

On the feasibility of classification-based product package authentication



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- Proposed authentication scheme

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- Dataset structure

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- Classification approaches
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Motivation [1]

Counterfeited products



2013: *5% counterfeited products on EU level → faked medicals are a threat for the patients and cause an economic loss.*

*The Falsified Medicines Directive (FMD) should be implemented until 2018. The approached solution is relies on product **serialization** and tracking using unique numeric identifiers.*

Motivation [2]

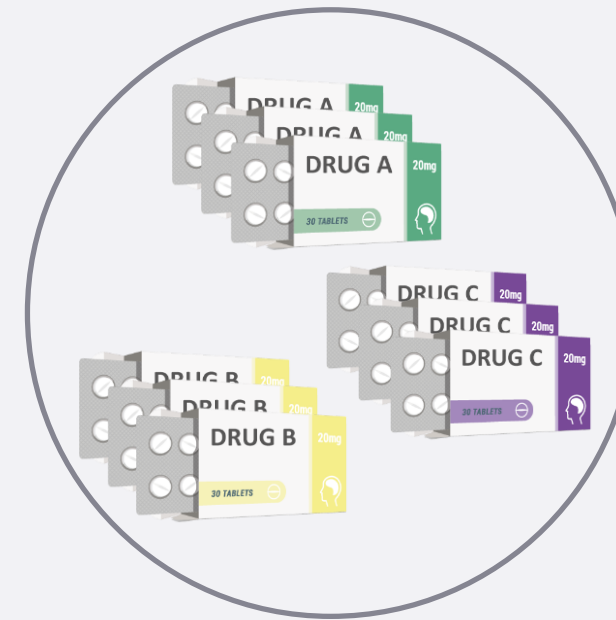
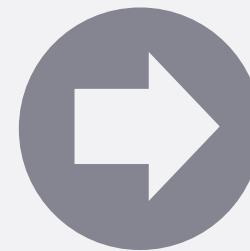
Paper-based PUFs



Previous literature showed that the fibre structure of paper or packaging material is positional highly unique and enables to identify single instances.

Basic idea

move from serialization to classification



serialization

Individualize each instance of a product using unique identifiers or PUF-based approaches, e.g. fibre fingerprints

classification

Use intrinsic or extrinsic features which are constant across all instances but different to features from other products.

Pre-requirements

Fundamental research questions



serialization

Individualize each instance of a product using unique identifiers or PUF-based approaches, e.g. fibre fingerprints

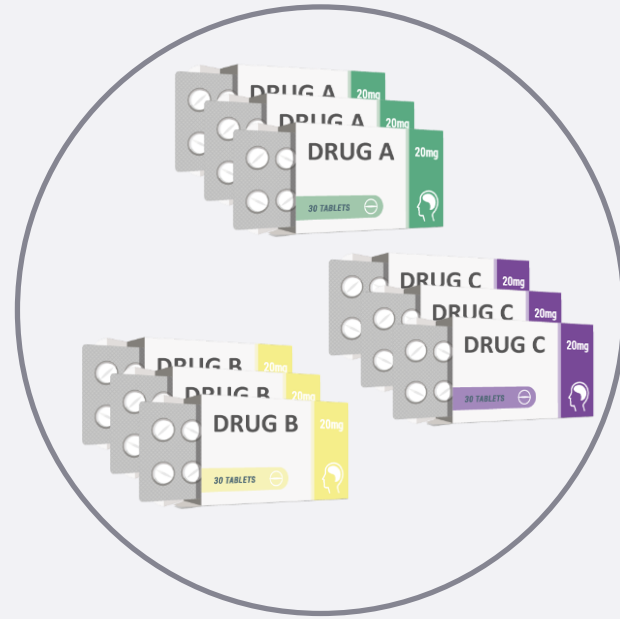
It is clear that the fibre-structure is locally unique.

Uniqueness



Pre-requirements

Fundamental research questions



classification

Use intrinsic or extrinsic features which are constant across all instances but different to features from other products.

01

It is not clear

- If the fibre structure shows constant features across different regions and
- if those features are discriminative enough to distinguish between different types of paper or packagings.

Instance
invariance

?

02

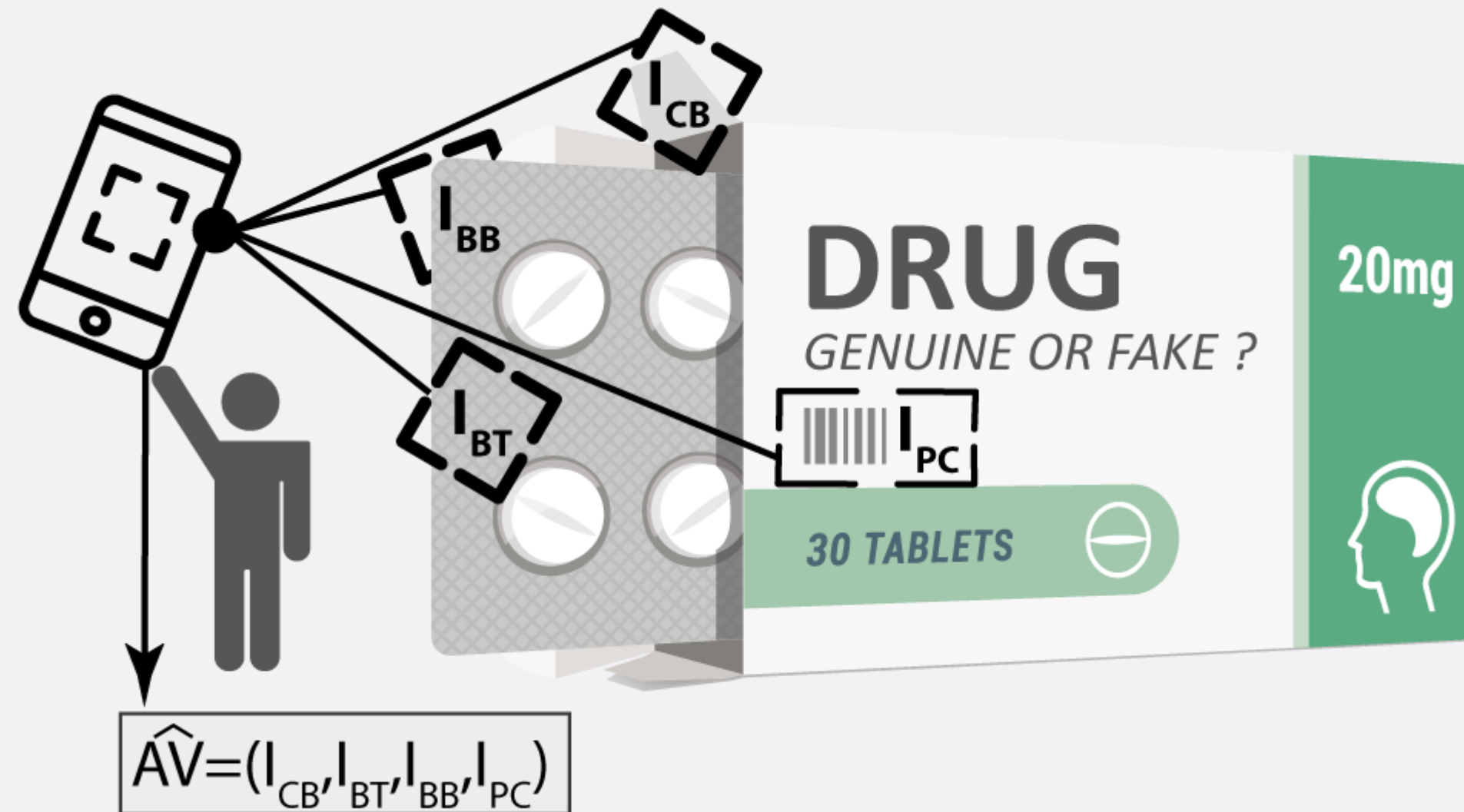
It is not clear

- how the texture and the computed features vary between different instances of a product.

Instance
generalisation

Drug packaging authentication system

Basic concept [1/3]



Capture packaging **modalities**:

CB = Cardboard

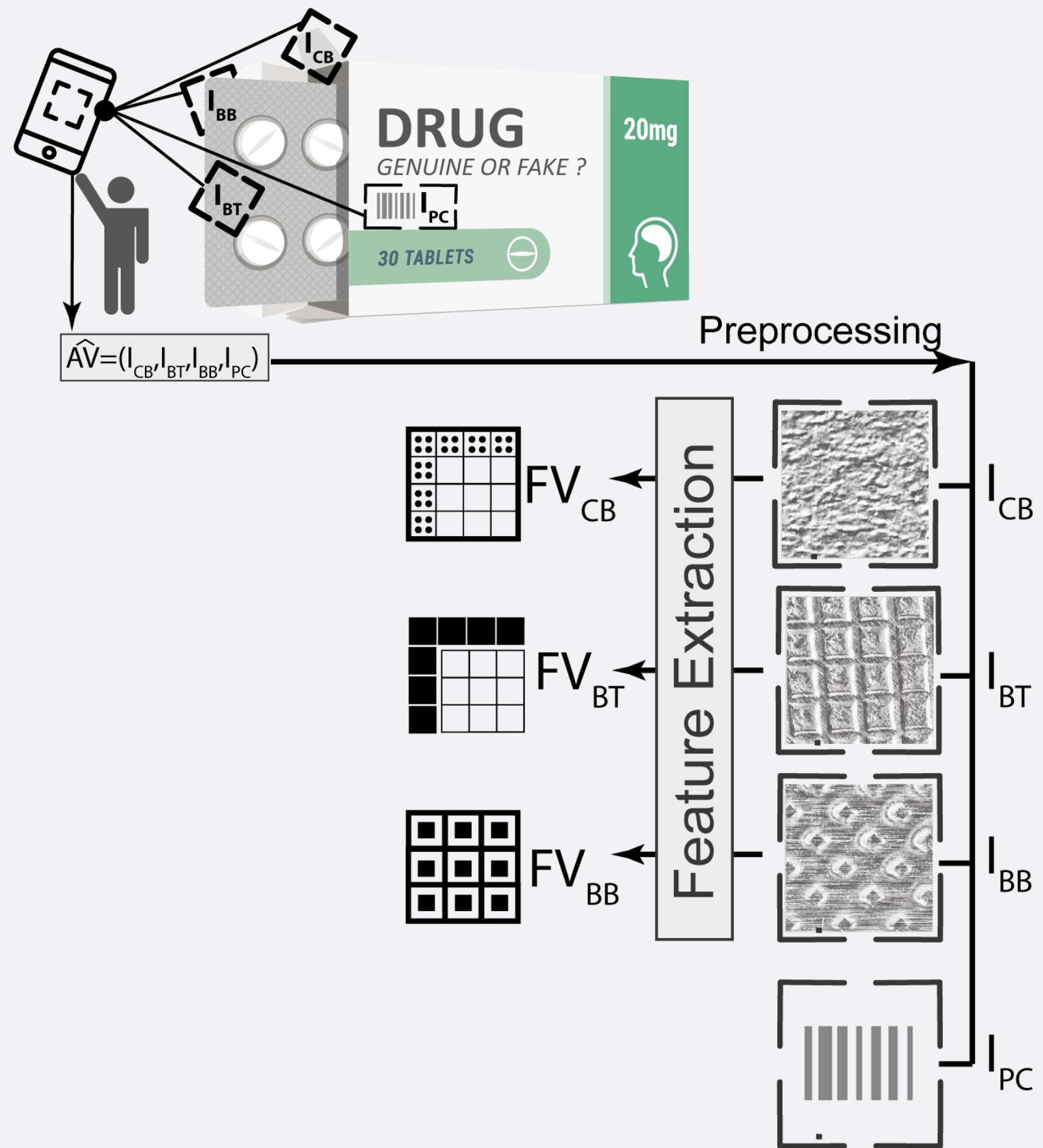
BB = Blister Bottom

BT = Blister Top

& the product code (PC)

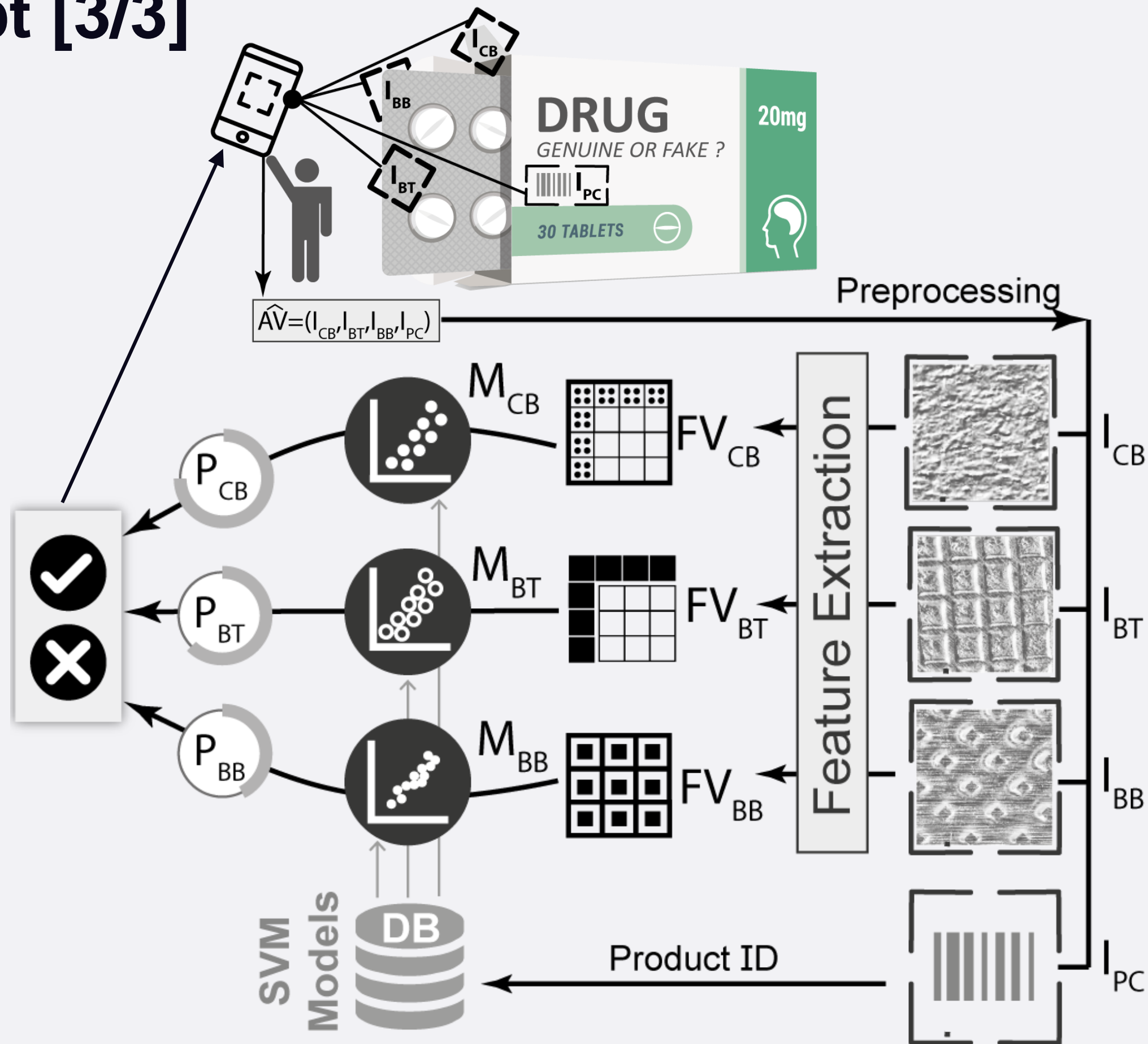
Drug packaging authentication system

Basic concept [2/3]



Drug packaging authentication system

Basic concept [3/3]



Drug packagings texture database

Acquisition details [1/2]



Sample collection

Packages were collected in different pharmacies in Salzburg.



Sorting & Labelling

All packages were sorted and each drug was assigned an identifier and the available instances were numbered.

- **Drugs #45**
- **Producers #28**
- **1 to 15 instances per drug**



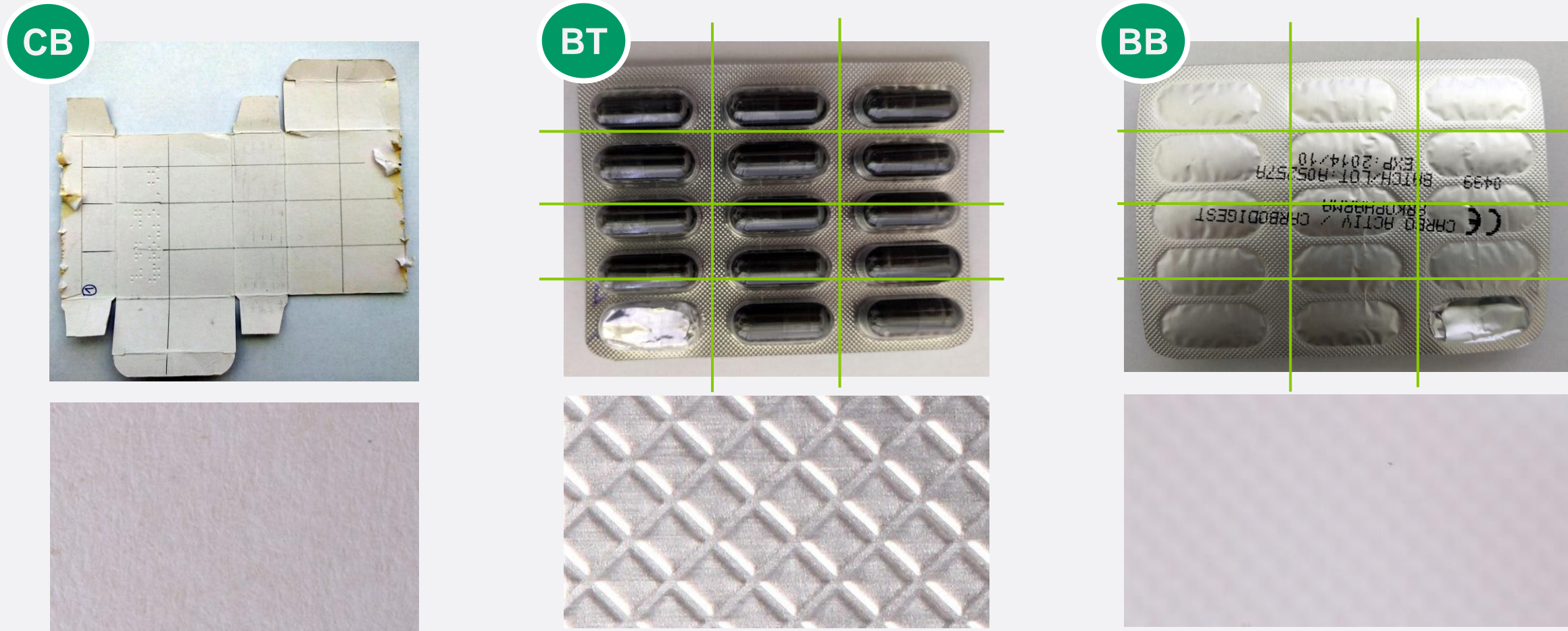
Image Acquisition

Images were captured in a controlled environment using a Canon 70D with a 100 mm macro lens and a flashlight.

The image distance was approximately 28cm.

Drug packagings texture database

Acquisition details [2/2]



Non-overlapping

Capture non-overlapping sections of each instance and modality



Cropping

The final images are of arbitrary size and show textural information of the modality.

Classification pipeline

Scenarios



CLASS or PACKAGE

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CLASS

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generalisation

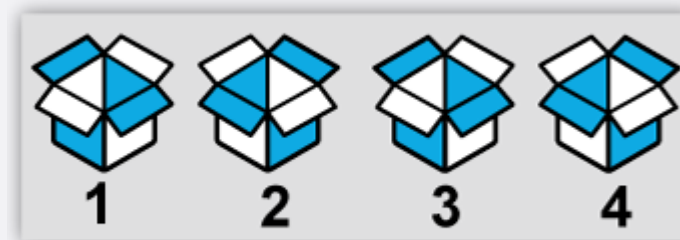
PACKAGE

Classification pipeline

Data selection

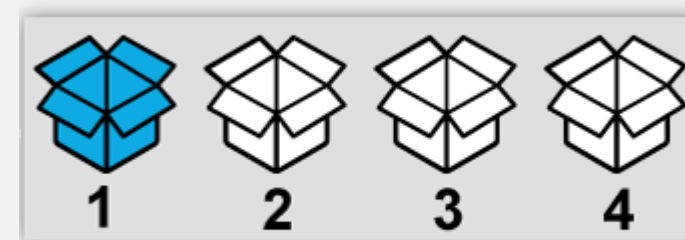


CLASS



$$|1+2+3+4| = k$$

PACKAGE



$$|1|, |2|, |3|, |4| = k$$

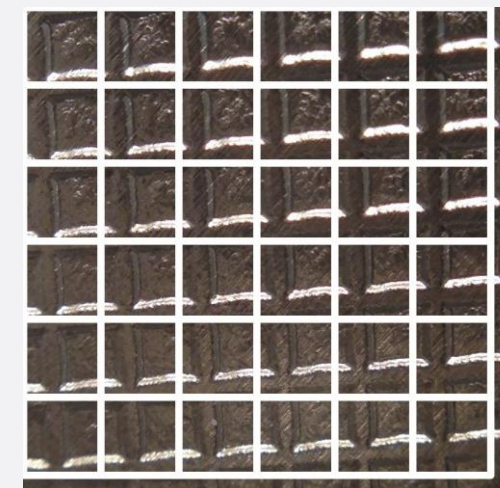
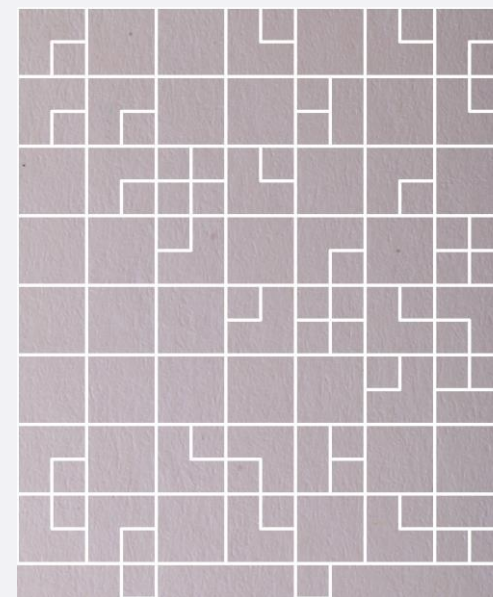
CLASS or PACKAGE

Keypoint selection:

k – image patches, with a predefined size, are selected for each modality in a scenario specific manner.

Patch sizes:

128x128, 256x256



Classification pipeline

Feature Extraction



CLASS or PACKAGE

Keypoint selection

Low-level features:

Each selected patch is contrast enhanced (CLAHE) and a set of feature vectors are computed.

e.g. LBP & variants, HOG, DTCWT



Classification pipeline

Classification approaches



CLASS or PACKAGE

Keypoint selection

Low-level features

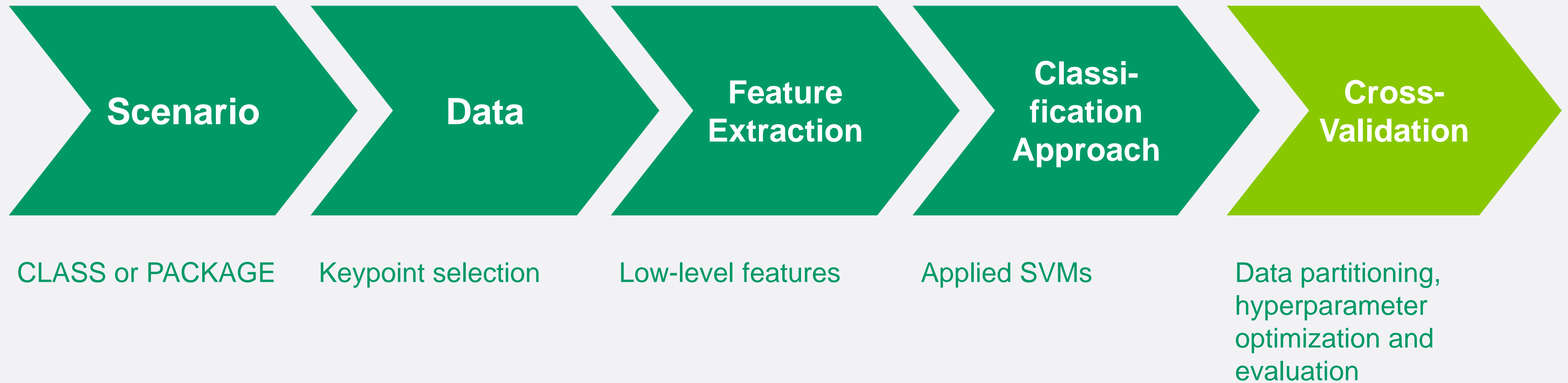
Applied SVMs:

- LIBSVM:
 - **ONE-CLASS**
 - **BINARY C-SVC**
- LIBSVM Extension:
 - **WSVM**

For all a radial basis function (RBF) kernel is utilized.

Classification pipeline

Cross-fold validation [1/4]



Classification pipeline

Cross-fold validation [2/4]



Parameters

Drugs #45	$D = \{d_1, \dots, d_{45}\}$
Drug manufacturers	$DM = \{dm_1, \dots, dm_{28}\}$
Packaging modality	$M = \{CB, BB, BT\}$
Feature Extraction M.	$FE = \{fe_1, \dots, fe_n\}$
Classification Scenario	$CS = \{CLASS, PACKAGE\}$

$$CC = (d \in D, m \in M, fe \in M, cs \in CS)$$

Classification Configuration

$$FV_{CC} = \{ FV_{(d_1, m, fe, cs)}, \dots, FV_{(d_{45}, m, fe, cs)} \}$$

CC specific Feature Vector Sets



Nested cross-validation using a specific classification approach

Positive Training Data

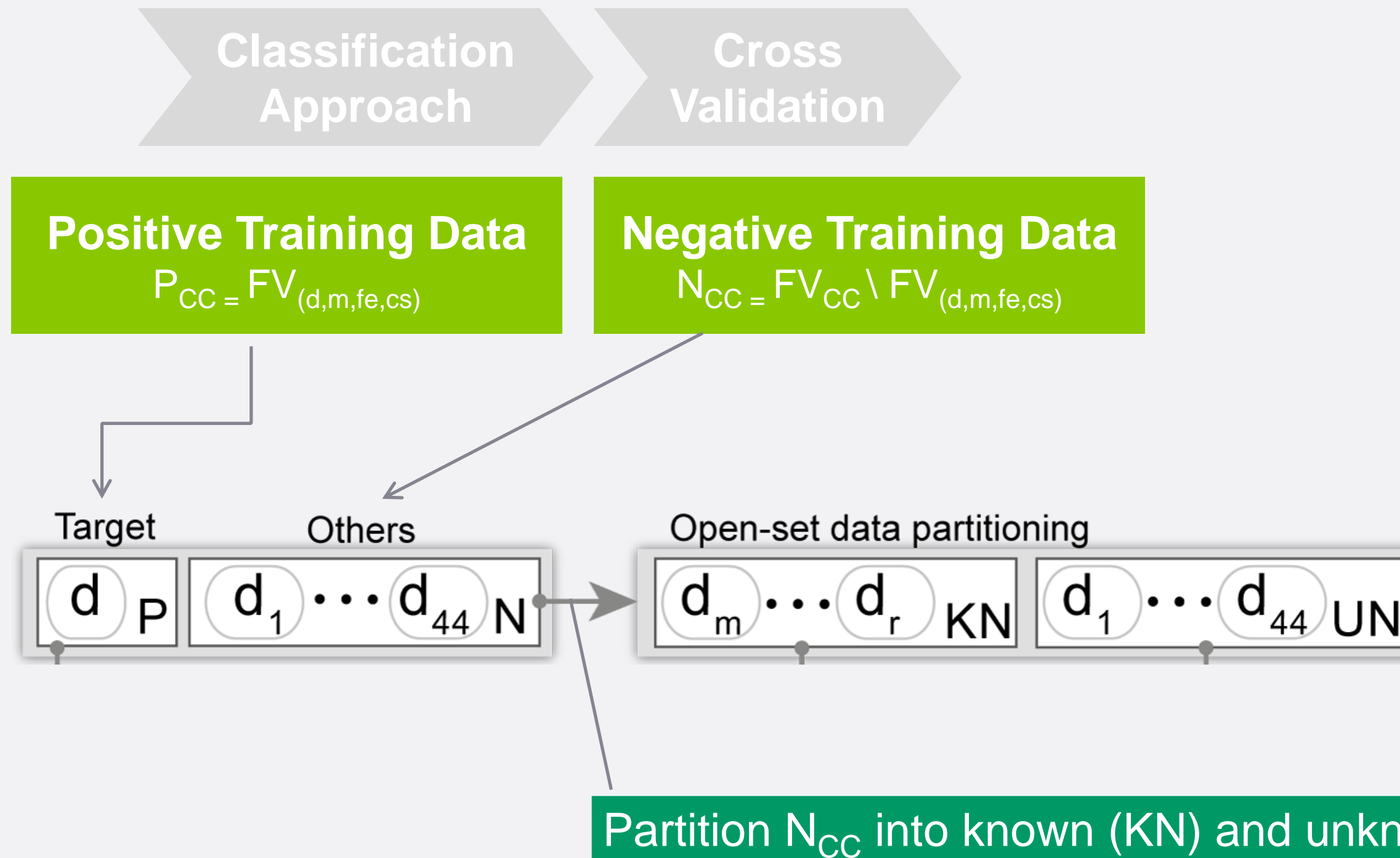
$$P_{CC} = FV_{(d, m, fe, cs)}$$

Negative Training Data

$$N_{CC} = FV_{CC} \setminus FV_{(d, m, fe, cs)}$$

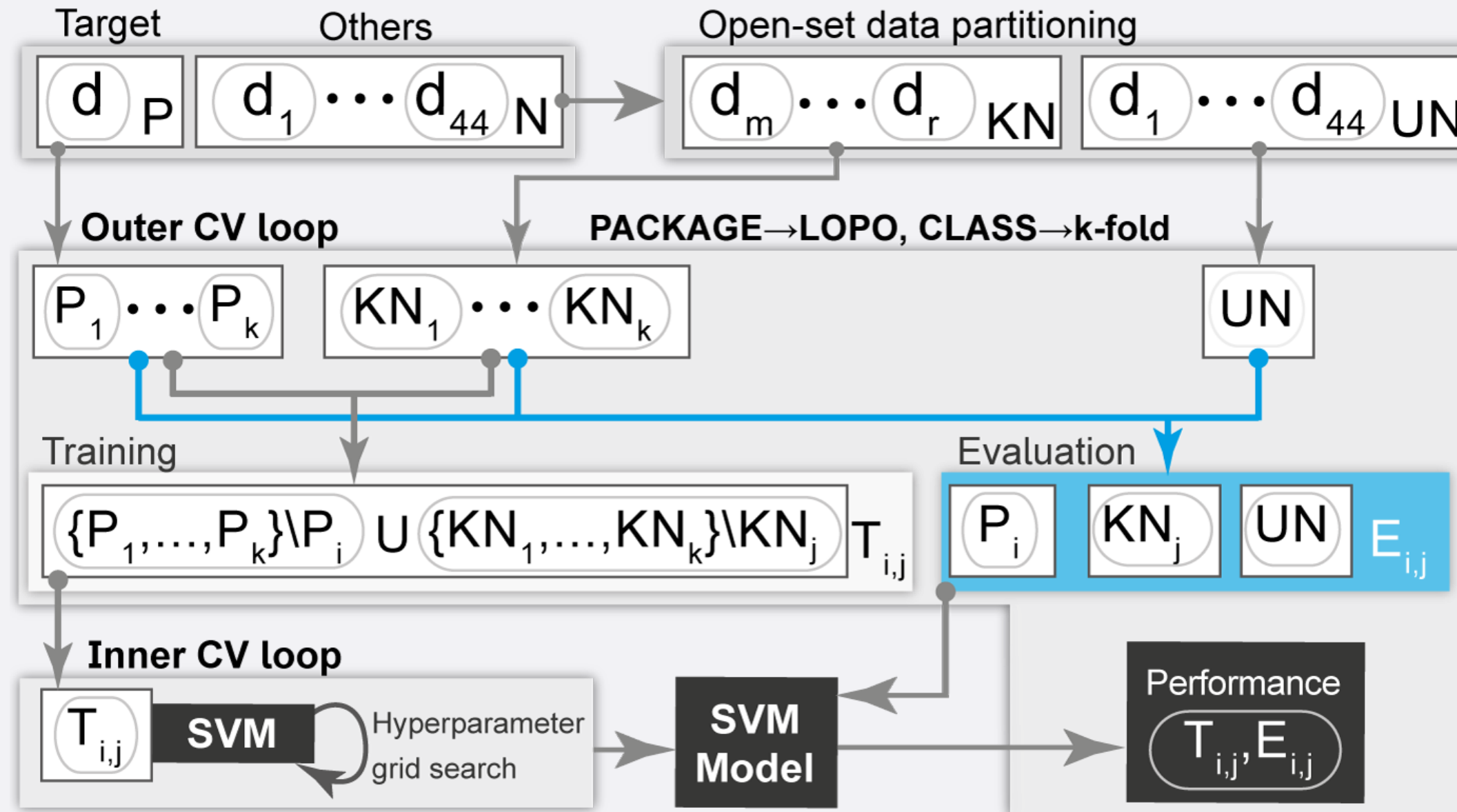
Classification pipeline

Cross-fold validation [3/4]



Classification pipeline

Cross-fold validation [4/4]



Optimize classifier using known negatives in the inner CV loop

In case of binary SVMs, a subset of the known negatives is not used for training; i.e. only for evaluation in order to address the open-set problem in the inner CV loop. The SVM parameters and a probability threshold achieving the highest F-Measure are determined.

Experiments

Research questions - Reminder

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Instance
invariance

CLASS

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Instance
generalisation

PACKAGE

Classification Performances – Overview:

All CCs were computed and the best CLASS and PACKAGE configurations (=highest averaged F-Measure and StDev) for different CC parameters and modalities were determined.

- **CLASS:** averaged results over all 45 drugs
- **PACKAGE:** averaged results for drugs with at least 5 instances.

Experiments

Results overview

01

02

page 24

page 23

A

B

CC	CLASS						PACKAGE					
	128×128			256×256			128×128			256×256		
	CB	BT	BB	CB	BT	BB	CB	BT	BB	CB	BT	BB
ONE-CLASS	<i>LTP</i> 0.83 ±7.9	<i>LTP</i> 0.9 ±6.2	<i>LTP</i> 0.92 ±5.8	<i>LTP</i> 0.91 ±4.4	<i>LTP</i> 0.85 ±13.6	<i>LBP</i> 0.87 ±13.5	<i>LBP</i> 0.81 ±8.7	<i>LBP</i> 0.86 ±6.3	<i>LTP</i> 0.84 ±11.3	<i>LTP</i> 0.85 ±9.1	<i>LBP</i> 0.88 ±5.0	<i>LBP</i> 0.85 ±7.1
BINARY	<i>LTP</i> 0.88 ±6.9	<i>LiLBP</i> 0.94 ±3.2	<i>LTP</i> 0.93 ±4.1	<i>LTP</i> 0.91 ±5.2	<i>LiLBP</i> 0.92 ±9.0	<i>LTP</i> 0.93 ±5.0	<i>LTP</i> 0.82 ±9.5	<i>LTP</i> 0.92 ±3.7	<i>LTP</i> 0.87 ±8.9	<i>LTP</i> 0.85 ±5.5	<i>LTP</i> 0.94 ±5.7	<i>LiLBP</i> 0.87 ±10.0
WSVM	<i>LTP</i> 0.86 ±7.6	<i>LTP</i> 0.93 ±4.1	<i>LTP</i> 0.93 ±4.3	<i>LiLBP</i> 0.88 ±6.0	<i>LTP</i> 0.88 ±7.6	<i>MFS</i> 0.88 ±9.1	<i>LTP</i> 0.85 ±8.2	<i>LTP</i> 0.91 ±4.2	<i>LiLBP</i> 0.85 ±9.2	<i>LiLBP</i> 0.83 ±8.5	<i>LTP</i> 0.89 ±8.7	<i>LiLBP</i> 0.84 ±10.1

CLASS results show high mean F-Measures over 0.9, indicating that textures from all three modalities show constant but highly discriminative features. This enables to recognize the same drug class and to distinguish it from others.

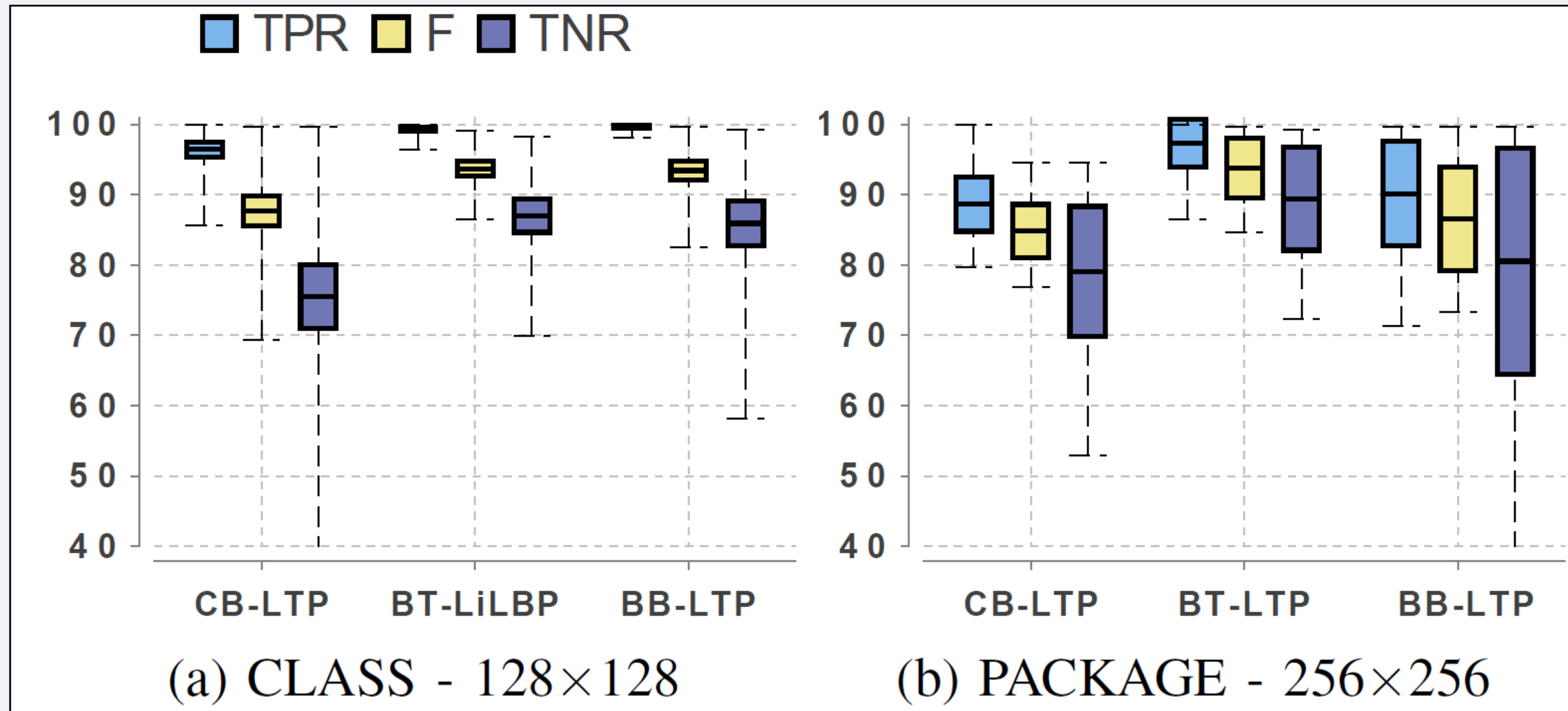
Instance invariance

Package results show that textural features are constant across different instances for all three modalities. This is a basic requirement for a classification-based authentication system.

Instance generalisation

Experiments

Details [A]



CLASS vs. PACKAGE
(Binary C-SVC) performance comparison.

Class Accuracy / True Positive Rate:

$$TPR = \frac{TP}{TP + FN}$$

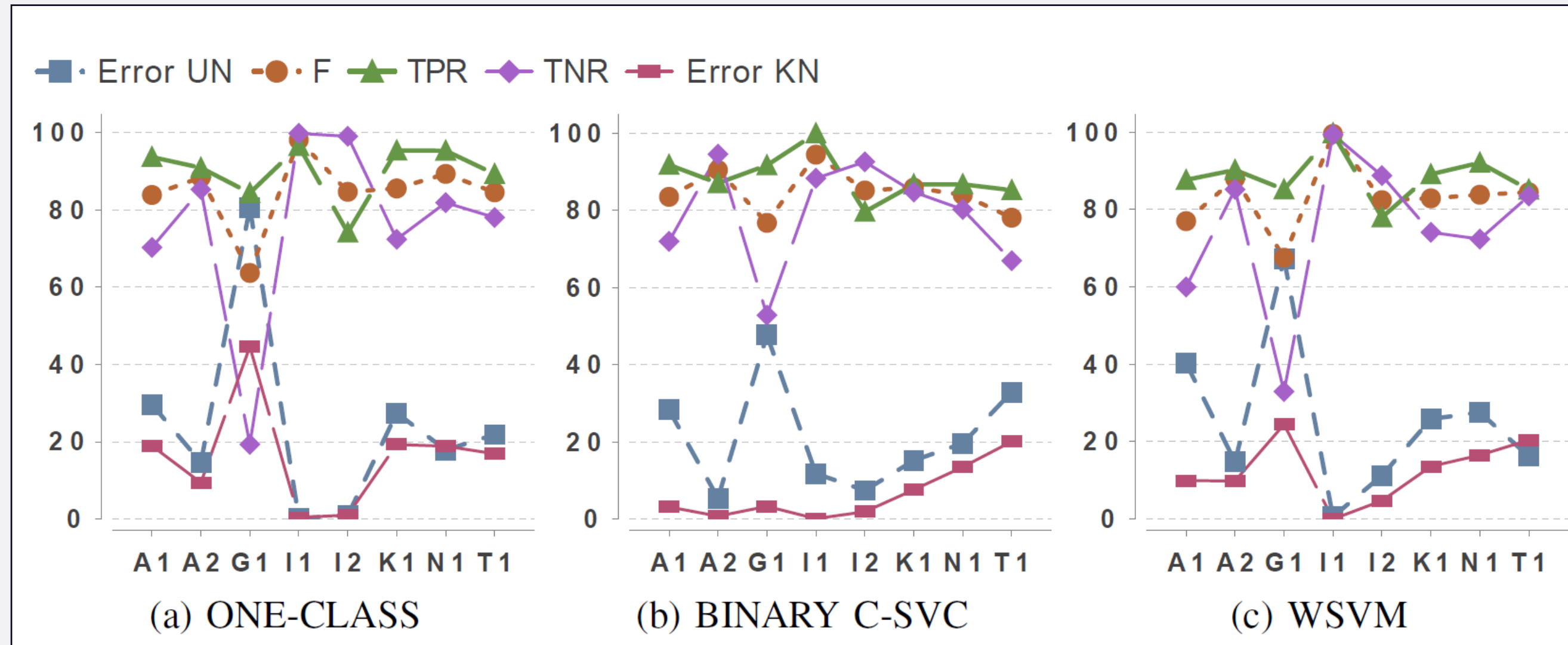
Others Accuracy / True Negative Rate

$$TNR = \frac{TN}{TP + FP}$$

Performance (*StDev*) decreases in case of the PACKAGE scenario which reflects a real world package-authentication setup.

Experiments

Details [B]



PACKAGE (256x256) - SVM performance comparison for **CB** and all target drugs (=8) with more than 5 instances.

X-Axis: Target Drug IDs: (e.g. A1 = manufacturer A + drug number 1).

Error UN = False classified unknown negatives.

Error KN = False classified known negatives.

Error KN (data seen in training) is lower than Error UN (=data not used for training = open world).

Thank you

Conclusions and Outlook



Instance **invariance**

Textural features of drug packaging material are constant and highly discriminative.



Instance **generalisation**

Experiments indicate that a classifier can be trained with a set of known instances and is able to authenticate unseen instances.



TODO

Real world data from faked packages is required. Use high-level features, feature encoding and feature fusion techniques. Investigate error sources, i.e. probably other drugs from the same manufacturer use the same packaging material.