hr estimation

December 12, 2024

1 Heart Rate Estimation

The goal of this exercise is to estimate heart rate from PPG and acceleration signals. We signals from the PPG-DaLiA dataset (https://archive.ics.uci.edu/ml/datasets/PPG-DaLiA). It includes PPG and acceleration signals as well as reference heart rate computed from an ECG signal.

These signals were collected during various activities but we focus on two of them: sitting and walking.

First, we import all the packages we will need, define some global variables, and seed the random number generators.

```
[1]: %matplotlib widget
     import copy
     import functools
     import itertools
     import logging
     import operator
     import pathlib
     import warnings
     import IPython.display
     import joblib
     import matplotlib.pyplot as plt
     import numpy as np
     import pandas as pd
     import pytorch_lightning as pl
     import torch
     DATA_FILE = pathlib.Path('../data/ppg_dalia.pkl')
     LOG_DIRECTORY = pathlib.Path('../logs/hr_estimation')
     # Disable logging for PyTorch Lightning to avoid too long outputs.
     logging.getLogger('pytorch_lightning').setLevel(logging.ERROR)
```

```
# Seed random number generators for reproducible results.
pl.seed_everything(42)
```

Seed set to 42

[1]: 42

Then, we load the PPG and acceleration signals as well as the reference heart rate. The signals are already pre-processed with the following steps:

- Band-pass filtering between 0.4 and 4.0 Hz (24 240 bpm).
- Resampling to 25 Hz.

We also define the window length and shift length used to compute the reference heart rate.

```
[2]: FS = 25.0 # Sampling frequency of the PPG and acceleration signals in Hertz.

WINDOW_LENGTH = 8.0 # Window duration in seconds used to compute the reference_

sheart rate.

SHIFT_LENGTH = 2.0 # Shift between successive windows in seconds.

WINDOW_SIZE = round(FS * WINDOW_LENGTH)

SHIFT_SIZE = round(FS * SHIFT_LENGTH)

records = joblib.load(DATA_FILE)

subjects = set(record['subject'] for record in records)

print(f'Window length: {WINDOW_LENGTH} s (n = {WINDOW_SIZE})')

print(f'Shift length: {SHIFT_LENGTH} s (n = {SHIFT_SIZE})')

print(f'Number of records: {len(records)}')

print(f'Number of subjects: {len(subjects)}')
```

```
Window length: 8.0 s (n = 200)
Shift length: 2.0 s (n = 50)
Number of records: 29
Number of subjects: 15
```

Here are two examples of PPG and acceleration signals. One recorded when the subject is sitting and one recorded when the subject is walking.

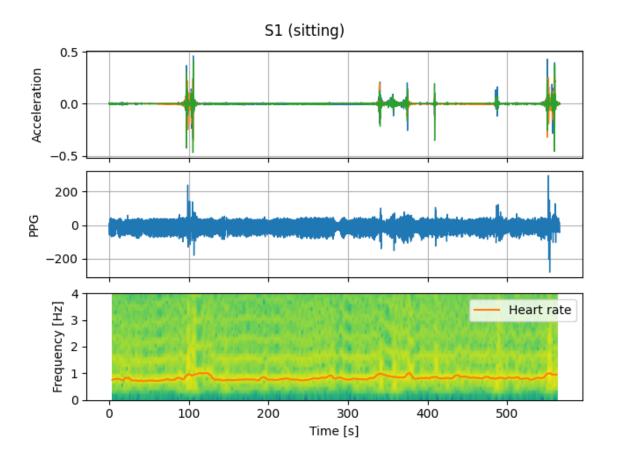
Each figure includes three plots: the tri-axis acceleration signals, the PPG signal, and a spectrogram of the PPG signal with the reference heart rate on top.

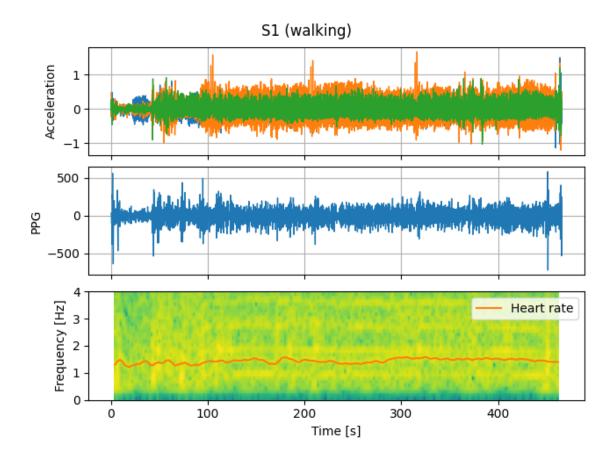
```
[3]: def plot_signals(record):
    signals = record['signals']
    hr = record['hr']

    fig, axes = plt.subplots(3, 1, sharex='all', constrained_layout=True)
    plt.suptitle(f'{record["subject"]} ({record["activity"]})')

    plt.sca(axes.flat[0])
    plt.plot(signals['time'].to_numpy(),
```

```
signals[['acc_x', 'acc_y', 'acc_z']].to_numpy(),
             linewidth=1)
   plt.grid()
   plt.ylabel('Acceleration')
   plt.sca(axes.flat[1])
   plt.plot(signals['time'].to_numpy(), signals['ppg'].to_numpy(),
             linewidth=1)
   plt.grid()
   plt.ylabel('PPG')
   plt.sca(axes.flat[2])
   plt.specgram(signals['ppg'].to_numpy(), Fs=FS, NFFT=WINDOW_SIZE,
                 noverlap=WINDOW_SIZE - SHIFT_SIZE)
   plt.plot(hr['time'].to_numpy(), hr['hr'].to_numpy() / 60.0,
             color='tab:orange', label='Heart rate')
   plt.ylim(0.0, 4.0)
   plt.xlabel('Time [s]')
   plt.ylabel('Frequency [Hz]')
   plt.legend(loc='upper right')
plot_signals(records[0])
plot_signals(records[1])
```





By zooming on the PPG signal, it is clear that walking cause a degradation in signal quality.

We will try to estimate the heart rate on sliding windows of the PPG and acceleration signals. To make things easier, we use the same window length and shift between windows as the reference heart rate.

So the next step is to extract sliding windows from all the records. We also extract the corresponding subject identifier for splitting the windows into subsets for training, validation, and testing.

In addition, we also prepare windows that include only the PPG signal (first channel).

```
[4]: def extract_windows(record):
    x = record['signals'][['ppg', 'acc_x', 'acc_y', 'acc_z']].to_numpy()
    n = x.shape[0]

windows = []
    for start in range(0, n - WINDOW_SIZE + 1, SHIFT_SIZE):
        end = start + WINDOW_SIZE
        windows.append(x[start:end].T)

windows = np.stack(windows)
    targets = record['hr']['hr'].to_numpy()
```

```
return windows, targets
def extract_all_windows(records):
   windows = []
   targets = []
   subjects = []
   activities = []
   for record in records:
        x, y = extract windows(record)
       windows.append(x)
       targets.append(y)
        subjects.extend(itertools.repeat(record['subject'], x.shape[0]))
        activities.extend(itertools.repeat(record['activity'], x.shape[0]))
   windows = np.concatenate(windows, axis=0)
   targets = np.concatenate(targets)[:, None]
    subjects = np.array(subjects)
   activities = np.array(activities)
   return windows, targets, subjects, activities
ppg_acc_windows, targets, subjects, activities = extract_all_windows(records)
ppg_windows = ppg_acc_windows[:, :1, :]
print(f'Shape of PPG and accleration windows: {ppg_acc_windows.shape}')
print(f'Shape of PPG windows: {ppg_windows.shape}')
```

```
Shape of PPG and accleration windows: (7420, 4, 200) Shape of PPG windows: (7420, 1, 200)
```

We have 7420 windows with 1 or 4 channels and that each window includes 200 samples (8 seconds at 25 Hz).

Next, we split the windows for training, validation, and testing by subjects. The test set includes 9 subjects, the validation set 3 subjects, and the test set 3 subjects.

```
[5]: def split_subjects(subjects):
    val_subjects = ('S10', 'S11', 'S12')
    test_subjects = ('S13', 'S14', 'S15')

    i_val = np.flatnonzero(np.isin(subjects, val_subjects))
    i_test = np.flatnonzero(np.isin(subjects, test_subjects))
    i_train = np.setdiff1d(np.arange(subjects.size), np.union1d(i_val, i_test))

    assert not (set(subjects[i_train]) & set(subjects[i_val]))
    assert not (set(subjects[i_train]) & set(subjects[i_test]))
```

```
assert not (set(subjects[i_val]) & set(subjects[i_test]))
assert (set(subjects[i_train]) | set(subjects[i_val]) |
set(subjects[i_test])) == set(subjects)

return i_train, i_val, i_test

i_train, i_val, i_test = split_subjects(subjects)

print(f'Subject used for training : {pd.unique(subjects[i_train])}')
print(f'Subject used for validation : {pd.unique(subjects[i_val])}')
print(f'Subject used for testing : {pd.unique(subjects[i_test])}')
```

```
Subject used for training : ['S1' 'S2' 'S3' 'S4' 'S5' 'S6' 'S7' 'S8' 'S9'] Subject used for validation : ['S10' 'S11' 'S12'] Subject used for testing : ['S13' 'S14' 'S15']
```

To make training more stable, we scale the windows such that they have approximately unit variance.

```
[6]: def compute_scaling(windows):
    sigma = np.mean(np.std(windows, axis=-1, keepdims=True), axis=0)
    return 1.0 / sigma

ppg_acc_alpha = compute_scaling(ppg_acc_windows[i_train])
ppg_acc_windows *= ppg_acc_alpha
ppg_alpha = compute_scaling(ppg_windows[i_train])
ppg_windows *= ppg_alpha
```

Question 1

The windows are scaled but not centered. Why?

Answer

We can see in the cell below that the mean of the signal is already zero. Therefore there is no need to center the signal. When filtering (Band-Pass) the signal, the offset (zero frequency) is removed and the signal is centered.

```
[24]: ppg_windows.mean(axis=(0, 2))
```

```
[24]: array([-6.8200391e-05])
```

Now we define the convolutional neural network (CNN) we will use to estimate heart rate. It is composed of convolutional layers to extract features and dense layers to estimate heart rate. The convolutional layers can optionally include batch normalization and the dense layers dropout.

```
[7]: class CnnModel(torch.nn.Module):

def __init__(self,
```

```
input_shape,
             output_shape,
             n_convolutional_layers=1,
             kernel_size=5,
             n_initial_channels=16,
             use_normalization=False,
             n_dense_layers=1,
             n_units=128,
             dropout=0.0):
    super().__init__()
    self.input_shape = input_shape
    self.output_shape = output_shape
    self.n_convolutional_layers = n_convolutional_layers
    self.kernel_size = kernel_size
    self.n_initial_channels = n_initial_channels
    self.use_normalization = use_normalization
    self.n_dense_layers = n_dense_layers
    self.n_units = n_units
    self.dropout = dropout
    self.layers = self._build_layers()
@property
def input_size(self):
    return functools.reduce(operator.mul, self.input_shape)
@property
def output_size(self):
    return functools.reduce(operator.mul, self.output_shape)
def _build_layers(self):
    layers = self._build_convolutional_layers()
    layers.extend(self._build_dense_layers())
    return torch.nn.Sequential(*layers)
def _build_convolutional_layers(self):
    layers = []
    n_output_channels = self.input_shape[0]
    for i in range(self.n convolutional layers):
        n_input_channels = n_output_channels
        n_output_channels = self.n_initial_channels * 2 ** i
        layers.append(torch.nn.Conv1d(
            in_channels=n_input_channels,
            out_channels=n_output_channels,
            kernel_size=self.kernel_size,
            padding='same',
        ))
```

```
if self.use_normalization:
            layers.append(torch.nn.BatchNorm1d(n_output_channels))
        layers.append(torch.nn.ReLU())
        if i < self.n_convolutional_layers - 1:</pre>
            layers.append(torch.nn.MaxPool1d(kernel_size=2))
        else:
            layers.append(torch.nn.AdaptiveAvgPool1d(1))
    layers.append(torch.nn.Flatten())
    return layers
def _build_dense_layers(self):
    sizes = [self.n_initial_channels
             * 2 ** (self.n_convolutional_layers - 1)]
    sizes.extend(itertools.repeat(self.n_units, self.n_dense_layers - 1))
    sizes.append(self.output_size)
    layers = []
    for i in range(self.n_dense_layers - 1):
        layers.append(torch.nn.Linear(sizes[i], sizes[i + 1]))
        layers.append(torch.nn.ReLU())
        if 0.0 < self.dropout < 1.0:</pre>
            layers.append(torch.nn.Dropout(self.dropout))
    layers.append(torch.nn.Linear(sizes[-2], sizes[-1]))
    return layers
def forward(self, x):
    return self.layers(x)
```

We also define a class to specify how the model should be trained and evaluated and a few utility functions. It is also here that we select the mean squared error (MSE) as the loss function to optimize the parameters. We also compute the mean absolute error (MAE) as an additional metric to monitor training.

```
[8]: class Regressor(pl.LightningModule):

    def __init__(self, config):
        super().__init__()
        self.save_hyperparameters()
        self.config = config
        self.model = CnnModel(**self.config['model'])
        self.example_input_array = torch.zeros((1,) + self.model.input_shape)

    def configure_optimizers(self):
        return torch.optim.Adam(self.parameters(), **self.config['optimizer'])
```

```
def forward(self, x):
        return self.model(x)
    def training_step(self, batch, batch_idx):
        return self._run_step(batch, 'train')
    def validation_step(self, batch, batch_idx):
        self._run_step(batch, 'val')
    def test_step(self, batch, batch_idx):
        self._run_step(batch, 'test')
    def predict_step(self, batch, batch_idx, dataloader_idx=0):
        x, y = batch
        return self.model(x)
    def _run_step(self, batch, subset):
        x, y = batch
        z = self.model(x)
        mse = torch.nn.functional.mse_loss(z, y)
        mae = torch.nn.functional.l1_loss(z, y)
        self.log dict({
            f'{subset}_mse': mse,
            f'{subset} mae': mae,
        }, on_step=False, on_epoch=True, prog_bar=True)
        return mse
def build_loader(*tensors, batch_size=100, shuffle=False, n_workers=0):
    dataset = torch.utils.data.TensorDataset(*map(torch.Tensor, tensors))
    return torch.utils.data.DataLoader(
        dataset=dataset,
        batch_size=batch_size,
        shuffle=shuffle,
        num_workers=n_workers,
    )
def train_model(config, windows, targets, i_train, i_val, n_epochs, name):
    train_loader = build_loader(windows[i_train], targets[i_train],__
 ⇒shuffle=True)
    val_loader = build_loader(windows[i_val], targets[i_val])
    regressor = Regressor(config)
    print(pl.utilities.model_summary.ModelSummary(regressor, max_depth=-1))
    with warnings.catch_warnings():
        warnings.simplefilter('ignore')
```

```
trainer = pl.Trainer(
            default_root_dir=LOG_DIRECTORY,
            logger=pl.loggers.TensorBoardLogger(LOG_DIRECTORY, name),
            enable_model_summary=False,
            max_epochs=n_epochs,
        trainer.fit(regressor, train_loader, val_loader)
    return regressor
def compute_metrics(targets, predictions):
    targets = targets.ravel()
    predictions = predictions.ravel()
    return {
        'count': targets.size,
        'mse': np.mean((targets - predictions) ** 2),
        'mae': np.mean(np.abs(targets - predictions)),
    }
def evaluate_model(model, windows, targets, i_train, i_val, i_test):
    loader = build_loader(windows, targets)
    with warnings.catch_warnings():
        warnings.simplefilter('ignore')
        trainer = pl.Trainer(
            default_root_dir=LOG_DIRECTORY,
            logger=False,
            enable_progress_bar=False,
            enable_model_summary=False,
        predictions = trainer.predict(model, loader)
    predictions = np.vstack([p.numpy() for p in predictions])
    metrics = []
    for subset, indices in (('train', i_train), ('val', i_val), ('test', _
 →i_test)):
        metrics.append({
            'subset': subset,
            **compute_metrics(targets[indices], predictions[indices]),
        })
    return pd.DataFrame(metrics)
```

We also start TensorBoard to monitor training.

If you prefer to view TensorBoard in a separate window, you can open http://localhost:6006/ in your web browser.

```
[28]: %reload_ext tensorboard %tensorboard --logdir ../logs/hr_estimation --port 6007
```

<IPython.core.display.HTML object>

We are finally to define the first model configuration we will train. It includes the following layers:

- Input windows (input_size = 200, input channels = 1 or 4)
- Convolutional layer (kernel size = 5, output size = 200, output channels = 16)
- ReLU activation
- Max pooling (output size = 100, output channels = 16)
- Convolutional layer (kernel size = 5, output size = 100, output channels = 32)
- ReLU activation
- Max pooling (output size = 50, output channels = 32)
- Convolutional layer (kernel size = 5, output size = 50, output channels = 64)
- ReLU activation
- Max pooling (output size = 25, output channels = 64)
- Convolutional layer (kernel size = 5, output size = 25, output channels = 128)
- ReLU activation
- Global averaging pooling (output size = 128)
- Dense layer (output size = 128)
- ReLU activation
- Dense layer (output size = 128)
- ReLU activation
- Dense layer (output size = 1)

We use the same configuration for PPG only and PPG and acceleration windows (only the input shape changes).

We also define the number of epochs to use for training.

```
[29]: | ppg_cnn_config = {
          'model': {
              'input_shape': ppg_windows.shape[1:],
              'output_shape': targets.shape[1:],
              'n_convolutional_layers': 4,
              'kernel_size': 5,
              'n_initial_channels': 16,
              'use_normalization': False,
              'n dense layers': 3,
              'n_units': 128,
              'dropout': 0.0,
          },
          'optimizer': {
              'lr': 0.0001,
          },
      }
      ppg_acc_cnn_config = copy.deepcopy(ppg_cnn_config)
      ppg_acc_cnn_config['model']['input_shape'] = ppg_acc_windows.shape[1:]
```

```
n_{epochs} = 30
      print('PPG CNN config')
      IPython.display.display(ppg_cnn_config)
      print()
      print('PPG ACC CNN config')
      IPython.display.display(ppg_acc_cnn_config)
     PPG CNN config
     {'model': {'input_shape': (1, 200),
       'output_shape': (1,),
       'n_convolutional_layers': 4,
       'kernel_size': 5,
       'n_initial_channels': 16,
       'use_normalization': False,
       'n_dense_layers': 3,
       'n_units': 128,
       'dropout': 0.0},
      'optimizer': {'lr': 0.0001}}
     PPG ACC CNN config
     {'model': {'input_shape': (4, 200),
       'output_shape': (1,),
       'n_convolutional_layers': 4,
       'kernel_size': 5,
       'n_initial_channels': 16,
       'use_normalization': False,
       'n_dense_layers': 3,
       'n_units': 128,
       'dropout': 0.0},
      'optimizer': {'lr': 0.0001}}
     We are ready to train the first two models with these configurations.
[30]: ppg_cnn = train_model(
          config=ppg_cnn_config,
          windows=ppg_windows,
          targets=targets,
          i_train=i_train,
          i_val=i_val,
          n_epochs=n_epochs,
          name='ppg_cnn',
      )
      ppg_acc_cnn = train_model(
          config=ppg_acc_cnn_config,
```

sizes	· 		· 		· 						· 	
0 model 1]		CnnModel		87.2 K	١	train	l	[1,	1, 2	00]		[1,
<pre>1 model.layers 1]</pre>		Sequential		87.2 K	١	train	l	[1,	1, 2	00]		[1,
2 model.layers.0 16, 200]	I	Conv1d	١	96	١	train	l	[1,	1, 2	[00		[1,
•	I	ReLU		0	1	train	l	[1,	16,	200]	١	[1,
4 model.layers.2 16, 100]	I	MaxPool1d	١	0	١	train	l	[1,	16,	200]		[1,
5 model.layers.3 32, 100]	I	Conv1d	١	2.6 K	١	train	l	[1,	16,	100]		[1,
6 model.layers.4 32, 100]	1	ReLU	1	0	١	train	l	[1,	32,	100]	١	[1,
	I	MaxPool1d	١	0	١	train	I	[1,	32,	100]		[1,
8 model.layers.6 64, 50]	I	Conv1d	١	10.3 K	١	train	I	[1,	32,	50]		[1,
	I	ReLU	١	0	١	train	I	[1,	64,	50]	١	[1,
10 model.layers.8 64, 25]	I	MaxPool1d	١	0	١	train	I	[1,	64,	50]	I	[1,
	I	Conv1d	١	41.1 K	١	train	l	[1,	64,	25]		[1,
12 model.layers.10 128, 25]	I	ReLU		0		train	l	[1,	128,	25]		[1,
13 model.layers.11 128, 1]	I	AdaptiveAvgPool1d	١	0	I	train	l	[1,	128,	25]	I	[1,
14 model.layers.12 128]	I	Flatten	١	0	١	train	l	[1,	128,	1]		[1,
15 model.layers.13 128]	I	Linear	١	16.5 K	I	train	I	[1,	128]		l	[1,
16 model.layers.14 128]	I	ReLU	I	0	I	train	I	[1,	128]		I	[1,
17 model.layers.15 128]	I	Linear	١	16.5 K	١	train	I	[1,	128]		I	[1,

```
18 | model.layers.16 | ReLU | 0 | train | [1, 128] | [1,
1287
19 | model.layers.17 | Linear | 129 | train | [1, 128] | [1,
17
______
87.2 K
        Trainable params
        Non-trainable params
87.2 K Total params
0.349
        Total estimated model params size (MB)
        Modules in train mode
20
        Modules in eval mode
Sanity Checking: |
                        | 0/? [00:00<?, ?it/s]
Training: | | 0/? [00:00<?, ?it/s]
Validation: | 0/? [00:00<?, ?it/s]
Validation: |
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| 0/? [00:00<?, ?it/s]

Validation: |

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Validation: |
                | 0/? [00:00<?, ?it/s]
                        | Params | Mode | In sizes | Out
  Name
              | Type
______
0 | model
          1]
1 | model.layers | Sequential | 87.5 K | train | [1, 4, 200] | [1,
17
2 | model.layers.0 | Conv1d | 336 | train | [1, 4, 200] | [1,
16, 200]
                      | 0 | train | [1, 16, 200] | [1,
3 | model.layers.1 | ReLU
16, 200]
4 | model.layers.2 | MaxPool1d | 0 | train | [1, 16, 200] | [1,
16, 100]
5 | model.layers.3 | Conv1d | 2.6 K | train | [1, 16, 100] | [1,
32, 100]
6 | model.layers.4 | ReLU | 0 | train | [1, 32, 100] | [1,
32, 100]
7 | model.layers.5 | MaxPool1d | 0 | train | [1, 32, 100] | [1,
32, 50]
8 | model.layers.6 | Conv1d | 10.3 K | train | [1, 32, 50] | [1,
64, 50]
9 | model.layers.7 | ReLU | 0 | train | [1, 64, 50] | [1,
64, 50]
10 | model.layers.8 | MaxPool1d | 0 | train | [1, 64, 50] | [1,
64, 25]
11 | model.layers.9 | Conv1d | 41.1 K | train | [1, 64, 25] | [1,
128, 25]
                      12 | model.layers.10 | ReLU
128, 25]
13 | model.layers.11 | AdaptiveAvgPool1d | 0 | train | [1, 128, 25] | [1,
14 | model.layers.12 | Flatten | 0 | train | [1, 128, 1] | [1,
15 | model.layers.13 | Linear | 16.5 K | train | [1, 128] | [1,
128]
```

```
16 | model.layers.14 | ReLU | 0 | train | [1, 128] | [1,
1287
17 | model.layers.15 | Linear | 16.5 K | train | [1, 128]
                                                              | [1,
128]
                           | 0
18 | model.layers.16 | ReLU
                                          | train | [1, 128]
                                                              Ι [1.
1287
19 | model.layers.17 | Linear | 129 | train | [1, 128]
                                                              | [1,
17
______
87.5 K
        Trainable params
        Non-trainable params
87.5 K
        Total params
0.350
        Total estimated model params size (MB)
        Modules in train mode
20
        Modules in eval mode
0
Sanity Checking: |
                      | 0/? [00:00<?, ?it/s]
Training: | | 0/? [00:00<?, ?it/s]
Validation: | | 0/? [00:00<?, ?it/s]
Validation: | 0/? [00:00<?, ?it/s]
Validation: | 0/? [00:00<?, ?it/s]
Validation: | | 0/? [00:00<?, ?it/s]
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Validation: |
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                  | 0/? [00:00<?, ?it/s]
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Validation: |
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Validation: |
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Validation: |
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                        | 0/? [00:00<?, ?it/s]
Validation: |
                        | 0/? [00:00<?, ?it/s]
Validation: |
Validation: |
                       | 0/? [00:00<?, ?it/s]
                        | 0/? [00:00<?, ?it/s]
Validation: |
```

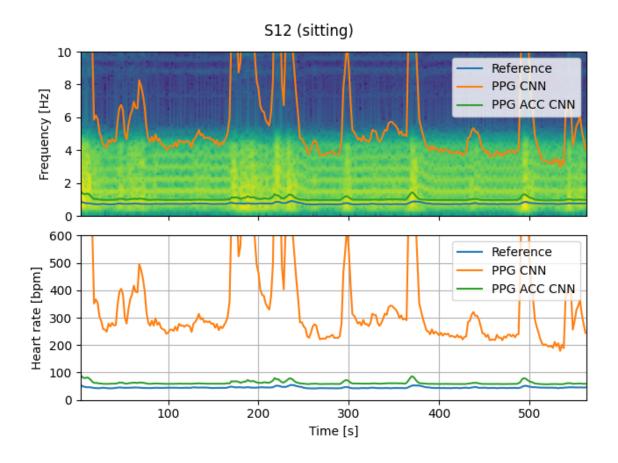
Based on metrics shown in TensorBoard, does using the acceleration signals in addition to the PPG signal help to improve performance?

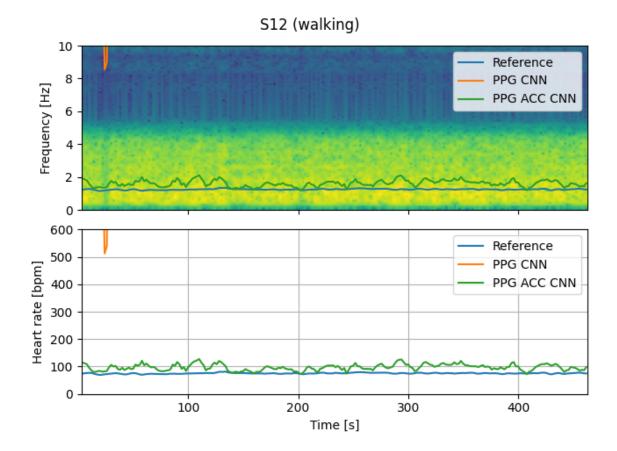
Answer Yes, for the acceleration model, we found a RMSE of 278.6604 for the validation set, while 534.7772 for the normal one. This shows that using the acceleration signals indeed improve the model performances. It can also be seen in the plot below that the estimation is more accurate for the predictions with acceleration signals.

We plot the heart rate predicted by these two models for two records from the validation set: one where the subject is sitting and on where the subject is walking.

```
[12]: def apply model(model, record, alpha):
          windows, targets = extract_windows(record)
          if alpha.shape[0] == 1:
              windows = windows[:, :1, :]
          windows *= alpha
          loader = build_loader(windows, targets)
          with warnings.catch_warnings():
              warnings.simplefilter('ignore')
              trainer = pl.Trainer(
                  default_root_dir=LOG_DIRECTORY,
                  logger=False,
                  enable progress bar=False,
                  enable_model_summary=False,
              )
              predictions = trainer.predict(model, loader)
          predictions = np.vstack([p.numpy() for p in predictions])
          return targets.ravel(), predictions.ravel()
```

```
def plot_results(record, models, limits=(0.0, 600.0)):
   predictions = {}
   for name, (model, alpha) in models.items():
        _, predictions[name] = apply_model(model, record, alpha)
   signals = record['signals']
   hr = record['hr']
   limits = np.array(limits)
   fig, axes = plt.subplots(2, 1, sharex='all', constrained_layout=True)
   plt.suptitle(f'{record["subject"]} ({record["activity"]})')
   plt.sca(axes.flat[0])
   plt.specgram(signals['ppg'].to_numpy(), Fs=FS, NFFT=WINDOW_SIZE,
                 noverlap=WINDOW_SIZE - SHIFT_SIZE)
   plt.plot(hr['time'].to_numpy(), hr['hr'].to_numpy() / 60.0,
             label='Reference')
   for name, prediction in predictions.items():
       plt.plot(hr['time'].to_numpy(), prediction / 60.0, label=name)
   plt.ylim(limits / 60.0)
   plt.ylabel('Frequency [Hz]')
   plt.legend(loc='upper right')
   plt.sca(axes.flat[1])
   plt.plot(hr['time'].to_numpy(), hr['hr'].to_numpy(), label='Reference')
   for name, prediction in predictions.items():
       plt.plot(hr['time'].to_numpy(), prediction, label=name)
   plt.ylim(limits)
   plt.grid()
   plt.xlabel('Time [s]')
   plt.ylabel('Heart rate [bpm]')
   plt.legend(loc='upper right')
models = {
    'PPG CNN': (ppg_cnn, ppg_alpha),
    'PPG ACC CNN': (ppg_acc_cnn, ppg_acc_alpha),
}
plot_results(records[21], models)
plot_results(records[22], models)
```





What can you say about the predictions computed with the two models?

Answer As already mentioned in Question 2, the prediction with the model including acceleration signals is way more accurate. We can see in Figure 3 that this prediction sticks more to the reality, while the PPG without acceleration does not make sense at all.

Next, we modify the configuration of the model using both PPG and acceleration to add batch normalization after each convolution layer. We do not do the same for the model using only PPG since it does not work at all.

```
[31]: ppg_acc_norm_cnn_config = copy.deepcopy(ppg_acc_cnn_config)
    ppg_acc_norm_cnn_config['model']['use_normalization'] = True

    print('PPG ACC norm CNN config')
    IPython.display.display(ppg_acc_norm_cnn_config)

PPG ACC norm CNN config

{'model': {'input_shape': (4, 200),
    'output_shape': (1,),
    'n_convolutional_layers': 4,
    'kernel size': 5,
```

```
'use_normalization': True,
       'n_dense_layers': 3,
       'n_units': 128,
       'dropout': 0.0},
      'optimizer': {'lr': 0.0001}}
     We train a model with this new configuration.
[32]: ppg_acc_norm_cnn = train_model(
         config=ppg_acc_norm_cnn_config,
         windows=ppg_acc_windows,
         targets=targets,
         i_train=i_train,
         i_val=i_val,
         n_epochs=n_epochs,
         name='ppg_acc_norm_cnn',
     )
                                             | Params | Mode | In sizes
        | Name
                         | Type
                                                                            | Out
     sizes
                                           | 88.0 K | train | [1, 4, 200] | [1,
     0 | model
                         CnnModel
     17
     1 | model.layers | Sequential | 88.0 K | train | [1, 4, 200] | [1,
     1]
     2 | model.layers.0 | Conv1d
                                             | 336
                                                      | train | [1, 4, 200] | [1,
     16, 200]
     3 | model.layers.1 | BatchNorm1d
                                                      | train | [1, 16, 200] | [1,
                                       | 32
     16, 200]
     4 | model.layers.2 | ReLU
                                                      | train | [1, 16, 200] | [1,
                                             1 0
     16, 200]
     5 | model.layers.3 | MaxPool1d
                                             1 0
                                                      | train | [1, 16, 200] | [1,
     16, 100]
                                             | 2.6 K | train | [1, 16, 100] | [1,
     6 | model.layers.4 | Conv1d
     32, 100]
     7 | model.layers.5 | BatchNorm1d
                                             | 64
                                                      | train | [1, 32, 100] | [1,
     32, 100]
     8 | model.layers.6 | ReLU
                                                      | train | [1, 32, 100] | [1,
                                             | 0
     32, 100]
     9 | model.layers.7 | MaxPool1d
                                             1 0
                                                      | train | [1, 32, 100] | [1,
     32, 50]
     10 | model.layers.8 | Conv1d
                                             | 10.3 K | train | [1, 32, 50] | [1,
     64, 50]
     11 | model.layers.9 | BatchNorm1d
                                           | 128
                                                      | train | [1, 64, 50] | [1,
     64, 50]
                                                      | train | [1, 64, 50] | [1,
     12 | model.layers.10 | ReLU
                                             10
     64, 50]
```

'n_initial_channels': 16,

```
13 | model.layers.11 | MaxPool1d | 0 | train | [1, 64, 50] | [1,
64, 25]
14 | model.layers.12 | Conv1d | 41.1 K | train | [1, 64, 25] | [1,
128, 25]
15 | model.layers.13 | BatchNorm1d | 256 | train | [1, 128, 25] | [1,
128, 25]
16 | model.layers.14 | ReLU | 0 | train | [1, 128, 25] | [1,
128, 25]
17 | model.layers.15 | AdaptiveAvgPool1d | 0 | train | [1, 128, 25] | [1,
128, 1]
18 | model.layers.16 | Flatten | 0 | train | [1, 128, 1] | [1,
128]
19 | model.layers.17 | Linear
                                | 16.5 K | train | [1, 128] | [1,
128]
                                     | train | [1, 128] | [1,
20 | model.layers.18 | ReLU | 0
1287
21 | model.layers.19 | Linear | 16.5 K | train | [1, 128] | [1,
128]
22 | model.layers.20 | ReLU
                         1287
23 | model.layers.21 | Linear | 129 | train | [1, 128] | [1,
1]
_____
88.0 K Trainable params
0 Non-trainable params
88.0 K Total params
0.352 Total estimated model params size (MB)
24 Modules in train mode
       Modules in eval mode
Sanity Checking: | 0/? [00:00<?, ?it/s]
Training: | | 0/? [00:00<?, ?it/s]
Validation: | 0/? [00:00<?, ?it/s]
Validation: | 0/? [00:00<?, ?it/s]
Validation: | | 0/? [00:00<?, ?it/s]
Validation: | 0/? [00:00<?, ?it/s]
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Validation: |
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Validation: |
Validation: |
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Validation: |
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Validation: |
```

| 0/? [00:00<?, ?it/s]

Validation: |

```
Validation: |
                        | 0/? [00:00<?, ?it/s]
                        | 0/? [00:00<?, ?it/s]
Validation: |
Validation: |
                        | 0/? [00:00<?, ?it/s]
                        | 0/? [00:00<?, ?it/s]
Validation: |
Validation: |
                        | 0/? [00:00<?, ?it/s]
Validation: |
                        | 0/? [00:00<?, ?it/s]
```

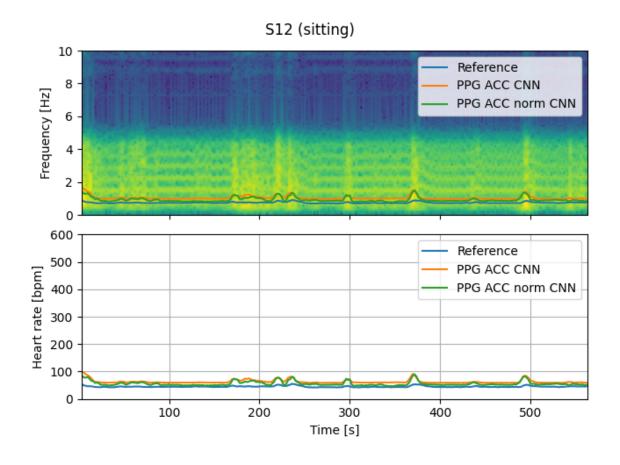
What is the effect of batch normalization on the training procedure and on the performance metrics?

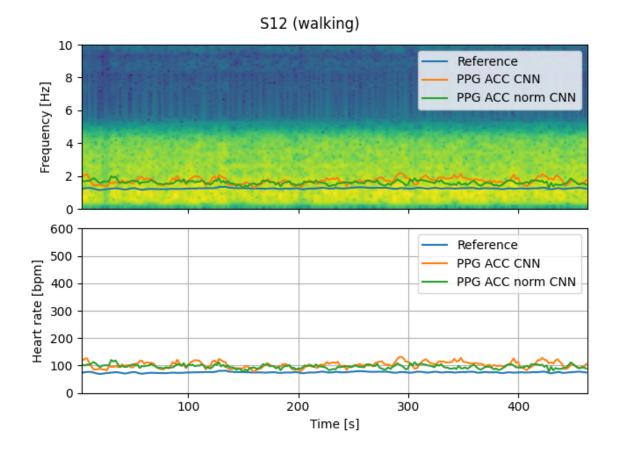
Answer Using batch normalization helps the training loss to converge faster as it reduces the variance inside each batch. Furthermore, inspecting training and validation RMSE show that BN helps achieving lower values.

We plot the predicted heart rate with respect to the reference for these new models (and we drop the one that did not work at all).

```
[33]: models = {
    'PPG ACC CNN': (ppg_acc_cnn, ppg_acc_alpha),
    'PPG ACC norm CNN': (ppg_acc_norm_cnn, ppg_acc_alpha),
}

plot_results(records[21], models)
plot_results(records[22], models)
```





Visually, is there a large difference between the CNN models with and without batch normalization on these examples?

Anser No, visually, no clear difference can be observed between the two predictions.

Finally, we try another configuration where we add dropout after each dense layer (except the last one which is the output layer).

```
[35]: ppg_acc_norm_dropout_cnn_config = copy.deepcopy(ppg_acc_norm_cnn_config)
    ppg_acc_norm_dropout_cnn_config['model']['dropout'] = 0.5

    print('PPG ACC norm dropout CNN config')
    IPython.display.display(ppg_acc_norm_dropout_cnn_config)

PPG ACC norm dropout CNN config

{'model': {'input_shape': (4, 200),
    'output_shape': (1,),
    'n_convolutional_layers': 4,
    'kernel_size': 5,
    'n_initial_channels': 16,
    'use_normalization': True,
```

```
'dropout': 0.5},
     'optimizer': {'lr': 0.0001}}
    And we train a new model with this configuration.
[36]: ppg_acc_norm_dropout_cnn = train_model(
        config=ppg_acc_norm_dropout_cnn_config,
        windows=ppg_acc_windows,
        targets=targets,
        i_train=i_train,
        i_val=i_val,
        n_epochs=n_epochs,
        name='ppg_acc_norm_dropout_cnn',
       | Name
                       | Type
                                        | Params | Mode | In sizes
                                                                    | Out
                     0 | model
    1]
    1 | model.layers | Sequential | 88.0 K | train | [1, 4, 200] | [1,
    2 | model.layers.0 | Conv1d
                                  | 336
                                                | train | [1, 4, 200] | [1,
    16, 200]
    3 | model.layers.1 | BatchNorm1d | 32
                                              | train | [1, 16, 200] | [1,
    16, 200]
    4 | model.layers.2 | ReLU
                               1 0
                                               | train | [1, 16, 200] | [1,
    16, 200]
    5 | model.layers.3 | MaxPool1d | 0
                                                | train | [1, 16, 200] | [1,
    16, 100]
    6 | model.layers.4 | Conv1d | 2.6 K | train | [1, 16, 100] | [1,
    32, 100]
    7 | model.layers.5 | BatchNorm1d | 64
                                                | train | [1, 32, 100] | [1,
    32, 100]
    8 | model.layers.6 | ReLU
                               | 0
                                                | train | [1, 32, 100] | [1,
    32, 100]
    9 | model.layers.7 | MaxPool1d | 0
                                                | train | [1, 32, 100] | [1,
    32, 50]
    10 | model.layers.8 | Conv1d
                                      | 10.3 K | train | [1, 32, 50] | [1,
    64, 50]
    11 | model.layers.9 | BatchNorm1d | 128
                                                | train | [1, 64, 50] | [1,
    64, 50]
    12 | model.layers.10 | ReLU
                                       1 0
                                                | train | [1, 64, 50] | [1,
```

'n_dense_layers': 3,
'n_units': 128,

64, 50]

64, 25]

13 | model.layers.11 | MaxPool1d | 0

| train | [1, 64, 50] | [1,

```
14 | model.layers.12 | Conv1d | 41.1 K | train | [1, 64, 25] | [1,
128, 25]
15 | model.layers.13 | BatchNorm1d | 256 | train | [1, 128, 25] | [1,
128, 25]
16 | model.layers.14 | ReLU | 0 | train | [1, 128, 25] | [1,
128, 25]
17 | model.layers.15 | AdaptiveAvgPool1d | 0 | train | [1, 128, 25] | [1,
128, 1]
18 | model.layers.16 | Flatten | 0 | train | [1, 128, 1] | [1,
1287
19 | model.layers.17 | Linear | 16.5 K | train | [1, 128] | [1,
128]
20 | model.layers.18 | ReLU
                               128]
21 | model.layers.19 | Dropout | 0 | train | [1, 128] | [1,
1287
22 | model.layers.20 | Linear
                               | 16.5 K | train | [1, 128] | [1,
128]
23 | model.layers.21 | ReLU
                        1287
24 | model.layers.22 | Dropout | 0 | train | [1, 128] | [1,
128]
25 | model.layers.23 | Linear | 129 | train | [1, 128] | [1,
______
88.0 K Trainable params
0 Non-trainable params
88.0 K
       Total params
0.352 Total estimated model params size (MB)
26
       Modules in train mode
       Modules in eval mode
Sanity Checking: |
                     | 0/? [00:00<?, ?it/s]
Training: | | 0/? [00:00<?, ?it/s]
                 | 0/? [00:00<?, ?it/s]
Validation: |
Validation: |
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Validation: |
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              | 0/? [00:00<?, ?it/s]
Validation: |
                 | 0/? [00:00<?, ?it/s]
```

```
Validation: |
                        | 0/? [00:00<?, ?it/s]
                        | 0/? [00:00<?, ?it/s]
Validation: |
Validation: |
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                        | 0/? [00:00<?, ?it/s]
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Validation: |
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Validation: |
                        | 0/? [00:00<?, ?it/s]
```

Does using dropout help to reduce overfitting compare to the same model without dropout?

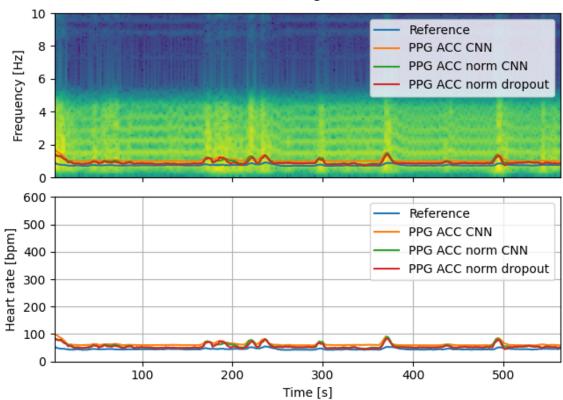
Answer Yes, applying dropout reduces overfitting. This can be observed since MSE and MAE are slightly higher for the training set when applying Dropout. This can be explained since dropout randomly selects nodes at each steps, preventing nodes to be too specific to the data. Nevertheless, this difference remains small.

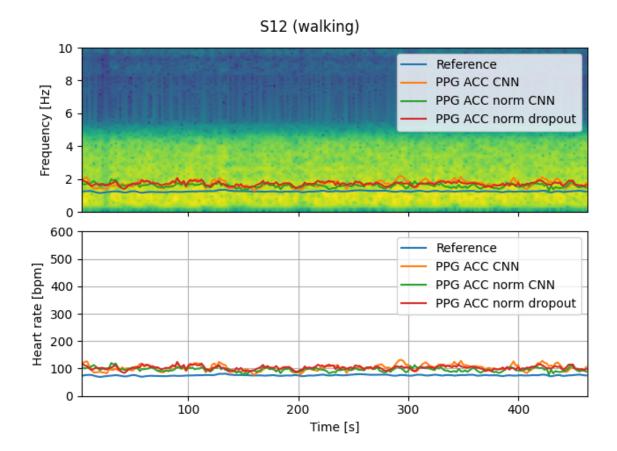
Again, we plot the predicted heart rate for two records from the validation set.

```
[37]: models = {
    'PPG ACC CNN': (ppg_acc_cnn, ppg_acc_alpha),
    'PPG ACC norm CNN': (ppg_acc_norm_cnn, ppg_acc_alpha),
    'PPG ACC norm dropout': (ppg_acc_norm_dropout_cnn, ppg_acc_alpha),
}
```

plot_results(records[21], models)
plot_results(records[22], models)

S12 (sitting)





Finally, we compute performance metrics (MSE and MAE) for all models on the subsets for training, validation, and testing.

```
[19]: models = {
          'PPG CNN': ppg_cnn,
          'PPG ACC CNN': ppg_acc_cnn,
          'PPG ACC norm CNN': ppg_acc_norm_cnn,
          'PPG ACC norm dropout': ppg_acc_norm_dropout_cnn,
      }
      metrics = []
      for name, model in models.items():
          if 'ACC' in name:
              windows = ppg_acc_windows
          else:
              windows = ppg_windows
          df = evaluate_model(model, windows, targets, i_train, i_val, i_test)
          df.insert(0, 'model', name)
          metrics.append(df)
      metrics = pd.concat(metrics, axis=0, ignore_index=True)
      metrics = metrics.set_index(['model', 'subset'])
```

	count		mse			\
subset	train val	test	train	val	test	
model						
PPG CNN	4360 1530	1530 253	.105415	519.009044	199.510891	
PPG ACC CNN	4360 1530	1530 127	.893587	263.449593	198.234142	
PPG ACC norm CNN	4360 1530	1530 27	.097020	230.597111	71.887220	
PPG ACC norm dropout	4360 1530	1530 30	.607524	240.611002	60.324805	
	mae					
subset	train	val	tes	st		
model						
PPG CNN	12.369010	19.012989	12.44398	80		
PPG ACC CNN	8.779760	13.099361	12.04571	.0		
PPG ACC norm CNN	3.844666	11.438116	6.17752	27		
PPG ACC norm dropout	4.044759	11.505437	5.49847	'3		

What is the best model in terms of MSE and MAE? What can you say about batch normalizaton and dropout?

Answer Based on the table, the best mse, as well as the mae on the test set is achieved for the model with batch normalization and dropout. This highlights the theoretical aspects of both methods, effectively preventing overfitting. But regarding the validation set, the model without dropout seems to perform better. With this difference, it is therefore difficult to ensure that dropout really helps in this case.