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A PPG (Photoplethysmography) signal is an optical technique used to measure blood volume changes in the microvascular tissue. It is commonly used for heart rate monitoring and oxygen saturation (SpO₂) measurement.

Working Principle of PPG Signal:

- 1. **Light Emission & Absorption:** An LED emits light onto the skin, which is either absorbed or reflected by the blood vessels.
- 2. **Detection:** A photodetector measures the amount of reflected or transmitted light, which varies with blood flow.
- 3. **Signal Processing:** The detected variations are converted into a PPG waveform, which represents the pulsatile changes in blood volume with each heartbeat.

Good ppg signal:



Fig-01: Good Ppg signal

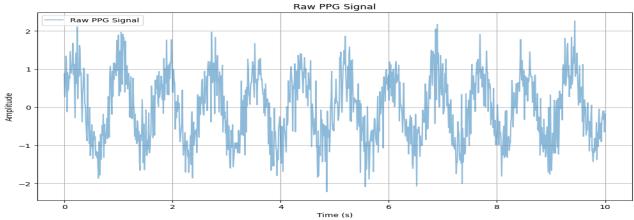
Features of PPG (Photoplethysmography) Signal

1.Raw ppg signal: A raw PPG (Photoplethysmogram) signal represents the variation in light absorption due to changes in blood volume in the microvascular bed of tissue. It's typically measured using a photodetector, like in smartwatches or pulse oximeters, and provides information about heart rate, blood oxygen levels, and other cardiovascular parameters. The raw PPG signal consists of **AC Component** and **DC Component**.

Raw ppg signal CODE

import numpy as np import pandas as pd import matplotlib.pyplot as plt from scipy.signal import find_peaks, butter, filtfilt

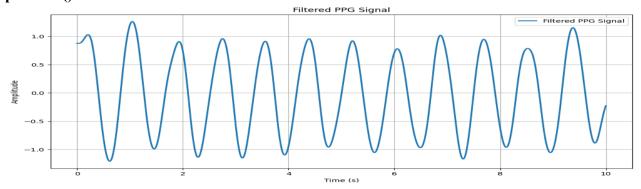
```
# Generate sample PPG signal
np.random.seed(0)
time = np.linspace(0, 10, 1000)
ppg signal = np.sin(2 * np.pi * 1.2 * time) + 0.5 * np.random.normal(size=len(time))
# Define the lowpass filter function
def butter lowpass filter(data, cutoff, fs, order=5):
  nyquist = 0.5 * fs
  normal cutoff = cutoff / nyquist
  b, a = butter(order, normal cutoff, btype='low', analog=False)
  y = filtfilt(b, a, data)
  return y
# Plot 1: Raw PPG Signal
plt.figure(figsize=(12, 6))
plt.plot(time, ppg signal, label="Raw PPG Signal", alpha=0.5)
plt.title("Raw PPG Signal")
plt.xlabel("Time (s)")
plt.ylabel("Amplitude")
plt.grid()
plt.legend() plt.show()
```



2. Filtered PPG signal:

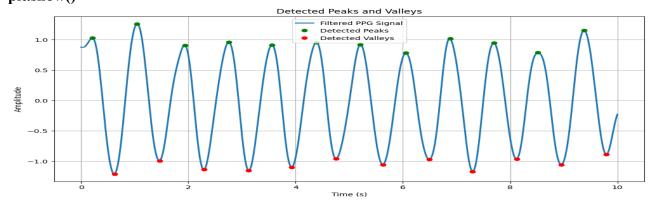
```
# Filter settings
fs = 100 # Sampling frequency in Hz
cutoff = 3 # Cutoff frequency in Hz
filtered_ppg = butter_lowpass_filter(ppg_signal, cutoff, fs)
# Plot 2: Filtered PPG Signal
plt.figure(figsize=(12, 6))
plt.plot(time, filtered_ppg, label="Filtered PPG Signal", linewidth=2)
```

```
plt.title("Filtered PPG Signal")
plt.xlabel("Time (s)")
plt.ylabel("Amplitude")
plt.grid()
plt.legend()
plt.show()
```



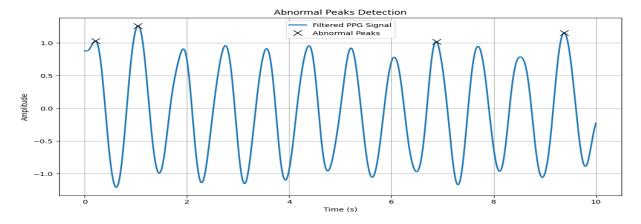
3.Peaks And Value:

```
# Peak and valley detection
peaks, _ = find_peaks(filtered_ppg, height=0.5, distance=fs//2)
valleys, _ = find_peaks(-filtered_ppg, height=0.5, distance=fs//2)
# Plot 3: Peaks and Valleys
plt.figure(figsize=(12, 6))
plt.plot(time, filtered_ppg, label="Filtered PPG Signal", linewidth=2)
plt.plot(time[peaks], filtered_ppg[peaks], "go", label="Detected Peaks")
plt.plot(time[valleys], filtered_ppg[valleys], "ro", label="Detected Valleys")
plt.title("Detected Peaks and Valleys")
plt.xlabel("Time (s)")
plt.ylabel("Amplitude")
plt.grid()
plt.legend()
plt.show()
```



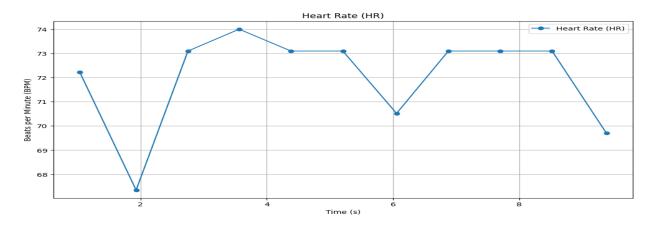
4. Abnormal Peaks:

```
#Abnormal peaks detection
peak_heights = filtered_ppg[peaks]
abnormal_peaks = peaks[peak_heights > 1] # Threshold for high spikes
# Plot 4: Abnormal Peaks
plt.figure(figsize=(12, 6))
plt.plot(time, filtered_ppg, label="Filtered PPG Signal", linewidth=2)
plt.plot(time[abnormal_peaks], filtered_ppg[abnormal_peaks], "kx", label="Abnormal
Peaks", markersize=10)
plt.title("Abnormal Peaks Detection")
plt.xlabel("Time (s)")
plt.ylabel("Amplitude")
plt.grid()
plt.legend()
plt.show()
```



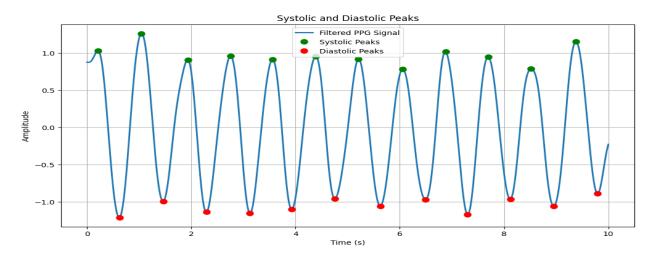
5. Heart Rate (HR)

```
# Heart Rate (HR) calculation
peak_times = time[peaks]
hr = 60 / np.diff(peak_times) # Beats per minute
# Plot 5: Heart Rate (HR)
plt.figure(figsize=(12, 6))
plt.plot(peak_times[1:], hr, label="Heart Rate (HR)", marker='o')
plt.title("Heart Rate (HR)")
plt.xlabel("Time (s)")
plt.ylabel("Beats per Minute (BPM)")
plt.grid()
plt.legend()
plot.show()
```



6. Systolic & Diastolic Peaks

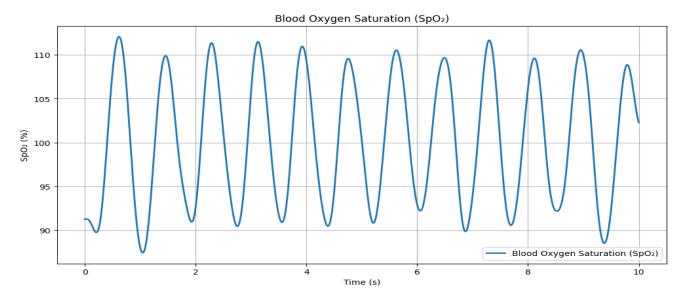
```
# Systolic & Diastolic Peaks
systolic peaks = peaks
diastolic peaks = valleys
# Plot 6: Systolic & Diastolic Peaks
plt.figure(figsize=(12, 6))
plt.plot(time, filtered ppg, label="Filtered PPG Signal", linewidth=2)
plt.plot(time[systolic peaks], filtered ppg[systolic peaks], "go", label="Systolic Peaks",
markersize=8)
plt.plot(time[diastolic peaks], filtered ppg[diastolic peaks], "ro", label="Diastolic Peaks",
markersize=8)
plt.title("Systolic and Diastolic Peaks")
plt.xlabel("Time (s)")
plt.ylabel("Amplitude")
plt.grid()
plt.legend()
plt.show()
```



7 Blood Oxygen Saturation (SpO₂).

```
# Blood Oxygen Saturation (SpO<sub>2</sub>) simulation
# Assuming SpO<sub>2</sub> is inversely proportional to the amplitude of the PPG signal spo<sub>2</sub> = 100 - (filtered_ppg * 10) # Simulated SpO<sub>2</sub> values

# Plot 7: Blood Oxygen Saturation (SpO<sub>2</sub>)
plt.figure(figsize=(12, 6))
plt.plot(time, spo<sub>2</sub>, label="Blood Oxygen Saturation (SpO<sub>2</sub>)", linewidth=2)
plt.title("Blood Oxygen Saturation (SpO<sub>2</sub>)")
plt.xlabel("Time (s)")
plt.ylabel("SpO<sub>2</sub> (%)")
plt.grid()
plt.legend()
plt.show()
```



8. Pulse Transit Time (PTT)

```
# Pulse Transit Time (PTT) simulation
# Assuming PTT is the time difference between systolic and diastolic peaks
ptt = np.zeros(len(systolic_peaks))
for i in range(len(systolic_peaks)):
    if i < len(diastolic_peaks):
        ptt[i] = time[diastolic_peaks[i]] - time[systolic_peaks[i]]

# Plot 8: Pulse Transit Time (PTT)
plt.figure(figsize=(12, 6))
plt.plot(time[systolic_peaks[:len(ptt)]], ptt, label="Pulse Transit Time (PTT)", marker='o')
plt.title("Pulse Transit Time (PTT)")</pre>
```

plt.xlabel("Time (s)")
plt.ylabel("PTT (s)")
plt.grid()
plt.legend()
plt.show()

