

TMA4315: Compulsory exercise 2 Logistic regression and Poisson regression

Group 7: Henrik Syversveen Lie, Mikal Stapnes, Oliver Byhring

25.10.2018

Contents

Part 1: Logistic regression	1
a)	1
b)	2
c)	4
d)	8
Part 2: Poisson regression - Eliteserien 2018	10
a)	10
b)	10
c)	11
d)	14

Part 1: Logistic regression

a)

We let y_i be the number of successful ascents, and n_i be the total number of attempts (success + fail) of the i 'th mountain. We then do binary regression with the logit link to model the probability of success. This gives

1. Model for response: $Y_i \sim \text{Bin}(n_i, \pi_i)$ for $i = 1, \dots, 113$, with $E(Y_i) = \mu_i = n_i \pi_i$ and $\text{Var}(Y_i) = n_i \pi_i (1 - \pi_i)$
2. Linear predictor: $\eta_i = \mathbf{x}_i^T \boldsymbol{\beta}$
3. Link function: $\eta_i = \ln\left(\frac{\pi_i}{1 - \pi_i}\right)$, also known as logit link

where \mathbf{x}_i is a p dimensional column vector of covariates for observation i , and $\boldsymbol{\beta}$ is the vector of regression parameters.

Then, the likelihood can be written

$$L(\boldsymbol{\beta}) = \prod_{i=1}^G L_i(\boldsymbol{\beta}) = \prod_{i=1}^G f(y_i; \boldsymbol{\beta}) = \prod_{i=1}^G \binom{n_i}{y_i} \pi_i^{y_i} (1 - \pi_i)^{n_i - y_i}.$$

If we take the natural logarithm, we get the log likelihood function

$$l(\boldsymbol{\beta}) = \ln(L(\boldsymbol{\beta})) = \sum_{i=1}^G l_i(\boldsymbol{\beta}) = \sum_{i=1}^G [y_i \ln(\pi_i) + (n_i - y_i) \ln(1 - \pi_i) + \ln \binom{n_i}{y_i}] = \sum_{i=1}^G [\ln \binom{n_i}{y_i} + y_i \ln \left(\frac{\pi_i}{1 - \pi_i} \right) + n_i \ln(1 - \pi_i)]$$

To express the log likelihood as a function of $\boldsymbol{\beta}$ we first use the link between η_i and π_i . We use the response function $h(\eta) = g^{-1}(\eta)$, which is the inverse of the link function. $\pi_i = h(\eta_i) = \frac{\exp(\eta_i)}{1 + \exp(\eta_i)}$. The log-likelihood function can then be written:

$$l(\boldsymbol{\beta}) = \sum_{i=1}^G [\ln \binom{n_i}{y_i} + y_i \eta_i + n_i \ln \left(\frac{1}{1 + \exp(\eta_i)} \right)] = \sum_{i=1}^G [\ln \binom{n_i}{y_i} + y_i \eta_i - n_i \ln(1 + \exp(\eta_i))].$$

Finally we use that $\eta_i = \mathbf{x}_i^T \boldsymbol{\beta}$ and get

$$l(\boldsymbol{\beta}) = \sum_{i=1}^G [\ln \binom{n_i}{y_i} + y_i \mathbf{x}_i^T \boldsymbol{\beta} - n_i \ln(1 + \exp(\mathbf{x}_i^T \boldsymbol{\beta}))],$$

which is an expression for the log likelihood as a function of beta.

Since we know that this function is concave we can find the parameters, $\boldsymbol{\beta}$ that gives the maximum likelihood by taking the partial derivatives with respect to $\boldsymbol{\beta}$ to get the score function,

$$\mathbf{s}(\boldsymbol{\beta}) = \frac{\partial l(\boldsymbol{\beta})}{\partial \boldsymbol{\beta}} = \sum_{i=1}^G [y_i x_i - n_i \frac{x_i \exp \mathbf{x}_i^T \boldsymbol{\beta}}{1 + \exp \mathbf{x}_i^T \boldsymbol{\beta}}].$$

The parameters $\boldsymbol{\beta}$ that gives the maximum likelihood is those for which the score function equals the zero-vector.

$$\mathbf{s}(\boldsymbol{\beta}) = \frac{\partial l(\boldsymbol{\beta})}{\partial \boldsymbol{\beta}} = \mathbf{0}.$$

In practice, there is often no closed form solution, which means that we have to use numerical procedures to find the betas, such as Newton-Raphson or Fisher Scoring. The betas can be found iteratively by the formula

$$\boldsymbol{\beta}_{n+1} = \boldsymbol{\beta}_n + F^{-1}(\boldsymbol{\beta}_n) \mathbf{s}(\boldsymbol{\beta}_n),$$

where $F^{-1}(\boldsymbol{\beta}_n)$ is the inverse of the expected Fisher information.

b)

We load the dataset and fit a model as described in Part a) with success rate as response and height and prominence as predictors.

```
##
## Call:
## glm(formula = cbind(success, fail) ~ height + prominence, family = "binomial",
##      data = mount)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -6.2886  -0.8086   0.6893   1.4226   3.7456
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  1.369e+01  1.064e+00  12.861  < 2e-16 ***
## height      -1.635e-03  1.420e-04 -11.521  < 2e-16 ***
## prominence  -1.740e-04  4.554e-05  -3.821  0.000133 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
```

```
##
## Null deviance: 715.29 on 112 degrees of freedom
## Residual deviance: 414.68 on 110 degrees of freedom
## AIC: 686.03
##
## Number of Fisher Scoring iterations: 4
```

We want to interpret the model parameters by using the odds. We have from the link function that

$$\eta_i = \ln\left(\frac{\pi_i}{1 - \pi_i}\right).$$

Our linear predictor is $\eta_i = \mathbf{x}_i^T \boldsymbol{\beta}$. If we insert this, assuming intercept and two covariates, we get

$$\ln\left(\frac{\pi_i}{1 - \pi_i}\right) = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2}.$$

We can now take $\exp(\cdot)$ of both sides, yielding

$$\frac{\pi_i}{1 - \pi_i} = \exp(\beta_0) \cdot \exp(\beta_1 x_{i1}) \cdot \exp(\beta_2 x_{i2}).$$

The left hand side is what we call the odds. If we now increase covariate j by one unit and keep the other constant, the odds will be multiplied by $\exp(\beta_j)$. Coefficient estimates for both height and prominence are negative. This means that for both coefficients $\exp(\beta_j) < 1$, and an increase in **height** or **prominence** will give a decrease in the odds and thus a decrease in success probability of climbing the mountain. For example, increasing **height** by one meter will decrease the odds by multiplying it with $\exp(\beta_1) = \exp(-0.001635) = 0.998$.

We can interpret the intercept in the following way. For a mountain with zero height, and zero prominence, the odds of climbing it will be $\frac{\pi_i}{1 - \pi_i} = \exp(\beta_0) = \exp(13.685845) = 878389$, or $\pi_i \simeq 1$. Although the intercept in this case is just theoretical (there would be no point in climbing a mountain of zero height), it makes sense because climbing a mountain of zero height should have probability 1.

Now we want to discuss the significance of the parameters. We do a Wald-test for each of the parameters. The Wald-test has hypotheses

$$H_0 : \beta_j = 0 \quad \text{vs.} \quad H_1 : \beta_j \neq 0.$$

Asymptotically ($n \rightarrow \infty$), the parameter estimates are normally distributed with mean $\boldsymbol{\beta}$ and variance $F^{-1}(\hat{\boldsymbol{\beta}})$, or the inverse of the expected Fisher information matrix evaluated at the ML estimate. Equivalently,

$$\hat{\boldsymbol{\beta}} \approx N_p(\boldsymbol{\beta}, F^{-1}(\hat{\boldsymbol{\beta}})).$$

We denote c_{jj} for diagonal element j of $F^{-1}(\hat{\boldsymbol{\beta}})$ and get that for each coefficient β_j ,

$$\frac{\beta_j - \hat{\beta}_j}{\sqrt{c_{jj}}} \sim N(0, 1).$$

We can test the significance of the β_j using the test statistics

$$z_j = \frac{\hat{\beta}_j - 0}{\sqrt{c_{jj}}}.$$

We read the values of the test statistics from the summary above, and get $z_0 = 12.861$, $z_1 = -11.521$, $z_2 = -3.821$. This gives p-values of $p_0 < 2 \cdot 10^{-16}$, $p_1 < 2 \cdot 10^{-16}$, $p_2 = 0.000133$. All these p-values are below any reasonable significance level, so we reject the null-hypothesis for all parameters. This means that all the parameters are significant.

Next, we want to use a likelihood ratio test (LRT), which compares the likelihood of two models. We want to test the significance of the above model, so we use the deviance. We have hypotheses

$$H_0 : \text{"Our model fits the data well"}, \quad H_1 : \text{"Our model does not fit the data well"}.$$

The deviance compares the candidate model with a saturated model. The saturated model is the model that provides a perfect fit to our data, so in the saturated model, $\tilde{\pi}_i = y_i$. We then get the deviance,

$$D = -2(\ln L(\text{candidate model}) - \ln L(\text{saturated model})) = -2(l(\hat{\pi}) - l(\tilde{\pi})).$$

Asymptotically, D is distributed as χ^2 with $G - p = 110$ degrees of freedom. We calculate the deviance and the p-value.

```
## deviance: 414.6842
```

```
## p-value: 0
```

Because the p-value is below any reasonable significance level, we reject the null-hypothesis. So our candidate model does not fit the data well. Although all the parameters are significant, this test indicates that we haven't included enough parameters to make a good model. The model could be improved if we included additional covariates, e.g. weather, amount of avalanches, "fitness" of climber, etc.

Asymptotically, the parameter estimates are normally distributed with mean β and variance $F^{-1}(\hat{\beta})$, or the inverse of the expected Fisher information matrix evaluated at the ML estimate. Or equivalently

$$\hat{\beta} \approx N_p(\beta, F^{-1}(\hat{\beta})).$$

As mentioned in the Wald-test. The parameteres are asymptotically normal with mean β and variance c_{jj} . The `r` printout gives us the coefficient estimates and the estimated standard deviation $\sqrt{c_{jj}}$, of parameter j . From this which we can construct a confidence interval. We can make a 95% confidence interval for the height parameter β_{height} . We have 95% confidence that

$$\beta_{\text{height}} \in [\hat{\beta}_{\text{height}} - z_{0.025}\sqrt{c_{jj}}, \hat{\beta}_{\text{height}} + z_{0.025}\sqrt{c_{jj}}],$$

where $\sqrt{c_{jj}}$ can be read off as the standard deviation of the height parameter. Inserting values from the `r` printout, we get

$$\beta_{\text{height}} \in [-0.001635 - 1.96 \cdot 0.000142, -0.001635 + 1.96 \cdot 0.000142] = [-0.00191332, -0.00135668] = [\hat{\beta}_L, \hat{\beta}_H].$$

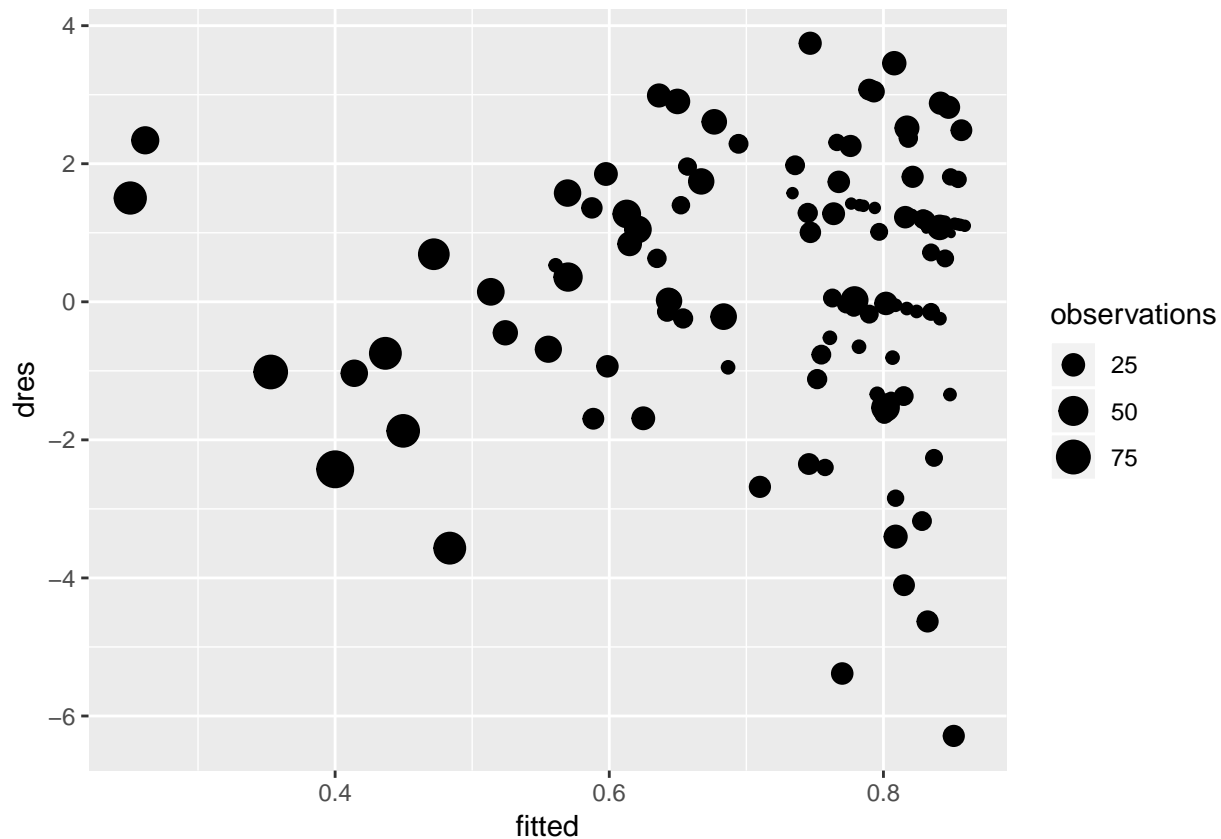
Finally we can take the $\exp(\cdot)$ function of both limits of the interval, and we get that

$$\exp(\beta_{\text{height}}) \in [\exp(\hat{\beta}_L), \exp(\hat{\beta}_H)] = [0.9981, 0.9986].$$

In this way, we have found a 95% confidence interval for $\exp(\beta_{\text{height}})$. This means that we can say with 95% confidence that an increase in height of 1 meter means that the odds will be multiplied by $\exp(\beta_{\text{height}}) \in [0.9981, 0.9986]$.

c)

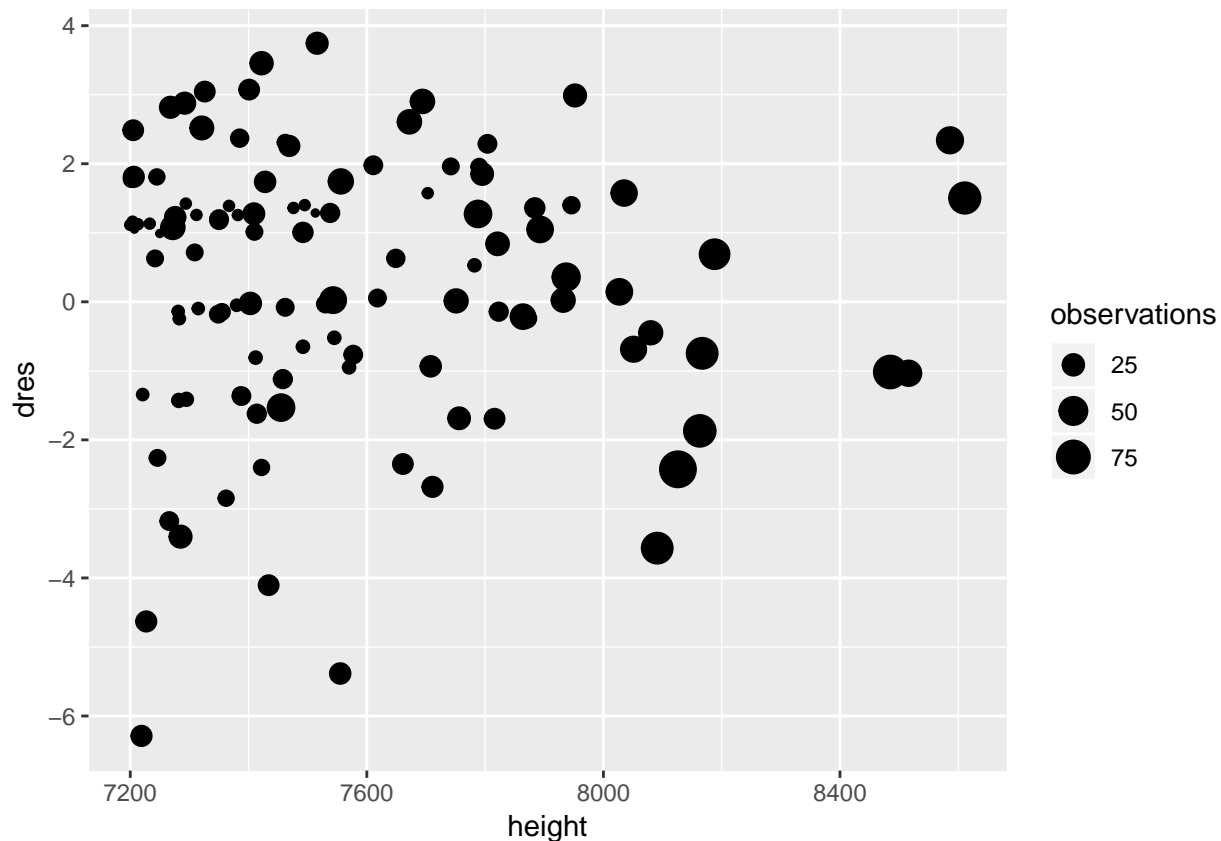
We plot the deviance residuals against the fitted values



The dots indicate a single covariate pattern (here a mountain or several mountains with identical height and prominence). The x -axis indicate the fitted probability of ascent success while the y -axis indicate the deviance residual. The size of the dots indicate the amount of observations (both failed and successful attempts).

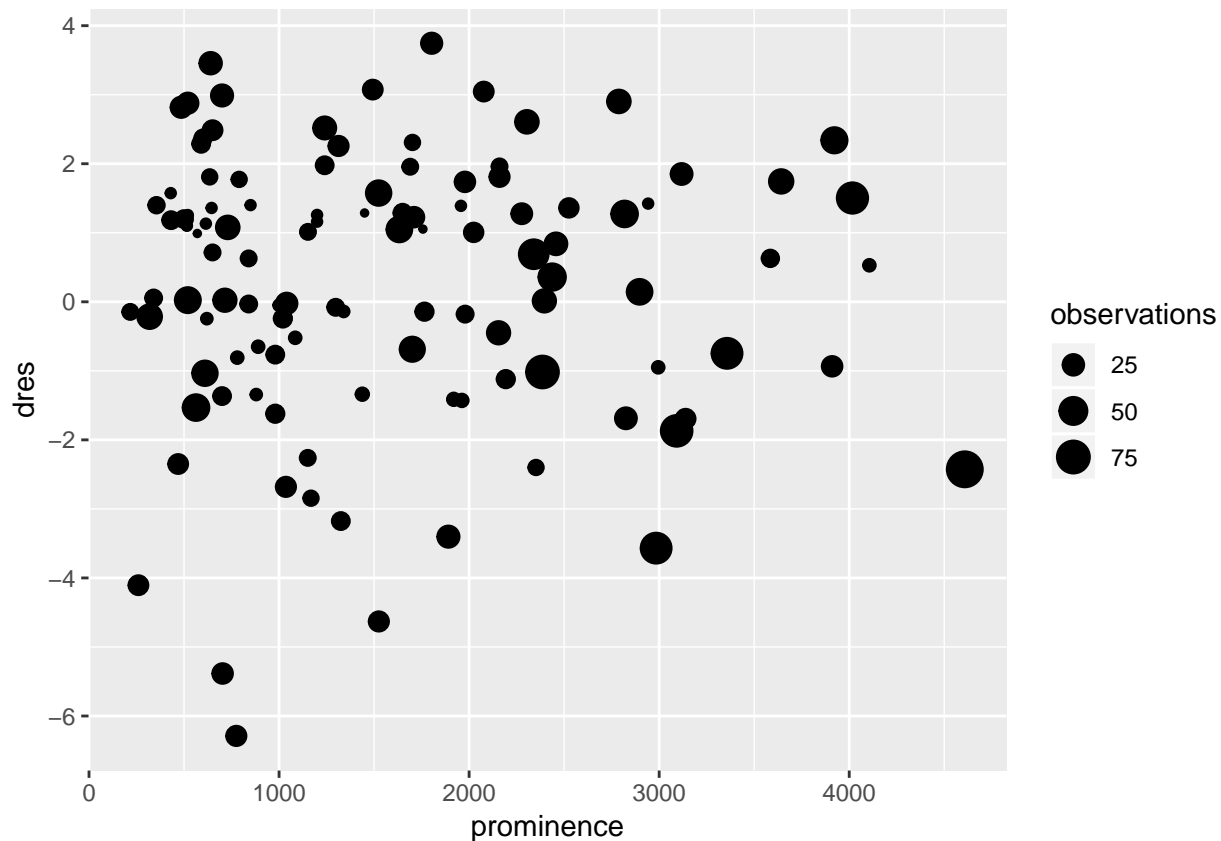
We see some indication of a larger spread of the residuals towards higher probabilities and some non-centering around lower probabilities, but observe no obvious structure. We investigate whether either of these trends are associated with a particular covariate by plotting the residuals as a function of **height** and **prominence**.

```
ggplot(df, aes(x = height, y = dres, size = observations)) + geom_point()
```



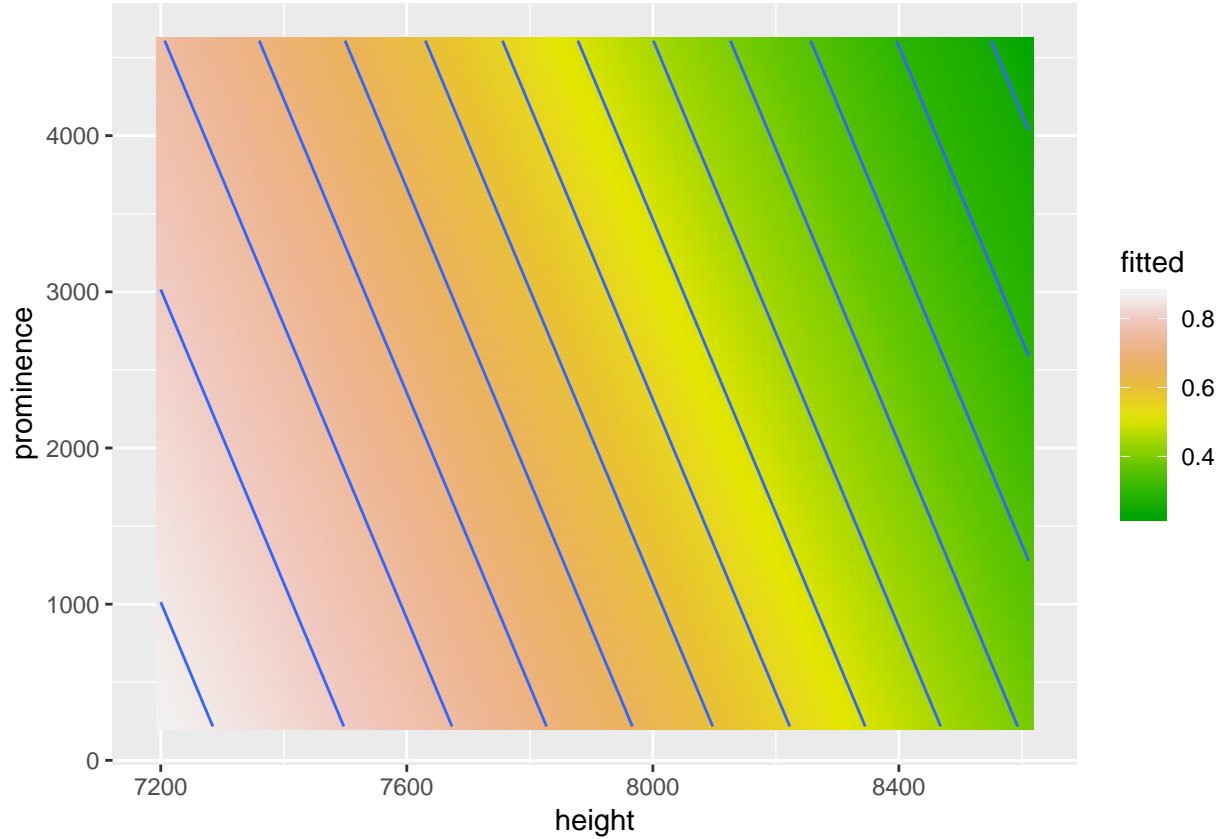
The x -axis now indicates height. We see some larger spread at lower altitudes, but note that there are only a small amount of summits with height above 8000m and it is quite natural to observe fewer outliers. We also observe a slight tendency towards negative residuals at greater altitudes with the exception of the two highest summits. This might indicate that there is some covariate that separates the summits above 8km. This further supports our claim that the model can be improved with the inclusion of additional covariate, e.g. some measure of summit infrastructure (amount of base camps).

```
ggplot(df, aes(x = prominence, y = dres, size = observations)) + geom_point()
```



The x -axis now indicates prominence. We see no apparent structure in the data.

Structure in the data would indicate that our model assumptions are unreasonable, and an obvious trend would motivate us to discard our model. However, the lack of structure does not necessarily indicate that our model is good. It will therefore be important to validate the model predictions using observations excluded from the model data.



We plot the predicted probability of ascent success (marked by color) as a function of **height** and **prominence**. The probability decreases with increasing **height** and **prominence**. The linear level curves are indicated by blue lines. We observe that the level curves are steep, indicating that **height** has a stronger model effect than **prominence**. This makes sense, as

$$\frac{\hat{\beta}_{\text{height}}}{\hat{\beta}_{\text{prominence}}} = 9.4$$

d)

Based on the model we have fitted, we want to estimate the probability of successfully ascending Mount Everest (height and prominence both equal 8848m), and Chogolisa (height 7665m and prominence 1624m). We denote Mt. Everest for x_1 and Chogolisa for x_2 . We use our fitted model to produce linear predictors η_i for both summits.

```
## Mt. Everest    Chogolisa
## -2.3237152    0.8678224
```

And we transform the linear predictors to probabilities using the response function $h(\eta) = \exp(\eta)/(1 + \exp(\eta))$,

```
## Mt. Everest    Chogolisa
## 0.08917783    0.70429238
```

We also want to make a 95% confidence interval for the probability of successfully ascending both summits. We use the fact that the coefficients found by maximum likelihood asymptotically will be normal distributed with mean β and variance $F^{-1}(\hat{\beta})$, $\hat{\beta} \sim N(\beta, F^{-1}(\hat{\beta}))$. We then get that the linear predictor, a linear

combination of the parameters β_j , is also normal distributed,

$$\widehat{LP} = \hat{\beta}_0 + x_{i1}\hat{\beta}_1 + x_{i2}\hat{\beta}_2 = x_i^T \hat{\beta} \sim N_1(x_i^T \beta, x_i F^{-1}(\hat{\beta}) x_i^T).$$

Which we can use to calculate the confidence intervals for the linear predictor η_i . We can then transform the linear predictor CI to the corresponding probability CI and get an interval for ascent success probability for Mount Everest and Chogolisa. As the estimated linear predictor is normally distributed, we get that

$$\frac{LP - \widehat{LP}}{\widehat{sd(LP)}} \sim N(0, 1).$$

We can then make a 95% confidence interval for the linear predictor, which will be

$$LP \in [\widehat{LP} - z_{0.025} \cdot \widehat{sd(LP)}, \widehat{LP} + z_{0.025} \cdot \widehat{sd(LP)}].$$

Then we take the $\exp()$ function of both limits of the interval, and we get that

$$\exp(LP) \in [\exp(\widehat{LP} - z_{0.025} \cdot \widehat{sd(LP)}), \exp(\widehat{LP} + z_{0.025} \cdot \widehat{sd(LP)})].$$

Earlier, we have found that

$$\frac{\pi_i}{1 - \pi_i} = \exp(LP),$$

which means that

$$\pi_i = \frac{\exp(LP)}{1 + \exp(LP)}.$$

We can then get the formula for a 95% confidence interval for the probability of a successful ascent by

$$\pi_{ME} \in \left[\frac{\exp(\widehat{LP} - z_{0.025} \cdot \widehat{sd(LP)})}{1 + \exp(\widehat{LP} - z_{0.025} \cdot \widehat{sd(LP)})}, \frac{\exp(\widehat{LP} + z_{0.025} \cdot \widehat{sd(LP)})}{1 + \exp(\widehat{LP} + z_{0.025} \cdot \widehat{sd(LP)})} \right],$$

where $\widehat{LP} = \hat{\beta}_I + \hat{\beta}_h \cdot x_{ME,h} + \hat{\beta}_p \cdot x_{ME,p}$ and $\widehat{sd(LP)} = \sqrt{x_i^T F^{-1}(\hat{\beta}) x_i}$.

For the linear predictor we get limits

##	2.5%	50 %	97.5%
## Mt. Everest	-2.8464382	-2.3237152	-1.8009922
## Chogolisa	0.7686346	0.8678224	0.9670102

Which is transformed to the probabilities

##	2.5%	50 %	97.5%
## Mt. Everest	0.05486572	0.08917783	0.1417303
## Chogolisa	0.68322546	0.70429238	0.7245232

The probability of reaching Mt. Everest is considerably smaller than for Chogolisa, with a predicted probability of 8.9% and a 95% confidence interval of [5.5%, 14.2%]. We note that the both the height and prominence of Mt. Everest are outside the data range used to fit the model. The prominence is almost double that of the greatest value in the data range. Thus we have no way to control if the model assumptions are valid for such large values of height and prominence. It is therefore not reasonable to use this model to estimate the probability of ascending Mt. Everest. In fact, common estimates for Mt. Everest success probability are $\sim 50\%$.

The model predicts the success rate for Chogolisa to be 70.4% with a 95% CI of [68.3%, 72.5%]. The height and prominence of Chogolisa lies inside the range of the data used to fit the model and it is reasonable to use the model to estimate this success rate. Even so, the observed data from Chogolisa, 20 successes and 2 failures, gives a success rate of $\frac{20}{22} = 0.91$, which is higher than the computed CI. Chogolisa is somewhat of an outlier, as the success rate of summits in our data set with height less than 8000m and prominence between 1000m and 2000m is 0.76. Nevertheless, this discrepancy indicates that our modelled response is highly variable and model predictions should be used with caution.

```
## [1] 0.7612245
```

Part 2: Poisson regression - Eliteserien 2018

a)

In this Part, we aim to simulate the remaining games in the Norwegian top division of football (Eliteserien). For each game, we assume that the score (the number of goals) of the home team is independent of the score of the away team. We assume that each team has a single parameter that measures its strength. We denote this strength parameter β_A for team A, β_B for team B, and so on.

Through watching football games, one could be made to believe that the goals scored by the away team in a football match is dependent on the goals scored by the home team and vice versa.

We therefore want to test if the assumption of independence between the goals scored by the home and away teams is reasonable. To do this, we first load the data set and make a contingency table of all the results, with the goals of the home team on the rows, and goals of the away team on the columns. We get the following contingency table.

```
##      0  1  2 3 4+
## 0    8 18  3 1  1
## 1   19 26 15 5  3
## 2   10 14 13 4  1
## 3   13 10  7 2  0
## 4+   8  7  3 1  0
```

We then want to test if the number of goals for home and away team are independent. We do this by conducting *Pearson's χ^2 test* on the contingency table. The test poses the following hypotheses

H_0 : The sampling distributions are independently chi-squared distributed, H_1 : They are not independently chi-squared distributed

We use the R function `chisq.test()` to compute the test statistic and the corresponding p-value.

```
##
## Pearson's Chi-squared test
##
## data:  contingency
## X-squared = 14.156, df = 16, p-value = 0.5871
```

We get a value of 14.156 for the test statistic, with a corresponding p-value of 0.5871. As this p-value is above any reasonable significance level, we keep the null hypothesis, and confirm that the goals scored by the home and away team are independent. This means that our assumption of independence is reasonable.

b)

Before we start simulating games, we want to construct the current standings in the Eliteserie based on all the results in our data set. By summing up the results from all games, we get the following table.

##	Team	Played	Won	Drawn	Lost	For	Against	GD	Points
## 1	Rosenborg	24	16	4	4	43	20	23	52
## 2	Brann	24	14	6	4	36	23	13	48
## 3	Molde	24	13	4	7	48	30	18	43
## 4	Haugesund	24	12	5	7	36	28	8	41
## 5	Ranheim_TF	24	11	5	8	38	40	-2	38
## 6	Vaalerenga	24	10	6	8	35	37	-2	36

```
## 7          Odd      24  9    7   8 35      29  6    34
## 8          Tromsøe   24 10    3  11 35      33  2    33
## 9          Sarpsborg08 24  9    5  10 39      34  5    32
## 10         Kristiansund 24  8    7   9 32      35 -3    31
## 11         Bodø/Glimt 24  6    9   9 28      30 -2    27
## 12         Stjøensvollene 24  6    8  10 38      38  0    26
## 13         Lillestrøm 24  6    7  11 26      37 -11   25
## 14         Stabæk     24  5    8  11 29      43 -14   23
## 15         Start      24  6    5  13 24      42 -18   23
## 16 Sandefjord Fotball 24  2    9  13 24      47 -23   15
```

c)

We now want to estimate the intercept, home advantage and strength parameters for each team. Then we produce a ranking based on the estimated strengths and compare with the rankings from b). To estimate the parameters, we create our own function `myglm` that performs the regression by maximum likelihood. The function uses the built in `optim` function to find the coefficients that maximizes the loglikelihood function (minimizes the negative of the loglikelihood, $-\ell(\beta)$).

```
##          Team      Strength
## 1      Rosenborg 0.366945548
## 2          Molde 0.279321007
## 3          Brann 0.225715115
## 4      Haugesund 0.141566217
## 5          Odd  0.099954079
## 6      Sarpsborg08 0.097625830
## 7          Tromsøe 0.060091773
## 8      Stjøensvollene 0.049639590
## 9          Vaalerenga 0.014445633
## 10     Kristiansund 0.012621369
## 11     Ranheim_TF 0.008439525
## 12     Bodø/Glimt 0.000000000
## 13     Lillestrøm -0.132589021
## 14     Stabæk    -0.148121316
## 15     Start     -0.225876528
## 16 Sandefjord Fotball -0.291815679

## [1] "Intercept: "
## [1] 0.1003129
## [1] "Home advantage: "
## [1] 0.4020541
##
## Call:
## glm(formula = goals ~ -1 + X, family = "poisson")
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.0205  -0.8748  -0.2014   0.5761   2.8679
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## XIntercept      0.100304   0.068489   1.465   0.1430
```

```
## XHomeAdvantage      0.402068    0.087521    4.594 4.35e-06 ***
## XRosenborg          0.366956    0.168373    2.179 0.0293 *
## XMolde              0.279264    0.168369    1.659 0.0972 .
## XLillestroem       -0.132857    0.168934   -0.786 0.4316
## XOdd               0.099975    0.166394    0.601 0.5480
## XHaugesund          0.141121    0.166320    0.848 0.3962
## XSandefjord_Fotball -0.291865    0.164767   -1.771 0.0765 .
## XRanheim_TF         0.008343    0.169495    0.049 0.9607
## XBrann              0.225678    0.165557    1.363 0.1728
## XSarpsborg08        0.097553    0.166444    0.586 0.5578
## XStabaek           -0.148047    0.168914   -0.876 0.3808
## XTromsoe            0.060348    0.166332    0.363 0.7167
## XStart             -0.225884    0.165079   -1.368 0.1712
## XVaalerenga         0.014465    0.169280    0.085 0.9319
## XKristiansund       0.012376    0.166170    0.074 0.9406
## XStroemsgodset      0.049657    0.166211    0.299 0.7651
## XBodoeGlimt         NA          NA          NA      NA
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
##      Null deviance: 499.35  on 384  degrees of freedom
## Residual deviance: 384.12  on 367  degrees of freedom
## AIC: 1135.3
##
## Number of Fisher Scoring iterations: 5
```

If we compare the results from our function `myglm` with the built-in function `glm`, we see that the regression coefficients are equal (to a precision of 4 digits).

We have set the strength of Bodø Glimt to zero, $\beta_{BodoeGlimt} = 0$. This means that the strength of Bodø Glimt is the “reference strength”, and the strength of every team is just the team’s strength compared to Bodø Glimt. This mean that all teams with $\beta_A > 0$ will be stronger than Bodø Glimt, and all teams with $\beta_A < 0$ will be weaker than Bodø Glimt. Also, a higher (lower) value of β_A indicates a stronger (weaker) team.

We get a coefficient for the intercept of $\beta_{Intercept} = 0.1003129$. This means that, if the teams are equally good (equal strength coefficient), one would expect the away team to score $\exp(\beta_{Intercept}) = 1.11$ goals on average. The coefficient for the home advantage is $\beta_{HomeAdvantage} = 0.4020541$. This means that, if the teams are equally good, one would expect the home team to score $\exp(\beta_{Intercept} + \beta_{HomeAdvantage}) = 1.65$ goals on average. This also means that teams are expected to score $\exp(\beta_{HomeAdvantage}) = 1.49$ times as many goals in home games as in away games.

Also, we see that Rosenborg has the significantly highest coefficient, and Sandefjord has the significantly lowest coefficient. This is what we would expect, seeing as they have been the best and worst team this season.

Now we want to compare the strength coefficient ranking with the actual ranking from the season so far. We therefore print the two rankings side by side.

##	Strength	Ranking
## 1	Rosenborg	Rosenborg
## 2	Molde	Brann
## 3	Brann	Molde
## 4	Haugesund	Haugesund
## 5	Odd	Ranheim_TF

## 6	Sarpsborg08	Vaalerenga
## 7	Tromsoe	Odd
## 8	Stroemsgodset	Tromsoe
## 9	Vaalerenga	Sarpsborg08
## 10	Kristiansund	Kristiansund
## 11	Ranheim_TF	BodoeGlimt
## 12	BodoeGlimt	Stroemsgodset
## 13	Lillestroem	Lillestroem
## 14	Stabaek	Stabaek
## 15	Start	Start
## 16	Sandefjord_Fotball	Sandefjord_Fotball

From the comparison, we get some really interesting results. Based on the strength ranking, we can say that teams that are higher on the actual ranking have “overachieved”, while teams that are lower on the actual ranking have “underachieved”.

Ranheim_TF is the stand out overachiever, placing in 11th on the strength ranking, and 5th on the actual ranking. One reason for this overachievement may be that Ranheim often win by only small scores, e.g. 1-0, while they lose with big scores, e.g. the scores on some of their losses were: 4-0, 4-0, 3-0, 4-1 and 3-1.

We also see that Brann and Molde have changed places on the actual ranking compared to the strength ranking. Again, this can be due to Brann winning by small scores, and losing by large scores, while Molde often wins by large scores and lose by small scores, e.g. some of Molde’s wins have been: 5-0, 4-0, 3-0, 5-1 and 5-1.

Other “overachievers” are Vålerenga and Bodø Glimt, while the “underachievers” are Odd, Tromsø, Sarpsborg 08 and Strømsgodset.

One final thought: If our explanation for why some teams “over”- and “underachieve” is correct (that they lose/win by small and large margins), then the strength ranking should be similar to a ranking based on goal difference. We therefore make a comparison between strength ranking and goal difference ranking.

##	Strength	RankingGD
## 1	Rosenborg	Rosenborg
## 2	Molde	Molde
## 3	Brann	Brann
## 4	Haugesund	Haugesund
## 5	Odd	Odd
## 6	Sarpsborg08	Sarpsborg08
## 7	Tromsoe	Tromsoe
## 8	Stroemsgodset	Stroemsgodset
## 9	Vaalerenga	Ranheim_TF
## 10	Kristiansund	Vaalerenga
## 11	Ranheim_TF	BodoeGlimt
## 12	BodoeGlimt	Kristiansund
## 13	Lillestroem	Lillestroem
## 14	Stabaek	Stabaek
## 15	Start	Start
## 16	Sandefjord_Fotball	Sandefjord_Fotball

We see that all teams are now in the same place, except for a permutation of the teams Vålerenga, Kristiansund, Ranheim and Bodø Glimt. All these teams except Kristiansund have a goal difference of -2, and Kristiansund has a goal difference of -3, so their goal differences are almost equal. All in all, there may be some truth to our explanation.

d)

Finally, we want to investigate rankings by means of simulation instead of comparing estimated strength. To do this, we use the estimated strengths of each team, the intercept and the home advantage, and simulate the remaining games in the current season 1000 times.

In each of the 1000 simulations, we get the goals for the home team in each match, by drawing a random variable from the poisson distribution with parameter $\lambda_H = \exp(\beta_{Intercept} + \beta_{HomeAdvantage} + \beta_{HomeTeam} - \beta_{AwayTeam})$. Similarly, the goals of the away team in each match is drawn from a poisson distribution with parameter $\lambda_A = \exp(\beta_{Intercept} - \beta_{HomeTeam} + \beta_{AwayTeam})$. When all matches are “played”, the final ranking is computed based on the current ranking plus the newly simulated games. The 1000 final rankings are stored in a .rds file. In this way we can load the results, instead of running the simulation multiple times.

After simulating the 48 remaining games 1000 times, we want to do some inference on the final results. We first investigate the “average final ranking”, that is, the average points, goals, wins etc. for each team. In addition, we look at how many times each team has placed in each place.

##	Team	Played	Won	Drawn	Lost	For	Against	GD	Points
## 1	Rosenborg	30	19.8	5.1	5.0	55.9	25.5	30.4	64.7
## 2	Brann	30	17.2	7.3	5.5	46.7	29.5	17.3	59.0
## 3	Molde	30	16.2	5.4	8.4	58.5	36.4	22.2	54.0
## 4	Haugesund	30	14.6	6.3	9.0	45.2	35.8	9.4	50.2
## 5	Ranheim_TF	30	13.2	6.4	10.4	46.2	48.6	-2.4	46.0
## 6	Vaalerenga	30	12.3	7.4	10.3	43.4	45.2	-1.9	44.3
## 7	Odd	30	11.2	8.3	10.4	43.4	37.8	5.6	42.1
## 8	Tromsø	30	12.4	4.4	13.2	43.6	41.2	2.4	41.6
## 9	Sarpsborg08	30	11.4	6.3	12.2	47.9	42.1	5.7	40.7
## 10	Kristiansund	30	10.4	8.5	11.1	40.8	42.8	-2.0	39.8
## 11	Stroemsgodset	30	8.6	9.4	12.0	47.2	45.6	1.6	35.3
## 12	Bodø/Glimt	30	8.1	10.4	11.5	35.8	39.0	-3.2	34.6
## 13	Lillestrøm	30	7.5	8.3	14.2	32.6	47.8	-15.2	30.7
## 14	Stabæk	30	6.8	9.4	13.8	36.2	52.5	-16.4	29.7
## 15	Start	30	7.2	6.3	16.6	29.8	53.5	-23.7	27.8
## 16	Sandefjord Fotball	30	3.2	10.1	16.7	29.9	59.7	-29.7	19.7

From the average rating, we see that Rosenborg win the Eliteserie by a margin of ≈ 6 points on average. Brann claims silver, and Molde bronze. Sandefjord ends bottom with a margin of ≈ 8 points, with Start also getting relegated and Stabæk claiming the play-off place. Ranheim (surprisingly) claim 5th place with a negative goal difference. We also see that the table is (almost) unchanged after simulating the remaining 48 games. The only difference is that Strømsgodset overtake Bodø Glimt.

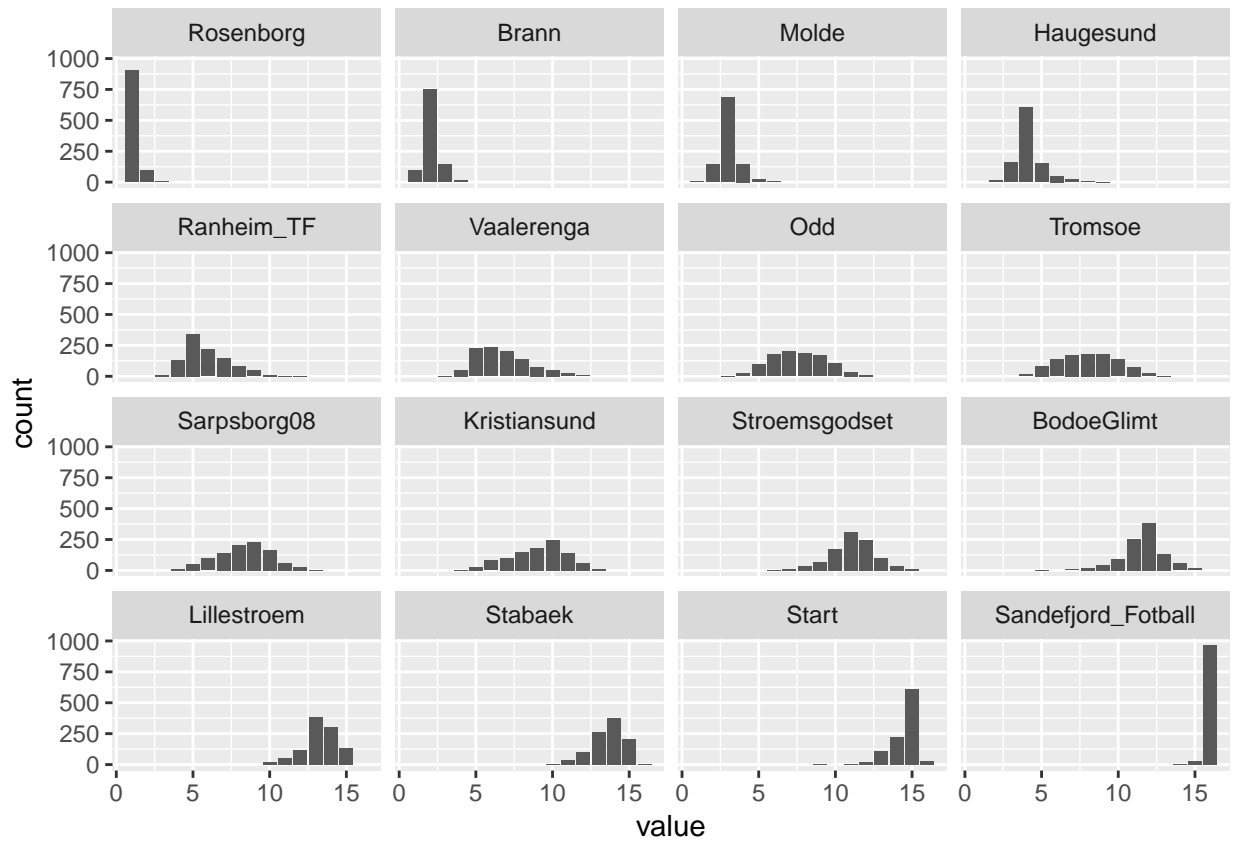
##	Team	One	Two	Three	Four	Five	Six	Seven	Eight	Nine	Ten
## 1	Rosenborg	901	95	4	0	0	0	0	0	0	0
## 2	Brann	94	750	145	11	0	0	0	0	0	0
## 3	Molde	5	142	682	146	19	6	0	0	0	0
## 4	Haugesund	0	13	157	606	151	50	19	3	1	0
## 5	Ranheim_TF	0	0	10	134	344	218	150	82	48	11
## 6	Vaalerenga	0	0	1	49	224	234	200	137	77	51
## 7	Odd	0	0	1	22	96	177	200	185	169	103
## 8	Tromsø	0	0	0	15	84	135	174	179	179	135
## 9	Sarpsborg08	0	0	0	14	55	96	141	207	230	168
## 10	Kristiansund	0	0	0	3	26	80	99	151	184	245
## 11	Stroemsgodset	0	0	0	0	0	4	10	37	69	173
## 12	Bodø/Glimt	0	0	0	0	1	0	7	19	42	89
## 13	Lillestrøm	0	0	0	0	0	0	0	0	0	17
## 14	Stabæk	0	0	0	0	0	0	0	0	0	8

## 15	Start	0	0	0	0	0	0	0	0	1	0
## 16	Sandefjord_Fotball	0	0	0	0	0	0	0	0	0	0
##	Eleven	Twelve	Thirteen	Fourteen	Fifteen	Sixteen					
## 1	0	0	0	0	0	0					
## 2	0	0	0	0	0	0					
## 3	0	0	0	0	0	0					
## 4	0	0	0	0	0	0					
## 5	2	1	0	0	0	0					
## 6	21	6	0	0	0	0					
## 7	35	12	0	0	0	0					
## 8	76	21	2	0	0	0					
## 9	62	26	1	0	0	0					
## 10	139	63	10	0	0	0					
## 11	312	245	101	39	10	0					
## 12	255	385	131	56	15	0					
## 13	55	116	381	301	130	0					
## 14	35	101	264	378	210	4					
## 15	8	24	110	223	608	26					
## 16	0	0	0	3	27	970					

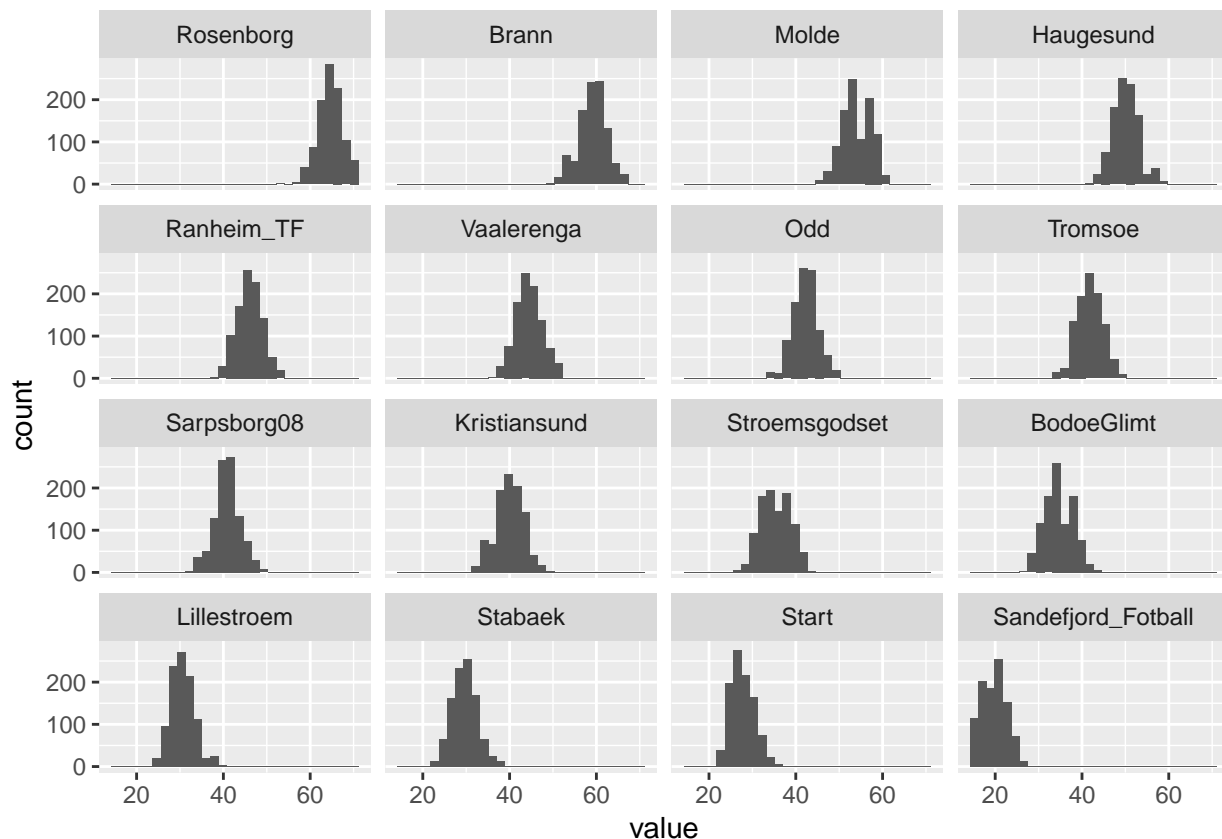
We see that Rosenborg win the Eliteserie with 90.1% probability, with Brann having 9.4% and Molde having 0.5% chance of winning. Also, Rosenborg get a medal in every simulation. Brann also claim a medal with 98.9% probability, with silver being the most likely result at 75% probability. Moreover, Molde gets bronze with 68.2% probability. Other teams with a chance of claiming a medal are Haugesund (17%), Ranheim (1%), Vålerenga (0.1%) and Odd (0.1%).

In the other end of the table, Sandefjord end bottom with 97% probability, managing a play-off place with only 0.3% probability. Start get relegated with 63.4% probability, manage play-off with 22.3% probability, and secure their place in the Eliteserie with a probability of 14.3%. Interestingly, in one simulation Start managed to get all the way up to 9th! Stabæk get relegated with a probability of 21.4%, managing play-off with probability 37.8%, and safe ground with probability 40.8%. Other teams in danger of relegation/play-off are Lillestrøm (13%/30.1%), Bodø Glimt (1.5%/5.6%) and Strømsgodset (1%/3.9%).

Based on the from these placings, we make barcharts for the placings of each team.



We also make a histogram of the points acquired by each team.

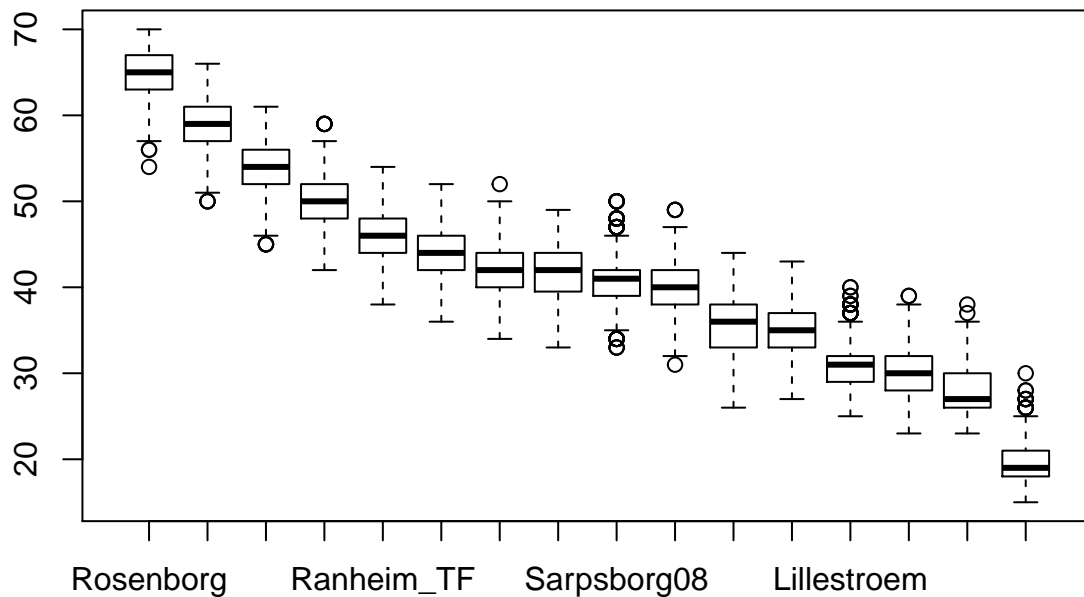


By the histogram of points, the distribution of points achieved in a season looks to be approximately normal. Seeing as the amount of points is a random variable, we can by the central limit theorem say that the mean of the points is normally distributed. We therefore find the average number of points for each team, as well as the standard deviation, and construct 90 % confidence intervals for the points of each team.

```
##           Teams mean  sd  low high
## 1      Rosenborg 64.7 2.8 60.0 69.3
## 2        Brann 59.0 3.1 53.9 64.1
## 3        Molde 54.0 3.1 49.0 59.1
## 4    Haugesund 50.2 3.0 45.4 55.1
## 5   Ranheim_TF 46.0 3.0 41.1 51.0
## 6   Vaalerenga 44.3 3.2 39.1 49.6
## 7         Odd 42.1 3.0 37.2 47.0
## 8     Tromsøe 41.6 3.1 36.5 46.6
## 9   Sarpsborg08 40.7 3.0 35.7 45.6
## 10  Kristiansund 39.8 3.2 34.6 45.0
## 11  Stroemsgodset 35.3 3.2 30.1 40.5
## 12   BodoeGlimt 34.6 3.0 29.7 39.6
## 13   Lillestroem 30.7 2.8 26.1 35.3
## 14     Stabaek 29.7 3.0 24.8 34.7
## 15        Start 27.8 2.9 23.1 32.4
## 16 Sandefjord_Fotball 19.7 2.6 15.5 24.0
```

From the confidence intervals, we see that Rosenborg will win or get silver. Brann may win, but will at worst get 4th. Meanwhile, all hope of staying in the division looks to be gone for Sandefjord.

At last we make a boxplot of the points for each team and interpret the results.



We see that the “boxes” (quartile 2 and 3) of Rosenberg, Brann, Molde and Haugesund are non-overlapping. This suggests that this will be the final standing among the top four in the table. Meanwhile, Sandefjord’s box is some distance behind Start’s box, emphasizing that Sandefjord will get relegated. Yet, there is still possibility for Start and Stabæk to avoid relegation/play-off.

Finally we include all code written in this document.

```
# Load the data
filepath <- "https://www.math.ntnu.no/emner/TMA4315/2018h/mountains"
mount <- read.table(file = filepath, header = TRUE, col.names = c("height",
  "prominence", "fail", "success"))
# Fit the model
fit = glm(cbind(success, fail) ~ height + prominence, data = mount, family = "binomial")
# Print summary
summary(fit)

# Calculate deviance, and print values from deviance test
D = deviance(fit)
cat("deviance: ", D)
cat("p-value: ", 1 - pchisq(D,110))

# Fit the residuals
library(ggplot2)
residual = residuals(fit, type = "deviance")
df = data.frame(fitted = fit$fitted.values, dres = residual, height = mount$height, prominence = mount$prominence)
df['observations'] = df['success'] + df['fail']
# Vs. fitted
```

```

ggplot(df, aes(x = fitted, y = dres, size = observations)) + geom_point()
# Vs. height
ggplot(df, aes(x = height, y = dres, size = observations)) + geom_point()
# Vs. prominence
ggplot(df, aes(x = prominence, y = dres, size = observations)) + geom_point()

# Make "Contour plot"
height = rep(seq(min(mount$height), max(mount$height), length.out = 100), each = 100)
prominence = rep(seq(min(mount$prominence), max(mount$prominence), length.out = 100), 100)

df2 = data.frame(fitted = exp(fit$coefficients[1] + fit$coefficients[2]*height + fit$coefficients[3]*prominence))
ggplot(df2, aes(x = height, y = prominence, fill = fitted)) + geom_raster() + geom_contour(aes(z = fitted))

# Intercept, height and prominence for Mount Everest and Chogolisa
x_ME <- c(1, 8848, 8848)
x_CH <- c(1, 7665, 1624)

# Linear predictors
eta_ME <- t(x_ME)%%fit$coefficients
eta_CH <- t(x_CH)%%fit$coefficients
# Std. deviations
std_ME <- sqrt(t(x_ME) %% vcov(fit) %% x_ME)
std_CH <- sqrt(t(x_CH) %% vcov(fit) %% x_CH)

# Limits for conf int of linear predictor
eta_limits_ME = c(eta_ME + qnorm(0.025) * std_ME, eta_ME, eta_ME + qnorm(0.975) * std_ME)
eta_limits_CH = c(eta_CH + qnorm(0.025) * std_CH, eta_CH, eta_CH + qnorm(0.975) * std_CH)

limits = rbind(eta_limits_ME, eta_limits_CH)
rownames(limits) = c('Mt. Everest', 'Chogolisa')
colnames(limits) = c('2.5%', '50 %', '97.5%')

limits[,2]

exp(limits[,2])/(1+ exp(limits[,2]))

limits
# Confidence intervals for the probabilities
prob = exp(limits)/(1 + exp(limits))
prob

smaller_summits = mount[mount$height <= 8000 & mount$prominence <= 2000 & mount$prominence >= 1000, ]
sum(smaller_summits['success']) / (sum(smaller_summits['success']) + sum(smaller_summits['fail']))

# Load results of matches
filepath <- "https://www.math.ntnu.no/emner/TMA4315/2018h/eliteserien2018"
eliteserie <- read.table(file = filepath, header = TRUE, colClasses = c("character",
    "character", "numeric", "numeric"))

# Initialize contingency table
contingency = matrix(rep(0,25),nrow = 5, ncol = 5)
rownames(contingency) = c("0","1","2","3","4+")
colnames(contingency) = c("0","1","2","3","4+")

```

```

# Compute values of the contingency table
for (i in 1:dim(eliteserie)[1]){
  if (eliteserie[i,3]<4 & eliteserie[i,4]<4){
    contingency[eliteserie[i,3]+1,eliteserie[i,4]+1] = contingency[eliteserie[i,3]+1,eliteserie[i,4]+1]
  } else if (eliteserie[i,3]<4 & !eliteserie[i,4]<4){
    contingency[eliteserie[i,3]+1,5] = contingency[eliteserie[i,3]+1,5] + 1
  } else if (!eliteserie[i,3]<4 & eliteserie[i,4]<4){
    contingency[5,eliteserie[i,4]+1] = contingency[5, eliteserie[i,4]+1] + 1
  } else{
    contingency[5,5] = contingency[5,5] + 1
  }
}
}
# Finally print the contingency table
print(contingency)

Xsq <- chisq.test(contingency)
print(Xsq)

# Initialize the current standings (ranking) for the teams
ranking = data.frame(
  Team = c("Rosenborg", "Molde", "Lillestroem", "Odd", "Haugesund", "Sandefjord_Fotball", "Ranheim_TF",
  stringsAsFactors = FALSE)

# Compute the standings by summing up over all matches
for (i in 1:dim(eliteserie)[1]){
  # First add number of matches played for both home team
  ranking[match(eliteserie[i,1], ranking[,1]),2] = ranking[match(eliteserie[i,1], ranking[,1]),2] + 1
  # and for the away team
  ranking[match(eliteserie[i,2], ranking[,1]),2] = ranking[match(eliteserie[i,2], ranking[,1]),2] + 1

  # Then add goals for and against home team
  ranking[match(eliteserie[i,1], ranking[,1]),6] = ranking[match(eliteserie[i,1], ranking[,1]),6] + eliteserie[i,3]
  ranking[match(eliteserie[i,1], ranking[,1]),7] = ranking[match(eliteserie[i,1], ranking[,1]),7] + eliteserie[i,4]
  # and for and against away team
  ranking[match(eliteserie[i,2], ranking[,1]),6] = ranking[match(eliteserie[i,2], ranking[,1]),6] + eliteserie[i,3]
  ranking[match(eliteserie[i,2], ranking[,1]),7] = ranking[match(eliteserie[i,2], ranking[,1]),7] + eliteserie[i,4]

  # Then count wins, draws and losses for home team
  ranking[match(eliteserie[i,1], ranking[,1]),3] = ranking[match(eliteserie[i,1], ranking[,1]),3] + as.numeric(eliteserie[i,3]>eliteserie[i,4])
  ranking[match(eliteserie[i,1], ranking[,1]),4] = ranking[match(eliteserie[i,1], ranking[,1]),4] + as.numeric(eliteserie[i,3]==eliteserie[i,4])
  ranking[match(eliteserie[i,1], ranking[,1]),5] = ranking[match(eliteserie[i,1], ranking[,1]),5] + as.numeric(eliteserie[i,3]<eliteserie[i,4])

  # Finally count wins, draws and losses for away team
  ranking[match(eliteserie[i,2], ranking[,1]),3] = ranking[match(eliteserie[i,2], ranking[,1]),3] + as.numeric(eliteserie[i,3]>eliteserie[i,4])
  ranking[match(eliteserie[i,2], ranking[,1]),4] = ranking[match(eliteserie[i,2], ranking[,1]),4] + as.numeric(eliteserie[i,3]==eliteserie[i,4])
  ranking[match(eliteserie[i,2], ranking[,1]),5] = ranking[match(eliteserie[i,2], ranking[,1]),5] + as.numeric(eliteserie[i,3]<eliteserie[i,4])
}
# Points is computed by 3 points for win, 1 for draw and 0 for loss
ranking[,9] = 3*ranking[,3] + ranking[,4]
# Goal difference is goals scored minus goals conceded
ranking[,8] = ranking[,6] - ranking[,7]
# We rank the teams first by points, then goal difference, finally by goals scored
ranking = ranking[order(ranking$Points, ranking$GD, ranking$For, decreasing = T),]

```

```

# Rename rows after sorting
row.names(ranking) <- seq(1,16)
# Display the standings
print(ranking)

# Initialize the design matrix X, which is a 2*192 x 18 matrix.
X = matrix(0, nrow = dim(eliteserie)[1]*2, ncol = 18)
colnames(X) = c("Intercept", "HomeAdvantage", "Rosenborg", "Molde", "Lillestroem", "Odd", "Haugesund",

# First we alter the design matrix for all "home games"
for (i in 1:dim(eliteserie)[1]){
  # Intercept
  X[i,1] = 1

  # Home team gets x_home = 1
  X[i,2] = 1

  # x_homeTeam is set to 1
  X[i, match(eliteserie[i,1], dimnames(X)[[2]])] = 1

  # x_awayTeam is set to -1
  X[i, match(eliteserie[i,2], dimnames(X)[[2]])] = -1
}

# Then alter design matrix for all "away games"
for (i in 1:dim(eliteserie)[1]){
  # Intercept
  X[dim(eliteserie)[1] + i,1] = 1

  # Away team, so x_home = 0

  # x_homeTeam is set to -1
  X[dim(eliteserie)[1] + i, match(eliteserie[i,1], dimnames(X)[[2]])] = -1

  # x_awayTeam is set to 1
  X[dim(eliteserie)[1] + i, match(eliteserie[i,2], dimnames(X)[[2]])] = 1
}

# And we now have the desired design matrix

# We then make the response vector (first all home goals, then all away goals)
goals = c(eliteserie[,3], eliteserie[,4])

# Finally we create the function that does the regression
myglm <- function(formula, data = list(), contrasts = NULL, ...){

  # Extract model matrix & responses
  mf <- model.frame(formula = formula, data = data)
  X <- model.matrix(attr(mf, "terms"), data = mf, contrasts.arg = contrasts)
  y <- model.response(mf)
  terms <- attr(mf, "terms")

  # Add code here to calculate coefficients, residuals, fitted values, etc...

```

```

# and store the results in the list est
est <- list(terms = terms, model = mf)

# Store call and formula used
est$call <- match.call()
est$formula <- formula

# Store information used for model fit
est$X <- X
est$y <- y

# Fit model (using optim function in r)
# Initial guess is just 0
est$initial <- rep(0,18)
# Function returns -maximum likelihood
est$fcn <- function(coeff){
  return(-y%*%X%*%coeff + sum(exp(X%*%coeff)))
}

# Use optim function to estimate coefficients
est$coefficients = optim(est$initial, est$fcn, method = "BFGS", hessian = TRUE)

# We only need to return the coefficients from the regression
return(est$coefficients)
}

# Conduct the regression
coeff <- myglm(goals~ -1 + X)

# Find relative strength by setting Bodø Glimt to 0
coeff$par[3:17] = coeff$par[3:17] - coeff$par[18]
coeff$par[18] = 0

# Create a strength ranking based on the calculated coefficients
strengthRanking = data.frame(
  Team = c("Rosenborg", "Molde", "Lillestroem", "Odd", "Haugesund", "Sandefjord_Fotball", "Ranheim_TF",
  Strength = coeff$par[3:18],
  stringsAsFactors = FALSE)

# Sort after strength
strengthRanking = strengthRanking[order(strengthRanking$Strength, decreasing = T),]
# Rename rows after sorting
row.names(strengthRanking) <- seq(1,16)

# Print the strength ranking
print(strengthRanking)
# Print intercept and home advantage
print("Intercept: ")
print(coeff$par[1])
print("Home advantage: ")
print(coeff$par[2])

```

```

# Compare with the results from the built in function glm
summary(glm(goals ~ -1 + X, family = "poisson"))

# Compare the calculated strength ranking with the current standings
compareableRanking = data.frame(
  Strength = strengthRanking[,1],
  Ranking = ranking[,1],
  stringsAsFactors = FALSE)
row.names(compareableRanking) <- seq(1,16)
print(compareableRanking)

# We make a ranking based on goal difference.
rankingGD = ranking[order(ranking$GD, ranking$For, decreasing = T),]
# And compare it to the strength ranking.
compareableRanking2 = data.frame(
  Strength = strengthRanking[,1],
  RankingGD = rankingGD[,1],
  stringsAsFactors = FALSE)
row.names(compareableRanking2) <- seq(1,16)
print(compareableRanking2)

# Load the unplayed matches
filepath <- "https://www.math.ntnu.no/emner/TMA4315/2018h/unplayed2018"
eliteserieUnplayed <- read.table(file = filepath, header = TRUE, colClasses = c("character",
  "character"))

# Add columns for goals in each match
eliteserieUnplayed$yh = rep(0,dim(eliteserieUnplayed)[1])
eliteserieUnplayed$ya = rep(0,dim(eliteserieUnplayed)[1])

# Initialize list to store final rankings
football <- list()

# Function to simulate one season
simseason <- function(){

  #Play the remaining matches
  eliteserieUnplayed[,3] = rpois(dim(eliteserieUnplayed)[1],exp(coeff$par[1]+coeff$par[2] + strengthRanking[match(eliteserieUnplayed[,1], ranking[,1]),1]))
  eliteserieUnplayed[,4] = rpois(dim(eliteserieUnplayed)[1],exp(coeff$par[1] - strengthRanking[match(eliteserieUnplayed[,1], ranking[,1]),1]))

  # Make the final ranking by starting with the matches already played
  rankingFinal <- ranking
  # Each team will play 30 times
  rankingFinal$Played = rep(30,16)

  for (i in 1:dim(eliteserieUnplayed)[1]){
    # Add goals for and against home team
    rankingFinal[match(eliteserieUnplayed[i,1], rankingFinal[,1]),6] = rankingFinal[match(eliteserieUnplayed[i,1], rankingFinal[,1]),6] + eliteserieUnplayed[i,3]
    rankingFinal[match(eliteserieUnplayed[i,1], rankingFinal[,1]),7] = rankingFinal[match(eliteserieUnplayed[i,1], rankingFinal[,1]),7] + eliteserieUnplayed[i,4]
    # and for and against away team
    rankingFinal[match(eliteserieUnplayed[i,2], rankingFinal[,1]),6] = rankingFinal[match(eliteserieUnplayed[i,2], rankingFinal[,1]),6] + eliteserieUnplayed[i,4]
    rankingFinal[match(eliteserieUnplayed[i,2], rankingFinal[,1]),7] = rankingFinal[match(eliteserieUnplayed[i,2], rankingFinal[,1]),7] + eliteserieUnplayed[i,3]
  }
}

```



```

# List the points for each team in each simulation
# And calculate number of times each team placed in each place:
# Average the final ranking over all the simulations:
for (i in 1:1000){
  cur = data[[i]]
  for (j in 1:16){
    # Add the placing of each team
    placings[i,match(cur[j,1],colnames(placings))] = j
    # Add the points of each team
    points[i,match(cur[j,1],colnames(points))] = cur[j,9]
    # Find number of placings for each team
    numPlace[match(cur[j,1], numPlace[,1]),j+1] = numPlace[match(cur[j,1], numPlace[,1]),j+1] + 1
    # Average the final ranking
    rankingAverage[match(cur[j,1], rankingAverage[,1]),3:9] = rankingAverage[match(cur[j,1], rankingAverage[,1]),3:9] + 1
  }
}

# Sort the table in the right order
rankingAverage = rankingAverage[order(rankingAverage$Points, rankingAverage$GD, rankingAverage$For, rankingAverage$Against)]
# Rename rows after sorting
row.names(rankingAverage) <- seq(1,16)

# Round to 1 decimal
rankingAverage[,2:9] = round(rankingAverage[,2:9],1)

print(rankingAverage)

# Print number of placings for each team
print(numPlace)

library("ggplot2")
library("reshape2")
# Plot the placings of each team in a barchart
placingsplot <- melt(placings)
ggplot(placingsplot, aes(x = value)) + geom_bar() + facet_wrap(~variable)

# Plot the points of each team in a histogram
pointsplot <- melt(points)
ggplot(pointsplot, aes(x = value)) + geom_histogram() + facet_wrap(~variable)

# Initialize data frame to store confidence intervals
confidence_intervals = data.frame(
  Teams = numPlace$Team, # All the teams
  mean = 0,
  sd = 0,
  low = 0,
  high = 0,
  stringsAsFactors = FALSE)

for (i in 1:16){
  confidence_intervals$mean[i] = mean(points[[i]])
  confidence_intervals$sd[i] = sd(points[[i]])
  confidence_intervals$low[i] = confidence_intervals$mean[i] - confidence_intervals$sd[i]*1.645
}

```

```
confidence_intervals$high[i] = confidence_intervals$mean[i] + confidence_intervals$sd[i]*1.645
}

# Round all values
confidence_intervals[,2:5] = round(confidence_intervals[,2:5],1)

print(confidence_intervals)

# Plot boxplot of points for each team to compare points of each team
boxplot(points)
```