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# REPLICATING ALGORITHM AN ADAPTIVE ALGORITHM FOR GREY IMAGE EDGE DETECTION BASED ON GREY CORRELATION ANALYSIS

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**Abstract:**

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# 1 Introduction

The purpose of this mini project in Image Processing and Computer Vision.

## 2 Method of analysis

First the dataset is analysed for stochastic transitivity violations, this covers:

Weak stochastic transitivity (WST) which holds if:

$$p_{ac} \geq 0.5, \text{ (violation if } p_{ac} < 0.5)$$

Moderate stochastic transitivity (MST) which holds if:

$$p_{ac} \geq \min(p_{ab}; p_{bc}), \text{ (violation if } p_{ac} < \min(p_{ab}; p_{bc}))$$

Strong stochastic transitivity (SST) which holds if:

$$p_{ac} \geq \max(p_{ab}; p_{bc}), \text{ (violation if } p_{ac} < \max(p_{ab}; p_{bc}))$$

This was tested using a script written in MATLAB which tested for all three types of transitivity violations. Generally there are no set rules for what too many violations are, but they give an indication of whether you should or should not continue with the making of your model. The first model used in this study was a BTL model which is one of the probabilistic choice models that the study group has tried before, but it was found that the fit was poorly and had a p-value of 0.0055 where the minimum value should be at least 0.1. The model used instead is preference tree, which is good for data with more than one attribute. Some different tree structures have been tried, and the one with the greatest fit is chosen, and the MATLAB script used is shown following page, where M is equal to the dataset shown in Table 1.

```

1 %Preference tree opgave
2 % Perceived health risk of drug
3 % N=48
4 M=[
5 0 28 35 10 4 7
6 20 0 18 2 0 3
7 13 30 0 3 1 0
8 38 46 45 0 1 17
9 44 48 47 47 0 44
10 41 45 48 31 4 0];
11
12 %Rusmidler:
13 %Alc Tob Can Ecs Her Coc
14
15 P = M/48;
16 WST=0;
17 MST=0;
18 SST=0;
19 it=0;
20
21 for a=1:6
22     for b=1:6
23         for c=1:6
24             it = it+1;
25             if a~=b && b~=c && a~=c
26                 if P(a,b)>=0.5 && P(b,c)>=0.5
27                     if P(a,c)<0.5
28                         WST=WST+1;
29                     end
30                     if P(a,c)<min(P(a,b),P(b,c))
31                         MST=MST+1;
32                     end
33                     if P(a,c)<max(P(a,b),P(b,c))
34                         SST=SST+1;
35                     end
36                 end
37             end
38         end
39     end
40 end
41
42 A = {[1];[2 9];[3 9];[4 7 8 9];[5 7 8 9];[6 8 9]};
43

```

```

44 [pModel, chistat, u, lL_eba, lL_sat, fit, cova] = fOptiPt(M,A);
45 pModel=pModel*10e-15;
46
47 err = 1.96*sqrt(diag(cova));
48 X = [1 2 3 4 5 6];
49 errorbar(u, err, 'o');
50 %set(gca, 'xtick', [1:6], 'xticklabel', {'Alc', 'Tob', 'Can', 'Ecs', '
    Her', 'Coc'});

```

### 3 Hypothesis

When conducting an experiment it is important to specify what exactly we want to measure. As it has already been mentioned, the focus of this study is the perception of the surface of the five dices in comparison with each other. As such, the H0 hypothesis can be formulated as:

*The surface of the dice are perceived equally for each die.*

Through hypothesis testing, it will take a p-value greater than 0.05 to accept this assumption. If the P-value is lesser than 0.05 there will be strong evidence against the H0 hypothesis, which means that the dices would be perceived differently.

### 4 Results

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### 5 Analysis

After constructing the script and executing it, the following findings were made:

0 WST

1 MST

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<sup>1</sup>FiXme Note: Write Results

## 5 SST

A p-value of 0.1414.

The p-value is above 0.1 which indicates that the preference tree model is not significant worse than a theoretical perfect model. In a preference tree model there is both scale values for the stimuli presented in the study, and the attributes the tree is made with. The scale values of the attributes is used to illustrate the perceived difference of how big a health risk the substance is, eg. longer branch bigger perceived risk. On Figure 1 the preference tree is illustrated with the scale values, and the numbers at each branch is a indicator for the belonging attribute. The green line is modified so it is three times shorter than it normally would be the red is also modified, but it is shortened ten times. This is done to ensure that all the trees proportions are viewable at the same time.

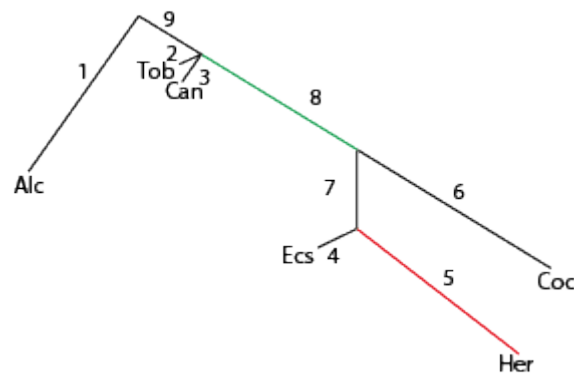
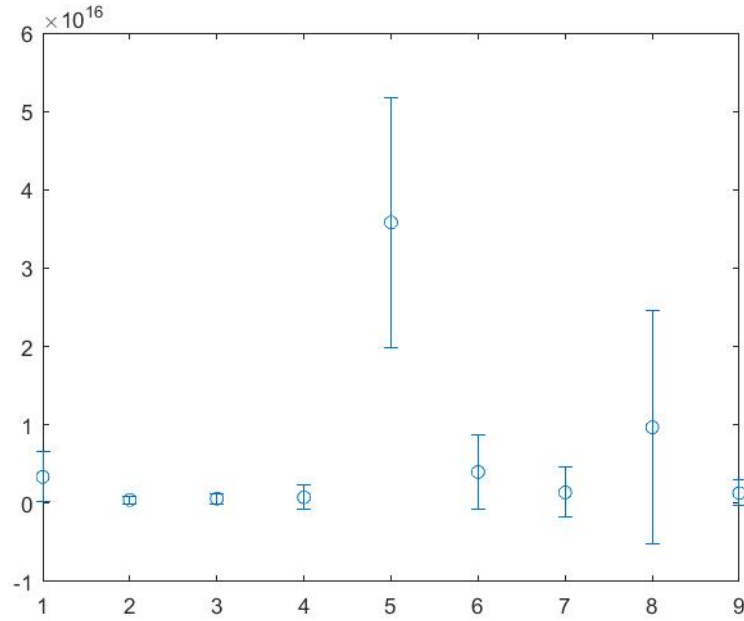


Figure 1

From the tree it's clear that the different attributes have different effects on how big the perceived health risk is. With these results it's found interesting to investigate how the attributes compare with each other. To compare the attributes the scale values for every branch of the tree is plotted, with the belonging 95% confidence interval, see Figure 2.



**Figure 2**

It's found that the only attribute that is significant different from the others is the cocaine, which is the one that have cocaine at the end.

## 6 Discussion

The Tree illustrated at Figure 1 uses attributes at every branch, which is compared in Figure 2. The structure of the tree is build up so the stimuli are sorted into different categories of attributes. With alcohol being the only drinkable substance it have it's own attribute, which is number 1. The other substances goes from the attribute number 9, which is the non drinkable substances. From 9 the attributes 2 and 3 goes down to tobacco and cannabis. Attribute 2 indicates social acceptable and legal substance, while attribute 3 indicates social acceptable but illegal substance. The 8'th attribute indicates non-social acceptable and illegal substances. The 6'th attribute indicates that the substance is a common party drug, hence cocaine, while the 7'th attribute is for more hard drug, where 4 goes to ecstasy is and 5 goes to heroin.

## 7 Conclusion

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