

PREDICTING THE SURVIVAL OF PASSENGERS IN THE TITANIC ACCIDENT

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Univariable logistic regression

Load the necessary library and the dataset

```
library(tidyverse)
```

```
titanic <- read_csv("C:/Users/PC/OneDrive/Desktop/Data Science/Datasets/titanic.csv")
```

```
titanic$Sex <- as.factor(titanic$Sex)
```

```
titanic$Sex <- as.numeric(titanic$Sex)
```

```
titanic$Pclass <- as.factor(titanic$Pclass)
```

Cross table for quick intuition

```
table(titanic$Survived, titanic$Pclass)
```

	1	2	3
0	80	97	372
1	134	87	119

- More people from the first class survived
- Many more people from the third class perished

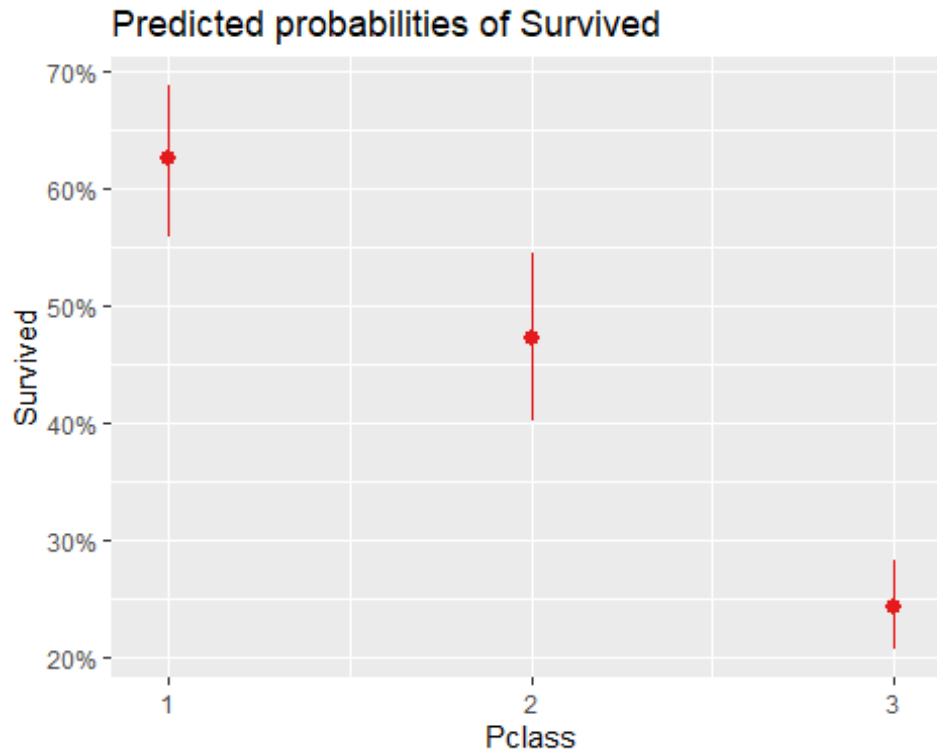
Run logistic regression with categorical predictors

```
m1 <- glm(Survived~Pclass, titanic, family = binomial)
```

Plot predictions

```
library(sjPlot)
```

```
plot_model(m1, type = "eff", terms = c("Pclass"))
```



- The probability of survival in class one is more than 60%
- The probability of survival in the third class is less than 25%
- The probability of survival in class two is less than 45 %

Get probabilities and odds ratios

```
library(emmeans)
```

```
emmeans(m1, pairwise~Pclass, type = "response",
infer = T)
```

```
$emmeans
```

Pclass	prob	SE	df	asympt.LCL	asympt.UCL	null	z.ratio	p.value
1	0.626	0.0331	Inf	0.559	0.688	0.5	3.651	0.0003
2	0.473	0.0368	Inf	0.402	0.545	0.5	-0.737	0.4612
3	0.242	0.0193	Inf	0.206	0.282	0.5	-10.822	<.0001

Confidence level used: 0.95

Intervals are back-transformed from the logit scale

Tests are performed on the logit scale

```
$contrasts
```

contrast	odds.ratio	SE	df	asympt.LCL	asympt.UCL	null	z.ratio
Pclass1 / Pclass2	1.87	0.382	Inf	1.16	3.01	1	3.056
Pclass1 / Pclass3	5.24	0.923	Inf	3.46	7.91	1	9.395
Pclass2 / Pclass3	2.80	0.509	Inf	1.83	4.29	1	5.684

```
p.value
0.0063
<.0001
<.0001
```

Confidence level used: 0.95

Conf-level adjustment: tukey method for comparing a family of 3 estimates

Intervals are back-transformed from the log odds ratio scale

P value adjustment: tukey method for comparing a family of 3 estimates

Tests are performed on the log odds ratio scale

- First class passengers are significantly more likely to survive than dying
- The probability of survival in third class is significantly lower than that of dying

Check

```
titanic2 <- titanic %>% slice(1:400)
m2 <- glm(Survived~Pclass, titanic2, family = binomial)
emmeans(m2, ~Pclass, infer = T, type = "response")
```

Pclass	prob	SE	df	asyp.LCL	asyp.UCL	null	z.ratio	p.value
1	0.576	0.0515	Inf	0.473	0.673	0.5	1.454	0.1460
2	0.434	0.0544	Inf	0.332	0.542	0.5	-1.204	0.2287
3	0.302	0.0306	Inf	0.246	0.365	0.5	-5.764	<.0001

Confidence level used: 0.95

Intervals are back-transformed from the logit scale

Tests are performed on the logit scale

Reverse odds ratios if needed

```
emmeans(m1, ~Pclass, type = "response") %>%
pairs(reverse = T, infer = T)
```

contrast	odds.ratio	SE	df	asyp.LCL	asyp.UCL	null	z.ratio
Pclass2 / Pclass1	0.535	0.1090	Inf	0.332	0.864	1	-3.056
Pclass3 / Pclass1	0.191	0.0337	Inf	0.126	0.289	1	-9.395
Pclass3 / Pclass2	0.357	0.0647	Inf	0.233	0.546	1	-5.684

```
p.value
0.0063
<.0001
<.0001
```

Confidence level used: 0.95

Conf-level adjustment: tukey method for comparing a family of 3 estimates

Intervals are back-transformed from the log odds ratio scale

P value adjustment: tukey method for comparing a family of 3 estimates

Tests are performed on the log odds ratio scale

- Reverse = T gives us odds ratios below one
- Passengers in 2nd class were 0.535 times as likely to survive compared to passengers in 1st class
- Passengers in 3rd class were 0.191 times as likely to survive compared to passengers in 1st class
- Passengers in 3rd class were 0.357 times as likely to survive compared to passengers in 2nd class

Get publication ready table

```
library(gtsummary)
```

```
fancy_table <- tbl_regression(
  m1, exponentiate = T, add_pairwise_contrasts = T) %>%
  add_significance_stars(
    hide_p = F, hide_se = T, hide_ci = F) %>%
  bold_p()
```

```
fancy_table
```

Characteristic	OR ^{1,2}	95% CI ²	p-value
Pclass			
Pclass2 / Pclass1	0.54**	0.33, 0.86	0.006
Pclass3 / Pclass1	0.19***	0.13, 0.29	<0.001
Pclass3 / Pclass2	0.36***	0.23, 0.55	<0.001

¹ *p<0.05; **p<0.01; ***p<0.001

How to produce model equations

```
library(equatiomatic)
```

```
extract_eq(m1)
```

$$\log \left[\frac{P(\text{Survived} = 1)}{1 - P(\text{Survived} = 1)} \right] = \alpha + \beta_1(\text{Pclass}_2) + \beta_2(\text{Pclass}_3)$$

How to get odds for categories

```
emmeans(m1, ~Pclass, infer = T) %>%  
  as_tibble() %>%  
  dplyr::select(Pclass, emmean) %>%  
  mutate(odds = exp(emmean)) %>%  
  mutate_if(is.numeric, ~round(., 2))
```

A tibble: 3 × 3

	Pclass	emmean	odds
	<fct>	<dbl>	<dbl>
1	1	0.52	1.67
2	2	-0.11	0.9
3	3	-1.14	0.32

- The odds of survival for 1st class passengers is 67% higher than those of dying
- 2nd class passengers have 10% lower odds of survival compared to their odds of dying
- 3rd class passengers face a significant 68% lower chance of survival relative to their odds of dying.

Contact logistic regression with numeric predictors

```
modela <- glm(Survived~Age, titanic, family = binomial)
```

Visualize predictions

```
library(sjPlot)  
summary(modela)
```

Call:

```
glm(formula = Survived ~ Age, family = binomial, data = titanic)
```

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	0.002310	0.153782	0.015	0.988018
Age	-0.013688	0.003958	-3.458	0.000544 ***

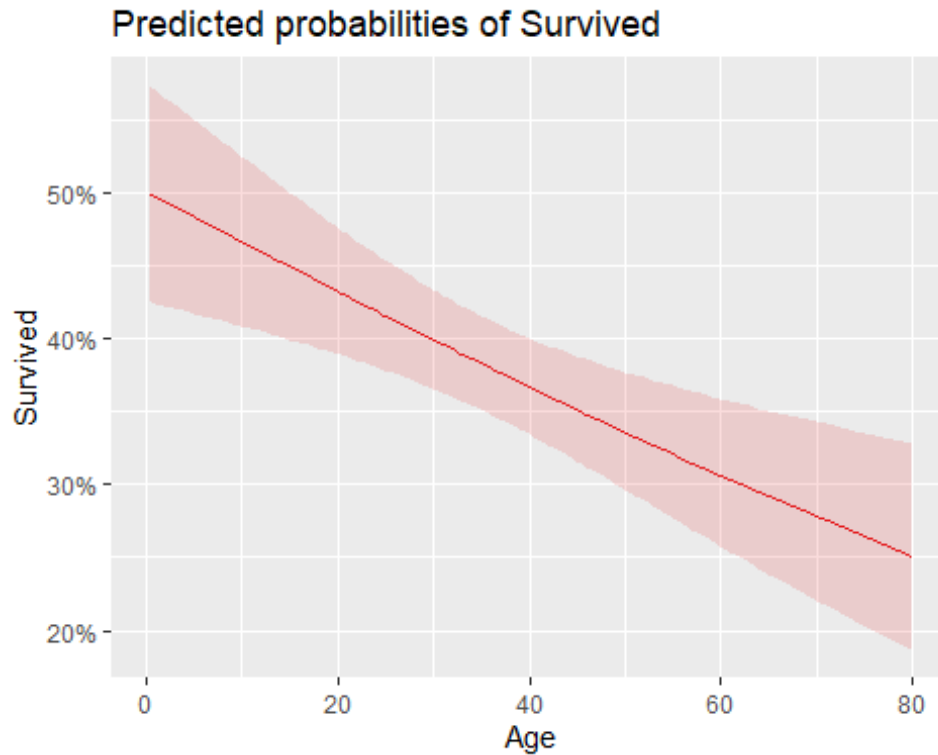
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 1182.8 on 888 degrees of freedom
Residual deviance: 1170.7 on 887 degrees of freedom
AIC: 1174.7

Number of Fisher Scoring iterations: 4

```
plot_model(modela, type = "pred", terms = "Age[all]")
```



- The probability of survival decreases with an increase in age
- Higher proportion of young people survived during the titanic accident as compared to older ones.

How to check for non-linearity

Use polynomial degrees based on age

```
modela1 <- glm(Survived~poly(Age, 2), data = titanic,  
               family = binomial)
```

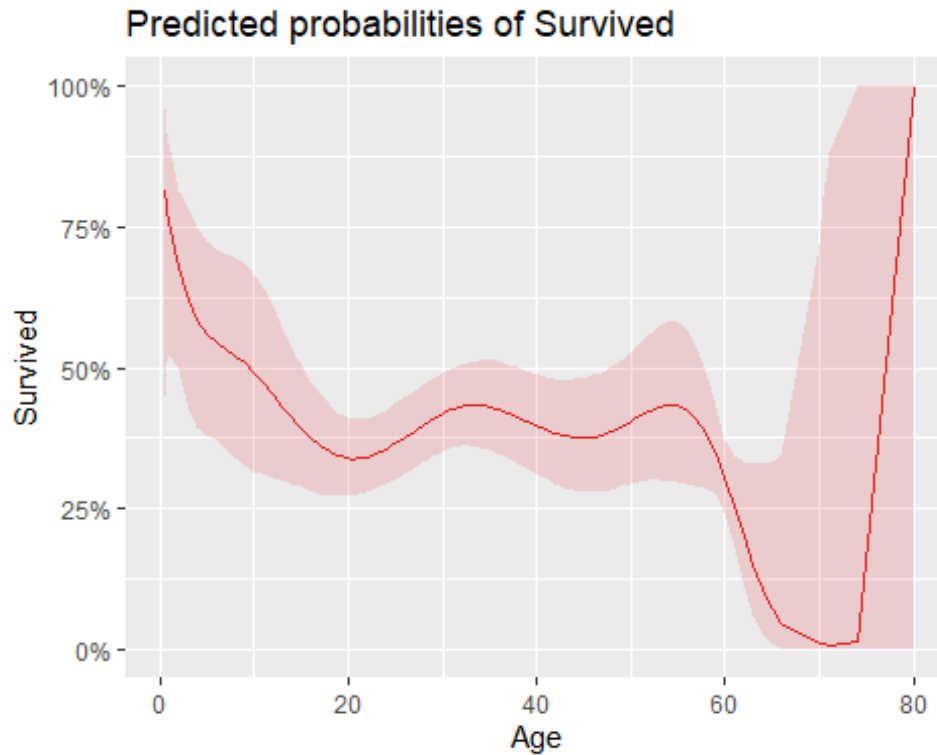
```
modela2 <- glm(Survived~poly(Age, 3), data = titanic,  
               family = binomial)
```

```
modela3 <- glm(Survived~poly(Age, 4), data = titanic,  
               family = binomial)
```

Dont overdo

```
modela4 <- glm(Survived~poly(Age, 10), data = titanic,  
               family = binomial)
```

```
plot_model(modela4, type = "pred", terms = "Age")
```



- It is difficult to interpret the survival rates of passengers due to complexity of the model.

```
summary(modela4)
```

Call:

```
glm(formula = Survived ~ poly(Age, 10), family = binomial, data = titanic)
```

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-0.50462	0.07648	-6.598	4.16e-11	***
poly(Age, 10)1	-8.28223	2.87775	-2.878	0.004	**
poly(Age, 10)2	0.96533	3.83039	0.252	0.801	
poly(Age, 10)3	-7.61989	7.14851	-1.066	0.286	
poly(Age, 10)4	4.66868	8.65548	0.539	0.590	
poly(Age, 10)5	3.66760	6.66164	0.551	0.582	
poly(Age, 10)6	6.87432	7.17499	0.958	0.338	
poly(Age, 10)7	7.23940	6.56334	1.103	0.270	
poly(Age, 10)8	3.12351	4.19513	0.745	0.457	
poly(Age, 10)9	-1.97385	3.24969	-0.607	0.544	
poly(Age, 10)10	-0.29224	3.01064	-0.097	0.923	

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 1182.8 on 888 degrees of freedom
Residual deviance: 1146.5 on 878 degrees of freedom
AIC: 1168.5

Number of Fisher Scoring iterations: 7

- Degrees more than 3 and 4 are rarely used due to the risk of overfitting, which makes the model overly complex and less interpretable as of the example above

Compare all the models we have created so far

`AIC(modela1, modela2, modela3, modela4)`

	df	AIC
modela1	3	1175.682
modela2	4	1167.686
modela3	5	1165.135
modela4	11	1168.470

- Select the model with the lowest aic. aic measures the relative quality of statistical models
- Here we will select the model with 3rd polynomial degree and move on

Another method used to choose polynomial degrees

`tab_model(modela4)`

Survived			
Predictors	Odds Ratios	CI	p
(Intercept)	0.60	0.51 – 0.74	<0.201
Age [1st degree]	0.00	0.00 – 63.93	0.304
Age [2nd degree]	2.63	0.00 – NA	0.801
Age [3rd degree]	0.00	0.00 – NA	0.001
Age [4th degree]	106.56	0.00 – NA	0.590
Age [5th degree]	39.16	0.15 – NA	0.582

Age [6th degree]	967.12	0.90 – NA	0.338
Age [7th degree]	1393.25	1.24 – NA	0.270
Age [8th degree]	22.73	0.09 – NA	0.457
Age [9th degree]	0.14	0.00 – 219448.52	0.544
Age [10th degree]	0.75	0.00 – 2800.19	0.923
<hr/>			
Observations	889		
R ² Tjur	0.038		

- Here we choose the polynomial degree that is statistically significant.

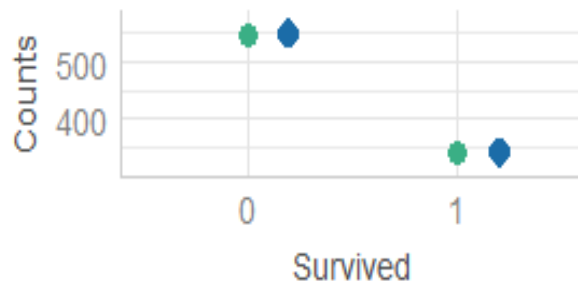
Check model assumptions

```
library(performance)
```

```
check_model(modela3)
```

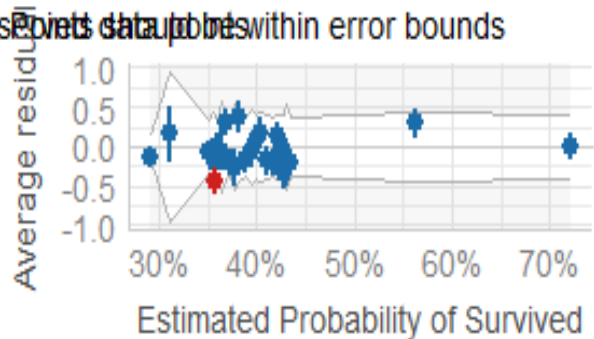
Posterior Predictive Check

Model-predicted intervals should include observed data points



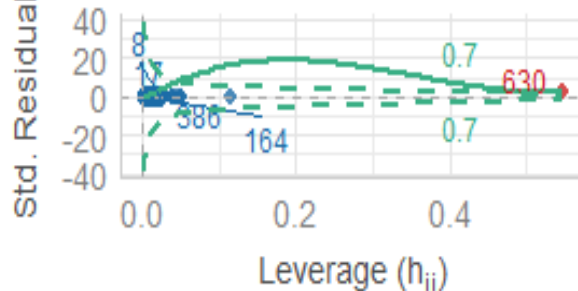
● Observed data ● Model-predicted data Within error bounds ● no ● yes

Binned Residuals



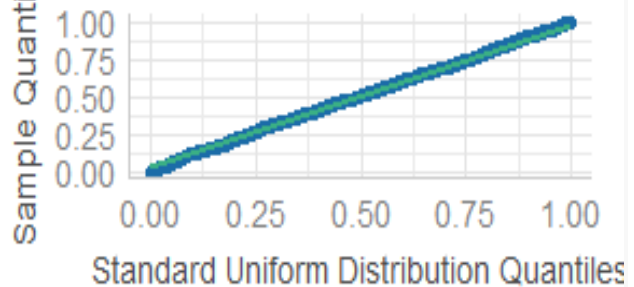
Influential Observations

Points should be inside the contour lines



Uniformity of Residuals

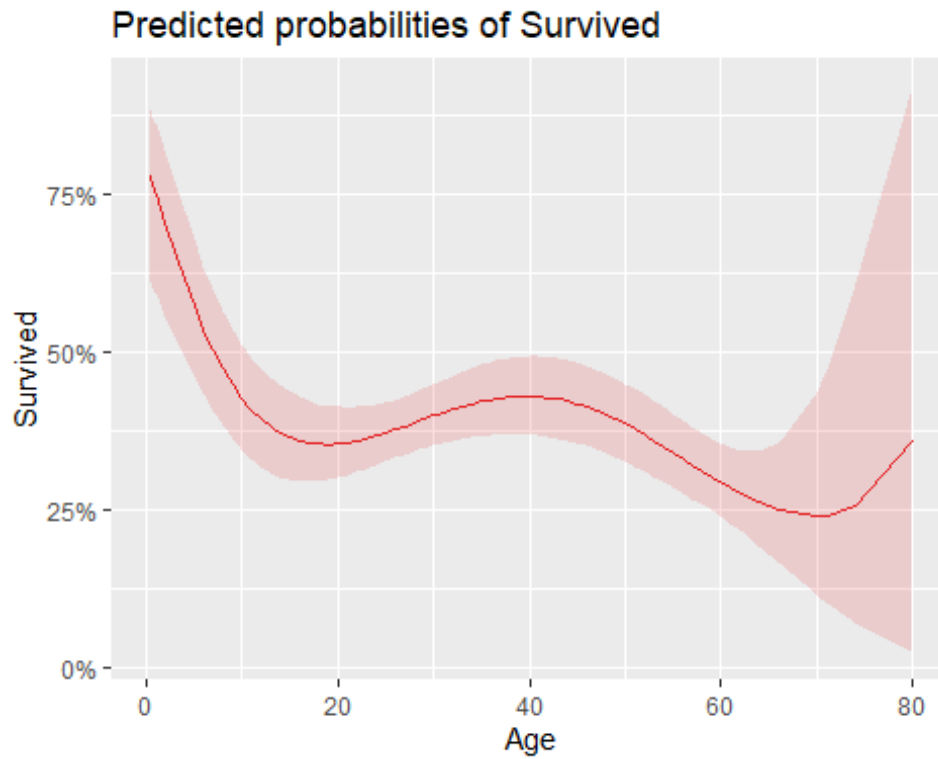
Dots should fall along the line



- The posterior predictive check involves comparing the models predicted intervals with the actual observed values. It helps assess how well the model aligns with the data
- Most residuals fall within the error bands with only one potential outlier
- The influential observational plot tells us that this outlier do not appear to be influential.
- The residuals exhibit a uniform distribution
- The model seems to be okay and we are good to go

Visualize predictions

```
plot_model(modela3, type = "eff", terms = "Age[all]")
```



- The plot clearly shows that babies and young children have the highest survival rate.
- Survival probability then decreases until around age 25 before gradually increasing to a peak at approximately 48 years old.
- After this, it declines again. This pattern indicates two turning points and essentially divides the data into three distinct areas.

Visualize particular predictions

```
library(emmeans)
b <- emmip(modela3, ~Age, CIs = T, type = "response",
            at = list(Age = c(1, 25, 48, 80)))+
  scale_y_continuous()
b
```

