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Extended Mosaic and Association Plots for Visualizing (Conditional) Independence

Achim Zeileis

David Meyer

Kurt Hornik

- ❄ The independence problem in 2-way contingency tables
 - ❖ Standard approach: χ^2 test
 - ❖ Alternative approach: max test
- ❄ Visualizing the independence problem
 - ❖ Association plots
 - ❖ Mosaic plots
- ❄ Extensions
 - ❖ Visualization & significance testing
 - ❖ HCL instead of HSV colors
 - ❖ Multi-way tables and conditional independence
 - ❖ Implementation in `grid`
- ❄ The `vcd` package

Standard approach:

- ❄ Analyze the relationship between two categorical variables based on the associated 2-way contingency table.
- ❄ Measure the discrepancy between observed frequencies $\{n_{ij}\}$ and expected frequencies under independence $\{\hat{n}_{ij}\}$ by the Pearson residuals:

$$r_{ij} = \frac{n_{ij} - \hat{n}_{ij}}{\sqrt{\hat{n}_{ij}}}.$$

- ❄ Use the Pearson X^2 statistic for testing:

$$X^2 = \sum_{ij} r_{ij}^2,$$

which has an asymptotic χ^2 distribution.

Alternative approach(es):

❄ There are many conceivable functionals $\lambda(\cdot)$ which lead to reasonable test statistics $\lambda(\{r_{ij}\})$.

❄ In particular:

$$M = \max_{ij} |r_{ij}|.$$

Then, every residual exceeding the critical value c_α violates the null hypothesis at level α .

❄ Instead of relying on unconditional limiting distributions, perform a permutation test, either by simulating or computing the conditional permutation distribution of $\lambda(\{r_{ij}\})$.

The independence problem



Relationship between hair color and eye color among 328 female students:

Hair color	Eye color				Total
	Brown	Blue	Hazel	Green	
Black	36	9	5	2	52
Brown	81	34	29	14	158
Red	16	7	7	7	37
Blond	4	64	5	8	181
Total	137	114	46	31	328

$$X^2 = 112.30 \quad p = 0$$

$$M = 6.76 \quad p = 0$$

The independence problem

Home and away goals in the Bundesliga in 1995:

Home goals	Away goals						
	0	1	2	3	4	5	6
0	26	16	13	5	0	1	0
1	19	58	20	5	4	0	1
2	27	23	20	5	1	1	1
3	14	11	10	4	2	0	0
4	3	5	3	0	0	0	0
5	4	1	0	1	0	0	0
6	1	0	0	1	0	0	0

$$X^2 = 46.07$$

$$p = 0.121$$

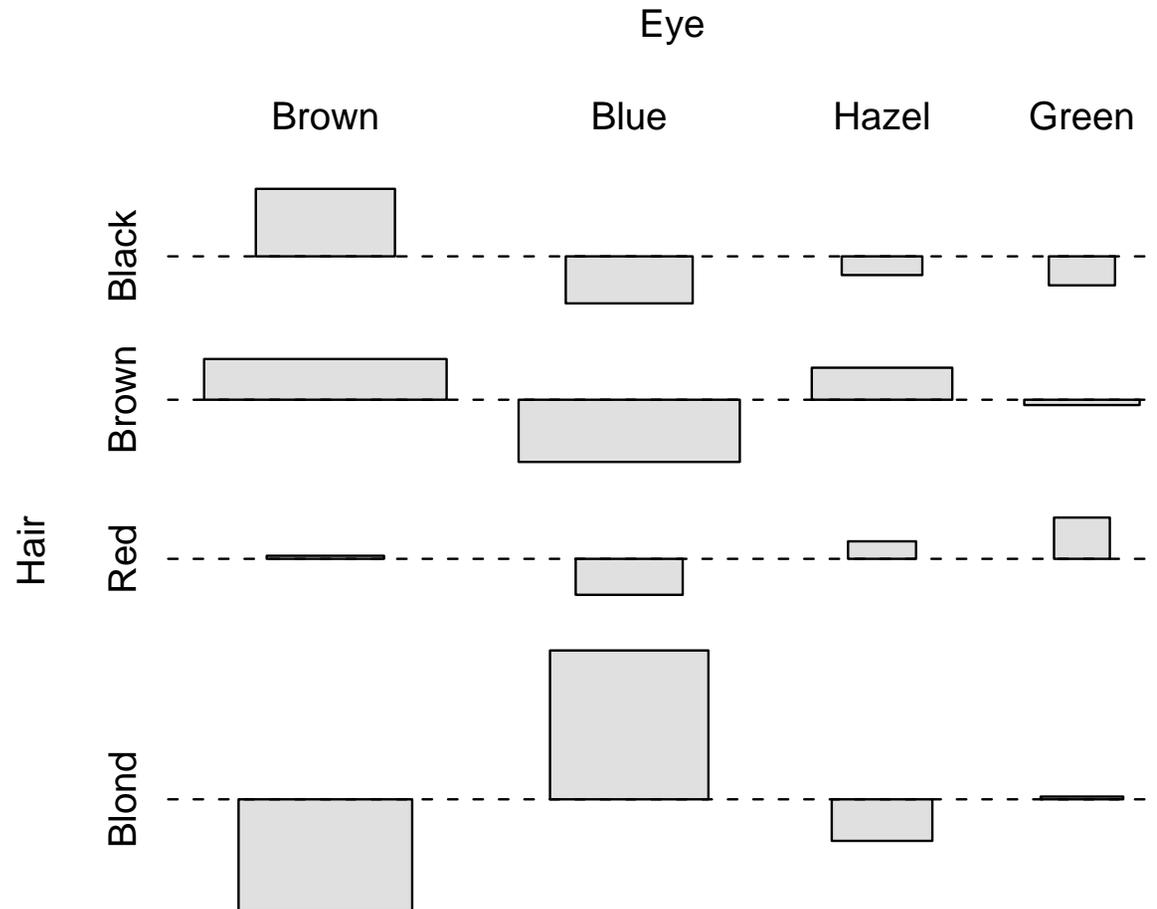
$$M = 2.87$$

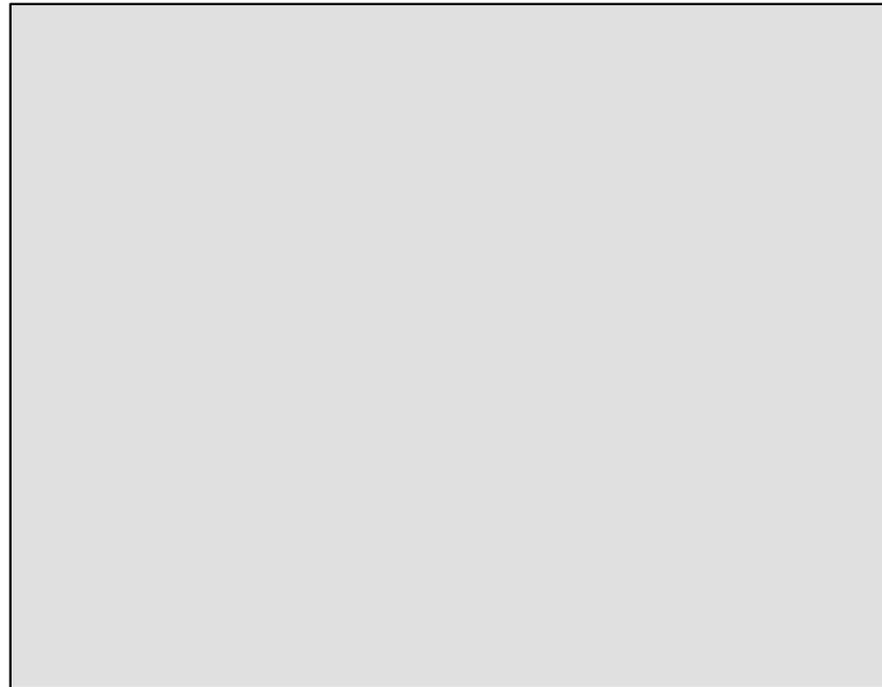
$$p = 0.355$$

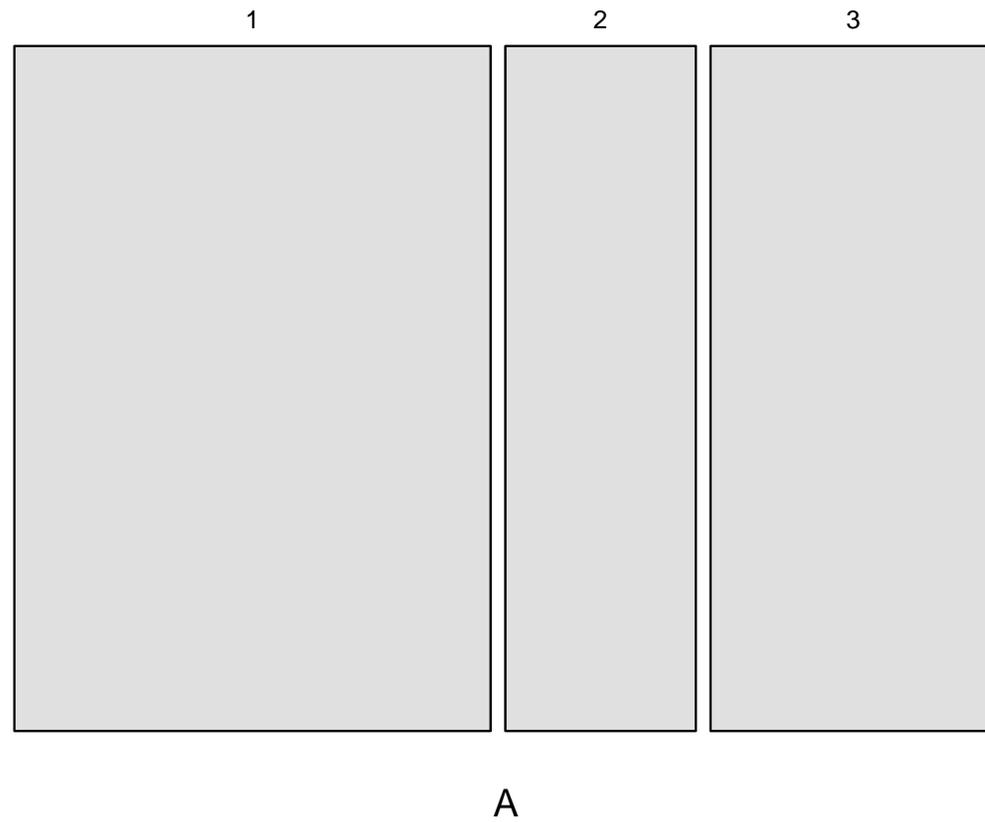
Association plot: display for the Pearson residuals $\{r_{ij}\}$ and the raw residuals $\{n_{ij} - \hat{n}_{ij}\}$ in an rectangular array.

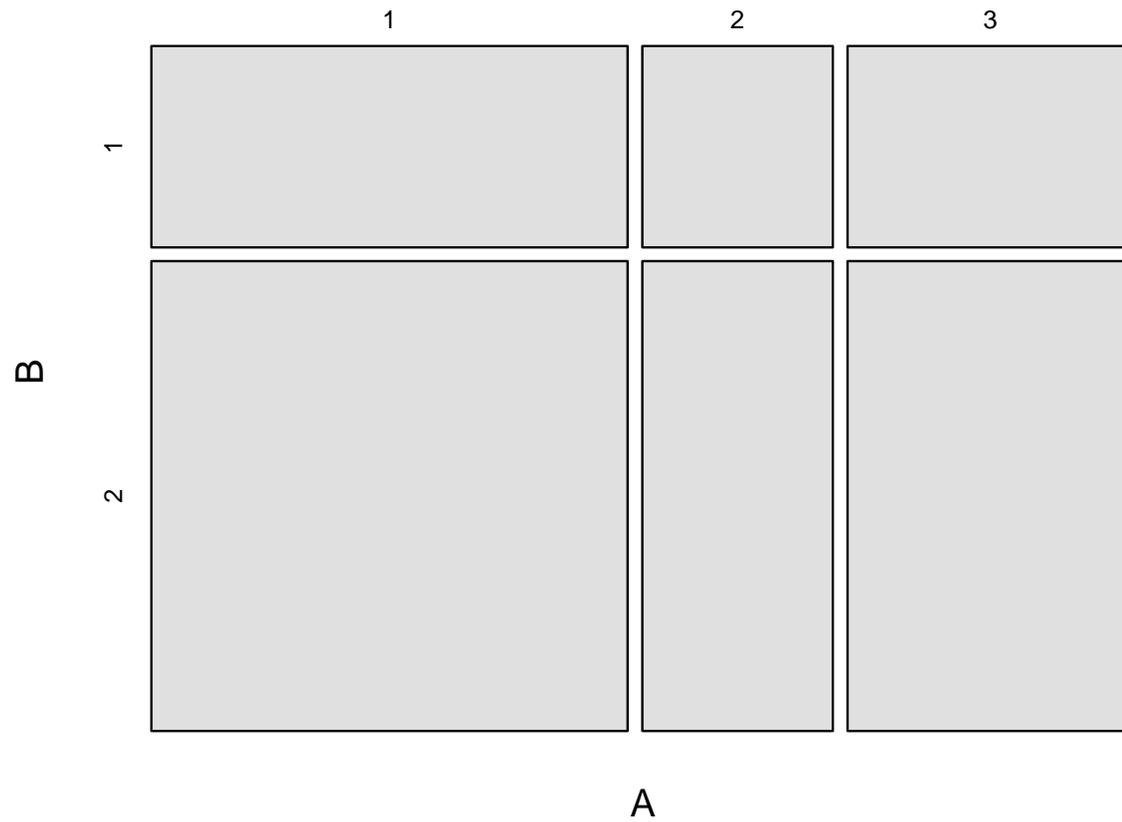
Mosaic plot: display in which the sizes of the mosaic tiles is proportional to the observed frequencies $\{n_{ij}\}$.

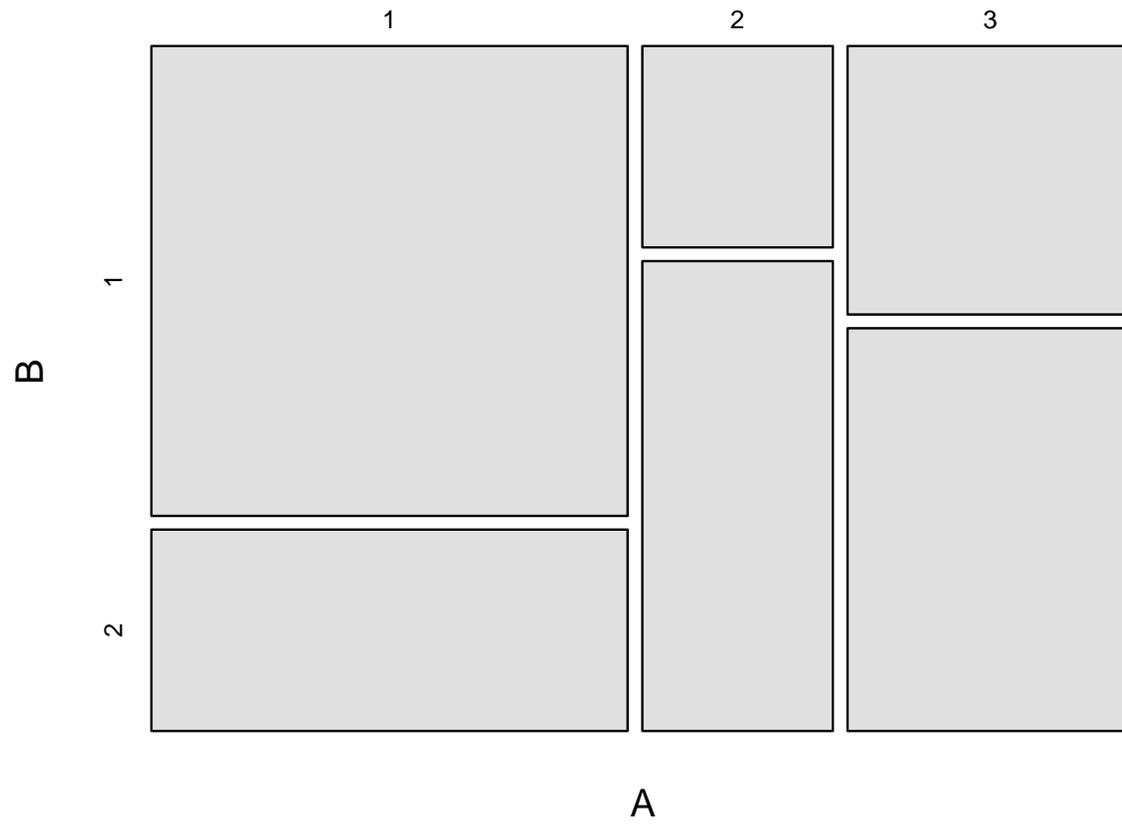
Visualization

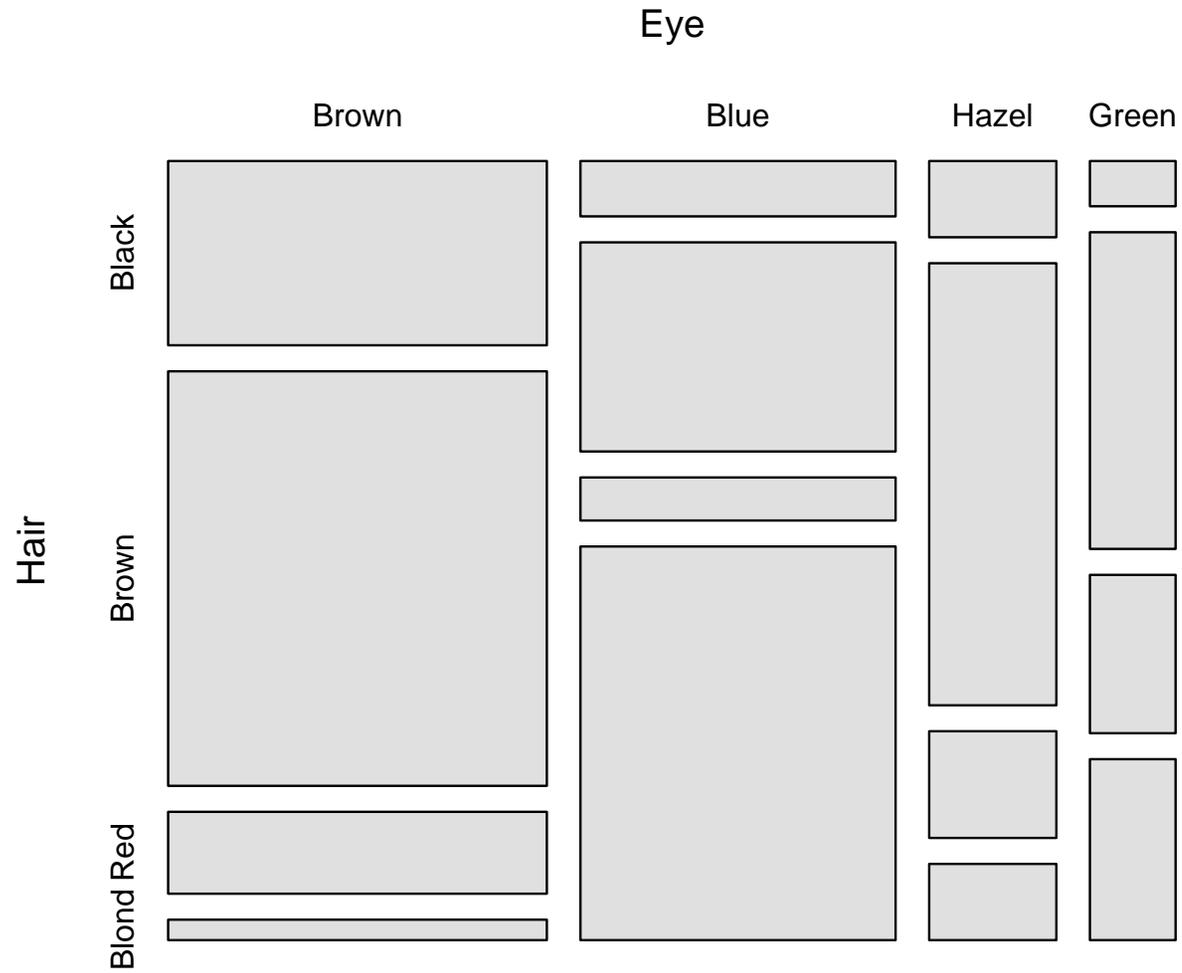












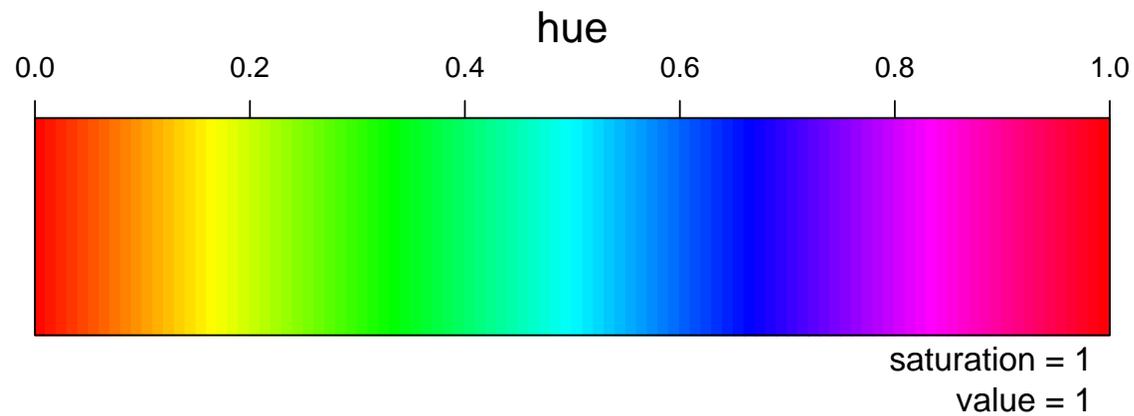
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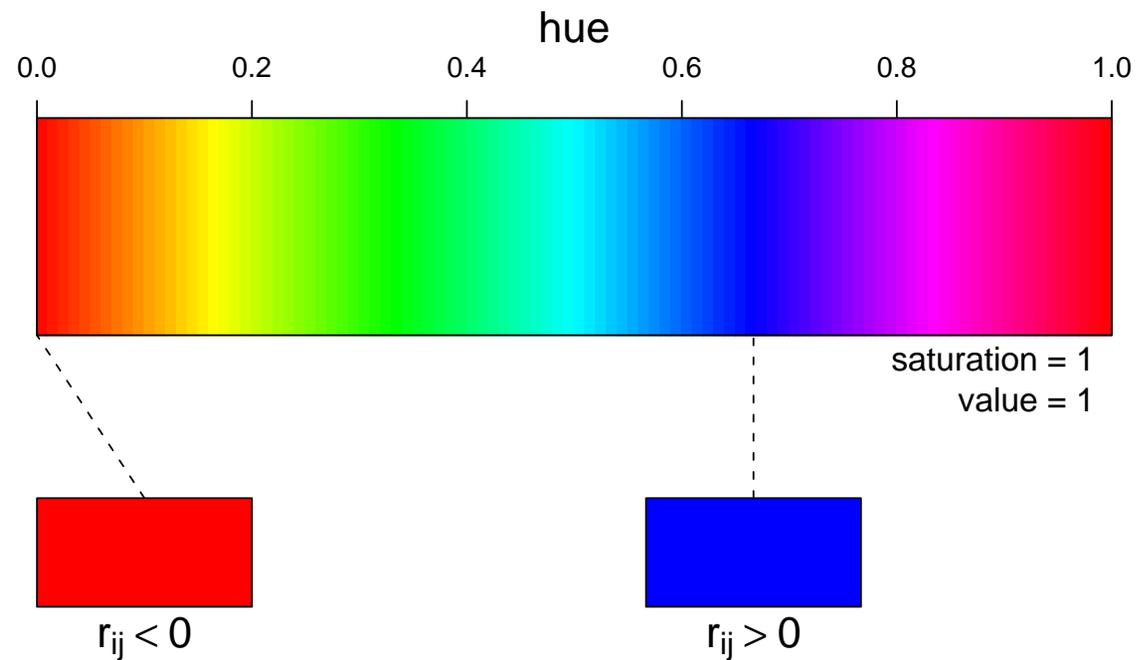
In R these are implemented based on HSV colors.

The HSV color space is one of the most common implementations of color in many computer packages. Hue, saturation and value range in $[0, 1]$.

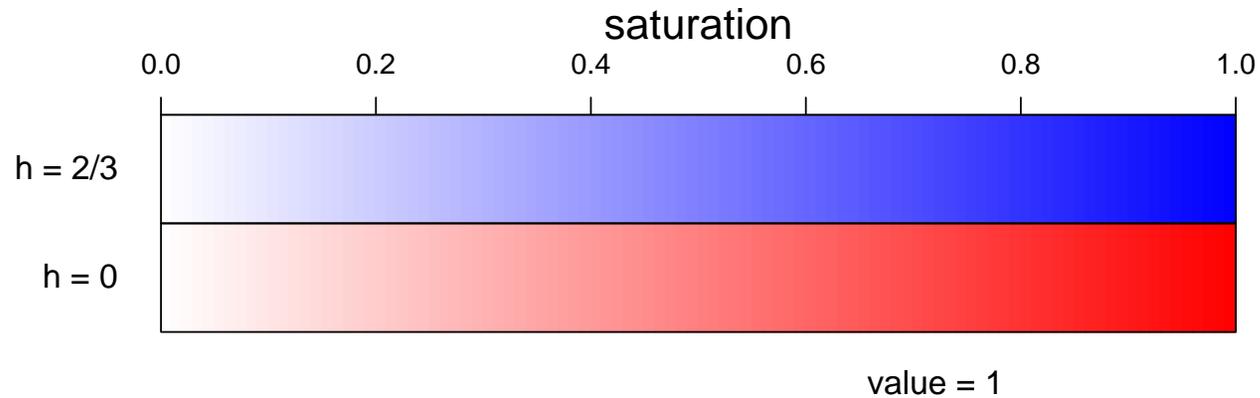
The hue is typically used to code the *sign* of the residuals.



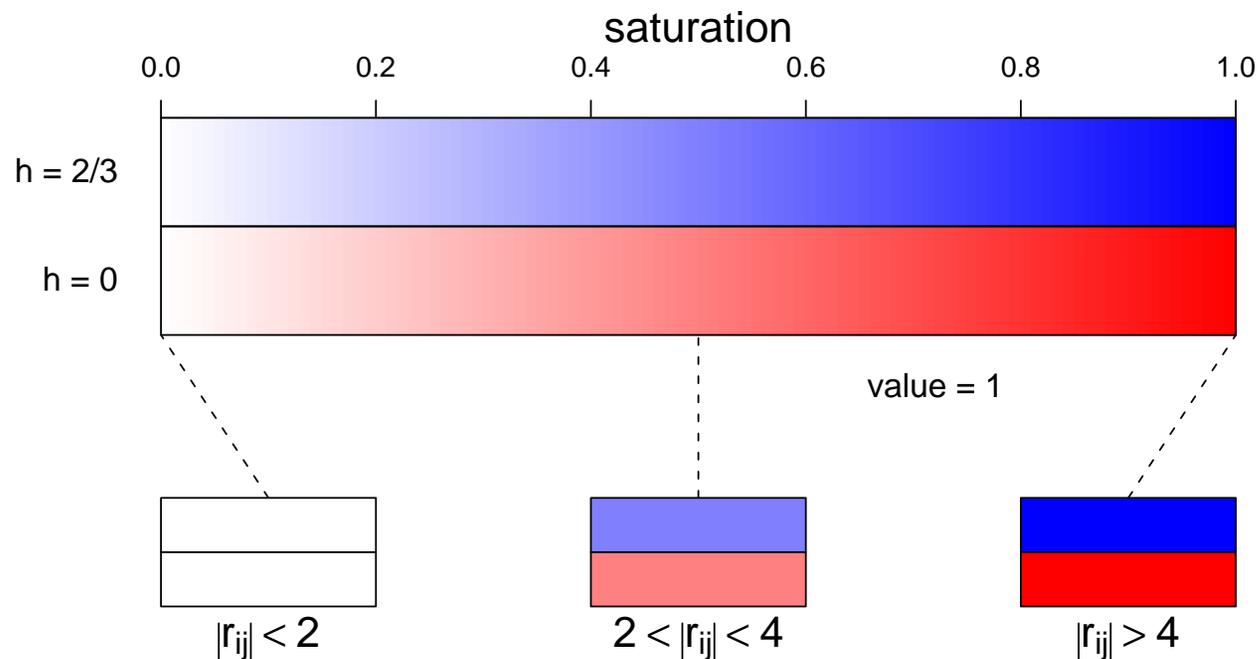
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Friendly's extended mosaic displays use the saturation to code the *absolute size* of the residuals.

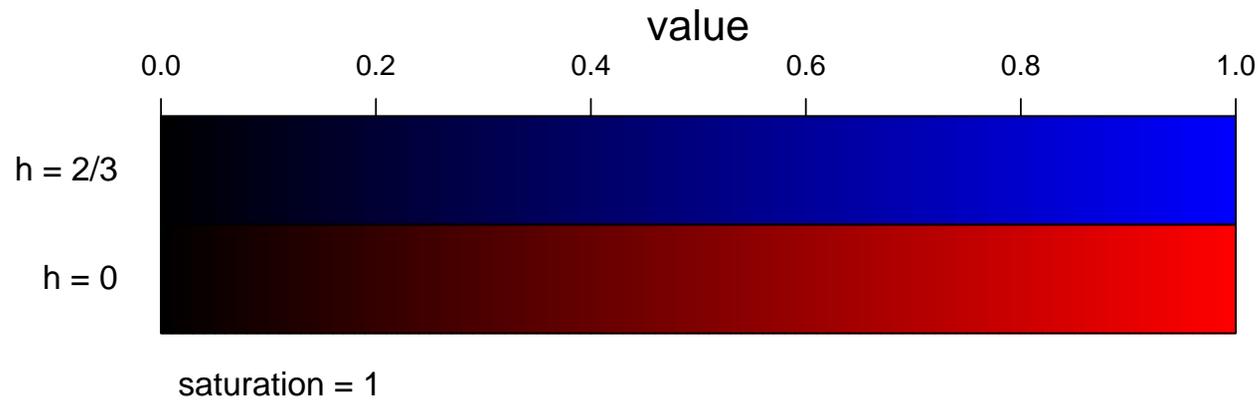


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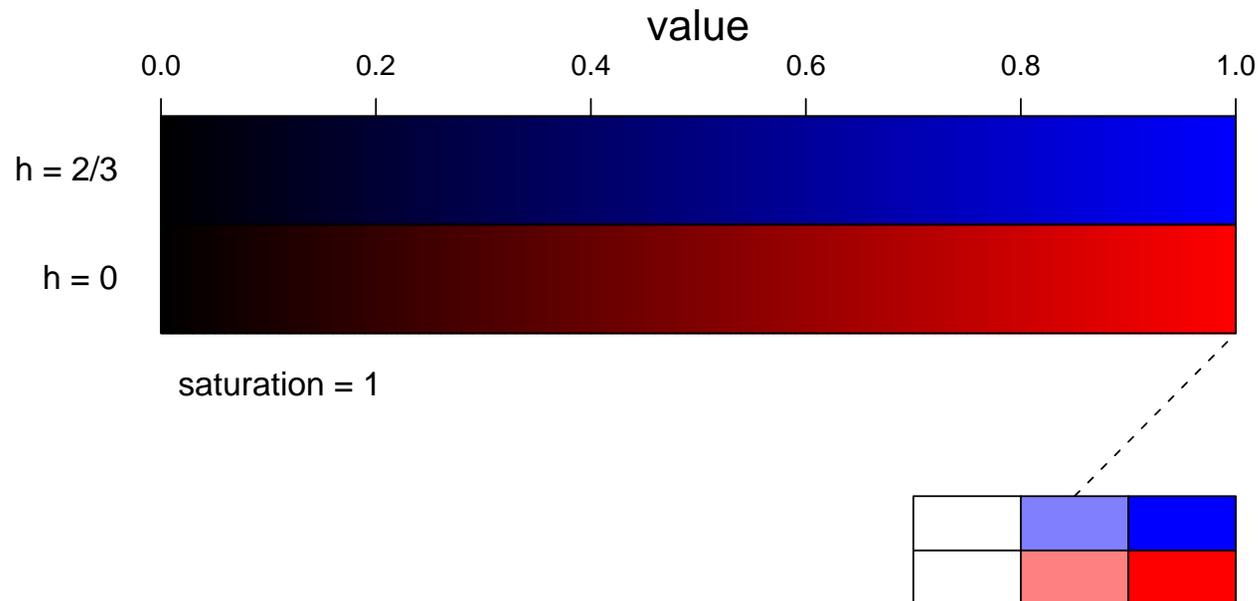
HSV colors

Value is currently not used for coding, always set to 1.

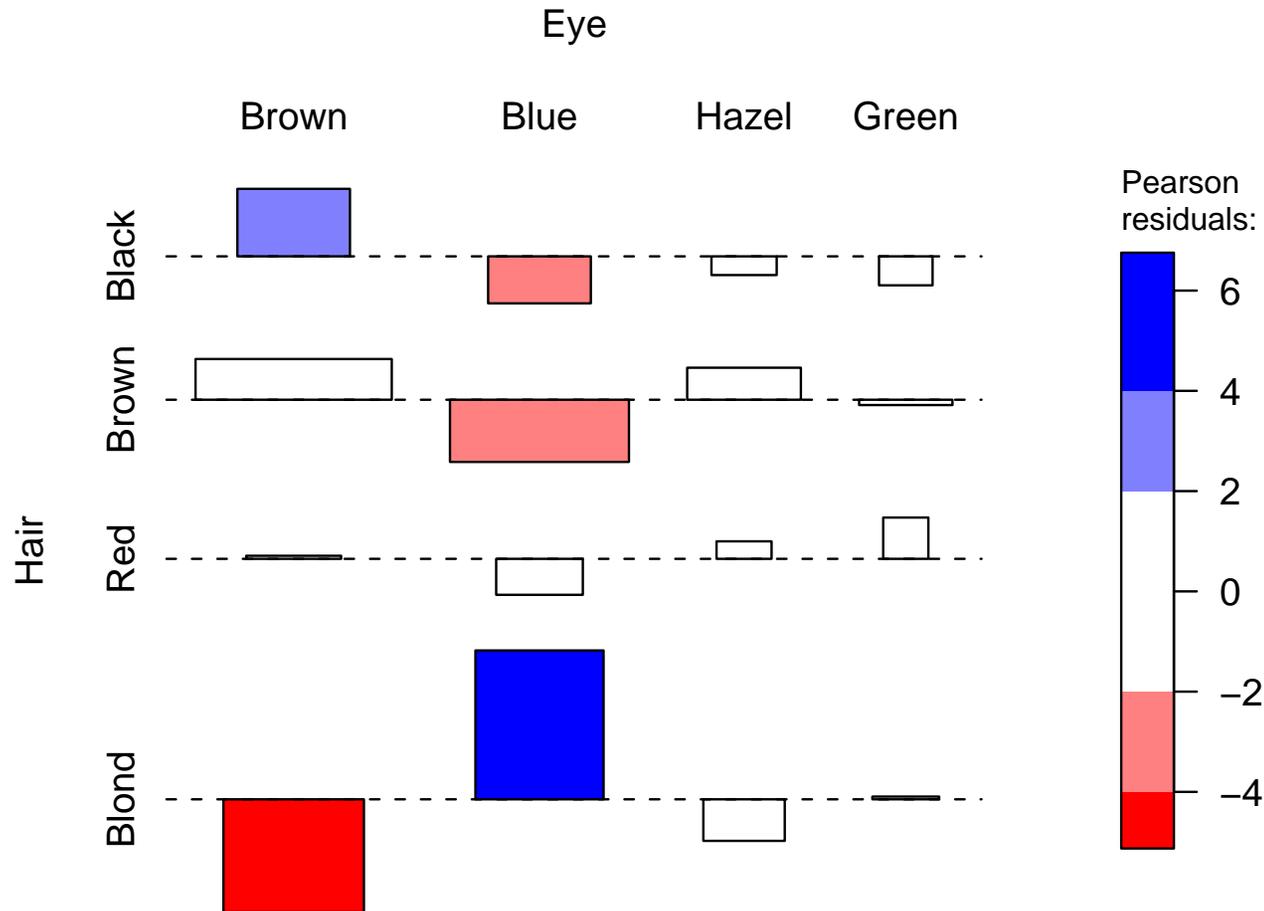


HSV colors

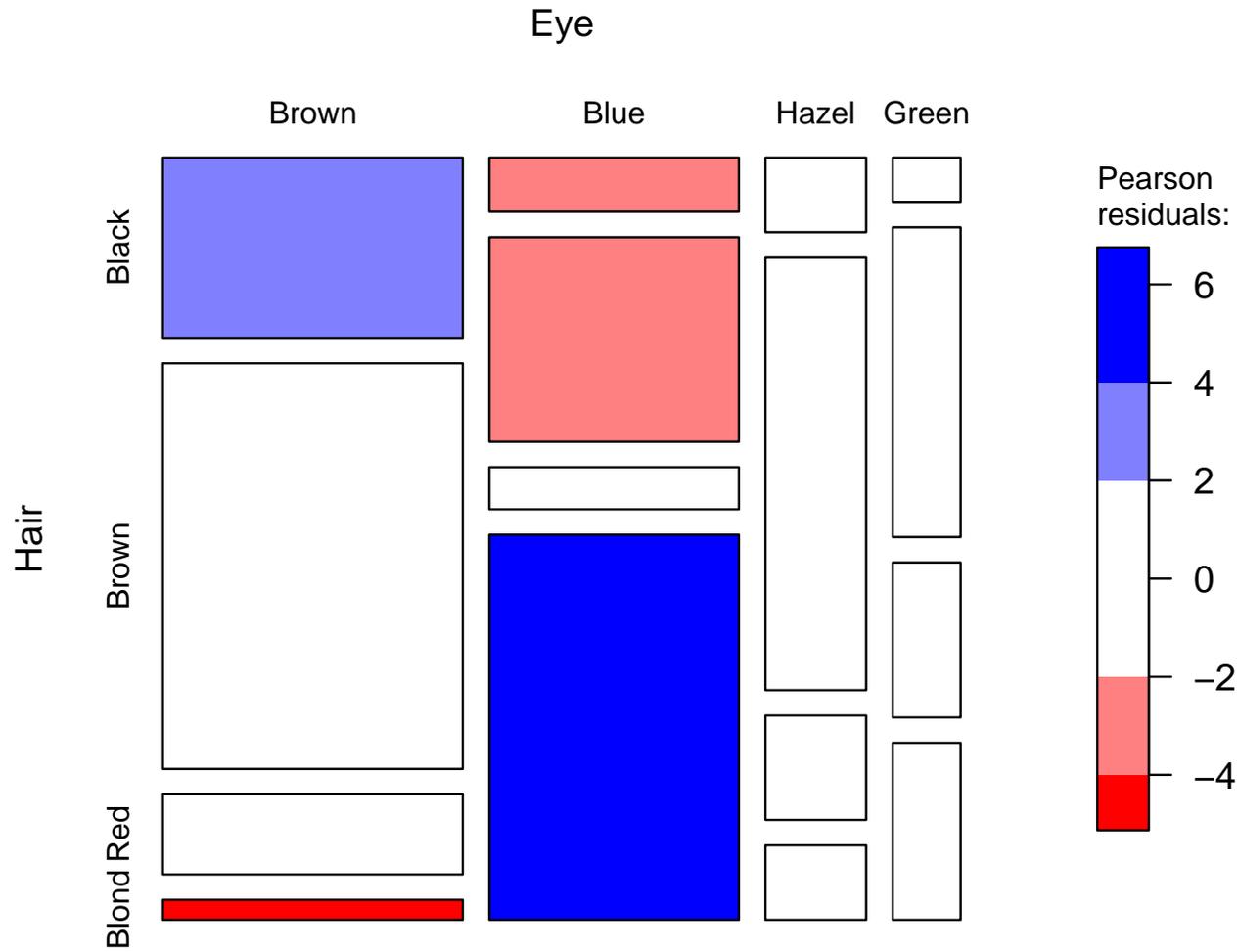
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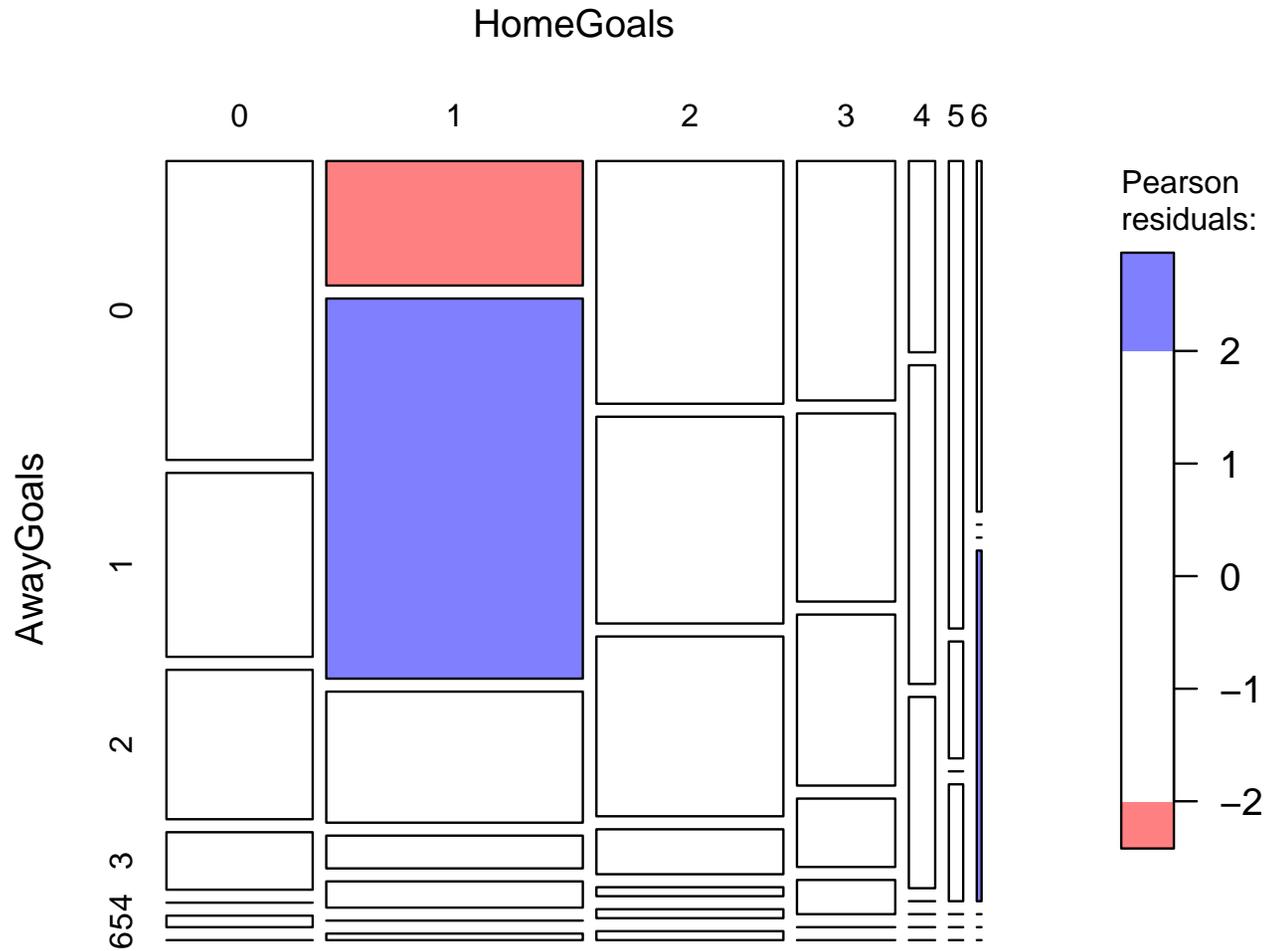
HSV colors



HSV colors



Visualization & testing



Intuition: colored cells convey the impression that there is significant dependence. Currently, this is *not* true.

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Approach 1: use the 90% and 99% critical values for the max statistic M instead of 2 and 4.

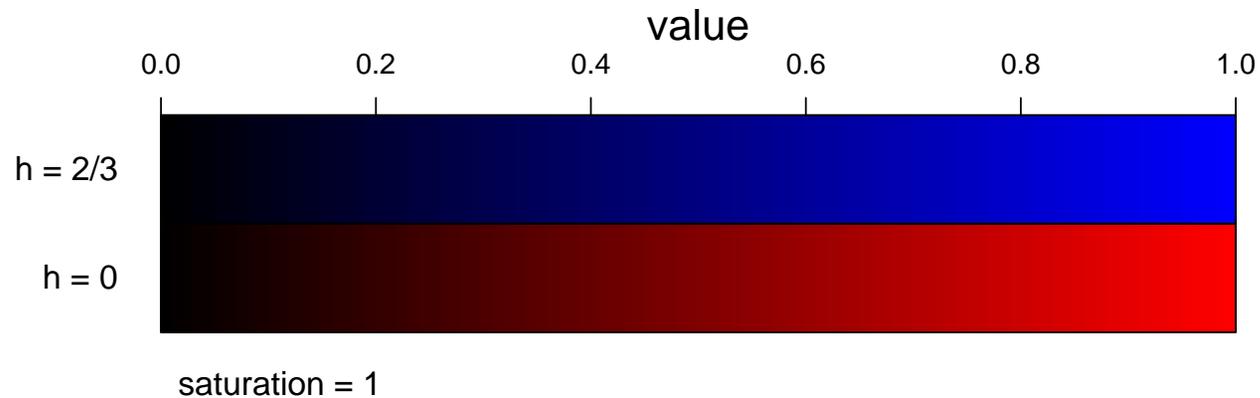
Advantage:

- ❄ color \Leftrightarrow significance
- ❄ highlights the cells which “cause” the dependence (if any).

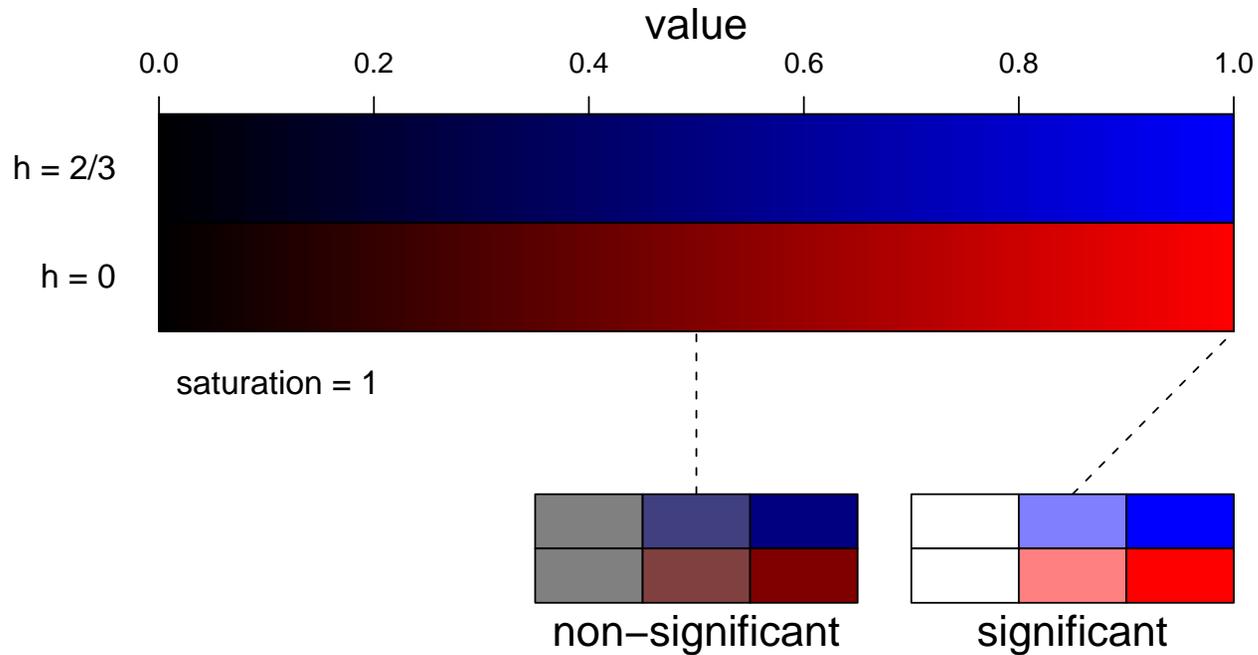
Disadvantage:

- ❄ does not work for the χ^2 test (or any other functional $\lambda(\cdot)$).

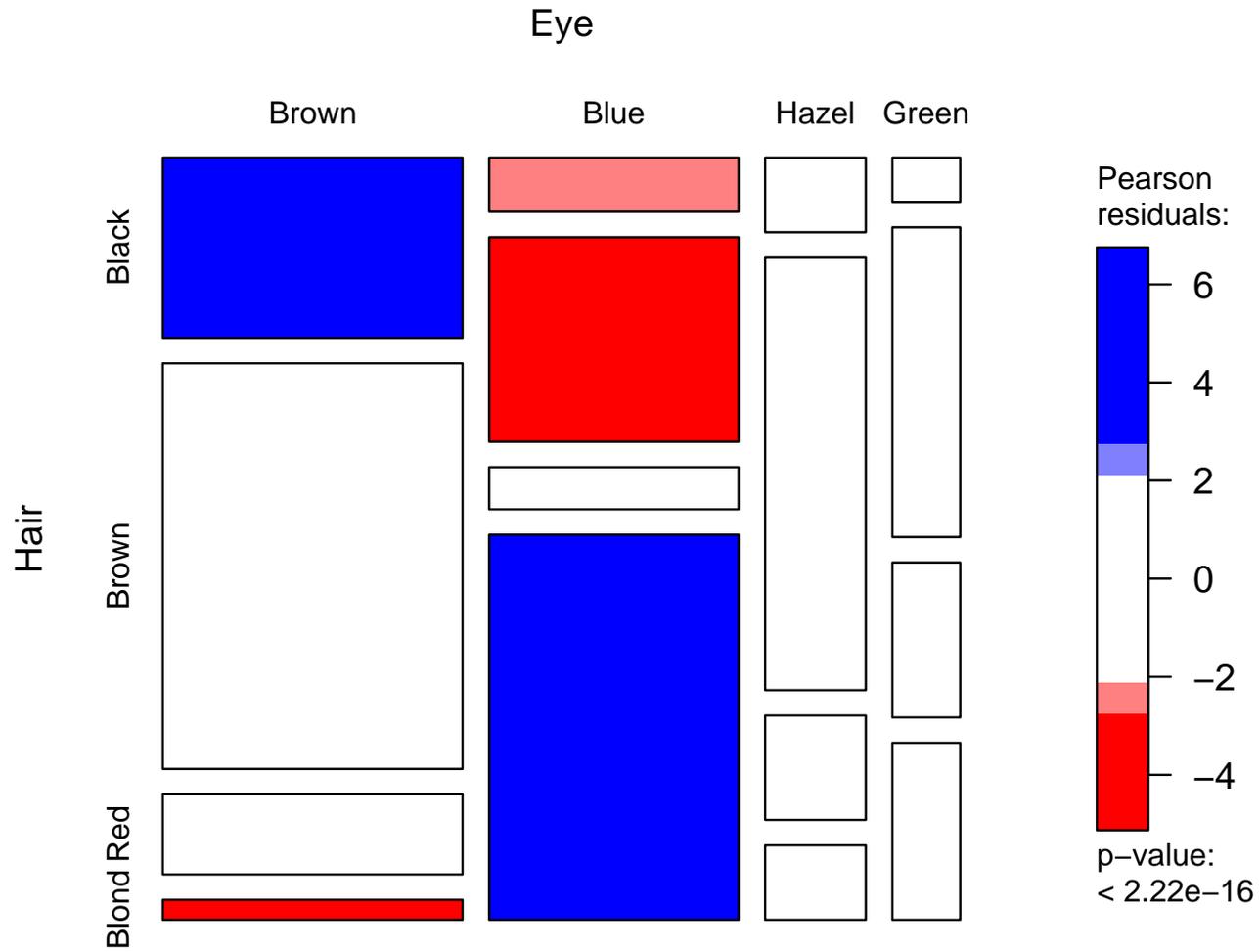
Approach 2: Use value to code the *result of a significance test* for independence.



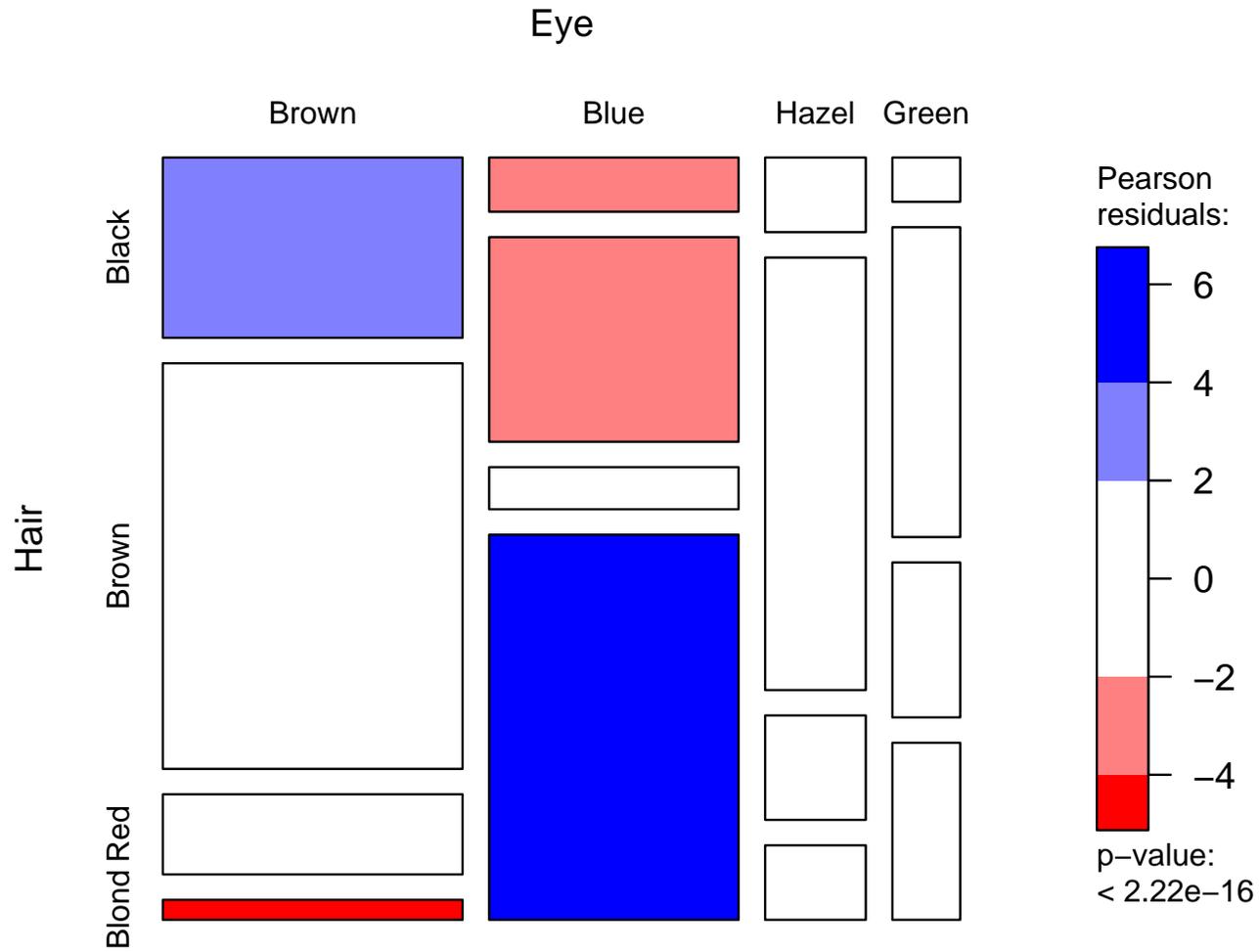
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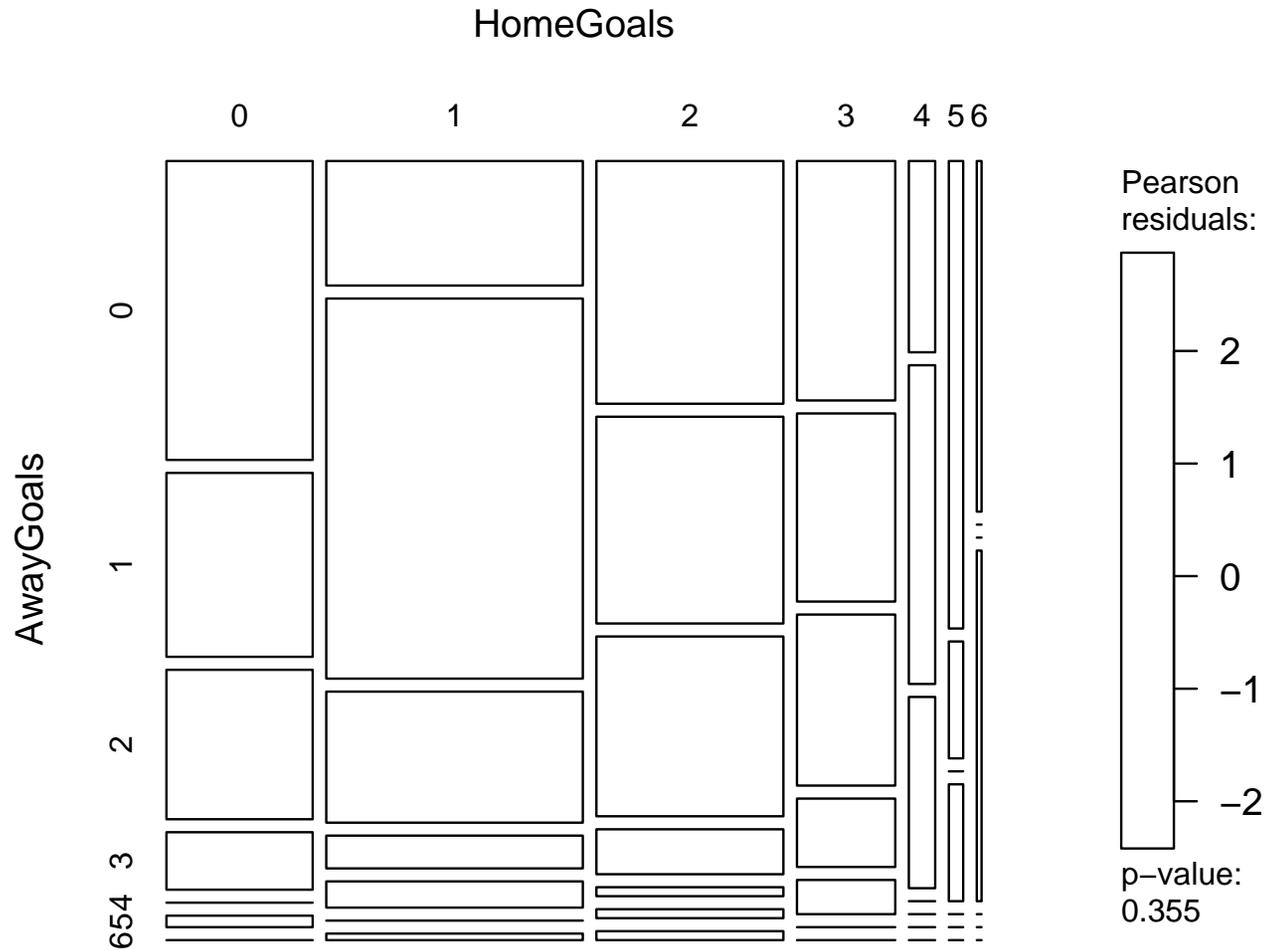
Visualization & testing



Visualization & testing



Visualization & testing



Disadvantages of HSV colors:

- ❄ device dependent,
- ❄ not copierproof,
- ❄ flashy colors good for drawing attention to a plot, but hard to look at.

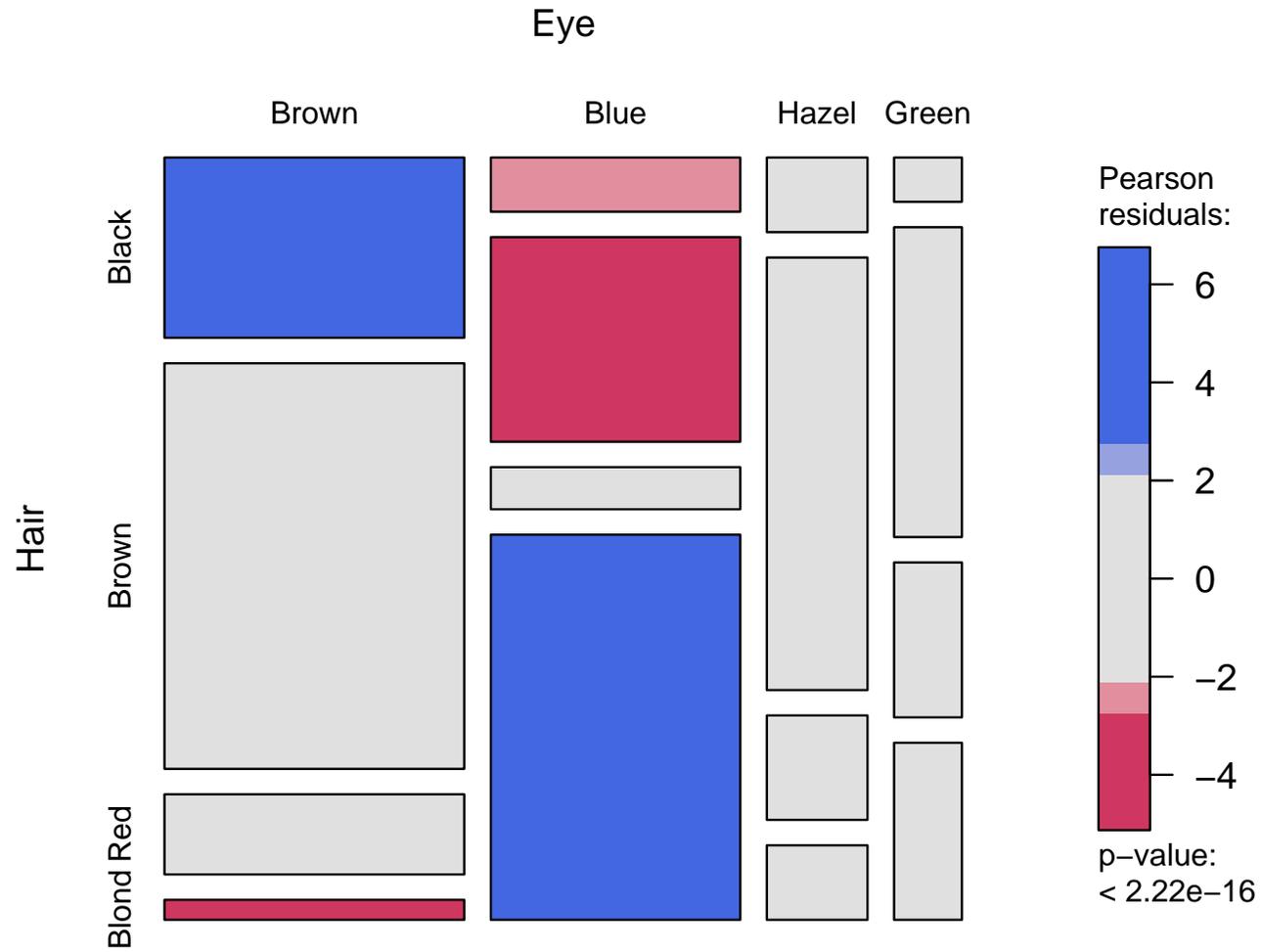
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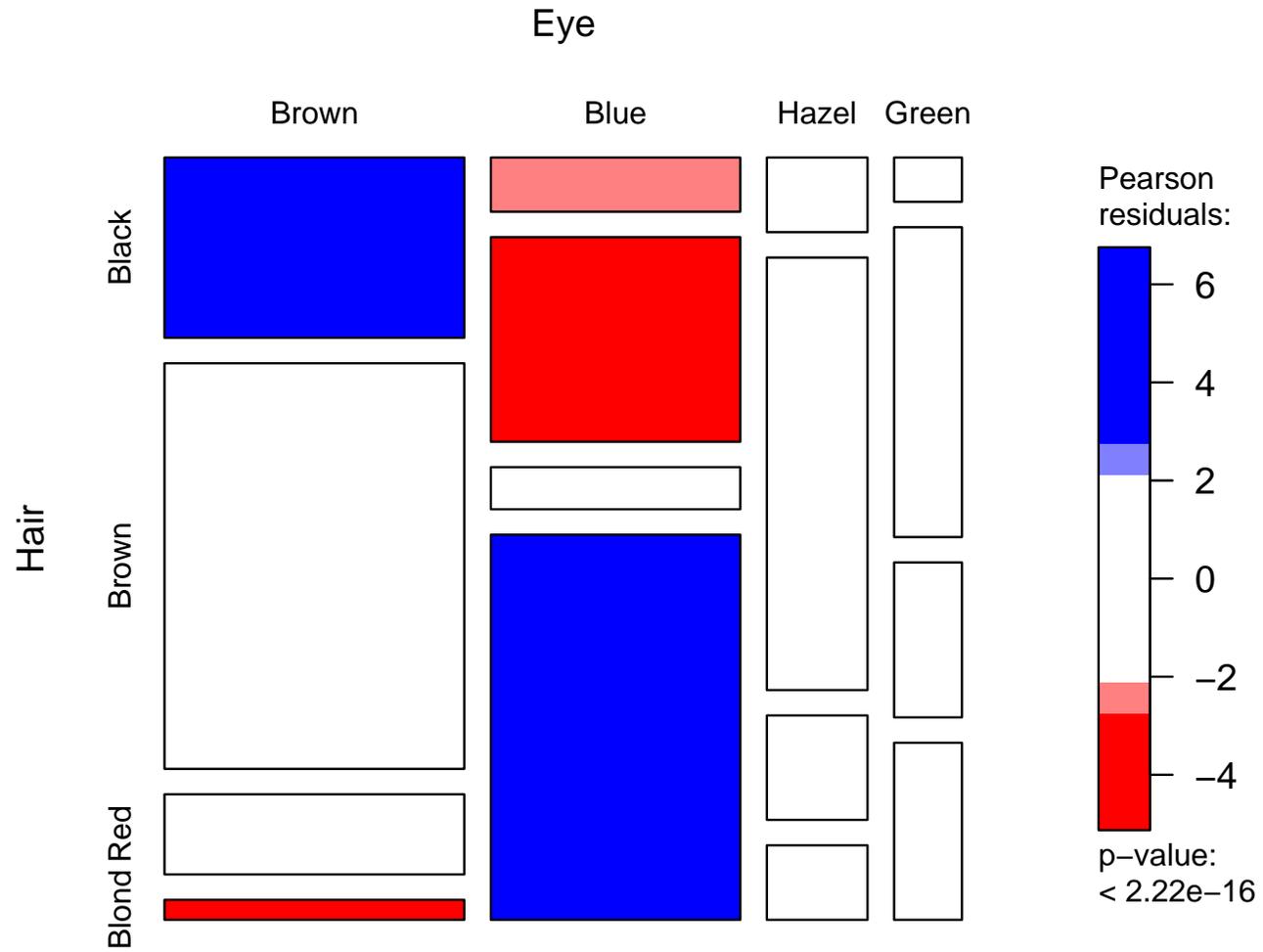
Alternative: use HCL colors instead (see Ihaka, 2003).

HCL colors are defined by hue (in $[0, 360]$), chroma and luminance (in $[0, 100]$). HCL space essentially looks like a double cone.

HCL colors



HCL colors



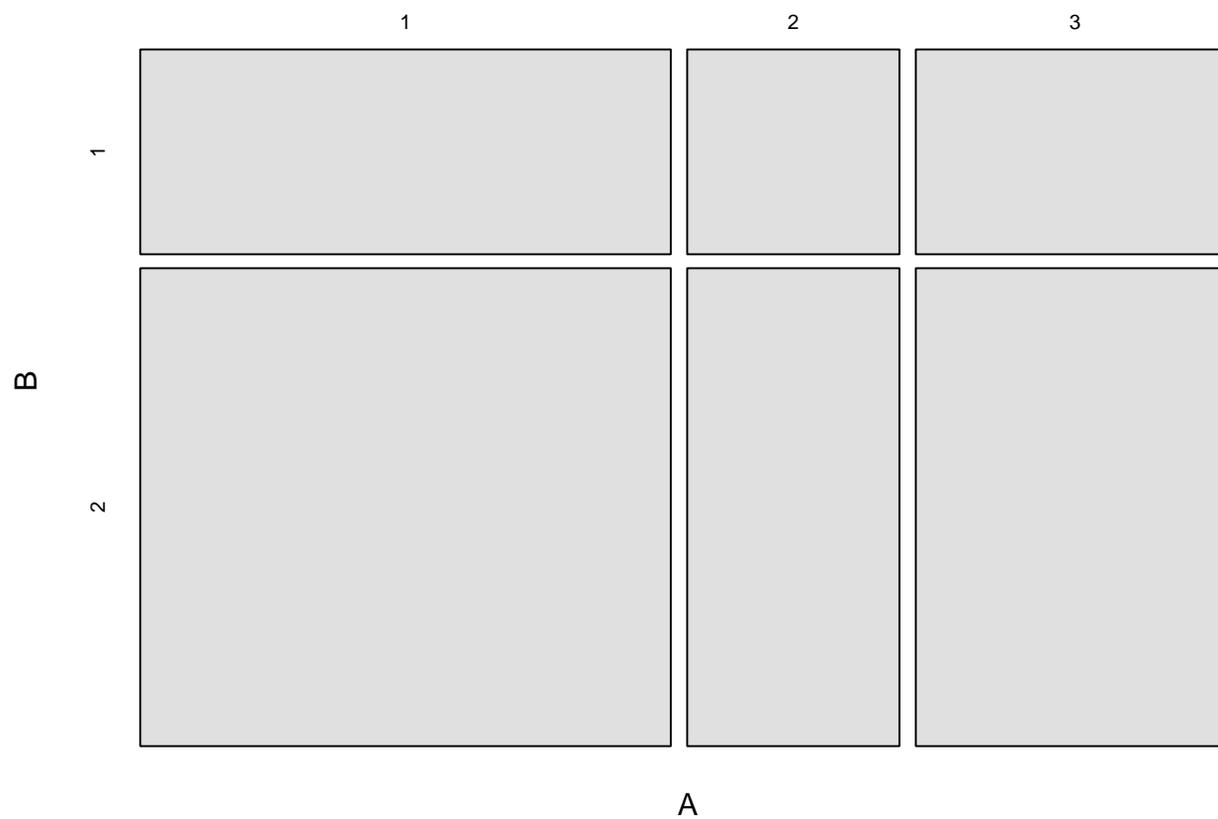
Principal idea of the mosaic plot:

❄ subdivision of tiles according to (conditional) probabilities

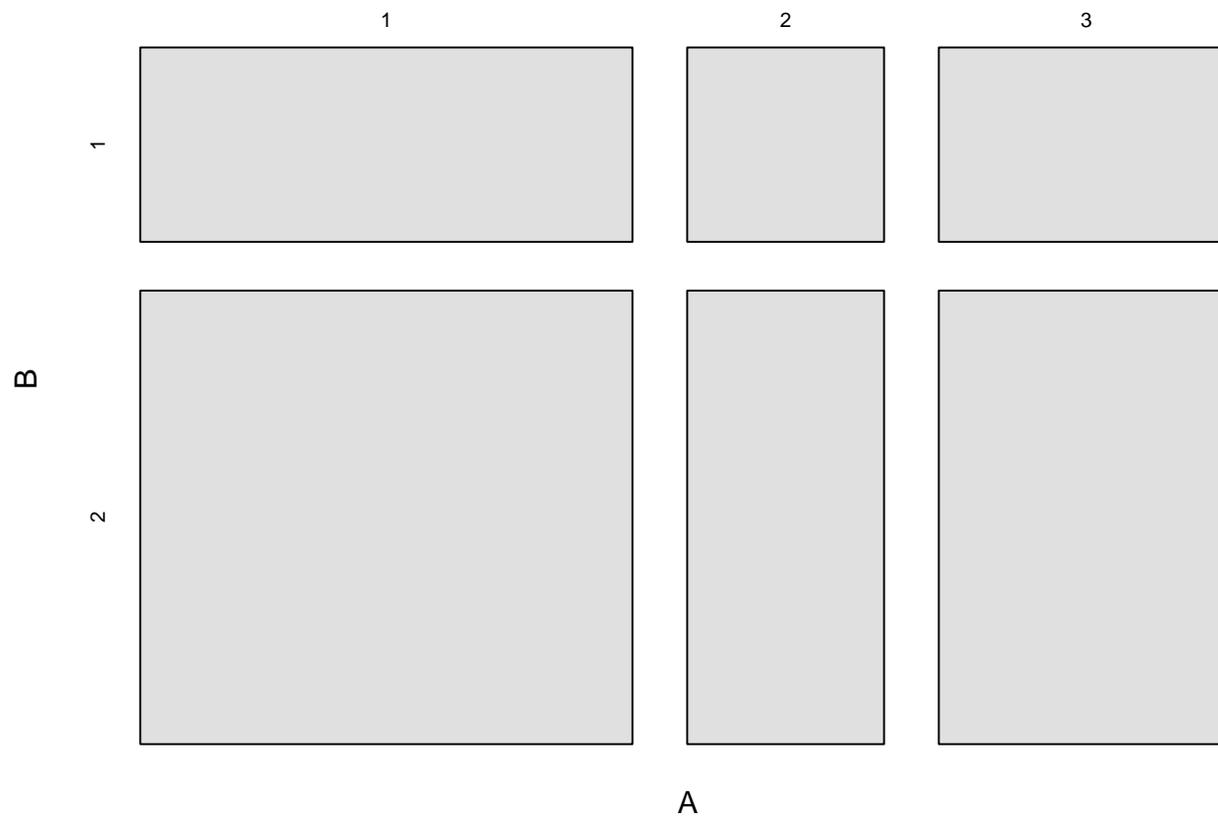
→ can also be used for n -way tables

The same idea does *not* apply to association plots.

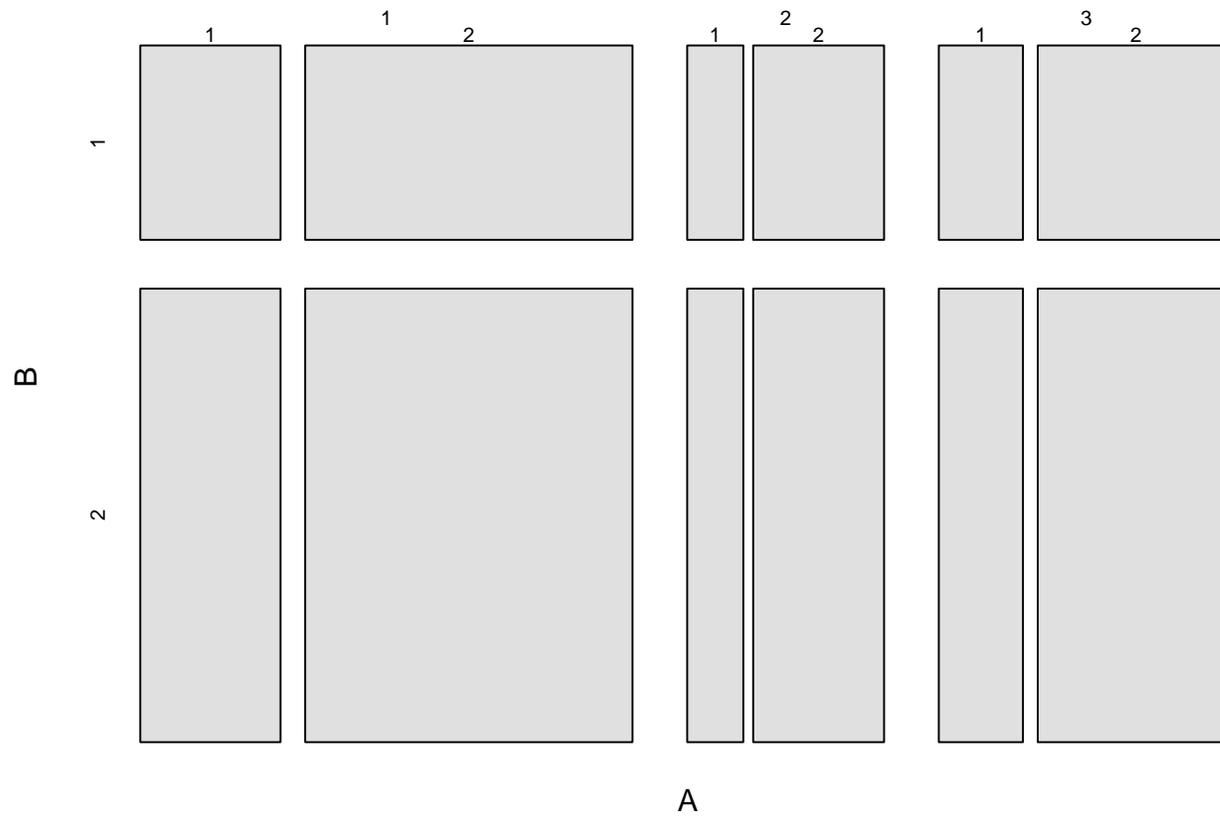
Multi-way tables



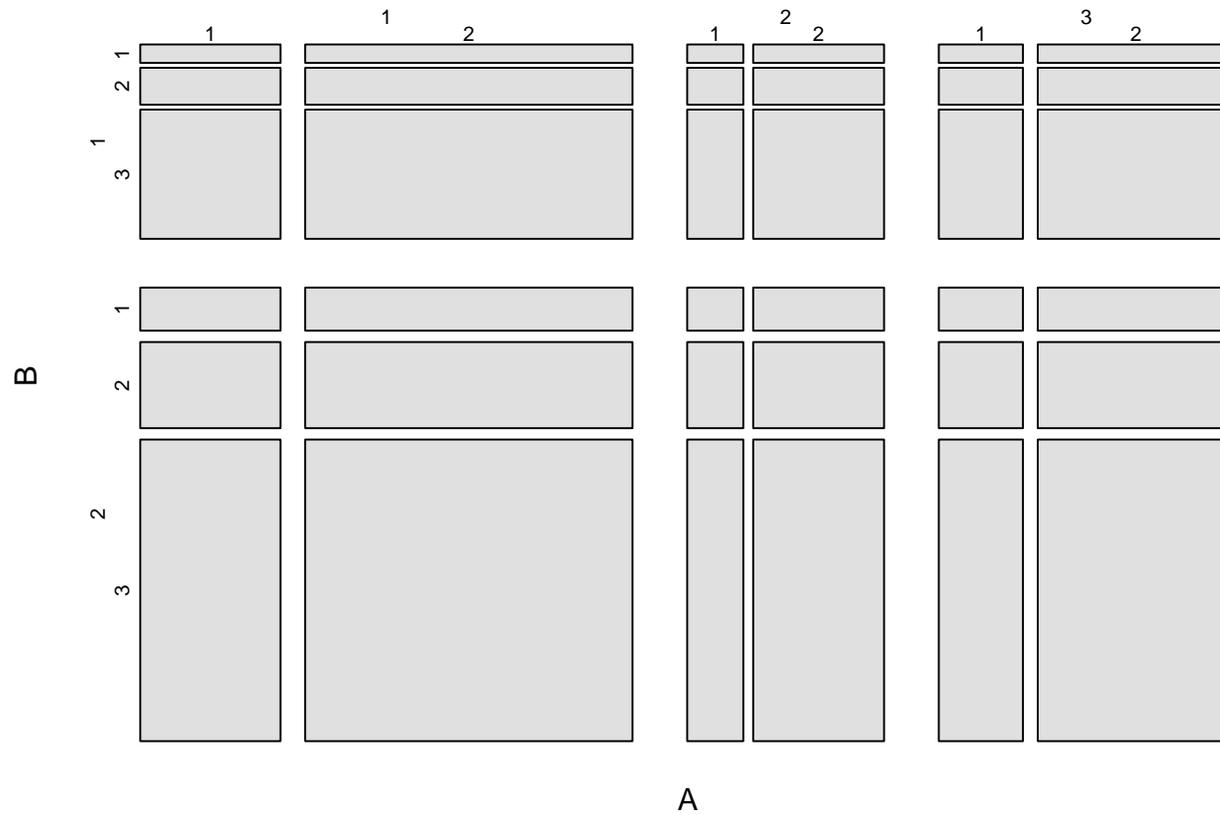
Multi-way tables



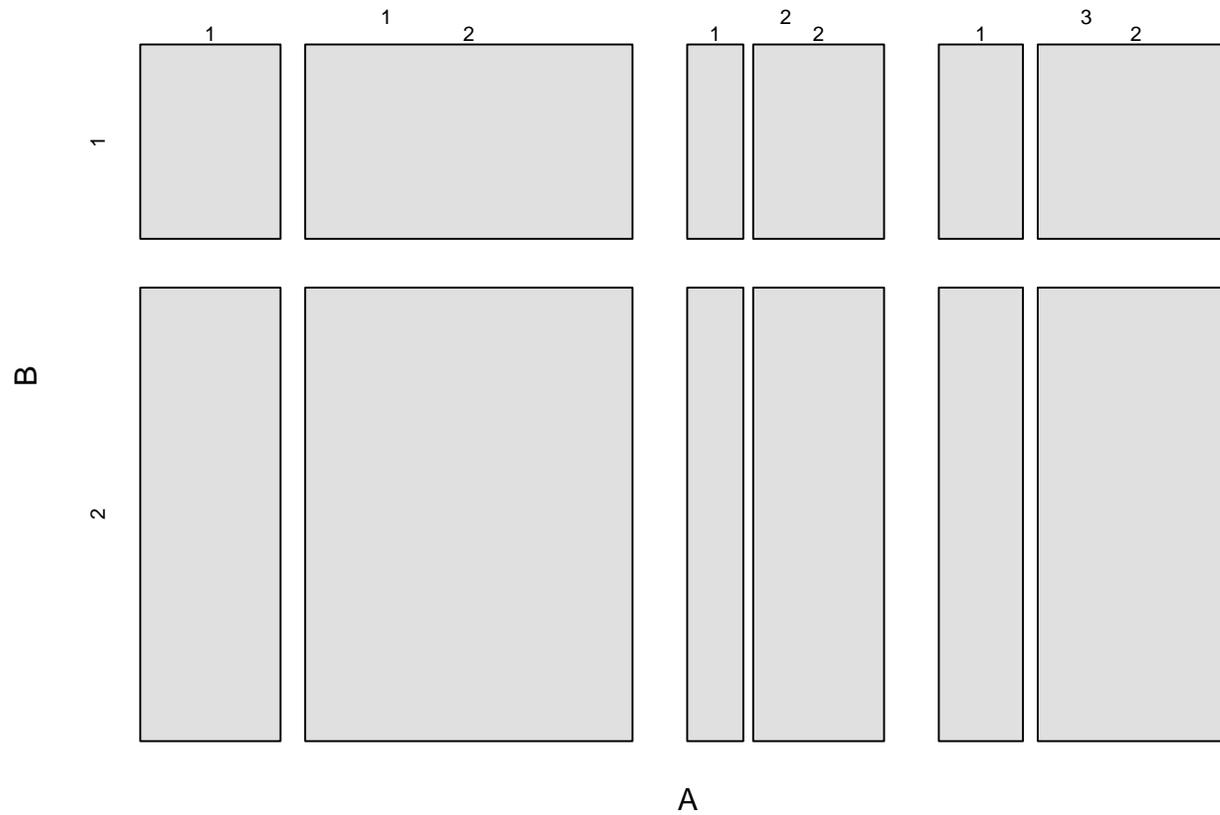
Multi-way tables



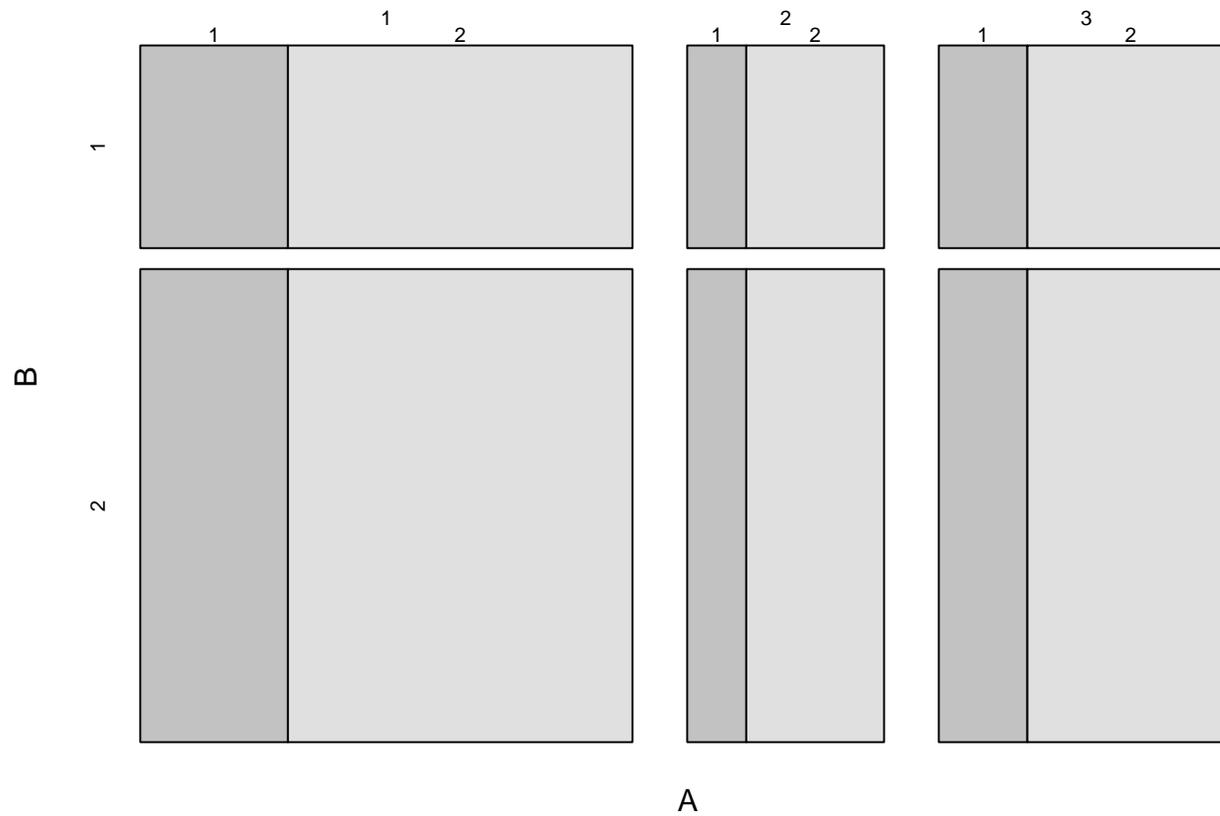
Multi-way tables



Multi-way tables

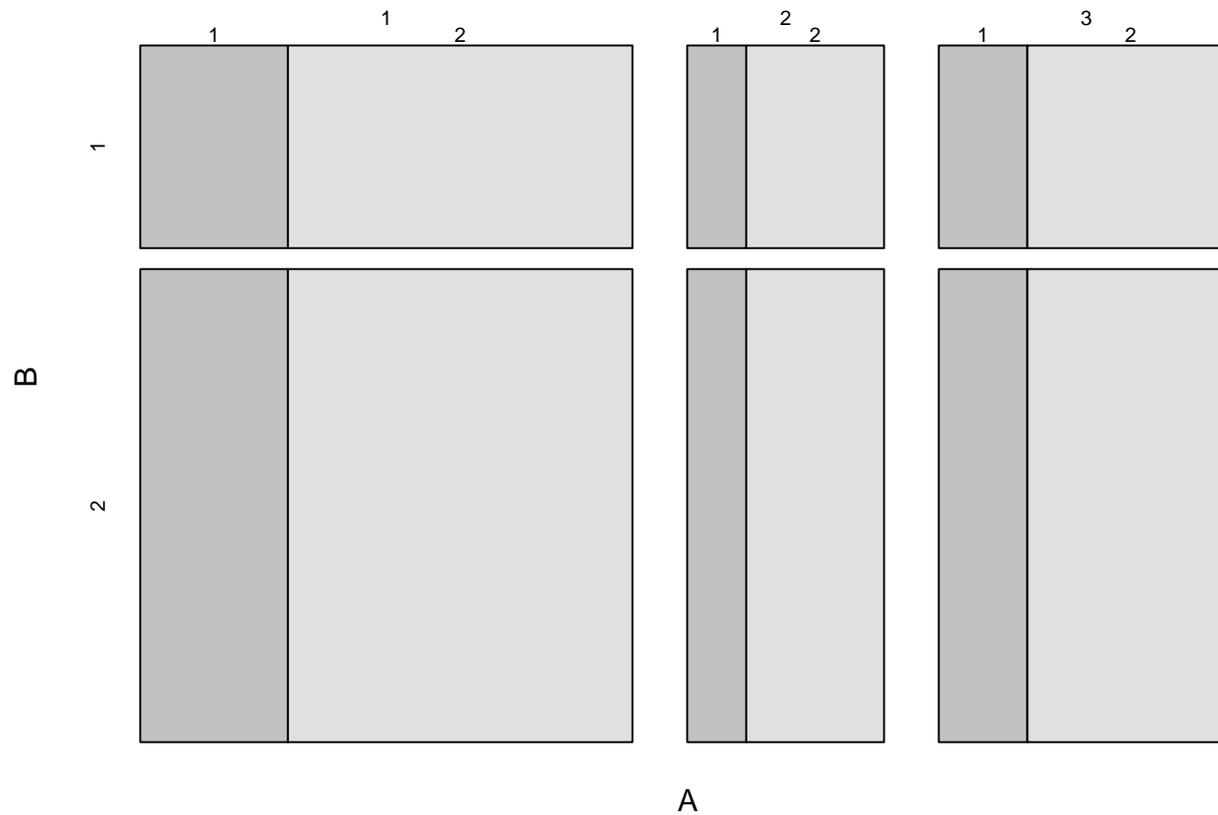


Multi-way tables



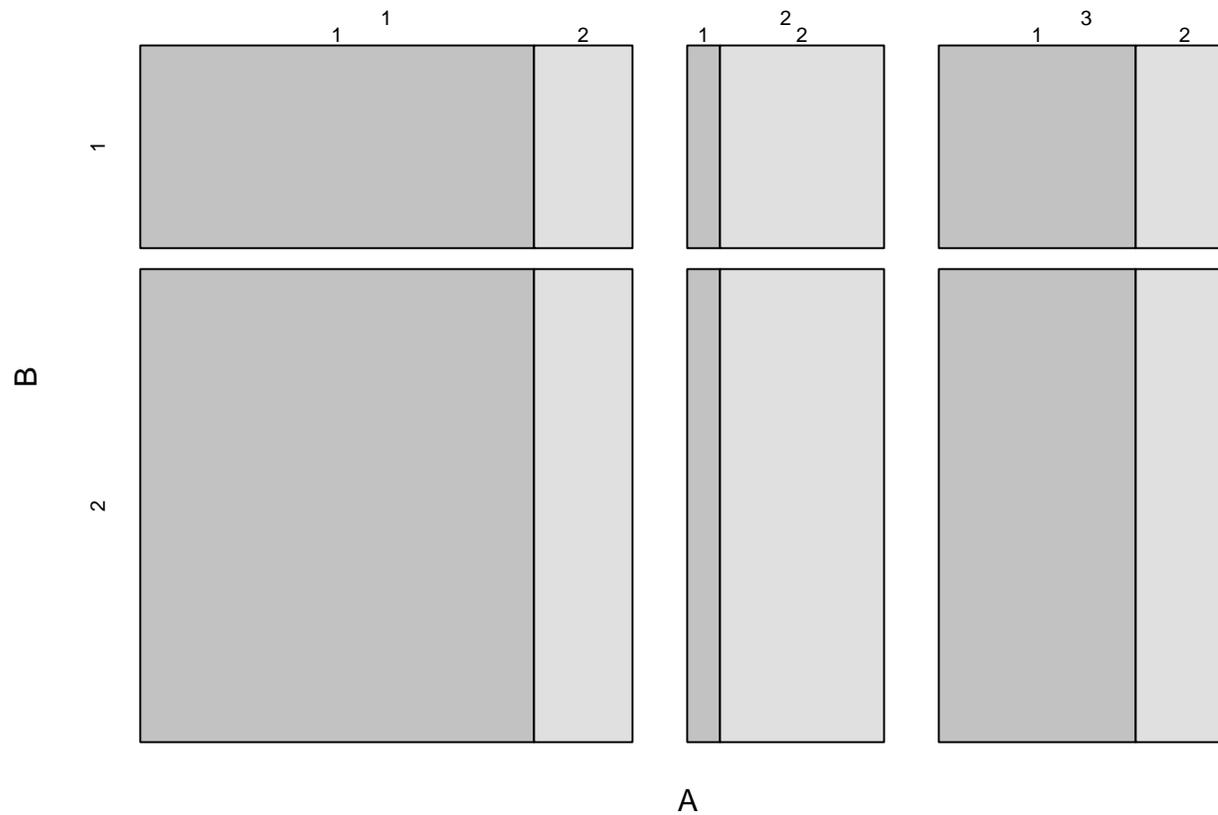
Multi-way tables

Complete independence: $A \perp\!\!\!\perp B \perp\!\!\!\perp C$



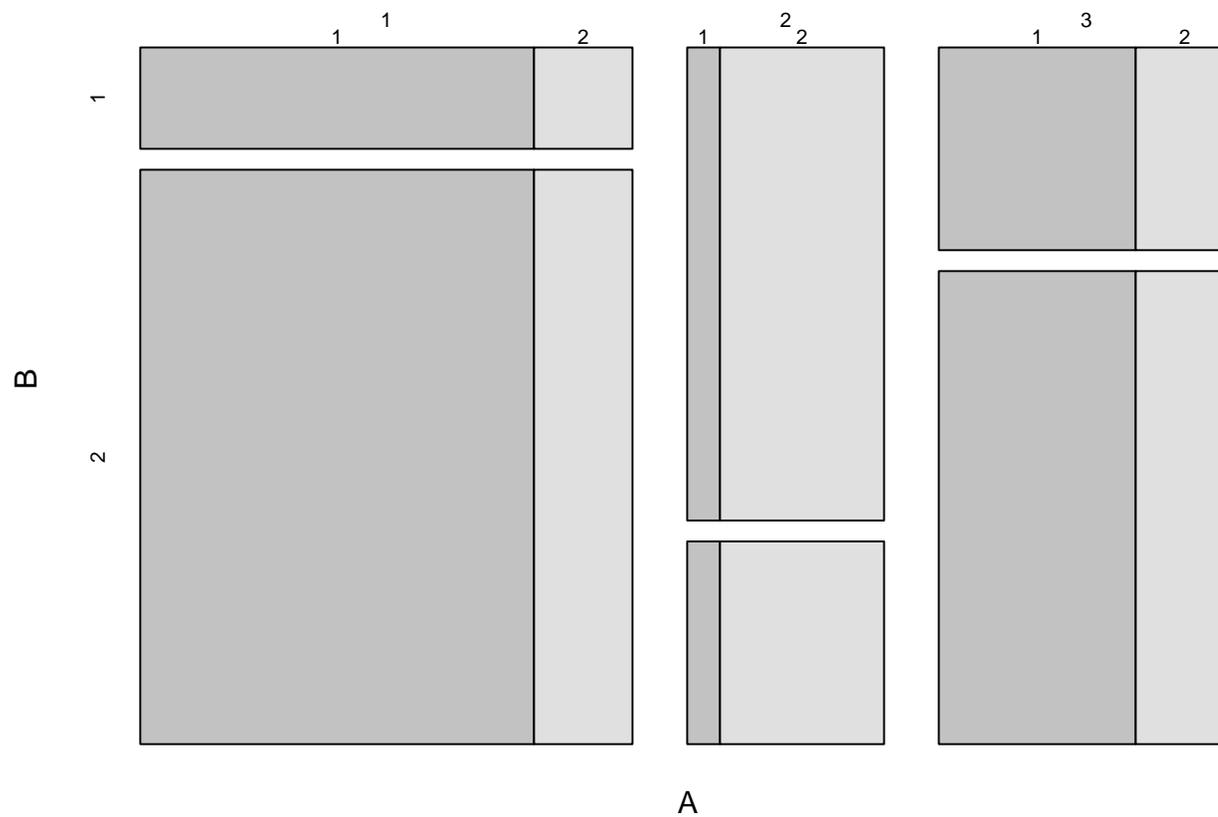
Multi-way tables

Joint independence: $(A, C) \perp\!\!\!\perp B$



Multi-way tables

Conditional independence: $B \perp\!\!\!\perp C \mid A$



Correspondence:

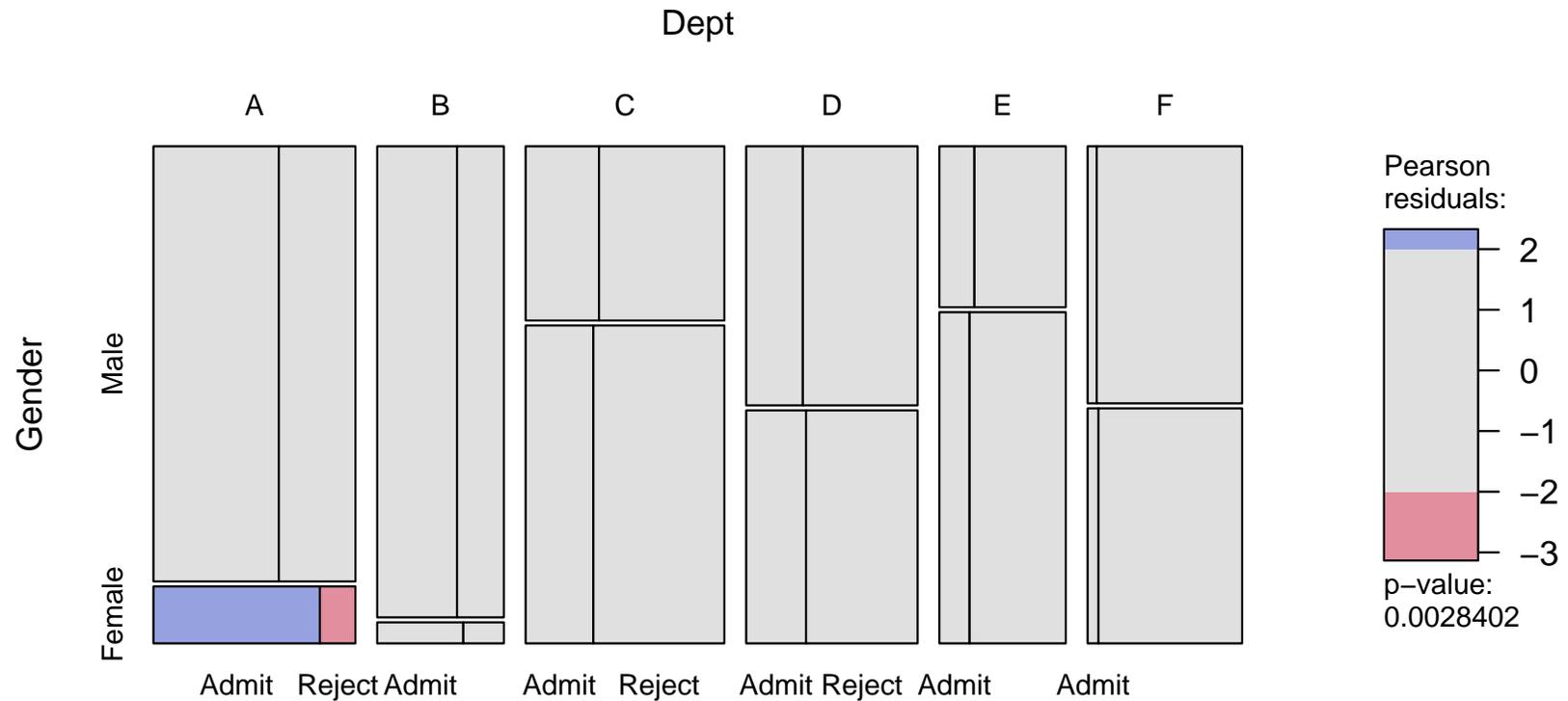
❄ conditioning in the model (→ shading of residuals)

❄ conditioning in the visual display

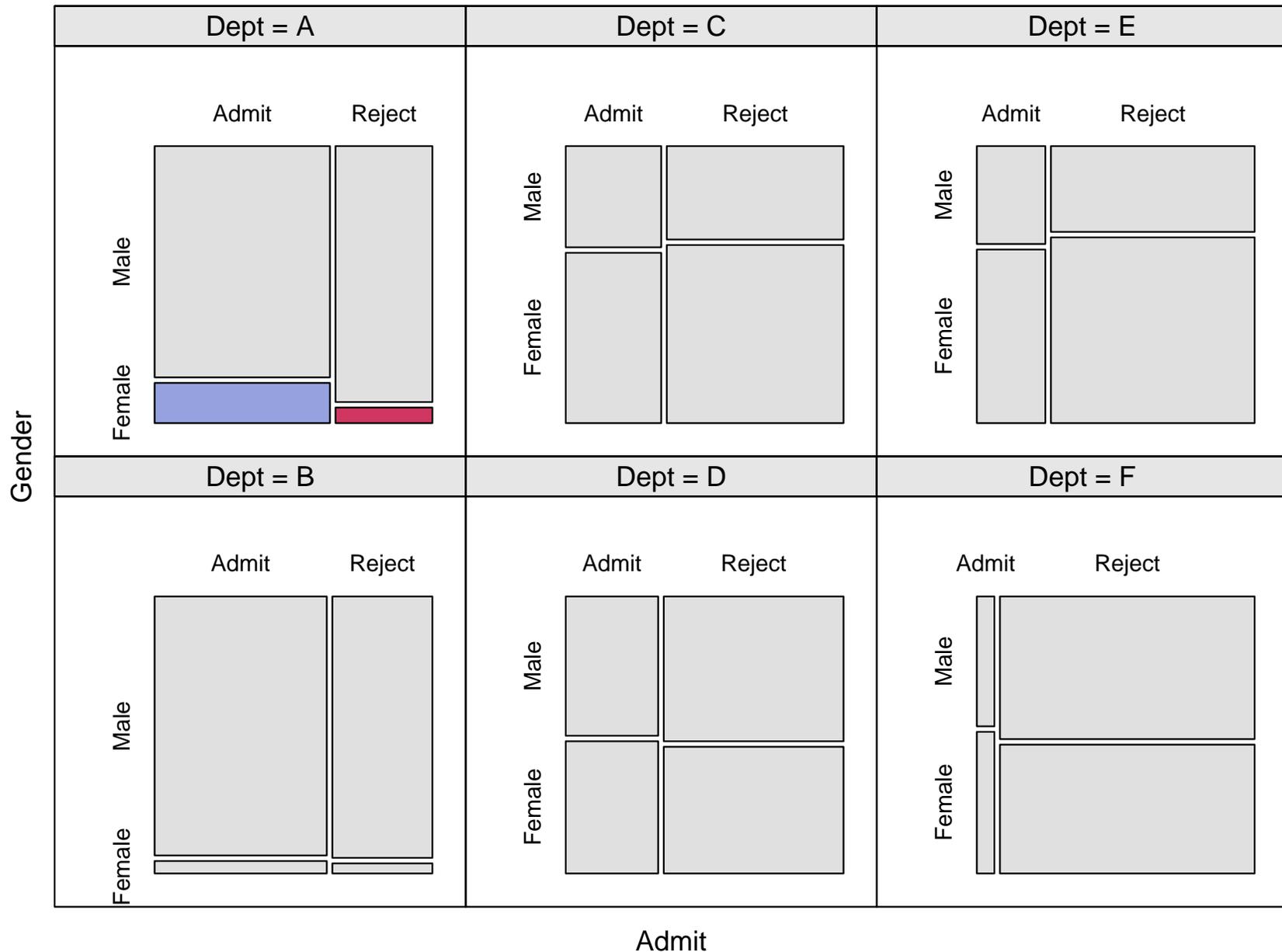
→ can also be done in Trellis-like layout

This idea *does* also work for association plots.

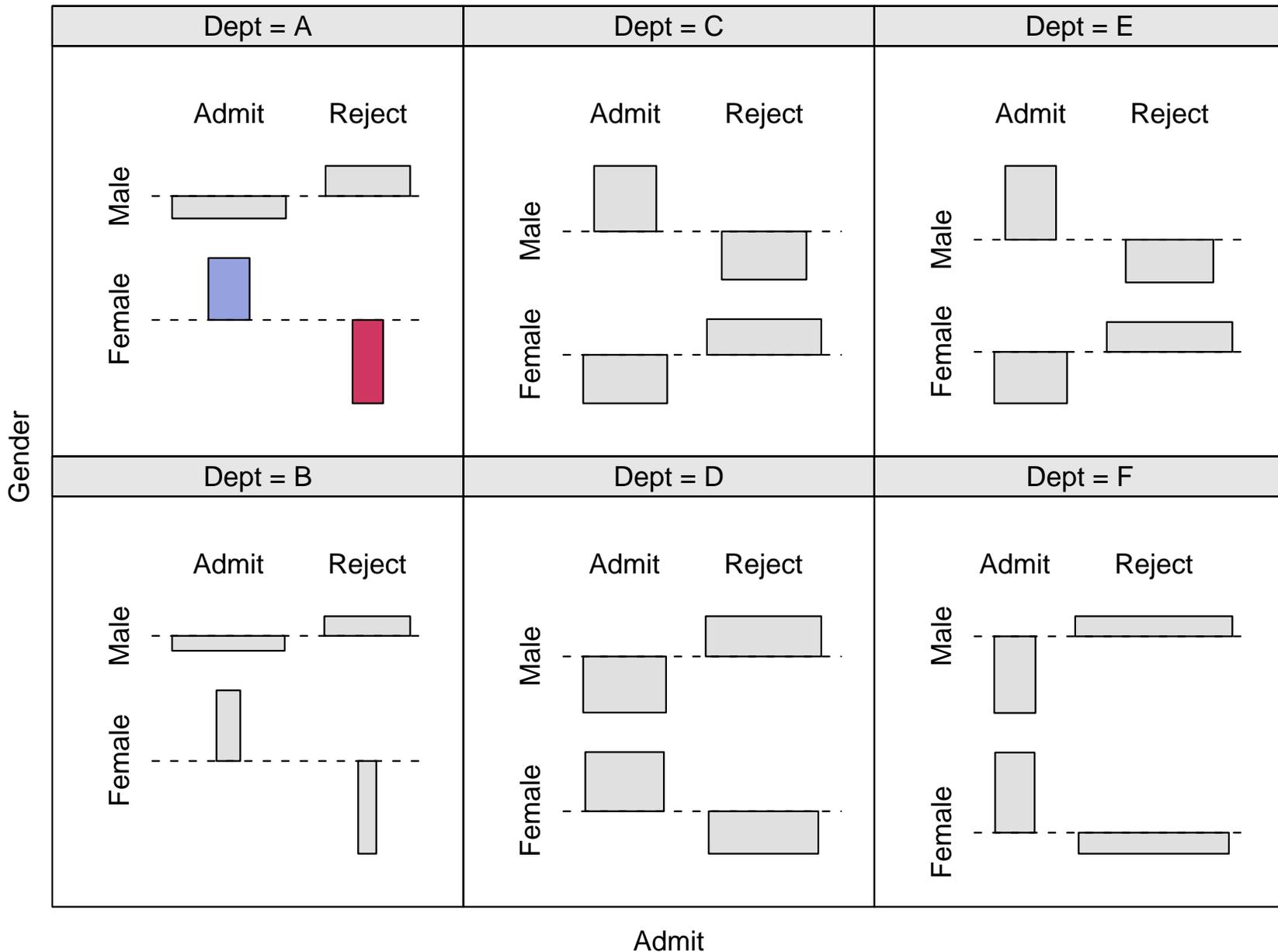
Multi-way tables



Multi-way tables



Multi-way tables



Multi-way tables



Conditioning in the plot:

```
R> assocplot(Admit ~ Gender | Dept, data = UCBAmissions)
```

Conditioning in the plot:

```
R> assocplot(Admit ~ Gender | Dept, data = UCBAmissions)
```

Conditioning in the model:

```
R> fm <- loglm(~(Admit + Gender) * Dept, data = UCBAmissions)
R> assocplot(fm)
```

The graphics engine `grid` overcomes the old R concept of plots with a plot region surrounded by a margin. `grid` is

- ❄ based on generic drawing regions (viewports),
- ❄ allows for plotting to relative coordinates,
- ❄ is also the basis for an implementation of Trellis graphics called `lattice`.

(see Murrell, 2002)

Thus, the new implementation of mosaic and association plots makes them easily reusable, e.g., in Trellis-like layouts.

Furthermore, graphics parameters for the rectangles, e.g.,

- ❄ fill color,
- ❄ line type,
- ❄ line color,

can be specified for each cell individually by the user. Each graphics parameter can be an object of the same dimensionality as the original table.

→ new shadings can easily be implemented.

New methods will be available in the package `vcd` for visualizing categorical data.

Currently only in development version. The released version is available from the Comprehensive R Archive Network

<http://CRAN.R-project.org/>

and it already offers some functionality for

- ❄ fitting & graphing of discrete distributions,
- ❄ plots for independence and agreement,
- ❄ visualization of log-linear models.