Dummy Baseline Model

This notebook creates a baseline model by extracting the first 1000 characters of the abstracts.

Drive already mounted at /content/drive; to attempt to forcibly remount, cal l drive.mount("/content/drive", force remount=True).

Load Validation Data

```
In [3]: INPUT_FILE_DIR = "/content/drive/MyDrive/BioLaySumm2024_main/extracted/"
        ELIFE_VAL_FILE_NAME = "eLife_val_extracted.csv"
        PLOS_VAL_FILE_NAME = "PLOS_val_extracted.csv"
        val_eLife_df = pd.read_csv(INPUT_FILE_DIR + ELIFE_VAL_FILE_NAME)
        val plos df = pd.read csv(INPUT FILE DIR + PLOS VAL FILE NAME)
In [6]: val_eLife_df['abstract'].apply(lambda x: x[:1000])
Out[6]: 0
               Cell-fate reprograming is at the heart of deve...
        1
               Klebsiella pneumoniae is a respiratory , blood...
               Plasmodium vivax relapse infections occur foll...
               When 2 Mha of Amazonian forests are disturbed ...
               Neurosphere formation is commonly used as a su...
        236
               Similarity between two individuals in the comb...
        237
               Early development of many animals is character...
        238
               Gene control systems sometimes interpret multi...
        239
               The number of neurotransmitter-filled vesicles...
        240
               HIV-1 Gag selects and packages a dimeric , uns...
        Name: abstract, Length: 241, dtype: object
```

Output Validation data

```
In [19]: OUTPUT_FILE_DIR = "/content/drive/MyDrive/BioLaySumm2024-evaluation_scripts/
    val_eLife_df['abstract'].apply(lambda x: x[:1000]).to_csv(OUTPUT_FILE_DIR+'e
    val_plos_df['abstract'].apply(lambda x: x[:1000]).to_csv(OUTPUT_FILE_DIR+'pl
```

Load test data

```
In [7]: ELIFE_TEST_FILE_NAME = "eLife_test_extracted.csv"
PLOS_TEST_FILE_NAME = "PLOS_test_extracted.csv"

test_eLife_df = pd.read_csv(INPUT_FILE_DIR + ELIFE_TEST_FILE_NAME)

test_plos_df = pd.read_csv(INPUT_FILE_DIR + PLOS_TEST_FILE_NAME)

In [8]: test_eLife_df
```

Out[8]:		Unnamed: 0	headings	keywords	id	abstract
	0	0	['Abstract', 'Introduction', 'Results and disc	['biochemistry and chemical biology', 'computa	elife- 81547- v1	Acylation of diverse carbohydrates occurs acro
	1	1	['Abstract', 'Introduction', 'Results', 'Discu	['computational and systems biology']	elife- 86176- v2	Honey bee ecology demands they make both rapid
	2	2	['Abstract', 'Introduction', 'Results', 'Discu	['genetics and genomics']	elife- 82210- v1	Biguanides , including the world's most prescr
	3	3	['Abstract', 'Introduction', 'Results', 'Discu	['microbiology and infectious disease', 'ecolo	elife- 83152- v2	Ecological relationships between bacteria medi
	4	4	['Abstract', 'Introduction', 'Results', 'Discu	['neuroscience']	elife- 83044- v2	Gamma oscillations are believed to underlie co
	•••					
	137	137	['Abstract', 'Introduction', 'Results', 'Discu	['tools and resources', 'genetics and genomics']	elife- 84831- v1	High-throughput transgenesis using synthetic D
	138	138	['Abstract', 'Introduction', 'Results', 'Discu	['structural biology and molecular biophysics'	elife- 82885- v1	To reach their final destinations , outer memb
	139	139	['Abstract', 'Introduction', 'Results', 'Discu	['neuroscience']	elife- 87902- v2	Midbrain dopamine (DA) neurons are key regul
	140	140	['Abstract', 'Introduction', 'Methods', 'Resul	['medicine', 'epidemiology and global health']	elife- 79615- v2	Mobile health (mHealth) interventions , whic
	141	141	['Abstract', 'Introduction', 'Methods', 'Resul	['medicine', 'epidemiology and global health']	elife- 80428- v2	Severe acute respiratory syndrome coronavirus

142 rows × 5 columns

Output Test Data

```
In [16]: OUTPUT_FILE_DIR = "/content/drive/MyDrive/BioLaySumm2024_main/test_submission
    test_eLife_df['abstract'].apply(lambda x: x[:1000]).to_csv(OUTPUT_FILE_DIR+'
    test_plos_df['abstract'].apply(lambda x: x[:1000]).to_csv(OUTPUT_FILE_DIR+'public line of the state o
```