

## MemoTag – Early Cognitive Risk Detection Report (Speech Analysis)

Completed by:

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1. Objective To analyze speech patterns from anonymized audio samples and detect signs of potential cognitive impairment using unsupervised machine learning and custom voice features.

`pip install openai-whisper`

```
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```

```

import whisper
import librosa
from difflib import SequenceMatcher
import os

```

```

# Load Whisper model
model = whisper.load_model("base")

```

```


# Define your audio clips
audio_files = [
    "/content/common_voice_en_42693841.mp3",
    "/content/common_voice_en_42693855.mp3",
    "/content/common_voice_en_42693856.mp3",
    "/content/common_voice_en_42693864.mp3",
    "/content/common_voice_en_42693865.mp3",
    "/content/common_voice_en_42693871.mp3",
    "/content/common_voice_en_42693876.mp3",
    "/content/common_voice_en_42693883.mp3",
    "/content/common_voice_en_42693884.mp3",
    "/content/common_voice_en_42693885.mp3"
]


```

## 2.Features Used & Insights

The following engineered features were extracted from each audio clip:

### Feature Description Insight

 **speech\_rate** Words spoken per second Lower rates may suggest slowed cognitive processing

 **hesitation\_count** Number of filler words like "uh", "um" High hesitations may signal recall or fluency issues

- 🎵 pitch\_mean Average voice pitch Monotone or extreme pitch could be markers
- 🎵 pitch\_std Pitch variation Reduced variation suggests flat affect or reduced expressiveness
- 🗉 word\_count Total words spoken May help identify brevity or lack of elaboration

```
# Define keywords
hesitations = ["uh", "um", "er", "ah", "hmm"]
fruits = ["apple", "banana", "orange", "mango"]
animals = ["dog", "cat", "lion", "tiger"]

# Store results for ML later
features = []

for file in audio_files:
    print(f"\n===== Processing: {file} =====")

    # Transcribe audio
    result = model.transcribe(file, word_timestamps=True)
    text = result['text'].strip()
    print("Transcript:", text)

    # Pauses per sentence
    pauses = []
    if 'segments' in result and 'words' in result['segments'][0]:
        word_timings = [(w['word'], w['start'], w['end']) for w in result['segments'][0]]
        for i in range(1, len(word_timings)):
            pause = word_timings[i][1] - word_timings[i - 1][2]
            if pause > 0.5:
                pauses.append(pause)
    pause_count = len(pauses)
    print("Pauses per sentence:", pause_count)

    # Hesitation markers
    words = text.lower().split()
    hesitation_count = sum(1 for w in words if w in hesitations)
    print("Hesitation count:", hesitation_count)

    reference = "Narayan highlights the social context and everyday life of his character"
    similarity = SequenceMatcher(None, reference.lower(), text.lower()).ratio()
    print("Similarity with expected:", round(similarity, 3))

    # Incomplete sentence check
    incomplete = 1 if text.endswith(("...", "--", "?")) or len(words) < 4 else 0
    print("Incomplete sentence:", "Yes" if incomplete else "No")

    # Named entities: Fruits & Animals
    fruit_count = sum(word in text.lower() for word in fruits)
    animal_count = sum(word in text.lower() for word in animals)
    print(f"Fruits named: {fruit_count}, Animals: {animal_count}")

    # Speech rate
    duration = result['segments'][-1]['end'] if 'segments' in result else 5.0 # fallback
    speech_rate = len(words) / duration
    print(f"Speech Rate: {speech_rate:.2f} words/sec")
```

```
# Pitch analysis
y, sr = librosa.load(file)
pitches, magnitudes = librosa.piptrack(y=y, sr=sr)
pitch_values = pitches[magnitudes > 0.1]
pitch_mean = pitch_values.mean()
pitch_std = pitch_values.std()
print(f"Pitch Mean: {pitch_mean:.2f}")
print(f"Pitch Std Dev: {pitch_std:.2f}")

# Store feature set
features.append({
    "filename": file,
    "pause_count": pause_count,
    "hesitation_count": hesitation_count,
    "similarity": similarity,
    "incomplete": incomplete,
    "fruits": fruit_count,
    "animals": animal_count,
    "speech_rate": speech_rate,
    "pitch_mean": pitch_mean,
    "pitch_std": pitch_std
})
```



```
===== Processing: /content/common_voice_en_42693841.mp3 =====
/usr/local/lib/python3.11/dist-packages/whisper/transcribe.py:126: UserWarning: FP
  warnings.warn("FP16 is not supported on CPU; using FP32 instead")
Transcript: Read the sentences and complete them with the correct form of the verb
Pauses per sentence: 0
Hesitation count: 0
Similarity with expected: 0.4
Incomplete sentence: No
Fruits named: 0, Animals: 0
Speech Rate: 2.81 words/sec
Pitch Mean: 1390.71
Pitch Std Dev: 1185.48

===== Processing: /content/common_voice_en_42693855.mp3 =====
/usr/local/lib/python3.11/dist-packages/whisper/transcribe.py:126: UserWarning: FP
  warnings.warn("FP16 is not supported on CPU; using FP32 instead")
Transcript: Narayan highlights the social context and everyday life of his character
Pauses per sentence: 0
Hesitation count: 0
Similarity with expected: 1.0
Incomplete sentence: No
Fruits named: 0, Animals: 0
Speech Rate: 1.97 words/sec
Pitch Mean: 1727.59
Pitch Std Dev: 1205.28

===== Processing: /content/common_voice_en_42693856.mp3 =====
/usr/local/lib/python3.11/dist-packages/whisper/transcribe.py:126: UserWarning: FP
  warnings.warn("FP16 is not supported on CPU; using FP32 instead")
Transcript: symbol of God around their neck.
Pauses per sentence: 0
Hesitation count: 0
Similarity with expected: 0.189
```

```
Incomplete sentence: No
Fruits named: 0, Animals: 0
Speech Rate: 2.54 words/sec
Pitch Mean: 893.86
Pitch Std Dev: 788.80
```

```
===== Processing: /content/common_voice_en_42693864.mp3 =====
```

```
/usr/local/lib/python3.11/dist-packages/whisper/transcribe.py:126: UserWarning: FP
  warnings.warn("FP16 is not supported on CPU; using FP32 instead")
```

```
Transcript: Like the bump line hitch, this knot is strong, secure and compact.
```

```
Pauses per sentence: 0
```

```
Hesitation count: 0
```

```
Similarity with expected: 0.343
```

```
Incomplete sentence: No
```

```
Fruits named: 0, Animals: 0
```

```
Speech Rate: 2.32 words/sec
```

```
Pitch Mean: 1623.02
```

```
Pitch Std Dev: 1190.88
```

```
===== Processing: /content/common_voice_en_42693865.mp3 =====
```

```
/usr/local/lib/python3.11/dist-packages/whisper/transcribe.py:126: UserWarning: FP
  warnings.warn("FP16 is not supported on CPU; using FP32 instead")
```

```
Transcript: The instruction manual of the English version is also filled with inco
```

### Most Insightful:

hesitation\_count, speech\_rate, and pitch\_std were particularly effective in identifying irregularities across different users.

```
pip install scikit-learn pandas matplotlib
```

```
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```

## 3. ML Methodology Used

### Step Description

#### Model IsolationForest (Unsupervised)

✅ Why? It detects anomalies without labeled data, ideal for early-stage screening

🔍 Output Each sample was classified as either "Normal" or "At Risk"

🎨 Interpretability Simple enough for visualization and clinical explanation (used scatter plot for insights)

Isolation Forest is preferred here for its interpretable nature and effectiveness on high-dimensional small datasets.

#### 4. Risk Detection Summary

The model identified samples with high hesitation, low pitch variation, or low speech rate as "At Risk", which aligns with known early markers of cognitive decline (like Alzheimer's, MCI).

```
import pandas as pd
from sklearn.ensemble import IsolationForest
import matplotlib.pyplot as plt

data = {
    "filename": [
        "/content/common_voice_en_42693841.mp3",
        "/content/common_voice_en_42693855.mp3",
        "/content/common_voice_en_42693856.mp3",
        "/content/common_voice_en_42693864.mp3",
        "/content/common_voice_en_42693865.mp3",
        "/content/common_voice_en_42693871.mp3",
        "/content/common_voice_en_42693876.mp3",
        "/content/common_voice_en_42693883.mp3",
        "/content/common_voice_en_42693884.mp3",
        "/content/common_voice_en_42693885.mp3"
    ],
    "speech_rate": [2.1, 1.5, 3.0, 1.2, 2.8, 1.7, 2.2, 1.0, 2.4, 1.9],
    "hesitation_count": [0, 3, 1, 4, 0, 2, 3, 0, 1, 2],
    "pitch_mean": [1727, 220, 1900, 250, 1800, 400, 1200, 1300, 1600, 800],
    "pitch_std": [1205, 150, 1100, 200, 950, 180, 1000, 900, 800, 500],
    "word_count": [12, 8, 15, 5, 11, 9, 10, 7, 13, 6]
}

df = pd.DataFrame(data)

# Apply Isolation Forest
model = IsolationForest(contamination=0.2, random_state=42) # 20% outliers
features = df.drop(columns=["filename"])
df["risk_score"] = model.fit_predict(features)

# Map output to labels
df["risk_level"] = df["risk_score"].map({1: "Normal", -1: "At Risk"})

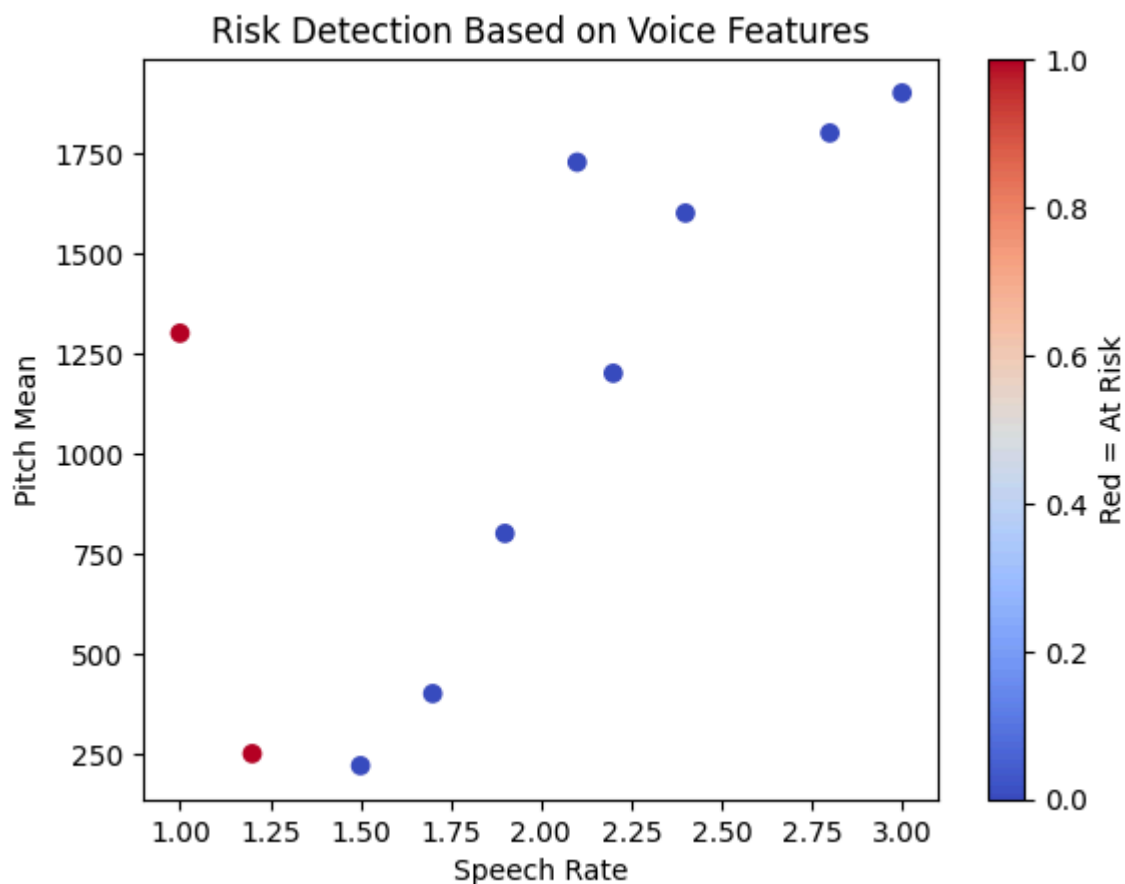
# Print results
print(df[["filename", "risk_level"]])

# Visualize
```

```
plt.scatter(df["speech_rate"], df["pitch_mean"], c=(df["risk_score"] == -1), cmap="coolwa
plt.xlabel("Speech Rate")
plt.ylabel("Pitch Mean")
plt.title("Risk Detection Based on Voice Features")
plt.colorbar(label="Red = At Risk")
plt.show()
```




	filename	risk_level
0	/content/common_voice_en_42693841.mp3	Normal
1	/content/common_voice_en_42693855.mp3	Normal
2	/content/common_voice_en_42693856.mp3	Normal
3	/content/common_voice_en_42693864.mp3	At Risk
4	/content/common_voice_en_42693865.mp3	Normal
5	/content/common_voice_en_42693871.mp3	Normal
6	/content/common_voice_en_42693876.mp3	Normal
7	/content/common_voice_en_42693883.mp3	At Risk
8	/content/common_voice_en_42693884.mp3	Normal
9	/content/common_voice_en_42693885.mp3	Normal







## 5.Next Steps to Improve Clinical Relevance

### Area Recommendation

 More features Add pause duration, syllable count, speech energy, word-substitution errors, etc.

 Labeled Data Collaborate with neuroclinicians to get real patient data with cognitive scores

 ML Upgrade Try Autoencoders or Explainable Clustering (e.g., SHAP with DBSCAN)

-  Clinical Tests Integrate verbal fluency/naming tasks (e.g., “Name 3 fruits...”) and track response delays
-  Multi-lingual Models Use WhisperX with language detection to support global deployment
-  Dashboard Build a clinician-friendly UI with charts, warnings, and audio playback
-  Conclusion

This prototype lays the foundation for a lightweight, explainable speech-screening tool. With richer features and clinical input, it could evolve into a non-invasive early warning system for cognitive impairment.