

What makes wine great?

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Task

- ▶ Prediction of quality of (white) wine (from 1, 2,... up to 10) using physicochemical variables.
- ▶ Actually only from 3 to 9 is observed.
- ▶ Data source: Cortez, Paulo, Cerdeira, A., Almeida, F., Matos, T., and Reis, J.. (2009). Wine Quality. UCI Machine Learning Repository. <https://doi.org/10.24432/C56S3T>.
- ▶ Support vector machine is used in their introductory paper.

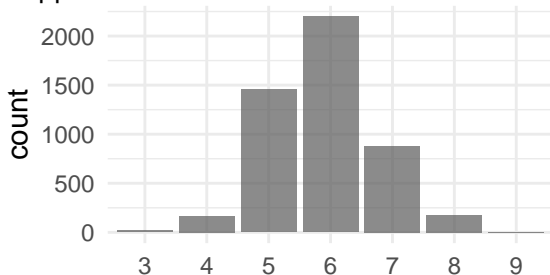


Figure 1: Histogram of quality

Data: Predictive variables

- ▶ Acidity: citric.acid, volatile.acidity
- ▶ Sweetness: residual.sugar
- ▶ Bitterness: sulphates
- ▶ Saltiness: chlorides
- ▶ Prevent oxidation and bacteria: total.sulfur.dioxide
- ▶ Literally interpretable: alcohol

Data: Predictive variables

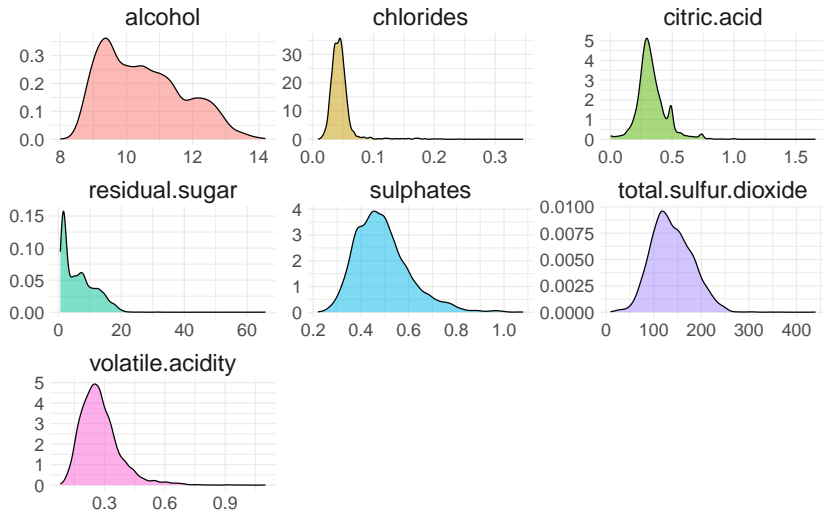


Figure 2: Density of predictive variables

How to model “quality”?

1. Categorical variable. $quality \in \{‘1’, \dots, ‘10’\}$
 - Classification
2. Continuous variable. $quality \in [1, 10]$
 - Linear Regression
3. Ordered Categorical variable. $quality \in \{1, \dots, 10\}$
 - Ordinal Regression

We want to retain ordered structure of data for interpretation.

⇒ Linear Regression (baseline) and Ordinal Regression

For following slides, y for *quality* and x for (vector of) predictive variables.

Regression

As a baseline model.

$$y \sim \text{Normal}(\eta, \gamma)$$

$$\eta = \mathbf{x}^T \boldsymbol{\beta}$$

$$\beta_j \sim \text{Normal}(0, \sigma_{\beta_j})$$

$$\gamma \sim \text{Half-normal}(0, \sigma_\gamma)$$

```
f <- quality ~ citric.acid + volatile.acidity +  
  residual.sugar + sulphates + chlorides +  
  total.sulfur.dioxide + alcohol
```

```
linear_reg <- brm(f,  
  data = d,  
  family = gaussian(),  
  prior = p_linear_reg)
```

Prior Specification

- ▶ Focus on “alcohol”: It takes from 8% to 14% (the range is 6%)
- ▶ The response takes from 3 to 9 (the range is 6)
- ▶ We don't expect the absolute value of coefficient to be larger than 1.
- ▶ Set weakly informative prior accordingly:
 $\beta_{alcohol} \sim \text{Normal}(0, 0.4)$



Figure 3: Distribution of (response / predictive) variables and prior distribution.

Prior Specification (cont)

We have

$$\begin{aligned}\beta_{\text{alcohol}} &\sim \text{Normal}(0, 0.4) \\ &:= \text{Normal}(0, \tau SD(y) / SD(\text{alcohol}))\end{aligned}$$

- ▶ We get scale free informativeness: $\tau \approx 0.5$
- ▶ Set prior for other variables as informative as coefficient for “alcohol”. (i.e., $\beta_j \sim \text{Normal}(0, \tau SD(y) / SD(x_j))$)

Regression: Result

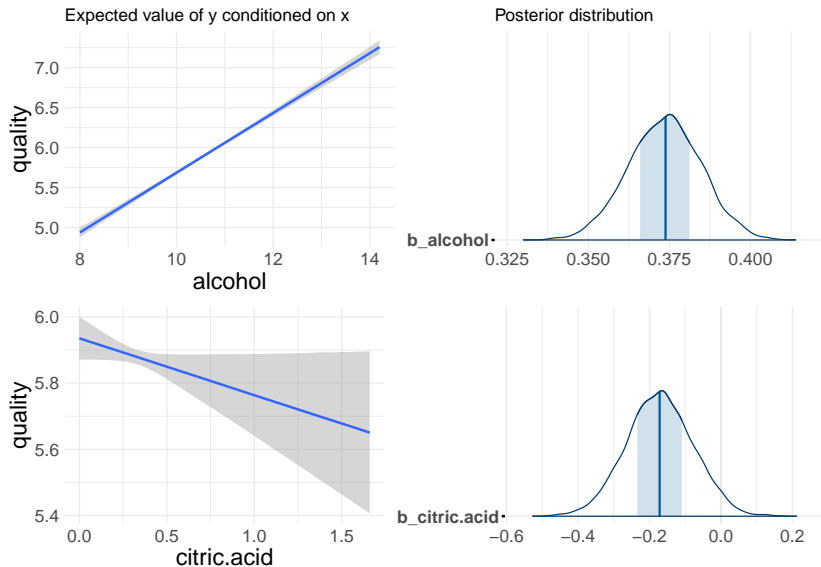


Figure 4: Result for linear regression (only for "alcohol" and "citric.acid")

Ordinal Regression: Cumulative Model

Consider a continuous latent variable \tilde{y} which determine the quality y through thresholds τ .

For $c = 2, \dots, C$:

$$\begin{aligned}Pr(y = c) &= Pr(y \leq c) - Pr(y \leq c - 1) \\&:= Pr(\tilde{y} \leq \tau_c) - Pr(\tilde{y} \leq \tau_{c-1}) \\ \tilde{y} &= \eta + \epsilon, \quad \epsilon \sim \text{Normal}(0, 1)\end{aligned}$$

Prior is set in the same way as regression (here assume $SD(\tilde{y}) = 1$)

```
cumlat <- brm(f,
  data = d,
  family = cumulative("probit"),
  prior = p_cumlat)
```

Cumulative model: Result

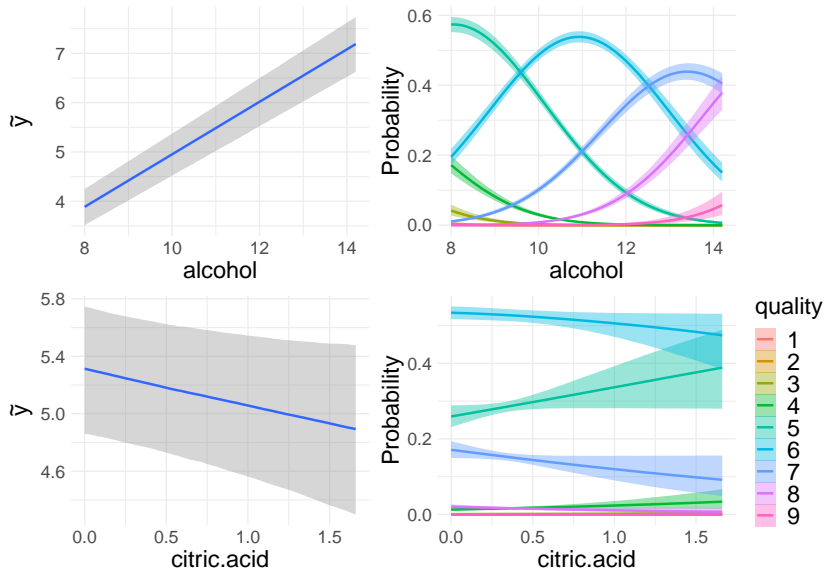


Figure 5: Result for cumulative model (only for "alcohol" and "citric.acid")

Cumulative model: Non-equidistant

Linear regression implicitly assume equidistant among categories (quality).

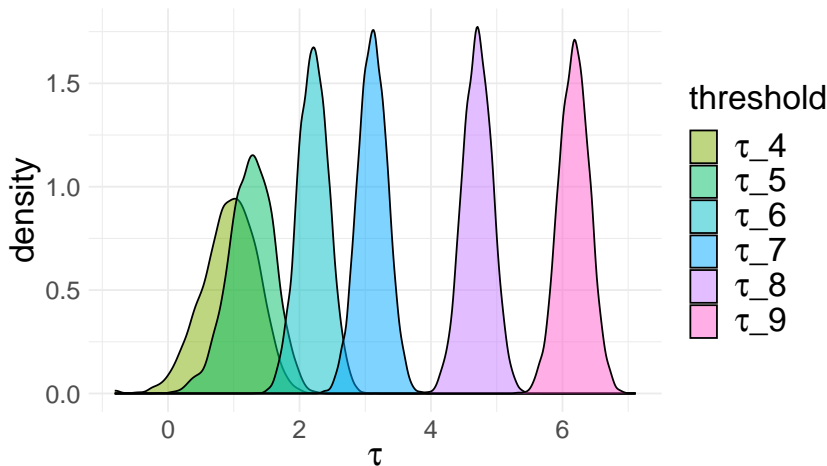


Figure 6: Posterior distribution for the thresholds τ_c

Model Comparison

Leave-one-out Cross Validation

```
loo_compare(linear_reg, cumlat)
```

	elpd_diff	se_diff
cumlat	0.0	0.0
linear_reg	-37.8	10.0

- ▶ Need to be carefully interpreted:
 - ▶ We modelled y differently.
- ▶ Cumulative model has lower ELPD.
 - ▶ We continue further analysis with cumulative model.

Adding non-linearity

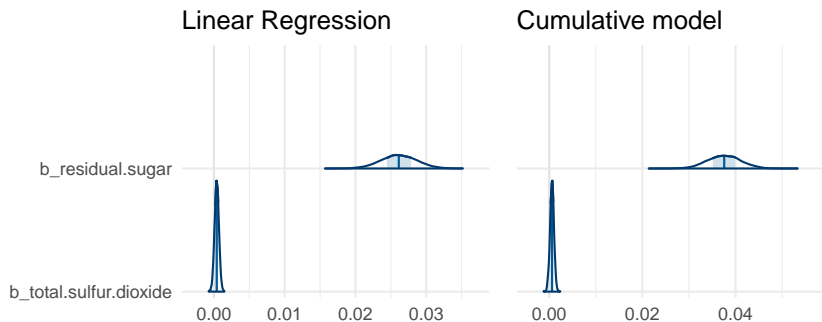


Figure 7: Posterior distributions from the two models.

- ▶ Coefficient for “residual.sugar” and “total.sulfur.dioxide” is concentrated in very small value or around zero.
- ▶ Might be due to non-linearity \Rightarrow use spline for the two variables.
- ▶ Does “optimal” value exist within the range of data we observed?

Spline: Result

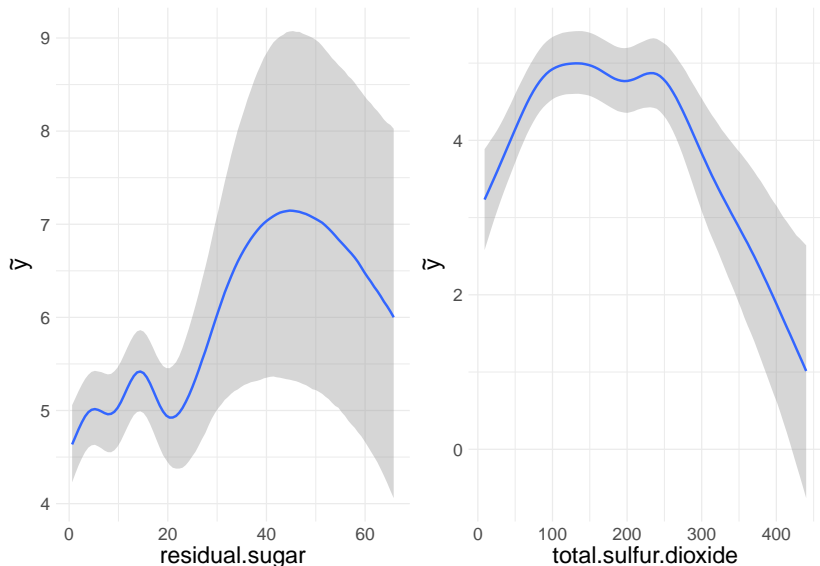


Figure 8: Result for cumulative model with spline (only variables with spline term)

Model Comparison

Leave-one-out CV

```
loo_compare(linear_reg, cumlat, cumlat_s)
```

	elpd_diff	se_diff
cumlat_s	0.0	0.0
cumlat	-91.9	15.4
linear_reg	-129.8	19.0

- Adding non-linearity improves ELPD.

Summary

- ▶ Positive effect of alcohol and negative effect of citric.acid on quality of wine.
- ▶ Ordinal Regression with Spline perform the best in terms of ELPD.
- ▶ Non-equidistant of quality
 - ▶ Lower quality wine tends to be more similar (quality 3, 4, and 5).
- ▶ Non-linear relationship between predictive variables and the quality.
 - ▶ The “optimal” values exist within the data range observed (total.sulfur.dioxide).
- ▶ Further analysis: more non-linearity and synergy effects.

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Appendix

Summary: Regression

```
Family: gaussian
Links: mu = identity; sigma = identity
Formula: quality ~ citric.acid + volatile.acidity + residual.sugar + sulphates + chlorides + total.sulfur
Data: d (Number of observations: 4898)
Draws: 4 chains, each with iter = 4000; warmup = 2000; thin = 1;
       total post-warmup draws = 8000
```

Population-Level Effects:

	Estimate	Est.Error	l-95% CI	u-95% CI	Rhat	Bulk_ESS
Intercept	2.20	0.15	1.90	2.50	1.00	8186
citric.acid	-0.17	0.09	-0.35	0.01	1.00	8908
volatile.acidity	-2.12	0.11	-2.34	-1.90	1.00	8456
residual.sugar	0.03	0.00	0.02	0.03	1.00	10690
sulphates	0.44	0.10	0.25	0.64	1.00	9337
chlorides	-0.87	0.54	-1.94	0.20	1.00	7324
total.sulfur.dioxide	0.00	0.00	-0.00	0.00	1.00	8627
alcohol	0.37	0.01	0.35	0.40	1.00	7269

	Tail_ESS
Intercept	6717
citric.acid	5856
volatile.acidity	5703
residual.sugar	6525
sulphates	5876
chlorides	5997
total.sulfur.dioxide	6665
alcohol	5937

Family Specific Parameters:

	Estimate	Est.Error	l-95% CI	u-95% CI	Rhat	Bulk_ESS	Tail_ESS
sigma	0.76	0.01	0.75	0.78	1.00	9825	5584

Draws were sampled using sampling(NUTS). For each parameter, Bulk_ESS and Tail_ESS are effective sample size measures, and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat = 1).

Summary: Cumulative model

```
Family: cumulative
Links: mu = probit; disc = identity
Formula: quality ~ citric.acid + volatile.acidity + residual.sugar + sulphates + chlorides + total.sulfur
Data: d (Number of observations: 4898)
Draws: 4 chains, each with iter = 4000; warmup = 2000; thin = 1;
       total post-warmup draws = 8000
```

Population-Level Effects:

	Estimate	Est.Error	l-95% CI	u-95% CI	Rhat	Bulk_ESS
Intercept[1]	0.94	0.42	0.08	1.67	1.00	3026
Intercept[2]	1.26	0.34	0.54	1.89	1.00	4158
Intercept[3]	2.21	0.23	1.75	2.66	1.00	5834
Intercept[4]	3.11	0.22	2.66	3.54	1.00	5747
Intercept[5]	4.70	0.22	4.26	5.14	1.00	5618
Intercept[6]	6.18	0.23	5.72	6.62	1.00	5395
Intercept[7]	7.34	0.24	6.88	7.79	1.00	5348
Intercept[8]	8.77	0.27	8.23	9.30	1.00	5629
citric.acid	-0.25	0.13	-0.51	0.01	1.00	5557
volatile.acidity	-3.11	0.16	-3.42	-2.79	1.00	5336
residual.sugar	0.04	0.00	0.03	0.04	1.00	6909
sulphates	0.63	0.14	0.37	0.90	1.00	5755
chlorides	-1.26	0.78	-2.79	0.29	1.00	6024
total.sulfur.dioxide	0.00	0.00	-0.00	0.00	1.00	8104
alcohol	0.53	0.02	0.50	0.57	1.00	4876

Tail_ESS

Intercept[1]	2646
Intercept[2]	3398
Intercept[3]	5648
Intercept[4]	5589
Intercept[5]	5566
Intercept[6]	5627
Intercept[7]	5466
Intercept[8]	5625
citric.acid	5392

Summary: Cumulative with Spline

Family: cumulative

Links: mu = probit; disc = identity

Formula: quality ~ s(residual.sugar) + s(total.sulfur.dioxide) + citric.acid + volatile.acidity + sulphates

Data: d (Number of observations: 4898)

Draws: 4 chains, each with iter = 4000; warmup = 2000; thin = 1;

total post-warmup draws = 8000

Smooth Terms:

	Estimate	Est.Error	l-95% CI	u-95% CI	Rhat	Bulk_ESS
sds(sresidual.sugar_1)	9.23	2.80	4.97	15.72	1.00	2470
sds(stotal.sulfur.dioxide_1)	3.47	1.18	1.78	6.37	1.00	3041
	Tail_ESS					
sds(sresidual.sugar_1)	4390					
sds(stotal.sulfur.dioxide_1)	4189					

Population-Level Effects:

	Estimate	Est.Error	l-95% CI	u-95% CI	Rhat	Bulk_ESS
Intercept[1]	0.34	0.45	-0.66	1.12	1.00	4096
Intercept[2]	0.68	0.36	-0.09	1.32	1.00	5887
Intercept[3]	1.73	0.22	1.31	2.16	1.00	6835
Intercept[4]	2.69	0.20	2.30	3.09	1.00	7458
Intercept[5]	4.33	0.20	3.93	4.74	1.00	7462
Intercept[6]	5.83	0.21	5.43	6.25	1.00	7700
Intercept[7]	7.02	0.22	6.60	7.45	1.00	7773
Intercept[8]	8.46	0.26	7.97	8.98	1.00	7144
citric.acid	-0.18	0.14	-0.44	0.08	1.00	9277
volatile.acidity	-3.04	0.16	-3.35	-2.72	1.00	8595
sulphates	0.60	0.14	0.33	0.87	1.00	8753
chlorides	-1.58	0.78	-3.12	-0.06	1.00	9246
alcohol	0.53	0.02	0.49	0.56	1.00	7822
sresidual.sugar_1	2.40	2.57	-2.60	7.39	1.00	5352
stotal.sulfur.dioxide_1	-0.06	2.36	-4.81	4.54	1.00	6721
	Tail_ESS					
Intercept[1]	1960					

Prior Summary: Regression

prior	class	coef	group	resp	dpar	nlpar	lb	ub
(flat)	b							
normal(0,0.36)	b	alcohol						
normal(0,20.268)	b	chlorides						
normal(0,3.659)	b	citric.acid						
normal(0,0.087)	b	residual.sugar						
normal(0,3.88)	b	sulphates						
normal(0,0.01)	b	total.sulfur.dioxide						
normal(0,4.393)	b	volatile.acidity						
normal(6, 5)	Intercept							
normal(0, 5)	sigma						0	
source								
default								
user								
user								
user								
user								
user								
user								
user								
user								
user								

Prior Summary: Cumulative

	prior	class	coef	group	resp	dpar	nlpar	lb
	(flat)	b						
	normal(0,0.406)	b	alcohol					
	normal(0,22.885)	b	chlorides					
	normal(0, 4.132)	b	citric.acid					
	normal(0,0.099)	b	residual.sugar					
	normal(0,3.88)	b	sulphates					
	normal(0,0.012)	b	total.sulfur.dioxide					
	normal(0,4.961)	b	volatile.acidity					
student_t(3, 0, 2.5)	Intercept							
normal(-2, 1)	Intercept		1					
normal(-1.43, 1)	Intercept		2					
normal(-0.86, 1)	Intercept		3					
normal(-0.29, 1)	Intercept		4					
normal(0.29, 1)	Intercept		5					
normal(0.86, 1)	Intercept		6					
normal(1.43, 1)	Intercept		7					
normal(2, 1)	Intercept		8					
ub	source							
	default							
	user							
	user							
	user							
	user							
	user							
	user							
	default							
	user							
	user							
	user							
	user							
	user							
	user							

Prior Summary: Cumulative with Spline

prior	class	coef	group	resp	dpar	nlpar
(flat)	b					
normal(0,0.406)	b	alcohol				
normal(0,22.885)	b	chlorides				
normal(0, 4.132)	b	citric.acid				
normal(0, 3)	b	sresidual.sugar_1				
normal(0, 3)	b	stotal.sulfur.dioxide_1				
normal(0,3.88)	b	sulphates				
normal(0,4.961)	b	volatile.acidity				
student_t(3, 0, 2.5)	Intercept					
normal(-2, 1)	Intercept		1			
normal(-1.43, 1)	Intercept		2			
normal(-0.86, 1)	Intercept		3			
normal(-0.29, 1)	Intercept		4			
normal(0.29, 1)	Intercept		5			
normal(0.86, 1)	Intercept		6			
normal(1.43, 1)	Intercept		7			
normal(2, 1)	Intercept		8			
student_t(3, 0, 2.5)	sds					
student_t(3, 0, 2.5)	sds	s(residual.sugar)				
student_t(3, 0, 2.5)	sds	s(total.sulfur.dioxide)				
lb	ub					
	source					
	default					
	user					
	user					
	user					
	user					
	user					
	user					
	user					
	default					
	user					
	user					
	user					

Cumulative Model (cont)

Other expression:

$$\begin{aligned}Pr(\tilde{y} \leq \tau_c) &= Pr(\eta + \epsilon \leq \tau_c) \\&= Pr(\epsilon \leq \tau_c - \eta) \\&= \Phi(\tau_c - \eta) \quad \Phi : \text{cdf of standard normal aka probit}\end{aligned}$$

Then we have:

$$\psi_c = \Phi(\tau_c - \eta) - \Phi(\tau_{c-1} - \eta)$$

Adding non-linearity with Spline

```
f_s <- quality ~ citric.acid + volatile.acidity +  
  sulphates + chlorides + alcohol +  
  s(residual.sugar) + s(total.sulfur.dioxide)  
  
cumlat_s <- brm(f_s,  
  data = d,  
  family = cumulative("probit"),  
  prior = p_cumlat_s)
```

- ▶ We are particularly interested in non-linearity of these two variables.
- ▶ Other variable could be non-linear.