
Wavelet-Based Masked Multiscale Reconstruction for PPG Foundation Models

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Abstract

We introduce Masked Multiscale Reconstruction (MMR), a self-supervised pre-training framework for photoplethysmography (PPG) signals that leverages the discrete wavelet transform. MMR is pretrained on ~ 18 M unlabeled 10-second PPG segments collected from over ~ 41 K smartwatch users largely in naturalistic field settings. The pretraining task is defined to randomly mask out subsets of wavelet coefficients derived from multi-resolution decomposition of raw PPG signals and train the encoder to reconstruct them. This enables the model to capture patterns across scales from fine-grained waveform morphology to long-term temporal dynamics crucial for diverse downstream tasks. On 10 of 13 health-related tasks, MMR trained on large-scale wearable PPG data outperforms or matches state-of-the-art open-source PPG foundation models and other self-supervised baselines. An ablation study of wavelet design further underscores the value of wavelet-based representations, paving the way toward robust and generalizable PPG foundation models.

1 Introduction

Photoplethysmography (PPG) has emerged as a key sensing modality in wearables, powering applications from cuffless blood pressure estimation [Song et al., 2019], arrhythmia detection [Bashar et al., 2019] to stress monitoring [Namvari et al., 2022]. Its ubiquity in consumer devices creates an opportunity for large-scale, continuous monitoring of cardiovascular health and the development of digital biomarkers [Charlton et al., 2022a, Lee and Akamatsu, 2025]. Traditional PPG models relied on handcrafted features or small datasets [Shao et al., 2021, Han et al., 2020]; however, recent foundation models such as [Abbaspourazad et al., 2024a, Pillai et al., 2024, Saha et al., 2025] have leveraged vast amounts of unlabeled datasets, establishing a new paradigm for large-scale representation learning.

While seminal works such as [Abbaspourazad et al., 2024a] explore patient-wise contrastive learning and [Pillai et al., 2024] introduce morphology-awareness through proxies like sVRI binning and SQI regression, these approaches remain primarily rooted in the time domain. Time-only representations overlook the spectral structure of PPG, where physiological rhythms unfold across multiple frequency bands. Standard Fourier methods attempt to capture these patterns but impose a stationarity assumption, leading to poor localization and resolution in non-stationary signals [Mallat, 2002]. Explicitly modeling the spectral domain is therefore beneficial for capturing the hierarchical, multi-resolution structure of PPG—rich information that purely temporal features or proxy objectives may fail to represent [Chen et al., 2025].

Wavelet decomposition [Daubechies, 1992] provides a natural way to analyze non-stationary signals in the time–frequency domain. By adaptively trading time and frequency resolution, wavelets capture short-lived fine details at higher frequencies while preserving coarse, long-term dynamics at lower frequencies. This multi-resolution view is critical, as physiological signals carry information across

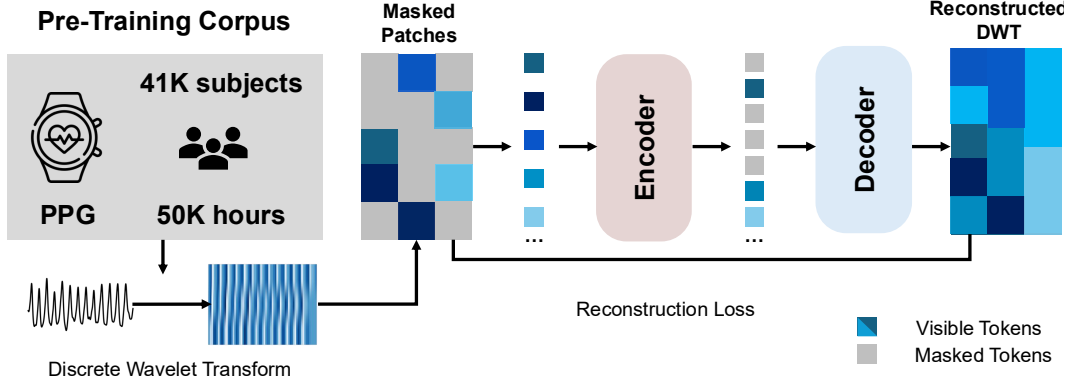


Figure 1: Masked Multiscale Reconstruction for Photoplethysmography (PPG) signals.

multiple scales, from local waveform morphology, which has been linked to vascular health [Charlton et al., 2022b], to longer-term rhythm dynamics, crucial for tasks such as heart rate variability [Namvari et al., 2022]. Motivated by these insights, we propose a masked multi-scale reconstruction framework for PPG, in which raw signals are decomposed into multiple resolution bands and a foundation model is trained to reconstruct masked coefficients across scales. To summarize, our contributions are: (i) We pretrain a large-scale *wavelet-based PPG foundation model* on $\sim 50\text{K}$ hours of PPG data with a masked multiscale reconstruction objective, enabling the model to capture rich time–frequency information across multiple scales. (ii) We demonstrate strong generalization across 13 diverse downstream tasks and provide detailed ablations that examine the impact of design choices such as wavelet family, decomposition scales, and patch size.

2 Background

Foundation models (FMs) for biosignals have recently shown strong promise, with large-scale self-supervised pretraining on ECG and PPG data from wearables demonstrating transferable representations across diverse downstream tasks [Abbaspourazad et al., 2024b, Pillai et al., 2024, Saha et al., 2025, Yang et al., 2023]. Yet most of these approaches treat signals purely in the time domain, overlooking spectral information that carries important physiological cues. A growing body of frequency-aware FMs [Zhang et al., 2022, Liu et al., 2023, Kara et al., 2024, Cheng et al., 2025, Fu and Hu, 2025, Duan et al., 2024] shows that explicitly modeling spectral content improves robustness and transferability. Wavelet analysis offers a natural way to capture both temporal localization and multi-resolution frequency structure, and has long been applied to PPG for denoising, feature extraction, and disease detection [Alafeef and Fraiwan, 2020, Singh et al., 2023, Shao et al., 2021]. Recent deep learning approaches incorporate wavelets in end-to-end pipelines, such as wavelet-informed tokenization [Masserano et al., 2024], and the combination of learnable wavelet decompositions with frequency-guided masking for biosignal foundation models [Chen et al., 2025]. Our work extends this line and introduces a multi-resolution masked pretraining framework for large-scale PPG data collected from smartwatches in real-world settings. By leveraging the fact that health tasks rely on information at multiple signal granularities, our approach provides more physiologically grounded and transferable representations (Refer Appendix A).

3 Method

PPG wavelet coefficients are patched, encoded, and reconstructed in a multi-scale framework for robust representation learning as shown in Fig. 1.

Discrete wavelet transform (DWT). The discrete wavelet transform (DWT) decomposes a signal into an approximation A_J and detail bands $\{D_j\}_{j=1}^J$ using paired low- and high-pass filters, with each level downsampled by half to provide joint time–frequency localization. At sampling rate f_s , the j -th level spans the frequency range $[f_s/2^{j+1}, f_s/2^j]$. We apply a level-4 Haar DWT (selected as

Table 1: Downstream Evaluation results across downstream tasks. Best scores are **bold**, second best underlined, with 95% CIs in gray brackets.

Classification - AUROC (\uparrow)	SimCLR	PaPaGei-P	PaPaGei-S	LSM	MMR
Hypertension - Lab	57.93 [54.2 – 61.2]	67.78 [64.7 – 71.2]	56.69 [52.4 – 59.9]	54.74 [51.0 – 58.4]	67.47 [63.9 – 70.9]
Hypertension	64.12 [57.4 – 71.1]	<u>62.10</u> [54.2 – 68.6]	61.46 [54.3 – 67.6]	54.28 [51.0 – 58.4]	60.69 [46.8 – 60.9]
PVC Detection	71.78 [71.1 – 72.5]	80.38 [79.7 – 80.9]	74.61 [73.9 – 75.3]	<u>72.29</u> [71.5 – 72.9]	82.47 [81.8 – 83.1]
HDL	41.12 [38.4 – 45.9]	49.71 [46.2 – 54.0]	33.43 [29.8 – 36.8]	<u>56.53</u> [52.9 – 59.9]	62.41 [58.1 – 66.7]
LDL	49.41 [46.5 – 52.5]	64.30 [61.1 – 67.5]	50.94 [47.7 – 54.0]	56.51 [53.1 – 59.5]	<u>61.50</u> [58.7 – 64.6]
Platelets	61.49 [59.0 – 63.9]	74.31 [72.0 – 76.8]	62.14 [59.4 – 64.7]	56.30 [53.7 – 58.8]	<u>66.18</u> [63.9 – 68.5]
Potassium	64.55 [62.2 – 66.3]	<u>81.20</u> [79.6 – 82.8]	71.53 [69.4 – 73.6]	67.34 [65.3 – 69.5]	82.85 [81.1 – 84.3]
Sodium	50.88 [46.7 – 54.8]	<u>60.71</u> [57.4 – 64.5]	50.65 [47.0 – 54.7]	49.04 [45.2 – 53.0]	71.90 [69.0 – 74.8]
Triglyceride	<u>51.51</u> [49.4 – 53.6]	44.36 [42.4 – 46.3]	50.63 [48.3 – 53.0]	55.23 [53.2 – 57.2]	47.50 [45.4 – 49.6]
Average	56.97 \pm 8.97	<u>64.98</u> \pm 11.93	56.89 \pm 11.74	58.02 \pm 6.76	66.99 \pm 10.47
Regression - MAE (\downarrow)					
Sys. BP (Lab)	<u>12.08</u> [11.6 – 12.6]	11.99 [11.5 – 12.5]	12.15 [11.6 – 12.7]	12.13 [11.6 – 12.6]	13.12 [12.6 – 13.6]
Dias. BP (Lab)	10.80 [10.4 – 11.4]	<u>10.38</u> [10.0 – 10.7]	10.78 [10.4 – 11.1]	10.72 [10.4 – 11.0]	9.66 [9.2 – 9.9]
Sys. BP	13.12 [11.9 – 14.4]	<u>12.93</u> [11.7 – 14.3]	13.04 [11.8 – 14.4]	13.12 [11.9 – 14.4]	12.80 [11.6 – 14.1]
Dias. BP	10.19 [9.3 – 11.0]	<u>10.28</u> [9.3 – 11.1]	10.30 [9.4 – 11.1]	10.37 [9.5 – 11.2]	<u>10.28</u> [9.4 – 11.1]
Average	11.55 \pm 1.14	11.40 \pm 1.12	11.57 \pm 1.09	11.59 \pm 1.10	<u>11.47</u> \pm 1.52

optimal setting based on ablations in Section 4.2) using PyWavelets [Lee et al., 2019], yielding one approximation and four detail subbands. The high-frequency subbands (2) dominated by noise and with close to zero coefficients are discarded. The remaining subbands are interpolated and arranged in order of increasing frequency to form a 2-D coefficient map of shape $[n_{\text{bands}}, \text{time}]$.

Masked Multiscale Reconstruction – MMR. We adopt a Vision Transformer (ViT) encoder–decoder within the masked autoencoder framework [He et al., 2022]. The 2-D wavelet coefficient map is divided into non-overlapping patches of size (1, 25) along the temporal axis, producing a sequence of tokens for each subband. Fixed 2-D sine–cosine positional embeddings are added to encode temporal and spectral structure. During pretraining, 75% of patches are randomly masked, and the decoder reconstructs the missing coefficients from the visible context. This Masked Multiscale Reconstruction (MMR) objective encourages the encoder to model dependencies across wavelet scales, enabling coarse bands to support fine-scale recovery and fine bands to refine coarse trends. Training minimizes mean-squared error (MSE) between reconstructed and original coefficients over the masked patches.

Experimental Setting We pretrain our encoder with unlabeled 10-second PPG segments collected from different Samsung Galaxy smartwatches, where the majority of the segments ($\sim 95\%$) are sampled at a low rate of 25 Hz (due to battery constraints in the wild). Each segment is upsampled, band-pass filtered, and z-score normalized before wavelet decomposition is applied. The dataset is split 80:20 into training and validation sets for pretraining. To evaluate generalization, the pretrained encoder is probed using random forest classifiers and regressors for a set of 13 downstream clinically motivated tasks. Classification tasks include the detection of hypertension, premature ventricular contractions (PVCs), and abnormal laboratory measures (e.g., high/low lipids, electrolytes, platelets), whereas for regression tasks, we perform the prediction of systolic and diastolic blood pressure in both field and laboratory data collection settings.

4 Results

We compare against various baselines such as SimCLR [Chen et al., 2020], masked auto encoding (time-domain) as done in LSM [Narayanswamy et al., 2024], and the open-source PaPaGei family (-S, -P variants) [Pillai et al., 2024]. We further analyze design choices through ablation studies.

4.1 Main Results

MMR performs competitively across the majority of downstream tasks, matching or surpassing strong baselines such as the PaPaGei family, LSM, and SimCLR. Across 13 classification and regression

tasks, MMR achieved the top score in 7 (PVC detection, hypertension-lab, diastolic BP, potassium, HDL, etc.) and came close in nearly all others (61.50 vs 64.30 for LDL, 10.28 vs 10.19 for Dias. BP. MAE), resulting in top-2 performance in 10/13 tasks overall. In classification, MMR achieved strong AUROC scores, including state-of-the-art performance for PVC detection and hypertension-lab. It also had the highest classification scores for abnormal lab results, such as high HDL, sodium, and potassium. Notably, MMR outperforms SimCLR and LSM (trained on the same wearable data as MMR) by up to +18 AUROC points (e.g., high potassium: 82.8 vs. 64.0) and by +10% for PVC detection, respectively. It also outperforms PaPaGei-S by margins of 10–15 AUROC points on several tasks. For regression, MMR achieved the lowest error in diastolic BP-lab, in the field setting, ranked second for diastolic BP, and led systolic BP regression. Across most tasks, PaPaGei-P remains a strong baseline, while the other variant lag significantly behind MMR. Importantly, these results were achieved using real-world wearable PPG data sampled at a low rate, compared to PaPaGei models trained on clean, high-frequency (125–500 Hz) clinical signals. Despite the lower sampling rate and higher noise inherent in field data, MMR not only competes but often surpasses clinical-data-trained models.

4.2 Ablation Studies

We ablated a 1M-sample subset, varying wavelet family, decomposition level, and patch size, and evaluated on two representative tasks (Table 2).

Wavelet Families: can exhibit distinct trade-offs in DWT. The Daubechies-4 (db4) provides stable performance across both tasks (Hypertension 63.8%, PVC 70.8%). Haar wavelet family achieves the highest PVC score (74.8%) while maintaining competitive Hypertension performance (64.1%). This can be attributed to Haar’s compact nature and sharp discontinuities, which can emphasize abrupt waveform changes that are critical for detecting ectopic beats [Yang et al., 2019]. By contrast, db4 and biorthogonal3.5 offer smoother base functions, which may miss key transients necessary for PVC detection.

Decomposition Level: governs the number of multi-resolution sub-bands/hierarchy available for analysis. The db4 wavelet attains 61.5% Hypertension accuracy at Level 4, increasing to 64.9% at Levels 7, suggesting that deeper decompositions with additional sub-bands yield more informative representations for classification [Singh et al., 2023, Attivissimo et al., 2023]. In contrast, PVC scores within the db4 family peak at Level 4 and decrease with deeper decompositions (70.0% at Level 7).

Patch size: regulates the granularity of the temporal context provided to the encoder. Smaller patches (25) achieve the best Hypertension score (64.9%), whereas a patch size of 50 offers a balanced trade-off, with reasonable Hypertension (63.4%) and strong PVC (71.1%). Large patches (100) reduce Hypertension to (62.9%), emphasizing that a longer window may average out subtle morphological patterns.

In summary, our experiments show that *different tasks benefit from distinct temporal and frequency scales*. This supports the hypothesis that multi-scale PPG information provides complementary cues, highlighting wavelet-based representations as an effective pretraining strategy for adaptive and robust physiological monitoring with PPG.

Table 2: AUROC scores for two tasks. Ablation of MMR across wavelet family, decomposition level, and patch size. Pretrained on 5% of the full dataset.

Configuration	Hypertension (avg. lab, field)	PVC
db4 – Level 4 – Patch 50	61.54	73.65
db4 – Level 6 – Patch 50	63.77	70.83
db4 – Level 7 – Patch 50	64.82	70.05
db4 – Level 6 – Patch 25	64.87	69.94
db4 – Level 6 – Patch 100	62.90	70.92
haar – Level 6 – Patch 50	64.11	74.84
bior3.5 – Level 6 – Patch 50	61.28	68.52

5 Discussion and Future Work

Pretraining on diverse smartwatch data with wavelet-based multi-scale reconstruction of PPG signals provides a strong foundation for robust physiological feature learning and downstream cardiovascular tasks. Future directions include dynamically learning or adapting to different levels of decomposition, incorporating frequency information through positional embeddings or cross-scale reconstruction where subbands are masked and reconstructed, and further analyzing the frequency–time components the model attends to for deeper interpretability. Advancing along these directions could yield richer and more generalizable PPG foundation models for real-world health applications.

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A Appendix

A.1 Extended Related Work

Self-supervised pretraining has emerged as the dominant paradigm for large-scale biosignal modeling [Lee et al., 2025]. For example, [Abbaspourazad et al., 2024b] trained foundation models on PPG and ECG from $\sim 141\text{K}$ Apple Watch users, demonstrating the value of contrastive learning at scale. In parallel, [Pillai et al., 2024] introduced PaPaGei, an open-source PPG foundation model trained on 20M unlabeled fingertip PPG segments that explicitly leverages waveform morphology, while [Saha et al., 2025] developed Pulse-PPG using 100 days of field data from 120 participants, showing improved efficiency and generalizability. Beyond single-modality PPG, multimodal biosignal foundations transfer representations across ECG, PPG, and other signals either via knowledge distillation [Abbaspourazad et al., 2024a] or unified embeddings [Yang et al., 2023]. Related work has also applied masked reconstruction on multivariate health time series, yielding strong generative and discriminative performance on tasks such as activity classification [Narayanswamy et al., 2024, Xu et al., 2025]. Together, these advances reflect a shift from task-specific models to general-purpose foundation models for biosignals.

While these foundation models highlight the value of large-scale self-supervision, most treat signals purely in the time domain. A growing body of work shows that explicitly incorporating spectral information provides a powerful inductive bias for robust and transferable representations. For instance, Time-Frequency Consistency [Zhang et al., 2022] proposed aligning time- and frequency-domain views via contrastive loss, while bioFAME [Liu et al., 2023] introduced a frequency-aware transformer encoder with multi-head spectral filters. Similarly, FreqMAE [Kara et al., 2024] leveraged temporal-shifting encoders to model spectral content in multimodal IoT data. More recent approaches, such as FAT [Cheng et al., 2025], FEI [Fu and Hu, 2025], and MF-CLR [Duan et al., 2024], further illustrate how spectral modeling can enhance time-series representation learning. These findings suggest that frequency-aware pretraining can serve as a complementary approach to large-scale training for physiological signals such as PPG.

Wavelet analysis provides a natural way to capture information at different temporal scales by decomposing signals into multi-resolution frequency bands. Earlier PPG studies applied discrete wavelet transforms (DWT) for denoising and handcrafted features, for example in respiratory rate estimation [Alafeef and Fraiwan, 2020], hypertension and diabetes detection [Singh et al., 2023], and peak stabilization pipelines [Shao et al., 2021]. More recently, deep learning models have incorporated wavelets end-to-end, such as wavelet-based tokenization for time-series foundation models [Masserano et al., 2024] and PhysioWave [Chen et al., 2025], which couples learned wavelet decompositions, frequency guided masking with Transformers for physiological signals such as ECG and EMG. Our work extends this line and introduces a multi-resolution masked pretraining framework for large-scale PPG data collected from smartwatches in real-world settings. By leveraging the fact that health tasks rely on information at multiple signal granularities, our approach provides more physiologically grounded and transferable representations.

A.2 Training Setup

We pretrain MMR using the AdamW [Loshchilov and Hutter, 2017] optimizer with a base learning rate of 1×10^{-4} , cosine decay schedule, and linear warmup over the first 10% of steps. Training is performed for $\sim 69\text{K}$ steps with a batch size of 512, weight decay of $1\text{e-}5$, and gradient clipping at 1.0, while adopting the same augmentations (i.e., time-flip, adding Gaussian noise, and stretching along the temporal axis) as LSM [Narayanswamy et al., 2024] to the PPG signal before wavelet decomposition. The backbone follows a ViT-Small configuration ($\sim 7\text{M}$ parameters) with 8 encoder blocks (hidden size 256, 4 heads, feedforward size 1024) and a lightweight decoder of 2 blocks (hidden size 192, 4 heads) used only during pretraining for reconstruction. Hyperparameter tuning was minimal, limited to a small grid search over 2–3 learning rates $\in \{1\text{e-}2, 1\text{e-}3, 1\text{e-}4\}$ and decay values $\in \{1\text{e-}3, 1\text{e-}4, 1\text{e-}5\}$, with sweeps and ablations run on a subset of the pretraining data ($\sim 1\text{M}$ data points). All experiments are conducted on 4 Tesla T4 GPUs (16GB each) with distributed data parallel (DDP) training in PyTorch [Paszke et al., 2019]. For the baselines, we use the pretrained weights for the PaPaGei family while we pretrain SimCLR and LSM on our pretraining data.

Table 3: Downstream datasets. Counts are (#positive / #negative) segments

Task	Setting	Train (pos/neg)	Test (pos/neg)
Hypertension	Lab (protocol)	746 / 3124	561 / 398
	Naturalistic (field)	542 / 373	140 / 104
PVC Detection	Wearable	14419 / 166270	4491 / 37947
<i>Laboratory Tests</i>			
HDL	Clinical reports	4117 / 3805	283 / 807
LDL	Clinical reports	2518 / 3155	779 / 613
Sodium	Clinical reports	3928 / 2700	1115 / 239
Potassium	Clinical reports	4755 / 5746	1689 / 835
Platelets	Clinical reports	3096 / 3590	1291 / 712

A.3 Dataset details

The data used in this research were collected with informed consent from all participants. Participants were informed that their wearable derived signals would be used for health-related research purposes. All data were fully de-identified prior to any analysis. We pretrain on data collected from various types of Samsung Galaxy smartwatches where PPG is sampled at different sampling frequencies (e.g., 100, 25 Hz). These datasets provide signals collected from diverse user groups and reflect real-world conditions, making them highly representative for PPG-based wearable applications. To ensure that there is no leakage of users across pretraining and test users, we follow strict user-wise splitting and create different evaluation sets settings as below:

- Hypertension-Lab: 63 users (50 train / 13 test) with protocolized data collection; segments: 746 positive / 3124 negative for training, and 561 positive / 398 negative for testing.
- Hypertension-Field: 915 train users / 244 test users; segments: 542 positive / 373 negative for training, and 140 positive / 104 negative for testing.
- PVC detection: 14,419 positive / 166,270 negative segments for training, and 4,491 positive / 37,947 negative segments for testing.
- Laboratory biomarkers (HDL, LDL, Sodium, Potassium): smaller user-sparse datasets (≈ 15 –30 users per task) with class imbalance; segment splits are provided in Table 3.

All analyses/tasks are performed at the segment level. To mitigate imbalance during probing of frozen embeddings, we downsample the majority class in the training (not in the test) split. Random forest classifiers and random forest regression models are used for probing with a 4-fold cross-validation. We also compute 95% confidence intervals via bootstrapping with 500 resampling runs similar to [Pillai et al., 2024].

Hypertension Classification We define hypertension as a binary classification task based on clinical guidelines: individuals are labeled as *Hypertensive* (label 1) if their systolic blood pressure is ≥ 130 mmHg or diastolic blood pressure is ≥ 80 mmHg, and *Normal* (label 0) otherwise. We apply buffer thresholds of ± 8 mmHg around the diagnostic cutoffs.

PVC Detection Premature Ventricular Contractions (PVCs) are early heartbeats originating in the ventricles [Cha et al., 2012]. They can indicate underlying cardiac conditions or increased risk of arrhythmias. We label high PVC burden as class 1 and low PVC burden as class 0.

Laboratory Tests For various laboratory tests (explained below as per [National Library of Medicine (US), 2020]), we adopt a binary classification scheme where high values are labeled as class 1 and class 0 otherwise.

- Sodium: Elevated sodium (hypernatremia) is linked to dehydration or adrenal gland/kidney dysfunction.
- Potassium: High potassium (hyperkalemia) may cause cardiac arrhythmias; low potassium (hypokalemia) is associated with muscle weakness, fatigue and rhythm disturbances.
- Platelets: Elevated platelet counts can signal inflammation or clotting risk.

- Low-Density Lipoprotein LDL: High LDL is a risk factor for peripheral artery disease and heart stroke.
- High-Density Lipoprotein HDL: High HDL helps lower heart disease and heart attack.
- Triglycerides: Elevated triglycerides increase cardiovascular risk and are often associated with metabolic syndrome