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Faculty of Computer Science, Institute of Systems Architecture

## Better Steganalysis (BEST) Final Report Reduction of Interfering Influence of Image Content on Steganalysis \*

Prof. Dr.-Ing. Andreas Westfeld

andreas.westfeld@htw-dresden.de

October 8, 2009

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## **Summary**

Steganography is the art and science of invisible communication. Its aim is the transmission of information embedded invisibly into carrier data. The goal of steganalysis is to discover steganographic alterations to carrier data.

The project BEST (Better Steganalysis) explored the influence of image content when detecting steganography. If this influence, which interferes with the detector statistics, is reduced, the reliability of steganalysis can be increased. Steganalysis can be regarded as an attempt to separate image (or carrier) content from embedded steganographic signals.

This project consisted of two phases. During its first phase, a new technique to scan elements of potential steganograms allowed a more fine-grained evaluation of the group size, a parameter of the RS attack by Fridrich et al., which affects the detection performance.

The results of Phase 2 include newly developed attacks for advanced JPEG steganography like MB2 that use targeted features to blockiness reduction artefacts. Phase 2 also comprises a new generic methodology to apply spatial domain attacks in the DCT domain.

All attacks are implemented in a software package called *steganographic work-bench* (swb), which is described in this report. The package also includes implementations of some previous state of the art attacks that have been used for evaluation.

## **Chapter 1**

### Introduction

Steganography is the ancient art and young science of hiding the communication in such a way that the existence of the private message cannot be detected. It works by hiding the secret message in a harmless carrier object. While cryptography merely ensures the confidentiality of the message content, steganography adds another layer of secrecy by concealing the fact that secret communication takes place.

Progress in steganography is tightly coupled with improvements of steganalysis, the detection of hidden information. In this report we introduce several techniques that aim to reduce the interfering influence of image content on steganalysis:

- · optimisation of the group size parameter for RS steganalysis,
- separation of error sources for measurement of the influence of image content,
- a generic methodology to apply spatial domain attacks to frequency domain,
- two new more reliable higher order statistical attacks on randomised Jsteg, one for small and one for large images, and
- a powerful attack on MB2 using targeted features to blockiness reduction artefacts.

## **Chapter 2**

# Methods, Assumptions, and Procedures

#### 2.1 A Steganographic Workbench

This report is intended to present the key ideas supported by the software in practical examples.

The software of this project, the *steganographic workbench* or swb package is implemented in R. Some parts are optimised for performance reasons in C.

R is a free programming language and computing environment that has become a de-facto standard among statisticians for the development of statistical software. Despite its strong points, and its large user base among statisticians, it is fair to say that R and its commercial cousin S-PLUS are not widely used in the signal processing community.

As far as our experience goes, R can significantly lower the threshold for experimental prototype implementations. While nested loops are necessary to process image files in C or C++, it is just one line of code in R due to the compact, clear and math-like syntax. Apart from the time saved while writing and reading the code, one source of its errors—the complexity—is suppressed.

To run the "steganographic workbench," an installed version of R is necessary (www.r-project.org). To install the swb package on Windows, start the R GUI environment and select "Install package(s) from local zip file..." in the "Packages" menu. Then choose the location of swb. zip and confirm.

On Unix systems this is accomplished by the shell command line

bash# R CMD INSTALL swb

where the package file swb.tgz must reside in the current working directory. You need supervisor rights to make packages available to all users. If you don't have these rights or want to install it only in your home directory, create and specify your own target for R libraries first:

bash\$ export R\_LIBS=~/.R

bash\$ mkdir -p \$R\_LIBS
bash\$ R CMD INSTALL swb

Once the swb package is installed, the user can load it into R using the command library (swb). This makes a the following collection of functions available:

ı	1	1	۲	٦
ı	/	١	L	J

Read or write image data from/to files.

read.gif<sup>1</sup>
read.jpg
read.pgm<sup>1</sup>
read.png
read.pnm<sup>1</sup>
write.png

#### Attacks/Feature Sets

Predict the use of steganography, estimate the length of the embedded message, or extract a feature vector in order to train a classifier.

abs.moments.attack

ca.attacks<sup>1</sup>
ca.estimate
ca.jphide<sup>1</sup>
ca.jsteg<sup>1</sup>
eca<sup>1</sup>
eca.jphide<sup>1</sup>
eca.jsteg<sup>1</sup>
gca.jphide<sup>1</sup>

gca.jsteg<sup>1</sup> gca.new.jsteg<sup>1</sup> f5attack<sup>1</sup> f5estimate<sup>1</sup>

hcfcom.attack jphide.normalise<sup>1</sup> jsteg.normalise

jpairs.attack irs.attack jspa.attack jws.attack

mb1.esorics.attack¹
mle.estimate
pairs.attack
rs.attack
spa.attack
qem
qem2

yu.attack<sup>1</sup> yu.estimate zp.attack<sup>1</sup> zp.estimate

wb.attack

f23<sup>1</sup> f274<sup>1</sup> f324<sup>1</sup>

ext.dct.features<sup>1</sup> ext.markov.features<sup>1</sup> merged.features<sup>1</sup>

#### **ROC Curve Handling**

Assess the quality (power/ accuracy) of attacks or plot an ROC curve.

assess.roc<sup>1</sup> eer<sup>1</sup> farid<sup>1</sup>

fpr.thresholds1

ker<sup>1</sup>
plot.roc<sup>1</sup>
prepare.roc
rho<sup>1</sup>
thresholds<sup>1</sup>
tor.thresholds<sup>1</sup>

#### **Embedding/Extraction**

Simulate the embedding of a message.

jphide.embed jsteg.embed lsb.embed lsb.extract ltsb.embed ltsb.extract pm1.embed pm1.extract

#### **Transformations**

DCT, quantisation, and scanning along a plane filling curve.

ndctq<sup>1</sup>
nidctq<sup>1</sup>
jndctq<sup>1</sup>
jnidctq<sup>1</sup>
scanpath
plot.scanpath

#### **Integer Operators**

Most of these operators are available in R, but require time-consuming conversions to and from floating point format. For performance reasons we implemented integer versions

in C.

iabs

<sup>&</sup>lt;sup>1</sup>There is currently no man page for functions that are rather for internal use in the swb package.

iand	iterative.mean.int <sup>1</sup>	tbl
iasr	iterative.mean <sup>1</sup>	jcalibrate <sup>1</sup>
ilsr	ixor	jhist
ior	ctbl <sup>1</sup>	jnzcoeff <sup>1</sup>
irange	dtbl <sup>1</sup>	
isign	mtbl <sup>1</sup>	

For some functions there is an online help available. For instance, the user can display the man page of read.jpg by entering ?read.jpg into R.

#### 2.2 Recursive Scanpaths

Many tools that detect steganographic payload extract a sequence of elements from the carrier media (e.g., images). In most cases the ordering of this sequence was just proposed by the ordering of the elements in the file. However, other sequences can lead to better detection performance. In addition we can turn a two-dimensional object like an image into a one-dimensional sequence object. Thus we can design detectors independently of an object's dimensionality.

Discrete objects of arbitrary complexity can be recursively scanned, if the recursion depth is chosen appropriately. We used recursive scanpaths to improve the detection power of the Pairs attack in images [23]. A space filling curve is a continuous map of a two-dimensional area (plane-filling) or a three-dimensional volume into a one-dimensional interval. David Hilbert, a German mathematician, invented a simple space-filling curve known as the Hilbert curve, which fills a square. The Hilbert curve can be encoded with the initial string L and the following string rewriting rules [20].

$$\begin{array}{cccc} L & \longrightarrow & +RF - LFL - FR + \\ R & \longrightarrow & -LF + RFR + FL - \\ F & \longrightarrow & \text{go one pixel forward} \\ + & \longrightarrow & \text{turn right} \\ - & \longrightarrow & \text{turn left} \end{array}$$

These rules terminate after a specific recursion depth.

Figure 2.1 shows the well-known Hilbert curve for the recursion depths 1, 2, and 3. The bold enumeration follows the curve and the italic follows the pixels in the file, which are stored row by row. The swb package provides a function scanpath that accepts the parameters width x and height y of the scanned image. The recursion depth d is internally determined for a Hilbert curve that is large enough to cover all pixels in the image:

$$d = \lceil \log_2 \max(x, y) \rceil \tag{2.1}$$

In most cases the Hilbert curve is able to cover more pixels than the actual image contains. The result of scanpath is a vector of  $x \cdot y$  indices<sup>2</sup>, which are used to

<sup>&</sup>lt;sup>2</sup>The return type is actually a matrix that is used as an index vector. The two dimensions given as parameters to the function are stored as a dimension attribute for convenience.

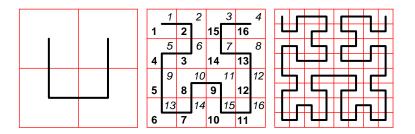


Figure 2.1: Hilbert curves for the recursion depths 1, 2, and 3

permute the pixels.

```
Example
lena <- read.png("lena.png")
sp <- scanpath(lena)
# or: sp<-scanpath(dim(lena)[1], dim(lena)[2], "hilbert1")
permuted.lena <- lena[sp]</pre>
```

In the permuted sequence of the image (i.e., pixels scanned along the Hilbert curve) almost all pixels are direct neighbours. This is also true if scanned row by row. However, also second order neighbours (every other pixel in the sequence) have a small mean distance (about  $\sqrt{2}$ ), while this distance is about 2 in the original row-by-row order in the file. As a consequence, close pixels in the permuted sequence show a stronger correlation than before. This property was used in the aforementioned improvement of the Pairs attack.

The swb package implements further scan methods, "hilbert1" is only the default. The remaining are listed in the online help (enter ?scanpath in the R command line). An example for a scan method can be displayed using the plot.scanpath method (e.g., plot.scanpath ("zigzag")).

#### 2.3 RS Group Sizes

RS analysis, which is another attack on LSB steganography, computes statistics on small disjoint groups of adjacent pixels[8]. Here it became evident that groups of pixels arranged in a square of  $m \times m$  deliver better results than slices of  $n \times 1$  pixels [8, 13]. This can be explained by the fact that locally close pixels differ less in colour or brightness than more distant ones. Certain types of systematically added randomness (e.g., LSB replaced by steganographic noise) are more easily detected in groups of closely related pixels with low variance in brightness. Unfortunately the quadratic shape scales in broad steps (group size  $g = 1, 4, 9, \ldots$ ; see Fig. 2.2), which might impede the group size to be adapted to the image size in an optimal way. Scanning pixels along a recursive scan path will allow us to refine the step size and close the gap for the groups of pixels. Therefore it could improve RS if the optimal is in the middle of one of the previous coarse steps in group size.

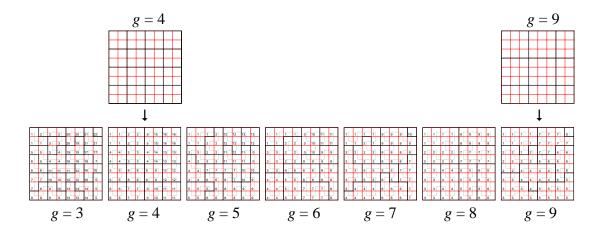


Figure 2.2: Conventional square shape of groups (above) compared to finer step size with proposed group shapes (below)

Along with refining the group size, recursive scanning can sequence elements: The pixels are no longer shaped in two dimensions. Therefore RS can work with slices of n pixels. In our experiments we consider all possible masks with  $n = 2 \dots 10$ . However, we cannot expect different results for isomorphic masks, e.g. for the mirrored masks 00101 and 10100. Likewise with the mirrored version of an image: RS will give a similar result for it. Not so for mask 01010. Although the bits are just shifted by one position, it is different if the bit to be flipped (1 in the mask) is at the end of the mask (influencing only one neighbour difference in the noise measure) or in the middle (influencing two differences). The rules to reproduce this set are:

- 1. include the binary pattern of ascending unsigned integers with  $n = 2 \dots 10$  bits unless a subsequent condition is fulfilled,
- 2. exclude masks where all bits are the same, since they are unusable for RS,
- 3. exclude a mask if it is a mirrored version of a mask that is already included,
- 4. exclude a mask if it is a version of a previously included mask that was shifted to the left by two or more positions.

Table 2.1 shows an example for RS masks of length n=4 that can be skipped in the experiments due to isomorphism. There are 776 different masks with n=2...10 bits; about one quarter of the workload is saved due to isomorphism.

#### 2.4 Separation of Error Sources

In this project we proposed some methods that estimate the length of the embedded message. We will average length estimates in order to separate different sources of errors. Böhme and Ker defined three error-measures [1], which are evaluated in our experiments.

Mask Drop-reason or "✓"

all equal (unusable)

√

shift □ left by 2 or more

shift □ left by 2 or more

shift □ left by 2 or more

minusable)

shift == left by 2 or more

Table 2.1: Drop isomorphic RS masks

A *cell* is a set of steganograms, produced when different messages of the same length are embedded into one particular carrier medium. To separate the error sources we measure the statistical dispersion (inter quartile range, IQR, difference between 0.75- and 0.25-quantile) and the central tendency (median, 0.5-quantile) of length estimations for each cell. From these two measures we derive three kinds of error (cf. Fig. 2.3):

all equal (unusable)

mirrored mirrored

- 1. The *within-image error* is the median of all cell IQRs. This kind of error is induced by the message and the secret key that is used for embedding. The distribution of the cell IQRs passed tests of normality.
- 2. The *between-image error* is the IQR of all cell medians. This is an image-specific error. The cell medians are rather Student-t distributed with  $\nu = 1 \dots 10$  degrees of freedom for the proposed set of attacks.
- 3. Finally, the *bias* is the median of the cell medians. This error can be precomputed for a given source of images and a particular length estimating attack.

#### 2.5 Aspects of High Performance Computing

The separation of error sources requires a large number of repetitions for each image in the test database. Apart from that, the experiments cover a number of independent

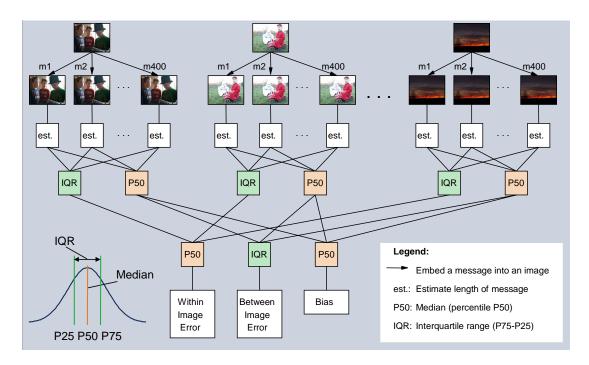


Figure 2.3: Separation of within-image error, between-image error, and bias

variations, such as five different image sizes  $(600 \times 840, 400 \times 560, 200 \times 280, 80 \times 112,$  and  $40 \times 56$  pixels), with quality q = 0.8. The medium sized images  $(200 \times 280)$  have been compressed at seven different qualities (q = 0.5, 0.6, 0.7, 0.8, 0.9, 0.95, 0.99). We applied 8 different embedding rates (1 %, 5 %, 10 %, 20 %, 40 %, 60 %, 80 %, and <math>100 % of the steganographic capacity).

We repeat the embedding with 400 random messages for over 600 images, at 11 combinations of size and quality, and at 8 embedding rates, detecting the result with 78 attacks. This results in about 1700 million length estimations.

Since 630 cell IQRs seem to follow a Gaussian distribution, the application of the median will reduce the error by a factor  $\sqrt{630}$  to a negligible level (4%). The reduction for Student-t distributed values is slightly more efficient.

If the steps to load an image, embed, and estimate the length of the embedded message required about one second, a first naive estimation will expect an execution time of 54 years (472,000 hours). Hence the goal is (a) to combine as many steps as possible, but also (b) generate executions steps that are as small as possible to allow fair scheduling without pushing up the costs for loading the runtime environment.

Let  $t_{\rm l}$  be the time to *load* an image into the system,  $t_{\rm e}$  the time for *embedding* a random message, and  $t_{\rm a}$  the time to execute an *attack* (length estimation). If we load an image only once for all related operations, and if we embed a message only once before applying the 78 attacks (cf. Figure 2.4), the necessary execution time is estimated as follows:<sup>3</sup>

$$t_{\text{total}} = 11 \cdot 630 \cdot (t_1 + (8 \cdot 400 \cdot (t_e + 78 \cdot t_a) + 78 \cdot t_a))$$

<sup>&</sup>lt;sup>3</sup>It is not necessary to repeat the attack for embedding rate 0.

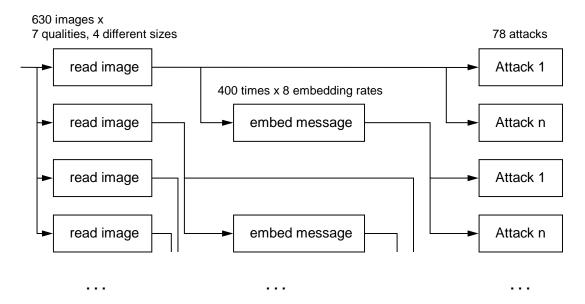


Figure 2.4: Pipeline for high performance computing

Different image sizes are weighted differently:

$$t_{\text{total}} = (1 + 0.48 + 7 \cdot 0.10 + 0.03 + 0.02) \cdot 630 \cdot (t_{\text{l}} + (8 \cdot 400 \cdot (t_{\text{e}} + 78 \cdot t_{\text{a}}) + 78 \cdot t_{\text{a}}))$$

With  $t_1$  = 0.123s,  $t_e$  = 0.335s, and  $t_a$  = 0.0577s, we expect an execution time of about 252 days. However, on a PC farm with about 2500 CPU cores at TU Dresden, this job was reduced to some hours wall time. Nevertheless, it was still challenging to aggregate and analyse all these results.

## 2.6 Applying Attacks from Spatial Domain in DCT Domain

During Phase 2 of BEST we applied spatial domain attacks in the DCT domain. In 2006 we published a paper on new histogram based attacks against Jsteg and JPhide (category attack). To be honest, we were not very confident about the contribution of this paper. Histogram based attacks have been used for a long time in the spatial domain. Such attacks (using first order statistics) have been outperformed in subsequent years by RS, SPA, WS. While the first histogram based attack (chi-square attack) has been used in both spatial domain and DCT domain, all the subsequent improvements were designed for the spatial domain only. RS, SPA, and WS were crying for a translation to the DCT domain. However, it was not obvious how to do so.

Apart from that, the chi-square attack can even be outperformed by other first order methods, e.g., dual statistics. Dual statistics have been ported to the DCT domain in the 2006 paper—a provoking result, since it was not only an improvement compared to the aged chi-square attack, but also outperforming contemporary blind attacks. This

result was followed by papers proposing blind attacks with an increased number of features that enabled a more reliable detection.

In this section we describe the functions in the swb package that have been used for the evaluation of spatial methods in the DCT domain.

#### 2.6.1 JPEG Image Handling

All these methods will operate on DCT coefficients. The function that reads a JPEG file and creates an R structure is called read.jpg. Its implementation uses the libjpeg. The

```
require(swb)  # load swb library
j <- read.jpg("lena.jpg") # store data from lena.jpg in j</pre>
```

The structure of its return value is modelled on Phil Sallee's JPEG toolbox for Matlab.<sup>4</sup> j contains the losslessly decompressed DCT coefficients along with the quantisation tables and other meta-information.

```
> str(j)
List of 14
 $ width
                : int 512
$ height
                : int 512
$ icolor
                : int 1
 $ ncomp
                : int 1
 $ jcolor
                : int 1
$ njpegcomp : int 1
 $ optimize.coding: logi FALSE
 $ comments
             : int 1
 $ coefficients :List of 1
  ..$: int [1:64, 1:4096] 47 2 1 0 0 0 0 2 0 ...
 $ qtables
                :List of 1
 ..$: int [1:8, 1:8] 5 3 3 5 7 12 15 18 4 4 ...
 $ ac.huff.tables : int 1
 $ dc.huff.tables : int 1
 $ info
                 :List of 6
  ..$ component.id : int 1
  ..$ h.samp.factor: int 2
  ..$ v.samp.factor: int 2
  ..$ quant.tbl.no : int 1
  ..$ dc.tbl.no
                 : int 1
  ..$ ac.tbl.no
                 : int 1
 $ progressive : int 0
```

 $<sup>^4</sup>$ During this project we also implemented his MB1 and MB2 methods in R and consequently needed a compatible input function

The above image "Lena" has 512 by 512 pixels. The structure of this JPEG image in j can be displayed using R's built-in function str. For JPEG files, the image is decomposed into disjoint blocks of  $8\times8$  pixels. Hence, the "Lena" image consists of  $64\times64=4096$  blocks, each of which is transformed using the DCT into 64 coefficients. Since this is a greyscale image, the file contains only one brightness channel. The coefficients of the first and only channel can be extracted by

```
coeffs <- j$coefficients[[1]]</pre>
```

#### 2.6.2 Scanpaths

<code>coeffs</code> is arranged in a  $64 \times 4096$  matrix. The first row contains all 4096 DC coefficients, rows 2...64 the AC coefficients. If we have a sorted image database with equally sized images, we can pre-calculate the index vectors for some scanpaths in advance:

```
d <- sqrt(dim(coeffs)[2]) # sqrt(4096)=64
HP <- scanpath(d, d)
HP2 <- scanpath(d, d, "hilbert2")
SL <- scanpath(d, d, "slalom")</pre>
```

There is also a function to embed a random message. The embedding method is Jsteg along a randomised embedding path. In the next example, we use 10 percent of the capacity (embedding rate 0.1) and store the resulting steganogram in s:

```
s <- jsteg.embed(coeffs, 0.1)</pre>
```

Before the set of steganalytic attacks is applied to the steganogram s, its elements can be reordered. If the matrix s is converted to a vector, the coefficients are in intra block order, i.e., the first 64 coefficients belong to the same  $8\times 8$  pixel block. Another option is the zigzag order that is used in the JPEG standard and easily achieved by permutation of the 64 columns in the matrix here. For this purpose the swb package has a global variable zigzagorder containing the appropriate permutation. R provides a function t () that transposes a matrix (turns rows into columns and vice versa). It can be used to switch from *intra* to *inter* block scan. In the transposed matrix, each 4096 elements belong to the same frequency, starting with the DC coefficients, and two consecutive coefficients belong to two different  $8\times 8$  pixel blocks.

```
x <- s
intra block scan
x <- s[zigzagorder,] zigzag order
x <- t(s) inter block scan row by row</pre>
```

In addition, the block ordering can be permuted using one of the pre-calculated index vectors for Hilbert 1 path  ${\tt HP}$ , Hilbert 2 path  ${\tt HP2}$ , or slalom  ${\tt SL}$ .

```
x \leftarrow t(s[,HP]) inter block scan Hilbert 1 path x \leftarrow t(s[,HP2]) inter block scan Hilbert 2 path x \leftarrow t(s[,SL]) inter block scan slalom path
```

we can remove the DC coefficients from each of the aforementioned scanpaths. Although DC coefficients are affected by Jsteg, some steganalytic methods perform better if only AC coefficients are used for the attack [15]. The first column of a matrix is easily deleted in R by a -1 in the column index. (s[-1,] is equivalent to s[2:64,1:4096], selecting all but the first column.)

```
x \leftarrow s[-1,] intra block scan (AC only)

x \leftarrow s[zigzagorder,][-1,] zigzag order (AC only)

x \leftarrow t(s[-1,]) inter block scan row by row (AC only)

x \leftarrow t(s[-1,HP]) inter block scan Hilbert 1 path (AC only)

x \leftarrow t(s[-1,HP2]) inter block scan Hilbert 2 path (AC only)

x \leftarrow t(s[-1,SL]) inter block scan slalom path (AC only)
```

#### 2.6.3 Histogram-Based Attacks

We will use three known histogram-based detection methods for randomised Jsteg to evaluate the reliability and precision of the methods that are proposed in the next subsection. The method of Zhang and Ping (ZP) divides the histogram into two interleaved groups [28], the attack of Yu et al. estimates the probability density function of the quantised values on a basis that is invariant to steganographic embedding [27]. The category attack (CA) compares equalising and complementary pairs of values with each other [15]. All three can estimate the length of the embedded message. For further details the reader is directed to the description in the respective original publications. As suggested in Section 2.5, reusable intermediate results are computed only once per image. Since the histogram is used in all three attacks, it is determined with a separate function jhist. The estimated embedding rate is stored in p.

```
hr <- jhist(x)
p <- zp.estimate(hr)
p <- yu.estimate(hr)
p <- ca.estimate(hr)</pre>
```

#### 2.6.4 Proposed Attacks

While there have been several proposals to use higher order statistics in the spatial domain, no specific method existed to the best of our knowledge for the DCT domain. In the following we will apply several techniques that are known from the spatial domain to (selected) DCT coefficients, which have been scanned along different paths (intrablock and interblock).

In this subsection the spatial domain attacks RS, Pairs, SPA, and WS [8, 10, 4, 7] are adapted to DCT coefficients [25]. In most cases it was easy to make the upper input limit of the attacks (usually fixed at 255 for maximum brightness) a more general parameter. For further details the reader is directed to the respective original publications.

There are two basic requirements for the input if attacks have been designed for brightness or colour intensity values.

- 1. The input should be non-negative. (However, DCT coefficients are positive and negative integers.)
- The input should not contain any values that are excluded from embedding. (However, the DCT values 0 and 1 are not used for embedding by randomised Jsteg.)

The function <code>jsteg.normalise</code> processes the sequence of coefficients to match the two requirements:

```
nc <- jsteq.normalise(x)</pre>
```

The modified attacks are applied directly to the normalised coefficients. An exception is the newly developed WB attack that also considers steganographically unused coefficients in its prediction. Hence it is applied to the coefficients in x. The JRS attack has a mask parameter. We used the two masks  $\blacksquare$  and  $\blacksquare$  in our experiments. The functions for JRS, JPairs, JSPA, JWS, and WB are used as follows (the estimated length will be stored in p):

```
p <- jrs.attack(nc, c(0,1))
p <- jrs.attack(nc, c(0,1,1,0))
p <- jpairs.attack(nc)
p <- jspa.attack(nc)
p <- jws.attack(nc)
p <- wb.attack(x)</pre>
```

## 2.7 Attacks on Some Advanced Embedding Methods

#### 2.7.1 Plus-Minus-One Embedding

One common embedding method is known as plus-minus-one embedding (PM1). Due to its simplicity it is not really "advanced." It is a rather simple derivative of the LSB replacement method. However, while the ubiquitous LSB embedding method can be detected with high reliability, PM1 requires advanced steganalytic methods to be detected.

To illustrate LSB embedding, let us consider greyscale images with pixel values in the range 0...255 as carrier medium. LSB steganography replaces the least significant bit of each pixel value in the image with the corresponding bit of the message to be hidden. An even-valued pixel will either retain its value or be incremented by one. However, it will never be decremented. The converse is true for odd-valued pixels. This asymmetry introduces a statistical anomaly in the histogram of brightness values: pairs of values (PoV) that consist of an even value and its successor tend to exhibit the same frequency if the image contains an LSB embedded message. This can be exploited for steganalytic purposes [26, 8, 4, 10, 7].

PM1, also known as LSB matching, overcomes this asymmetry: Whenever the LSB needs to be changed, the algorithm "flips a coin." If the obverse is up, the value is incremented, otherwise decremented. Although the change is never greater than one, this strategy can affect higher bits than just the LSB. For example 127 could be incremented to 128. This will change all eight bits.

PM1 embedding adjusts the least significant bit of carrier elements (e.g., pixel values) to match the message bit. Its result is compatible to LSB extraction, the embedded (and possibly encrypted) message is directly read from a sequence of least significant bits.

We looked at existing literature on PM1 detection and implemented some of those detectors. They performed less reliably than expected. Our initial goal to develop a targeted detector for PM1 seemed to move afar. We did not like to go the path to derive generic blind feature sets, but watched the development. The most promising feature set with a comprehensive evaluation was contributed by Giacomo Cancelli recently [2].

#### **Histogram Characteristic Function Center of Mass (HCFCOM)**

Harmsen and Pearlman noted that PM1 embedding acts as a low-pass filter for the histogram  $h_1$  of the image. Therefore the high frequency power in the histograms of PM1 steganograms is reduced. In other words, the FFT of the histogram  $H_1$ , also referred to as the histogram characteristic function (HCF), is likely to be significantly affected by PM1 embedding. In fact, its center of mass (COM), defined as

$$c_1(H_1) = \frac{\sum_{k=0}^{127} k |H_1(k)|}{\sum_{k=0}^{127} |H_1(k)|},$$

will be shifted toward the origin. The symmetric part of the FFT (128  $\leq k \leq$  255) is ignored here. This attack can be used with the swb package.

```
> require(swb)
> cover <- read.png("lena.png")$grey
> stego <- pm1.embed(cover)
> hcfcom.attack(cover)
[1] 8.326196
> hcfcom.attack(stego)
[1] 8.943867
```

This approach can be extended to multidimensional signals, e.g., RGB images, by using a multidimensional FFT and computing a multidimensional COM. Experimental results [14] have shown that the HCF strategy performs better with RGB images than with greyscale images.

Ker [14] suggested that this difference in performance is due to a lack of sparsity in the histogram of greyscale images. To address this issue, Ker proposed using a two dimensional adjacency histogram  $h_2(k, l)$ , which tabulates how often each pixel intensity is observed next to another.

#### Amplitude of Local Extrema (ALE)

The neighbourhood in the two dimensional adjacency histogram  $h_2(k, l)$  can be defined in four different ways: horizontal, vertical, main diagonal, and minor diagonal. [3]

#### **Wavelet Absolute Moments (WAM)**

One of the earlier feature sets that can be used for blind detection of PM1 embedding is called Wavelet Absolute Moments (WAM) [11]. These features can be extracted with the function abs.moments.attack.

```
> abs.moments.attack(stego)
Loading required package: rwt0

[1] 0.174671834 0.054164606 0.023816770 0.013684569 0.009844786 0.008620941
[7] 0.009000971 0.011008924 0.015468531 0.024383003 0.042012892 0.077243565
[13] 0.149049799 0.041256937 0.016352419 0.008503192 0.005523534 0.004302428
[19] 0.003876143 0.003922311 0.004359665 0.005234122 0.006699314 0.009043658
[25] 0.204717740 0.071622717 0.035275153 0.022848144 0.018702908 0.018731848
[31] 0.022321434 0.030907515 0.048688359 0.085451097 0.163569729 0.334627520
```

#### 2.7.2 Model-Based Steganography

Sallee modelled the marginal distribution of the DCT coefficients in JPEG images by the generalised Cauchy distribution [18]. In contrast to LSB steganography, the pairs of values are not equalised with his model-based approach. Instead, the embedded message is adapted to the generalised Cauchy distribution of each AC DCT subband in the JPEG carrier file. This adaptation is implemented as arithmetic decoding. Arithmetic coding transforms unevenly distributed bitstreams into shorter, uniform ones. Conversely, the arithmetic decoding can take a uniformly distributed bitstream (the message to be embedded) to produce a bitstream that is adapted to given probabilities of 0 and 1 according to the present generalised Cauchy distribution. In case the chosen distribution fits to the JPEG file, the first order statistics is preserved after embedding the adapted bitstream into the LSBs of the coefficients. This procedure is known as **MB1** today.

One weak property of MB1 is that block artefacts increase with growing size of the payload. **MB2** was developed to overcome this weakness [19]. It embeds the message in the same way as MB1 does but offers only half the capacity of MB1 to the user. The other half of the nonzero DCT coefficients are reserved for blockiness reduction. Early assessment by Fridrich showed good security compared to MB1 [6]. In later analyses, however, the tables have been turned [21, 17].

For the sake of simplicity we focus on the deblocking function and our most powerful feature set here. The reader is directed to our IWDW07 paper [22] for further information about other attacks on MB2 that we developed during this project.

Our key innovations behind MB2 detection are:

1. a gradient aware blockiness measure,

- 2. a classification of coefficient types regarding the blockiness reduction, and
- 3. a collection of suitable adjustment and calibration methods.

#### **Blockiness**

To reduce the blockiness, Sallee used a simple blockiness measure based on the difference of pixels at block boundaries. In Fig. 2.5a these pixels are marked grey.

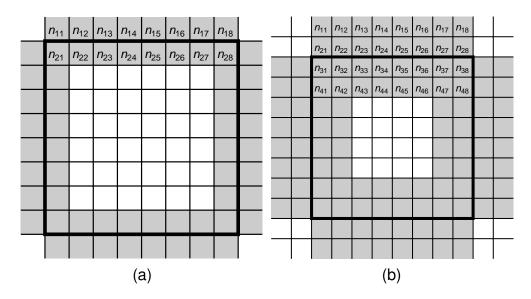


Figure 2.5: Pixels tested for (a) simple and (b) gradient aware blockiness

Each block has a north, west, south, and east boundary (with the pixels denoted by  $n_*$ ,  $w_*$ ,  $s_*$ , and  $e_*$ , respectively). The simple blockiness  $b_s$  is defined by the following sum of absolute differences:

$$b_{s} = \sum_{i=1}^{8} (|n_{1i} - n_{2i}| + |w_{1i} - w_{2i}| + |s_{1i} - s_{2i}| + |e_{1i} - e_{2i}|)$$

Corner pixels belong to several boundaries at the same time, e.g.,  $n_{28} = e_{21}$ .

This simple blockiness measure is suitable as optimisation goal to be minimised if the gradients at the block boundary were initially flat. If there is a smooth gradient across the block boundary, even in the absence of a steganographic message, e.g. a transition from dark to light, then the blockiness reduction will equalise the brightness at the boundary. This would introduce a stairstep in brightness, which is not only a visible artefact but also detectable using a **gradient aware** blockiness measure.

We developed an improved measure that respects the gradient within the block while measuring the difference at the boundary [24]. Figure 2.5b shows more pixels (marked grey) that are considered to estimate the boundary gradient. The mean of the differences of the surrounding should be equal to the difference at the boundary

(2.2). This leads to the new, gradient corrected version of the blockiness measure  $b_0$ (2.5).

$$n_{2i} - n_{3i} \approx \frac{(n_{1i} - n_{2i}) + (n_{3i} - n_{4i})}{2}$$

$$0 \approx n_{1i} - 3n_{2i} + 3n_{3i} - n_{4i}$$
(2.2)

$$0 \approx n_{1i} - 3n_{2i} + 3n_{3i} - n_{4i} \tag{2.3}$$

$$b_{g,n}^{(\lambda)} = \sum_{i=1}^{8} |n_{1i} - 3n_{2i} + 3n_{3i} - n_{4i}|^{\lambda}$$
 (2.4)

$$b_{g}^{(\lambda)} = b_{g,n}^{(\lambda)} + b_{g,w}^{(\lambda)} + b_{g,s}^{(\lambda)} + b_{g,e}^{(\lambda)}$$
 (2.5)

 $\lambda$  can be 1 or 2, depending on whether the absolute differences or a squared measure is desired.

#### **Coefficient Types**

The blockiness adjustment in MB2 leads to a differentiation of coefficients. The reason for this is that neither the coefficients used for embedding nor the coefficients that do not decrease the blockiness are altered during the blockiness adjustment. Regarding the blockiness reduction three sets of coefficient types can be defined. These three are the

**fixed coefficients** *F*, characterised by the fact that they cannot be altered because of model restriction even though they would decrease the blockiness if changed. The

different coefficients D are the ones that can be altered and if so they would decrease the blockiness. The remaining

indifferent coefficients / could be altered or not but if they are altered the blockiness would increase.

In order to categorise the coefficients a blockiness minimal image needs to be created. This is done with the function deblock\_delta in the MB2 algorithm, which is also used for blockiness reduction. It is advisable to separate the coefficient types further into the disjoint sets DC, AC1 and AC0 of coefficients. AC0 coefficients are the AC coefficients that are zero. The separation is useful because MB1 and MB2 do not use DC and AC0 coefficients for embedding. Having the blockiness minimal image at hand the indifferent coefficients can be enumerated. Doing this we get indifferent AC1, DC and AC0 coefficients. To separate the different from the fixed coefficients we need to take a closer look at the restrictions of the blockiness adjustment. MB1 and MB2 only modify coefficients within their bin in order to keep the low precision bins unchanged so the receiver of the steganogram can calculate the same model of the image. Thus, if a coefficient value needed to be altered into another low precision bin in order to decrease the blockiness it is a fixed coefficient because it must not be altered that way to keep the extraction of the model parameters correct. This again could happen with DC, AC1 and AC0 coefficients. The remaining coefficients are the different ones. They differ between the image and the blockiness minimal image but they could be altered without changing low precision bins and if so cause the blockiness to decrease.

In the case of MB1 a decrease of fixed and indifferent coefficients is expected while the different coefficients become more frequent. With MB2 the number of different coefficients should decrease because the longer the embedded message length the more coefficients are changed in order to decrease the blockiness, which can only be done by the different coefficients. Thus, the number of fixed coefficients increases, because different coefficients can only become indifferent or fixed. Since the blockiness is increasing during embedding the number of indifferent coefficients cannot increase significantly and so the fixed coefficients need to increase. Empirical analysis shows that the indifferent coefficients increase in most cases (92%) too.

#### **Adjustment and Calibration Methods**

We apply the calibration method described by Fridrich et al. [9]: The JPEG image *A* is dequantised and decompressed to spatial domain, cropped by a margin of 4 pixels from the left and from the top, then recompressed to the calibrated JPEG image *B* using the quantisation table of *A*. The cropping shifts the DCT block raster by half a block size.

Since the size of the payload embedded into the image under test is unknown, we produce the image at full embedding rate. However, in most cases the full embedding rate causes the MB2 blockiness adjustment to fail. For this reason we embed only 95 % of the capacity with both, MB1 and MB2 (separately, not on top of MB1, yielding two JPEG images).

#### **Feature Set**

The most powerful feature set  $\mathcal{F}$  that we developed includes five features, which we extract from six variants of the JPEG image. The resulting 30 features can be denoted as follows:

$$\mathcal{F} = \left\{ \begin{array}{c} b_{\mathrm{g}}^{(1)}, \\ b_{\mathrm{g}}^{(2)}, \\ |F_{\mathrm{ACO}}|, \\ |D|, \text{ and} \\ |I_{\mathrm{ACO}}| \end{array} \right\} \text{ extracted from } \left\{ \begin{array}{c} A, \\ B, \\ A+95 \% \text{ MB1 embedding,} \\ A+95 \% \text{ MB2 embedding, and} \\ B+95 \% \text{ MB1 embedding, and} \\ B+95 \% \text{ MB2 embedding.} \end{array} \right\}.$$

### **Chapter 3**

### **Results and Discussion**

#### 3.1 Possible sources of error

A meaningful comparison with previous state of the art is nearly impossible if the results are validated only by a single experiment carried out on the famous "Lena" image and the algorithm is vaguely described. However, even if the algorithm is precisely described and all parameters are listed in a table, the validity still depends on the data set that is needed to obtain the same results of the original paper.

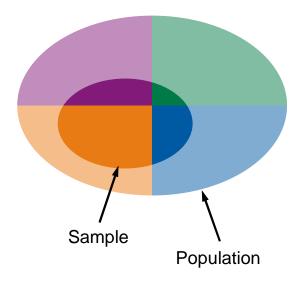


Figure 3.1: Coverage error of a sample, represented in a Venn diagram

A coverage error occurs when all members of the population do not have an equal (or known) probability of being included in the sample. If the sample is not carefully matched, the performance of a detector can considerably deviate in different evaluations.

Sampling error or estimation error is the error caused by observing a sample instead of the whole population, which is always the case in steganalysis. This type

of error decreases as the sample grows. High performance computing enables tests with larger image databases.

#### 3.2 Spatial Domain

In the original version of RS, Fridrich et al. divided the image into "disjoint groups of n adjacent pixels  $(x_1, \ldots, x_n)$ ." [8] The group size is variable. The masks that performed best in their experiments are groups of  $4 \times 1$  pixels with the mask  $\blacksquare$  as well as  $2 \times 2$  pixels with the mask  $\blacksquare$ .

Ker asked whether the groups should be disjoint or overlap [13]. In the similar case of Pairs Attack [10], where the homogeneity of colour cuts is measured, he argued conclusively that the groups of pairs must overlap so that every pair of adjacent pixels is considered. Also Dumitrescu used overlapping groups in their Sample Pairs Analysis (SPA) [4, 5]. However, Ker found lower reliability of RS for overlapping groups compared to their original disjoint version.

Ker experimented compared at least five linear masks and five square shaped masks [12]. He found that groups of  $3 \times 3$  pixels and the mask  $\blacksquare$  performed best, leading to a significant improvement compared to the standard mask  $\blacksquare$  [13]. Although we appreciate the way he set-up his image database, it was not possible for us to reproduce this improvement with our own image database. This might be a difference in implementation, since we got improved results for other  $3 \times 3$  masks, but it could also be a property of our image database that consists only of never compressed images—a typical covering error problem (cf. Section 3.1).

Standard RS determines the estimated length of the embedded message by solving a quadratic equation based on the cardinalities of regular and singular groups. RS fails if this equation has no real solution. Ker introduced a different measure that completely ignores singular groups, but is a relative difference based on the terms of regular groups only. This significantly improves the reliability of RS. Ker observed that the new mask with the larger group size provides no improvement when combined with the relative difference measure.

Only a handful of all possible masks has been tested in the literature. We guess that in terms of performance there are cross-dependencies between mask pattern, type of length estimation (relative difference or standard length estimation), and whether groups are disjoint or overlap.

We used 10,000 original images from the BOWS-2 image database. These images were taken with a number of different digital cameras. The shorter edge of the images was scaled to 512 pixels, then the center  $512 \times 512$  pixels were taken and converted to 8 bit greyscale. In contrast to the 10,000 images offered on the BOWS-2 webpage, the images used here do not contain any BOWS-2 watermark.

We considered 4 mask patterns with  $2\times 2$  pixels, 100 with  $3\times 3$  pixels, and 776 linear mask patterns with length  $2\dots 10$  pixels. The linear masks were applied three times: after scanning row by row and along a Hilbert 1 and 2 path. The attack was applied in four variants:

1. standard length estimation and disjoint groups (original),

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Table 3.1: $3 \times 3$ disjoin	nt groups (a)	with length	estimation	(b) with	relative
difference measure					

	(a)			(b)	
Rank	Mask	reliability	Rank	Mask	reliability
1		0.6789	1	H	0.7108
2		0.6759	2		0.7027
3		0.6751	3		0.7014
4	H	0.6747	4		0.7002
5		0.6709	5		0.6984
6		0.6690	6		0.6984
7		0.6659	7		0.6956
8		0.6651	8		0.6941
9		0.6597	9		0.6917
10		0.6594	10		0.6898
91		0.5486	91		0.5845
92		0.5484	92		0.5808
93		0.5362	93		0.5803
94		0.5339	94		0.5734
95		0.5323	95		0.5635
96		0.5229	96		0.5588
97		0.5168	97		0.5487
98		0.4915	98		0.5102
99		0.4737	99		0.5031
100		0.4495	100		0.4880

- 2. standard length estimation and overlapping groups,
- 3. relative difference measure and disjoint groups, and
- 4. relative difference measure and overlapping groups.

Each image was attacked twice, before and after embedding a random message that uses 3% of the capacity. This amounts to a total of almost **200 million RS attacks**, which will be presented in the following rank tables (Tables 3.1 . . . 3.7).

#### **3.2.1** Results for $3 \times 3$ Groups

Table 3.1 considers 100 mask patterns with  $3 \times 3$  pixels. All of the following rank tables present the first ten and last ten ranks. This ranking also depends on the image source and probably also on the actual embedded message. Table 3.1 presents the results for disjoint, non-overlapping groups. The attack seems to be more reliably for

"heavy" masks (more ones than zeros) than for "light." The reliability is between 1 (perfect separation) and 0 (when the area under the ROC curve is 0.5). The standard RS length estimation considers the cardinalities of both sets, the regular groups R and singular groups S to estimate the length. Since we are not interested in the precision, but only in the reliability in this case, we compare its performance to one of the relative difference  $(|R_{-m}| - |R_m|)/(|R_{-m}| + |R_m|)$ , where m is the mask used to determine if groups are regular.

The case for overlapping  $3 \times 3$  mask patterns is shown in Table 3.2. Apart from

Table 3.2:  $3 \times 3$  overlapping groups (a) with length estimation (b) with relative difference measure

	(a)				(b)	
Rank	Mask	reliability	•	Rank	Mask	reliability
1		0.7729	•	1		0.7899
2		0.7639		2	H	0.7851
3		0.7637		3		0.7811
4		0.7630		4		0.7798
5		0.7620		5		0.7798
6		0.7619		6		0.7790
7		0.7614		7	88	0.7786
8		0.7611		8	B	0.7783
9		0.7610		9		0.7779
10		0.7603		10	П	0.7772
91		0.7374		91		0.7495
92		0.7372		92		0.7484
93		0.7364		93		0.7478
94		0.7363		94		0.7426
95		0.7316		95		0.7422
96		0.7297		96		0.7405
97		0.7258		97		0.7384
98		0.7250		98		0.7246
99		0.7072		99		0.7139
100		0.7041		100		0.6693

the image margin, each pixel will be used in every position of the mask pattern here. We notice an improvement compared to the case of disjoint groups. The previous dependence between the weight of the mask and its reliability disappeared in the overlapping case. Also the difference between the two measures (standard RS length estimation; relative difference of R) is less distinctive. The mask  $\mathbb{H}$  selected by Ker [12] performed rather poor here. However, we verified the dominance of  $3 \times 3$  masks, which performed best in our experiments.

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#### **3.2.2** Results for $2 \times 2$ Groups

There are only four  $2 \times 2$  mask patterns (apart from isomorphic variants). Table 3.3 presents the ranking for their disjoint application, and Table 3.4 for overlapping. Again,

Table 3.3:  $2 \times 2$  disjoint groups (a) with length estimation (b) with relative difference measure

	(a)			(b)	
Rank	Mask	reliability	Rank	Mask	reliability
1		0.6729	1		0.6807
2		0.6532	2		0.6590
3		0.6153	3		0.6572
4		0.5692	4		0.5802

Table 3.4:  $2 \times 2$  overlapping groups (a) with length estimation (b) with relative difference measure

(a)				(b)	
Rank	Mask	reliability	Rank	Mask	reliability
1		0.7493	1		0.7354
2		0.7482	2		0.7248
3		0.7080	3		0.7184
4		0.7046	4		0.6697

like in the  $3 \times 3$  case, the attack works more reliably in the overlapping case. Also the dependence between heavy masks and stronger reliability could be true. However, while in the disjoint case the relative difference seems to be slightly more reliable compared to the standard RS length estimation, this is reversed in the overlapping case. Surprisingly, the heavy mask pattern performed better than the proposed by Fridrich et al. [8].

#### 3.2.3 Results for Groups with Linear Masks

This part of experiments considered linear mask that are usually applied row by row. We also applied these masks along the two different versions of the Hilbert curve (cf. Fig. 2.2). Indeed, the optimum group size for our image set was slightly above 4 pixels, but below 9 pixels. However, the desired gain by using recursive scanpaths was not found. The usual row by row scan exceeded the reliability of Hilbert paths, regardless the masks are applied disjoint or overlapping, and regardless the measure was standard length estimation or relative difference of *R*. Table 3.5 shows the results

Table 3.5: Linear disjoint groups,	scanned (a)	row by row	(b) along a Hilbert 1
curve (c) along a Hilbert 2 curve			

	(a) (b)			(c)				
Rank	Mask	reliability	Rank	Mask	reliability	Rank	Mask	reliability
1		0.6956	1		0.6548	1		0.6790
2		0.6922	2		0.6527	2		0.6746
3		0.6879	3		0.6521	3		0.6744
4		0.6865	4		0.6511	4		0.6739
5		0.6860	5		0.6506	5		0.6680
6		0.6844	6		0.6492	6		0.6666
7		0.6833	7		0.6490	7		0.6666
8		0.6830	8		0.6456	8		0.6663
9		0.6828	9		0.6438	9		0.6647
10		0.6808	10		0.6433	10		0.6645
767		0.4913	767		0.4678	767		0.4790
768		0.4894	768		0.4659	768		0.4789
769		0.4871	769		0.4578	769		0.4752
770		0.4834	770		0.4565	770		0.4738
771		0.4760	771		0.4537	771		0.4734
772		0.4735	772		0.4441	772		0.4693
773		0.4500	773		0.4308	773		0.4229
774		0.4366	774		0.4181	774		0.4147
775		0.4255	775		0.4050	775		0.4126
776		0.3895	776		0.3879	776		0.4003

for the particular scanpaths, standard RS length estimation, and disjoint application of groups. Again we can see an advantage for heavy mask patterns. There is no significant difference between the favourite that was experimentally determined by Fridrich and the best reliability that we found in our experiments for linear masks and conventional row by row scanpath.

Table 3.6 presents the reliability for the relative difference measure of *R*. This is a minor improvement against the standard RS length estimation.

We yielded a much stronger gain when groups overlap. As mentioned earlier the other scanpaths performed worse and are omitted in Table 3.7. We noticed a marginal performance gain for the relative difference. Interestingly, there is a characteristic pattern at the top that includes a repeated sequence 0-1-0-1. Overlapping linear groups, scanned row by row, evaluated by relative difference yield the second best reliability, right after the  $3\times3$  pixel masks.

Table 3.6: Linear groups with relative difference measure, scanned (a) row by row (b) along a Hilbert 1 curve (c) along a Hilbert 2 curve

	(a)		(b)				(c)		
Rank	Mask	reliability	Rank	Mask	reliability	Rank	Mask	reliability	
1		0.7139	1		0.6719	1		0.7068	
2		0.7111	2		0.6704	2		0.7007	
3		0.7093	3		0.6682	3		0.6934	
4		0.7084	4		0.6681	4		0.6920	
5		0.7083	5		0.6668	5		0.6911	
6		0.7065	6		0.6663	6		0.6908	
7		0.7064	7		0.6651	7		0.6897	
8		0.7058	8		0.6638	8		0.6895	
9		0.7056	9		0.6636	9		0.6881	
10		0.7044	10		0.6635	10		0.6879	
767		0.5297	767		0.5009	767		0.5176	
768		0.5261	768		0.4971	768		0.5175	
769		0.5212	769		0.4934	769		0.5150	
770		0.5179	770		0.4923	770		0.5128	
771		0.5157	771		0.4917	771		0.5069	
772		0.5148	772		0.4799	772		0.5017	
773		0.4931	773		0.4681	773		0.4659	
774		0.4765	774		0.4559	774		0.4549	
775		0.4679	775		0.4424	775		0.4524	
776		0.4296	776		0.4228	776		0.4388	

### 3.3 DCT Domain

#### 3.3.1 Impact of Image Size

For the comparison in Fig. 3.2 we downloaded 630 large TIFF images ( $1500 \times 2100$  pixels) from the NRCS database [16]. These images have been downsized (using pnmscale) to five different sizes ( $600 \times 840$ ,  $400 \times 560$ ,  $200 \times 280$ ,  $80 \times 112$ , and  $40 \times 56$  pixels), converted to greyscale, and JPEG compressed with quality q = 0.8. For the comparison in Fig. 3.2, we selected images with an embedding rate of 1 % of the capacity. In abundance of results we show only the best representative for each fundamental estimation method. This selection is based on the between-image error at an embedding rate of 1 % of the capacity ( $600 \times 840$ , JPEG quality 0.8). The between-image error is related to the detection power, since this kind of error dominates at low embedding rates, while the within-image error is negligible and the overall bias in the estimation is without influence to reliability. The representatives are the JPairs attack [25] applied to AC coefficients only, scanned interblock along a Hilbert 1 path, the JSPA attack [25] applied to AC coefficients only, scanned interblock row by row, the attack by Zhang and Ping [29], the category attack by Lee et al. [15],

Table 3.7: Overlapping linear groups, scanned row by row (a) with length estimation (b) with relative difference measure

	(a)				(b)	
Rank	Mask	reliability	•	Rank	Mask	reliability
1		0.7695	•	1		0.7732
2		0.7689		2		0.7732
3		0.7689		3		0.7732
4		0.7689		4		0.7732
5		0.7689		5		0.7732
6		0.7689		6		0.7726
7		0.7643		7		0.7726
8		0.7643		8		0.7726
9		0.7643		9		0.7726
10		0.7643		10		0.7726
767		0.7030		767		0.7090
768		0.7023		768		0.7090
769		0.7023		769		0.7072
770		0.7023		770		0.7072
771		0.7023		771		0.7072
772		0.7022		772		0.7072
773		0.7022		773		0.7072
774		0.7022		774		0.7072
775		0.7022		775		0.7072
776		0.7022		776		0.7071

the JWS attack [25] applied to AC coefficients only, scanned interblock row by row, the WB attack [25] applied to both AC and DC coefficients, scanned interblock along a slalom path, the JRS attack [25] with mask (0,1) applied to AC coefficients only, scanned intrablock in zigzag order, and finally the attack by Yu et al. [27]. As expected, the between-image error decreases with increasing image size. While the between-image error of the JPairs attack is smallest for images larger than  $200 \times 280$ , there are more suitable candidates for thumbnail images  $(40 \times 56)$ , where WB attack, category attack, and the attack by Yu et al. do a better job. Interestingly, the JWS attack, which is closely related to the front runner WB, has the biggest problem with thumbnail images. One possible explanation could be the increased variance in downscaled images.

## 3.3.2 Impact of JPEG Quality

The medium sized images  $(200 \times 280)$  have been compressed at seven different qualities (q = 0.5, 0.6, 0.7, 0.8, 0.9, 0.95, 0.99). Again, we only selected images with an

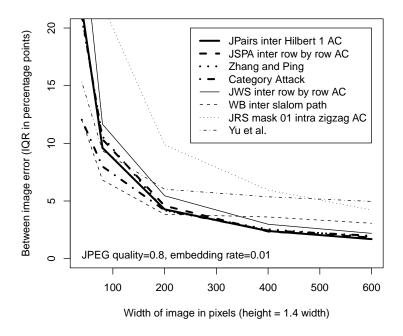


Figure 3.2: Between-image error as a function of the image size

embedding rate of 1 % of the image capacity here. Figure 3.3 shows the between-image error as a function of the JPEG quality. Except for the category attack and the attack by Zhang and Ping, this kind of error is increased for qualities above 0.9.

## 3.4 Steganalysis of Model-Based Steganography

For our experimental evaluation we downloaded 2300 large TIFF images (2100  $\times$  1500 pixels) and 630 further images (1500  $\times$  2100 pixels) from the NRCS database [16]. These images have been downsized to 840  $\times$  600 pixels by pnmscale's default method, converted to greyscale, and JPEG compressed with quality q=0.8. The rather small payload was 0.02 bits per nonzero coefficient for both, MB1 and MB2, using the original implementation by Phil Sallee.

The training was based on 2300 images and the classification was done by LDA based on 630 images. Table 3.8 shows the results, where "23" represents the attack by Fridrich with 23 DCT features [6], "274" the one with 274 mixed DCT and Markov features by Pevný and Fridrich [17], and "324" the set by Shi et al. [21] with the 324 Markov features. "81" and "193" is a segmentation of the 274 features, where "81" contains the calibrated averaged Markov features.  $\mathcal F$  is the proposed feature set, defined in Eq. 2.6 on page 29.

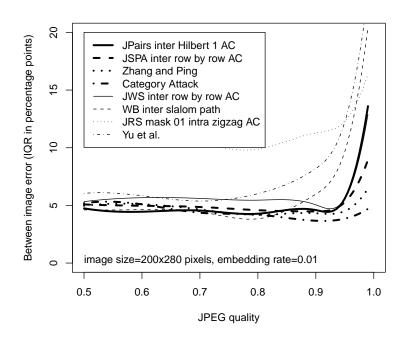


Figure 3.3: Between-image error as a function of the JPEG quality

Table 3.8: Detection reliability for feature combinations

		Additional	Number of features				
		features	23	324	274	81	193
Reliability	MB1	_	0.181	0.597	0.698	0.585	0.516
	IVIDI	${\cal F}$	0.596	0.793	0.791	0.743	0.708
	MB2	_	0.187	0.659	0.759	0.666	0.518
		${\cal F}$	0.873	0.924	0.937	0.919	0.909
False alarm rate at 50 % detection	MB1		0.348	0.133	0.077	0.133	0.136
	IVIDI	${\cal F}$	0.140	0.050	0.041	0.060	0.080
	MB2	_	0.370	0.105	0.042	0.086	0.147
		${\mathcal F}$	0.019	0.008	0.006	0.011	0.011

# **Chapter 4**

## **Conclusions**

In our comprehensive study of the RS attack we found that there is no single mask pattern that works best in all cases. Instead we found inter-dependencies between all parameters, e.g. mask shape (squared or linear), application (disjoint or overlapping groups), and the measure used for detection (length estimate or relative difference). The surprising result is that not a single change of a particular parameter makes an appreciable difference. However, if we change several of them at once, the detection power of RS grows significantly for mask patterns that did not receive attention in the literature so far, despite the fact that LSB replacement in spatial domain images is probably the best understood steganalysis problem today.

We also contributed to the "harder" problems in steganalysis, namely the detection of model-based steganography MB2. Our proposed feature set is based on coefficient types that can be derived from the blockiness adjustment of MB2. Used in combination with existing blind feature sets the false positive rate is reduced from 10 % to 1 % for a very low embedding rate (0.02 bits per non-zero coefficient).

We developed a methodology to apply higher order steganalytic attacks from the spatial domain in the transformed domain. Based on 1700 million attacks, we evaluated the performance of the proposed attacks under diverse image parameters (size, quality), and determined the most advisable schemes (WB for small images, JPairs for larger ones).

The implementation of our steganographic workbench will almost certainly be extended during the next years. It will shorten the development and evaluation cycles for new attacks and new steganographic embedding methods.

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# List of Symbols, Abbreviations, and Acronyms

**AC** alternating current (DCT subbands with nonzero video frequency)

**ALE** amplitude of local extrema, a feature set

BEST Better Steganalysis, project title
BEST2 Better Steganalysis, phase 2

BIE between image error
CA Category Attack
COM center of mass

**CPU** Central Processing Unit

**DC** direct current (DCT subbands with zero video frequency)

DCT discrete cosine transformFFT fast Fourier transformGPL General Public Licence

**HCF** histogram characteristic function

IQR inter quartile range

JPairs Pairs attack adopted to JPEG domain

JPEG Joint Photographic Experts Group (lossy image compression format)

JRS RS attack adopted to JPEG domain
JSPA SPA adopted to JPEG domain

JWS WS attack adopted to JPEG domain

LSB Least Significant Bit

LTSB Least Two Significant Bits

MB1 Model-based steganography (initial variant)MB2 Model-based steganography (with deblocking)

**Pairs** Pairs attack

**PM***k* Plus-Minus-*k* embedding

PNG Portable Network Graphics (lossless image compression format)

**ROC** Receiver Operating Characteristic

RS An attack based on *regular* and *singular* groups of pixels

SPA Sample Pairs analysis attack by Dumitrescu

**sqrt** square root

**WAM** wavelet absolute moments, a feature set

**WB** weighted non-steganographic boundary attack

WIE within image error
WPC Wet Paper Codes

WS An attack based on an estimation weighted statistics from surrounding

pixels by Fridrich and Goljan

Yu attack to Jsteg-like embedding using modified Cauchy model fitting by

Yu et al.

**ZP** simple attack by Zhang and Ping