The autoreload extension is already loaded. To reload it, use: %reload\_ext autoreload

# **KDE for Our Anomaly Detection Case Study**

#### A Few Tweaks

#### We will work with log probabilities

- This is what sklearn does by default
- ...And simplifies some operations
- I.e. products become sums

#### Additionally, we will work with negated (log) probabilities

- They can be interpreted as alarm signals
- ...Which is the customary approach in anomaly detection

#### Overall, our anomaly detection condition becomes:

$$-\log f(x,\theta) \ge \varepsilon$$

...Which is equivalent to the previous formulation

### **Training and Testing**

We will split our data in two segments

#### A training set:

- This will include only data about the *normal* behavior
- Ideally, there should be no anomalies here (we do not want to learn them!)
- We will use it to fit a KDE model

#### A test set:

To assess how well the approach can generalize

#### If the training set contains some anomalies

- Things may still be fine, as long as they are very infrequent
- ...Since we will still learn that they have low probability

### **Training and Testing**

In time series data sets are often split chronologically:

```
In [3]: train_end = pd.to_datetime('2014-10-24 00:00:00')
    util.plot_series(data, labels, test_start=train_end, figsize=figsize)
```

• Green: training set, orange: test set

### Fitting the Estimator

#### Now, we can separate the training set

...And estimate the bandwidth by using the rule of thumb:

```
In [4]: data_tr = data[data.index < train_end]
  q1 = data_tr['value'].quantile(0.25)
  q3 = data_tr['value'].quantile(0.75)
  sigma = data_tr['value'].std()
  m = len(data_tr)
  h = 0.9 * min(sigma, (q3-q1)/ 1.34) * m**(-0.2)
  print(f'The estimated bandwidth is {h:.3f}')</pre>
```

The estimated bandwidth is 1056.061

#### Then we fit a univariate KDE estimator (using scikit-learn)

```
In [5]: kde = KernelDensity(kernel='gaussian', bandwidth=h)
kde.fit(data_tr.values);
```

## Fitting the Estimator

Let's have a look at the estimate distribution:

```
In [6]: vmax = data['value'].max()
    xr = np.linspace(0, vmax, 100)
    util.plot_density_estimator_1D(kde, xr, figsize=figsize)
```

# Alarm Signal

We can now obtain (and plot) our alarm signal:

```
In [7]: | ldens = kde.score_samples(data.values) # Obtain log probabilities | signal = pd.Series(index=data.index, data=-ldens) # Build series with neg. putil.plot_series(signal, labels=labels, windows=windows, figsize=figsize) #
```

- Notice how the the process (inference) is noticeably slower than training
- Can you tell why this was the case?

### **Detecting Anomalies**

#### Let us pick a threshold (at random, for now) and try to detect some anomalies:

```
In [8]: thr = 13
    pred = pd.Series(signal.index[signal >= thr])
    print(pred.head(), f'#anomalies: {len(pred)}')

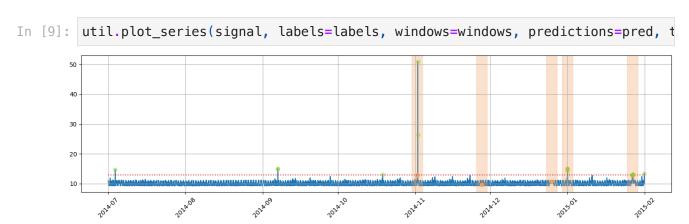
0    2014-07-03   19:00:00
    1   2014-09-06   22:30:00
    2   2014-09-06   23:00:00
    3   2014-10-18   23:30:00
    4   2014-11-02   01:00:00
    Name: timestamp, dtype: datetime64[ns] #anomalies: 15
```

- We just apply the filter signal >= thr to the signal index
- This yields a number of (predicted) anomalous timestamps
- Our module contains a function to perform this step:

```
def get_pred(signal, thr)
```

## **Detecting Anomalies**

Let us plot our predictions on the series:



- Not very good, but the threshold *is* random
- There are a many false positives, which are very common in anomaly detection

# **Metrics for Anomaly Detection**

How do we evaluate the quality of our system?

### **Metrics for Anomaly Detection**

**Evaluating the quality of an Anomaly Detection system can be tricky** 

- Usually, we do not need to match the anomalies exactly
- · Sometimes we wish to anticipate anomalies
- ...But sometimes we just want to detect them in past data

There is no "catch-all" metric, like accuracy in classification

#### It is much better to devise a cost model

- We evaluate the cost and benefits of our predictions:
- By doing this, we focus on the value for our customer

This is important for all industrial problems!

### A Simple Cost Model

We will use a simple cost model

Remember that our goals are:

- Analyzing anomalies
- Anticipating anomalies

First, we define:

- True Positives as windows for which we detect at least one anomaly
- False Positives as detected anomalies that do not fall in any window
- False negatives as anomalies that go undetected
- Advance as the time between an anomaly and when first we detect it

### A Simple Cost Model

They can be computed using a function in our module:

```
def get_metrics(pred, labels, windows)
```

### A Simple Cost Model

They can be computed using a function in our module:

```
def get_metrics(pred, labels, windows)
```

#### A Simple Cost Model

#### Then we introduce:

- A cost  $c_{alarm}$  for loosing time in analyzing false positives
- A cost  $c_{missed}$  for missing an anomaly
- A cost  $c_{late}$  for a late detection (partial loss of value)

Our cost model (simple, but serviceable) is then given by:

The cost with the current predictions is: 35

Of course, in a real problem the cost model should be discussed with the stakeholders

### **Threshold Choice**

## Refactoring the Cost Model

#### We will use this cost model for choosing our threshold

With this aim, we encapsulate the formula in a Python class in our module:

```
class ADSimpleCostModel:
    def __init__(self, c_alrm, c_missed, c_late):
        # Store cost parameters
        ...

def cost(self, signal, labels, windows, thr):
        # Compute the cost
```

### Refactoring the Cost Model

We can now evaluate our cost by:

- Instantiating the class with our values for  $c_{alarm}, c_{missed}, c_{late}$
- Calling the cost function with different threshold values

```
In [15]: cmodel = util.ADSimpleCostModel(c_alrm, c_missed, c_late)
  cost = cmodel.cost(signal, labels, windows, thr)
  print(f'The cost with the current predictions is: {cost}')
```

The cost with the current predictions is: 35

#### The process is rather efficient:

- · When we change the threshold
- ...We do not need to rebuild the alarm signal

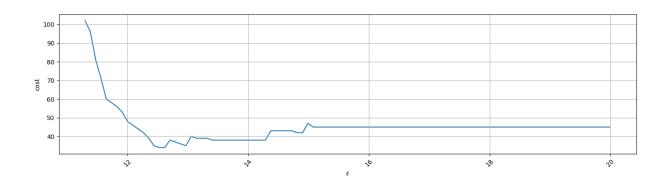
So, no need to repeat either training or inference

### Effect of the Threshold

We can now look at the effect of changing the threshold:

First, we'll do this for an *idealized problem* (i.e. over all the data)

```
In [16]: cmodel = util.ADSimpleCostModel(c_alrm, c_missed, c_late)
    thr_range = np.linspace(11.3, 20, 100)
    cost_range = pd.Series(index=thr_range, data=[cmodel.cost(signal, labels, wi
    util.plot_series(cost_range, figsize=figsize, xlabel=r'$\varepsilon$', ylabe
```



# Choosing the Threshold

#### Ideally, we wish to choose the best threshold

However, we have two problems:

- · We cannot really use all the data
- ...we need to keep some away to test generalization
- The threshold-to-cost function is non-smooth and non-differentiable

#### Luckily, both are easy to address

We can:

- Define a validation set to use when optimizing the threshold
- Use a simple line search approach, since  $\varepsilon$  is a scalar

#### **Define a Validation Set**

#### First, we define our validation set:

- It should contain some anomalies
- ...But some should be left our (for testing)

```
In [17]: val_end = pd.to_datetime('2014-12-10 00:00:00')
    util.plot_series(data, labels, val_start=train_end, test_start=val_end, figs
```

### Optimize the Threshold

We will use a line sampling approach:

We build the validation set and define a range of "sampled" thresholds:

```
In [18]: signal_opt = signal[signal.index < val_end]
    labels_opt = labels[labels < val_end]
    windows_opt = windows[windows['end'] < val_end]</pre>
```

Then, we use a function from our module to pick the best one:

```
def opt_thr(signal, labels, windows, cmodel, thr_range):
    costs = # compute costs as before
    best_idx = np.argmin(costs)
    return thr_range[best_idx], costs[best_idx]
```

### Validation Set?

Did you notice how we included the training data in the validation set?

Is that good or bad?

It's definitely not standard, but reasonable in this case!

The reason is that our "validation set"

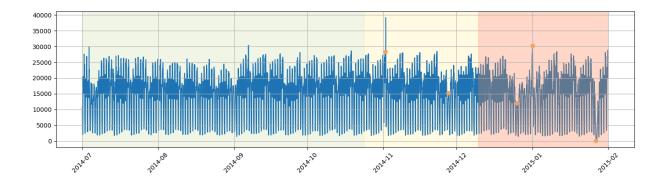
- ...Is not needed to ward off against overfitting
- ...But just to optimize one more parameter (i.e.  $\theta$ )

It is actually part of the training data

#### **Get Accustomed to Drift**

One more thing: see how the anomaly frequency changes over time?

```
In [19]: util.plot_series(data, labels, val_start=train_end, test_start=val_end, figs
```



- In time series, the data distribution often drifts over time
- When this happens, it is usually good to split the series in contiguous chunks
- I.e. the training, validation, and test set should be sequences

# Optimize the Threshold

#### Let's see the results of threshold optimization

```
In [20]: best_thr, best_cost = util.opt_thr(signal_opt, labels_opt, windows_opt, cmc
print(f'Best threshold: {best_thr:.3f}, corresponding cost: {best_cost:.3f}'
```

Best threshold: 15.079, corresponding cost: 15.000

The reported cost is on the training and validation set

#### For all the data (yes, we are cheating a bit) we have:

```
In [21]: ctst = cmodel.cost(signal, labels, windows, best_thr)
print(f'Cost on the whole dataset {ctst}')
```

Cost on the whole dataset 45

- This is, as expected, suboptimal
- ...But it works! We have our first complete anomaly detection approach