# Music Recommendation System

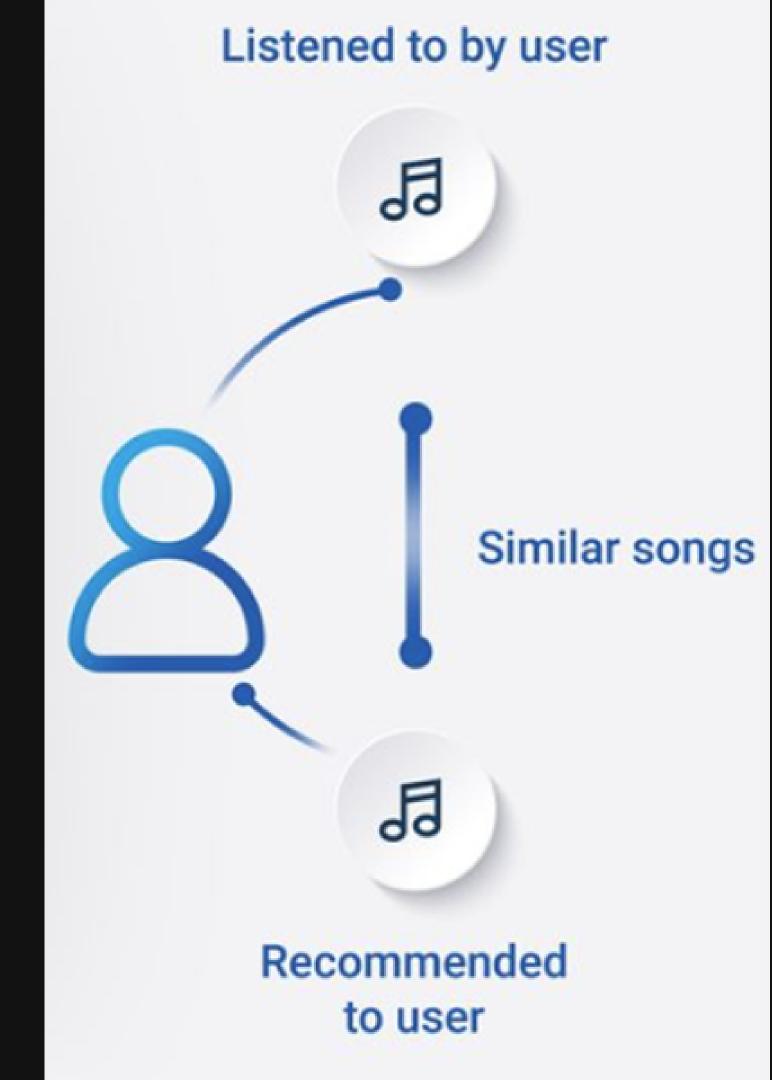
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# Team

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# Problem Statement

Building a Music Recommendation System using Similarity Finding Algorithms



# Motivation

Have you ever heard a song and wanted to listen similar songs?

- With commercial music streaming service which can be
  accessed from mobile devices, the availability of digital
  music currently is abundant compared to previous era.
   Sorting out all this digital music is a very timeconsuming and causes information fatigue. Therefore, it
  is very useful to develop a music recommender system
  that can search in the music libraries automatically and
  suggest suitable songs to users.
- By using music recommender system, the music provider can predict and then offer the appropriate songs to their users based on the characteristics of the music that has been heard previously.
- Our research would like to develop a music recommender system that can give recommendations based on similarity of features on audio signal.

# MP3S-32k Dataset

https://www.kaggle.com/datasets/dhrumil140396/mp3s32k



MP3 AUDIO FILES



1413 SONGS

### Project Pipeline

#### STEP 1

Converting the songs from mp3 to wav with Librosa and extraction of the peaks.

#### STEP 2

Extracting shingles from songs spectrogram

#### STEP 3

MinHashing with permutations on the shingles matrix.

#### STEP 4

Locality
sensitive
hashing to
divide the songs
in buckets.

#### STEP 5

Jaccard similarity calculation

#### STEP 6

Song recommendation

# MinHashing

- Minhashing involves compressing the large sets of unique shingles into a much smaller representation called "signatures".
- We then use these signatures to measure the similarity
- Although it is impossible for these signatures to give the exact similarity measure, the estimates are pretty close.
- The larger the number of signatures chosen, the more accurate the estimate is.

$$h(x) = (ax + b) \bmod c$$

- x is the row numbers of your original characteristic matrix.
- a and b are any random numbers smaller or equivalent to the maximum number of x, and they both must be unique in each signature.
- b- coefficient in signature
- c is a prime number slightly larger than the total number of shingle sets.

# LSH

The concept for locality-sensitive hashing (LSH) is that given the signature matrix of size n (row count), we will partition it into b bands, resulting in each band with r rows. This is equivalent to the simple math formula -n = br, thus when we are doing the partition, we have to be sure that the b we choose is divisible by n.

# Jaccard Similarity

- After creating shingle sets and characteristic matrix, we now need to measure the similarity between documents.
- We will make use of Jaccard Similarity for this purpose.
- For example, with two shingle sets as set1 and set2, the Jaccard Similarity will be:

| set1 ∩ set2 | | set1 ∪ set2 |

# Preprocessing

```
def unique_shingles(song_peaks):
    tot_shingles = list(song_peaks.values())
    shingles = []
    for i in tqdm(tot_shingles):
        shingles.append(i)

    shingles = np.hstack(shingles)
    shingles = np.array(list(dict.fromkeys(shingles))) # all unique peaks
    return shingles
```

```
def load_audio_peaks(audio, offset, duration, hop_size):
    d
    track, sr = librosa.load(audio, offset=offset, duration=duration)
    onset_env = librosa.onset.onset_strength(track, sr=sr, hop_length=hop_size)
    peaks = librosa.util.peak_pick(onset_env, 10, 10, 10, 10, 0.5, 0.5)
    return track, sr, onset_env, peaks
```

```
def convert_mp3_to_wav(audio:str) -> str:
    if audio[-3:] == "mp3":
        wav_audio = audio[:-3] + "wav"
        if not Path(wav_audio).exists():
            subprocess.check_output(f"ffmpeg -i {audio} {wav_audio}", shell=True)
        return wav_audio

    return audio
```

```
def onehot(peaks, shingles):
    return np.array([1 if x in peaks else 0 for x in shingles])

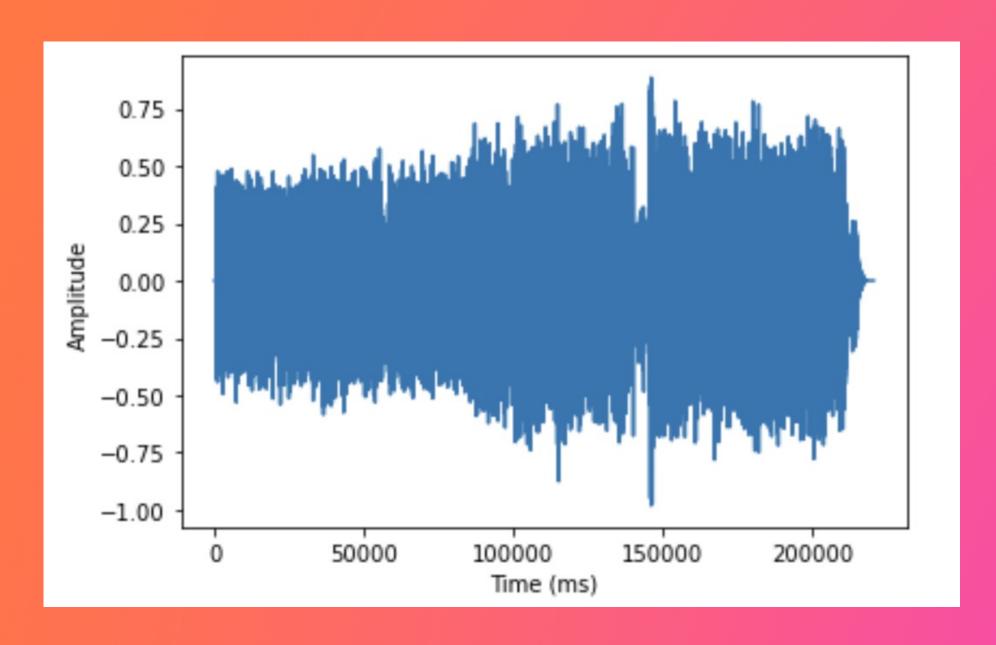
def shingles_matrix(shingles, song_peaks):
    matrix = np.zeros(len(shingles))

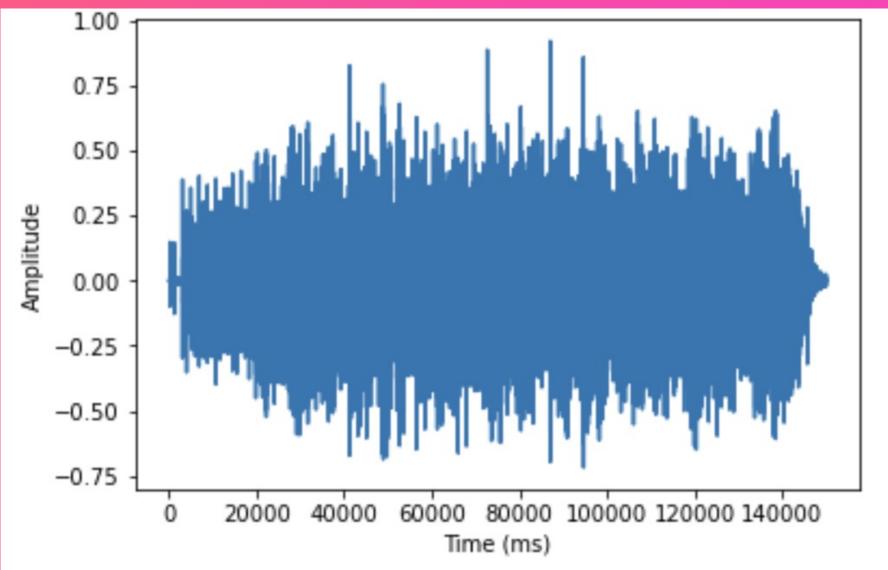
for v in tqdm(list(song_peaks.values())):
    matrix = np.vstack([matrix, onehot(v, shingles)])

matrix = np.delete(matrix, (0), axis=0)

return matrix
```

# Spectrograms





```
def save_pickle(element, path):
   with open(f"{path}", 'wb') as f:
        pickle.dump(element, f, pickle.HIGHEST_PROTOCOL)
def load_pickle(path):
   with open(f"{path}", 'rb',) as f:
        return pickle.load(f)
def convert_mp3_to_wav(audio:str) -> str:
   if audio[-3:] == "mp3":
       wav_audio = audio[:-3] + "wav"
        if not Path(wav_audio).exists():
                subprocess.check_output(f"ffmpeg -i {audio} {wav_audio}", shell=True)
        return wav_audio
    return audio
```

```
def load audio peaks(audio, offset, duration, hop size):
    track, sr = librosa.load(audio, offset=offset, duration=duration)
    onset env = librosa.onset.onset strength(track, sr=sr, hop length=hop size)
    peaks = librosa.util.peak_pick(onset_env, 10, 10, 10, 10, 0.5, 0.5)
    return track, sr, onset_env, peaks
def extract_peaks(song_path, rounded = False):
    song peaks = {}
    if rounded == True:
        for song in tqdm(song_path):
            tmp1, tmp2, onset, peaks = load_audio_peaks(str(song), OFFSET, DURATION, HOP_SIZE)
            song_peaks[' '.join(str(song).split('/')[-1][3:-4].split('_')).lower() + ' - ' + ' '.join(str(song).split('/')[0].spl
    else:
        for song in tqdm(song_path):
            tmp1, tmp2, onset, peaks = load_audio_peaks(str(song), OFFSET, DURATION, HOP_SIZE)
            song_peaks[' '.join(str(song).split('/')[-1][3:-4].split('_')).lower() + ' - ' + ' '.join(str(song).split('/')[0].spl
    return song_peaks
```

```
def unique_shingles(song_peaks):
   tot_shingles = list(song_peaks.values())
   shingles = []
   for i in tqdm(tot_shingles):
       shingles.append(i)
   shingles = np.hstack(shingles)
   shingles = np.array(list(dict.fromkeys(shingles))) # all unique peaks
   return shingles
def onehot(peaks, shingles):
    return np.array([1 if x in peaks else 0 for x in shingles])
def shingles_matrix(shingles, song_peaks):
    matrix = np.zeros(len(shingles))
    for v in tqdm(list(song_peaks.values())):
        matrix = np.vstack([matrix, onehot(v, shingles)])
    matrix = np.delete(matrix, (0), axis=0)
    return matrix
```

```
def hash matrix(matrix, shingles, song peaks):
   # we transpose the matrix in order to have the shingles on the rows and the songs on the columns
    df = pd.DataFrame(matrix.transpose(), index = range(len(shingles)), columns = list(song peaks.keys()))
    hash matrix = np.zeros(len(song peaks), dtype = int)
    # we permutate the rows of the matrix and for each column we look at the first non-zero value and store in a list
    # the corresponing raw index of that value, then by stacking the list at each permutation we get back the hash matrix
   for i in tqdm(range(nperm)):
        hash matrix = np.vstack([hash matrix, list(df.sample(frac = 1, random state = i).reset index(drop=True).ne(0).idxmax())]
        # .sample shuffles all the rows of the matrix
        \# .ne(x) looks for the values different from x
        # .idxmax finds the first index between all the indexes with non-zero values
    hash matrix = np.delete(hash matrix, (0), axis=0)
    hash mat = pd.DataFrame(hash matrix, index = range(1, nperm + 1), columns = list(song peaks.keys()))
    return hash mat
```

```
def db buckets(hash matrix, n bands):
    # first we have to decide a number of bands that is a divisor of the signature length in order to have equal slices of the si
    # of course the less bands we use the more discriminant the LSH will be
   rows = int(nperm/n bands)
    buckets = {}
    for song name, song hash in hash matrix.iteritems():
        song hash = list(song hash) # convert the columns of the dataframe from pandas series into lists
        for i in range(0, len(song hash), rows):
            bucket hash = tuple(song hash[i : i + rows]) # the hash of the bucket will be a tuple with number of elements = rows
            if bucket hash in buckets:
                buckets[bucket hash].add(song name) # if we already encountered that band we only add the song name
            else:
                buckets[bucket hash] = {song name} # otherwise we create a new key:value
    return buckets
```

```
def query_buckets(fingerprint, n_bands):
    # same as before but in this case fingerprint is a list and not a dataframe
    rows = int(len(fingerprint)/n_bands)
    # splitting the signature in nbands subvectors
    q_buckets = {}
    for i in range(0, len(fingerprint), rows):
        q_buckets[tuple(fingerprint[i : i + rows])] = 'query'
    return q_buckets
```

```
def shazamLSH(query, database, shingles, buckets):
    print('Im listening to your music, please dont make noise ...')
    score = (0, '')
    db keys = list(database.keys())
    buckets_keys = list(buckets.keys())
    query fingerprint = list(fingerprint(query, shingles, rounded=True)['query'])
    query_bands = query_buckets(query_fingerprint, 5)
    query keys = list(query bands.keys())
    # we compute the intersection between the query buckets and the database buckets
    common_bands = set(query_bands).intersection(set(buckets_keys))
    # we compute the jaccard only with the songs in the buckets of the intersection
    for band in common bands:
       for song in buckets[band]:
            jac = sklearn.metrics.jaccard score(query fingerprint, database[song], average = None)
            if score < (jac.any(), song):</pre>
                score = (jac.any(), song) # store the maximum score
    print('Maybe you were looking for this song: ', score[1], '\n-----\n')
```

### Result

# Thank You!