in
even in ves...
                     False [PRON, VERB, NOUN, ADP, DET, NOUN, PUNCT, ADV, ...
ves...
[nsubj, ROOT, dobj, prep, det, pobj, punct, ad... [(PronType=Rel),
(Number=Sing, Person=3, Tense...
                                        1.263158
2 3I7KR83SNADXAQ7HXK7S7305BYB9KD bible and they entered into the boat, and
were going...
                     0.109375 and they entered into the boat, and were
             sea
```

```
False [CCONJ, PRON, VERB, ADP, DET, NOUN, PUNCT,
going...
CCO... [cc, nsubj, ROOT, prep, det, pobj, punct, cc, ... [(ConjType=Cmp),
(Case=Nom, Number=Plur, Perso...
                                        1.437500
3 3BO3NEOQMOHK9ERYPNOGQIWCPC4IAQ bible Joseph laid up grain as the sand of
the sea. v...
                     0.160714 Joseph laid up grain as the sand of the sea,
             sea
                   False [PROPN, VERB, ADP, NOUN, ADP, DET, NOUN, ADP, ...
[nsubj, ROOT, prt, dobj, prep, det, pobj, prep... [(Number=Sing), (Tense=Past,
VerbForm=Fin), ()...
                            1.400000
4 3Y3CZJSZ9KTOW7IOKE38WZHHKSW5RH bible There will be a highway for the
                         0.000000 There will be a highway for the remnant
remnant that i... land
                        False [PRON, AUX, AUX, DET, NOUN, ADP, DET, NOUN,
that i...
PR... [expl, aux, ROOT, det, attr, prep, det, pobj, ... [(), (VerbForm=Fin),
(VerbForm=Inf), (Definite...
                                    1.277778
                               id corpus
sentence
                  token complexity
sentence_no_contractions contraction_expanded
pos_sequence
                                                   dep_sequence
morph_sequence morph_complexity binary_complexity
O 31HLTCK4BLVQ5B01AUR91TX9V9IVGH bible The name of one son was Gershom, for
Moses sai...
             foreign land
                             0.000000 The name of one son was Gershom, for
Moses sai...
                           False [DET, NOUN, ADP, NUM, NOUN, AUX, PROPN,
PUNCT,... [det, nsubj, prep, nummod, pobj, ROOT, attr, p... [(Definite=Def,
PronType=Art), (Number=Sing), ...
                                        1.520000
1 389A2A3040IXVY7G5B71Q9M43LEOCL bible unleavened bread, unleavened cakes
mixed with ... wheat flour
                               0.157895 unleavened bread, unleavened cakes
                             False [ADJ, NOUN, PUNCT, ADJ, NOUN, VERB, ADP,
mixed with ...
NOUN,... [amod, dep, punct, amod, appos, acl, prep, pob... [(Degree=Pos),
                                          1.200000
(Number=Sing), (PunctType=Comm)...
2 31N9JPQXIPIRX2A3S9NOCCFXO6TNHR bible However the high places were not taken
away; t... burnt incense
                          0.200000 However the high places were not taken
                         False [ADV, DET, ADJ, NOUN, AUX, PART, VERB, ADV,
away; t...
PU... [advmod, det, amod, nsubjpass, auxpass, neg, c... [(), (Definite=Def,
PronType=Art), (Degree=Pos...
                                     1.190476
3 3JVP4ZJHDPS081TGXL3N1CKZGQY0IN bible and he burnt incense of sweet spices
on it, as... burnt incense
                             0.250000 and he burnt incense of sweet spices on
                        False [CCONJ, PRON, VERB, NOUN, ADP, ADJ, NOUN,
it, as...
ADP,... [cc, nsubj, ROOT, dobj, prep, amod, pobj, prep... [(ConjType=Cmp),
(Case=Nom, Gender=Masc, Numbe...
                                        1.466667
4 3JAOYN9IHL25ZQAUV5EJZ4GHOKL33L bible The same day the king made the middle
                            0.214286 The same day the king made the middle of
of the c...
           bronze altar
the c...
                       False [DET, ADJ, NOUN, DET, NOUN, VERB, DET, NOUN,
A... [det, amod, npadvmod, det, nsubj, ccomp, det, ... [(Definite=Def,
PronType=Art), (Degree=Pos), (...
                                         1.352113
                               id corpus
sentence
             token complexity
sentence_no_contractions contraction_expanded
pos_sequence
                                                   dep_sequence
morph_sequence morph_complexity binary_complexity
```

```
O 3K8CQCU3KE19US5SN890DFPK3SANWR bible But he, beckoning to them with his
                          0.000000 But he, beckoning to them with his hand to
hand to be ...
                  hand
                     False [CCONJ, PRON, PUNCT, VERB, ADP, PRON, ADP, PRO...
[cc, nsubj, punct, advcl, prep, pobj, prep, po... [(ConjType=Cmp), (Case=Nom,
Gender=Masc, Numbe...
                             1.703704
1 3Q2T3FD00N86LCI41NJYV3PN0BW3MV bible If I forget you, Jerusalem, let my
                          0.197368 If I forget you, Jerusalem, let my right
hand ...
                       False [SCONJ, PRON, VERB, PRON, PUNCT, PROPN,
PUNCT,... [mark, nsubj, advcl, dobj, punct, npadvmod, pu... [(), (Case=Nom,
Number=Sing, Person=1, PronTyp...
                                         1.800000
2 3ULIZOH1VA5C32JJMKOTQ8Z4GUS51B bible the ten sons of Haman the son of
                            0.200000 the ten sons of Haman the son of
Hammedatha, t...
                    hand
                                      [DET, NUM, NOUN, ADP, PROPN, DET, NOUN,
Hammedatha, t...
                                True
ADP, P... [det, nummod, ROOT, prep, pobj, det, appos, pr... [(Definite=Def,
PronType=Art), (NumType=Card),...
                                         1.269231
3 3BFFODJK8XCEIOT3OZLBPPSRMZQTSD bible Let your hand be lifted up above your
adversar...
               hand
                       0.267857 Let your hand be lifted up above your
                          False [VERB, PRON, NOUN, AUX, VERB, ADP, ADP, PRON,
adversar...
... [ROOT, poss, nsubjpass, auxpass, ccomp, prt, p... [(VerbForm=Inf),
(Person=2, Poss=Yes, PronType...
                                        1.250000
4 3QREJ3J433XSBS8QMHAICCROBQ1LKR bible Abimelech chased him, and he fled
                           0.000000 Abimelech chased him, and he fled before
before him, ... entrance
                      False [PROPN, VERB, PRON, PUNCT, CCONJ, PRON, VERB,
... [nsubj, ROOT, dobj, punct, cc, nsubj, conj, pr... [(Number=Sing),
(Tense=Past, VerbForm=Fin), (C...
                                         1.652174
                               id corpus
sentence
                   token complexity
sentence_no_contractions contraction_expanded
pos_sequence
                                                   dep_sequence
morph_sequence morph_complexity binary_complexity
O 3UXQ63NLAAMRIP4WG4XPD98AOYOBLX bible for he had an only daughter, about
                                0.025000 for he had an only daughter, about
twelve year...
              only daughter
twelve year...
                             False [SCONJ, PRON, VERB, DET, ADJ, NOUN, PUNCT,
ADV... [mark, nsubj, ROOT, det, amod, dobj, punct, ad... [(), (Case=Nom,
Gender=Masc, Number=Sing, Pers...
                                         1.722222
1 3FJ2RVH25Z62TA3R8E1077EBUYU92W bible All these were cities fortified with
                              0.100000 All these were cities fortified with
high wall...
                high walls
high wall...
                           False [DET, PRON, AUX, NOUN, VERB, ADP, ADJ, NOUN,
P... [predet, nsubj, ROOT, attr, acl, prep, amod, p... [(), (Number=Plur,
PronType=Dem), (Mood=Ind, T...
                                      1.136364
2 3YO4AH2FPDK1PZHZAT8WAEBL70EQOF bible In the morning, 'It will be foul
                weather today
                                  0.125000 In the morning, 'It will be foul
weather today...
weather today...
                               False [ADP, DET, NOUN, PUNCT, PUNCT, PRON,
AUX, AUX,... [prep, det, pobj, punct, punct, nsubj, aux, RO... [(),
(Definite=Def, PronType=Art), (Number=Sin...
                                                    1.476190
3 3X52SWXEOX5Q3081YIOMX4V84QTCWZ bible Her young children also were dashed in
```

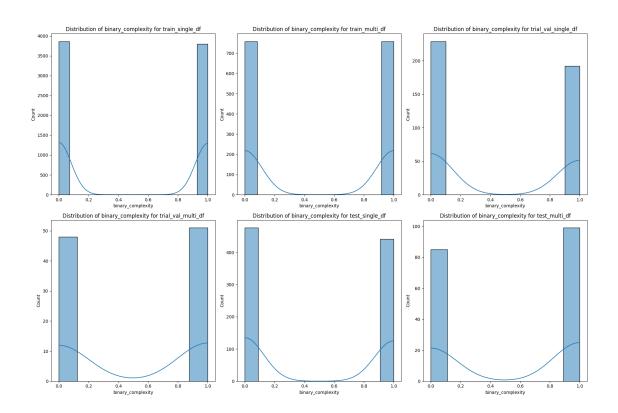
0.160714 Her young children also were dashed in

pieces ... young children

```
False [PRON, ADJ, NOUN, ADV, AUX, VERB, ADP, NOUN,
     pieces ...
     A... [poss, amod, nsubjpass, advmod, auxpass, ROOT,... [(Gender=Fem,
     Number=Sing, Person=3, Poss=Yes,...
                                                1.514286
     4 32K26U12DNONTREA84Q1V8UCIH2VD7 bible All king Solomon's drinking vessels
     were of go...
                       pure gold 0.178571 All king Solomon's drinking vessels
                                 False [DET, NOUN, PROPN, PART, NOUN, NOUN, AUX,
     were of go...
     ADP,... [det, compound, poss, case, compound, nsubj, c... [(), (Number=Sing),
     (Number=Sing), (), (Number...
                                          1.162791
[46]: # prompt: show me the distributions of binary_complexity for all 6 dataframes
      import matplotlib.pyplot as plt
      import seaborn as sns
      # Assuming your dataframes are named as in the provided code:
      dataframes = {
          "train_single_df": train_single_df,
          "train_multi_df": train_multi_df,
          "trial_val_single_df": trial_val_single_df,
          "trial_val_multi_df": trial_val_multi_df,
          "test_single_df": test_single_df,
          "test_multi_df": test_multi_df
      }
      # Create a figure and axes for the subplots
      fig, axes = plt.subplots(2, 3, figsize=(18, 12)) # Adjust figsize as needed
      # Iterate through dataframes and create distributions
      for i, (df_name, df) in enumerate(dataframes.items()):
       row = i // 3
        col = i % 3
       ax = axes[row, col]
       sns.histplot(df['binary_complexity'], kde=True, ax=ax) #kde=True adds a__
       ⇔kernel density estimate
       ax.set_title(f'Distribution of binary_complexity for {df_name}')
       ax.set_xlabel('binary_complexity')
```

plt.tight\_layout() # Adjust subplot params for a tight layout

plt.show()



## [49]: !ls -R /content/drive/MyDrive/266-final/data/266-comp-lex-master/

/content/drive/MyDrive/266-final/data/266-comp-lex-master/: fe-test-labels fe-train fe-trial-val test-labels train trial

/content/drive/MyDrive/266-final/data/266-comp-lex-master/fe-test-labels:

/content/drive/MyDrive/266-final/data/266-comp-lex-master/fe-train:

/content/drive/MyDrive/266-final/data/266-comp-lex-master/fe-trial-val:

/content/drive/MyDrive/266-final/data/266-comp-lex-master/test-labels:
lcp\_multi\_test.tsv lcp\_single\_test.tsv

/content/drive/MyDrive/266-final/data/266-comp-lex-master/train:
lcp\_multi\_train.tsv lcp\_single\_train.tsv

/content/drive/MyDrive/266-final/data/266-comp-lex-master/trial:
lcp\_multi\_trial.tsv lcp\_single\_trial.tsv

[50]: | tree /content/drive/MyDrive/266-final/data/266-comp-lex-master/

/content/drive/MyDrive/266-final/data/266-comp-lex-master/
fe-test-labels

```
fe-train
        fe-trial-val
        test-labels
           lcp_multi_test.tsv
           lcp_single_test.tsv
        train
           lcp_multi_train.tsv
           lcp_single_train.tsv
        trial
            lcp_multi_trial.tsv
            lcp_single_trial.tsv
     6 directories, 6 files
[51]: # prompt: save all 6 dataframes to fe-test-labels, fe-train, fe-trial-val which
       ware subdirectories located in /content/drive/MyDrive/266-final/data/
       \hookrightarrow 266-comp-lex-master/
      import os
      # Assuming your dataframes are named as in the provided code:
      dataframes = {
          "train_single_df": train_single_df,
          "train_multi_df": train_multi_df,
          "trial_val_single_df": trial_val_single_df,
          "trial_val_multi_df": trial_val_multi_df,
          "test_single_df": test_single_df,
          "test_multi_df": test_multi_df
      }
      base_dir = "/content/drive/MyDrive/266-final/data/266-comp-lex-master/"
      for df_name, df in dataframes.items():
          subdir = None
          if "train" in df_name:
            subdir = "fe-train"
          elif "trial_val" in df_name:
            subdir = "fe-trial-val"
          elif "test" in df_name:
            subdir = "fe-test-labels"
          if subdir:
            save_path = os.path.join(base_dir, subdir, f"{df_name}.csv")
            os.makedirs(os.path.dirname(save_path), exist_ok=True) # create directory_
       \hookrightarrow if it doesn't exist
            df.to_csv(save_path, index=False)
            print(f"Saved {df_name} to {save_path}")
```

Saved train\_single\_df to /content/drive/MyDrive/266-final/data/266-comp-lex-

```
master/fe-train/train_single_df.csv
     Saved train_multi_df to /content/drive/MyDrive/266-final/data/266-comp-lex-
     master/fe-train/train_multi_df.csv
     Saved trial_val_single_df to /content/drive/MyDrive/266-final/data/266-comp-lex-
     master/fe-trial-val/trial val single df.csv
     Saved trial_val_multi_df to /content/drive/MyDrive/266-final/data/266-comp-lex-
     master/fe-trial-val/trial val multi df.csv
     Saved test_single_df to /content/drive/MyDrive/266-final/data/266-comp-lex-
     master/fe-test-labels/test_single_df.csv
     Saved test_multi_df to /content/drive/MyDrive/266-final/data/266-comp-lex-
     master/fe-test-labels/test_multi_df.csv
[52]: | tree /content/drive/MyDrive/266-final/data/266-comp-lex-master/
     /content/drive/MyDrive/266-final/data/266-comp-lex-master/
        fe-test-labels
           test_multi_df.csv
           test_single_df.csv
        fe-train
           train_multi_df.csv
           train_single_df.csv
        fe-trial-val
           trial_val_multi_df.csv
           trial_val_single_df.csv
        test-labels
           lcp_multi_test.tsv
           lcp_single_test.tsv
           lcp_multi_train.tsv
           lcp_single_train.tsv
        trial
            lcp_multi_trial.tsv
            lcp_single_trial.tsv
     6 directories, 12 files
```

64

[]: