

Deviance Residuals:

Min	1Q	Median	3Q	Max
-2.0260	-0.5795	-0.4078	-0.2680	2.8794

Coefficients:

	Estimate	Std. Error	z value	Pr(> z )	
(Intercept)	-6.265e+00	6.454e-01	-9.707	< 2e-16	***
CMU.Hrs	5.280e-05	1.469e-05	3.593	0.000327	***
Fe	1.944e-01	4.025e-02	4.830	1.36e-06	***
Si	4.491e-01	1.249e-01	3.595	0.000324	***
Na	1.428e-01	3.996e-02	3.574	0.000351	***
ST	2.764e-02	1.266e-02	2.184	0.028994	*
PFc	1.541e+00	5.713e-01	2.697	0.007007	**
'Fe:Si'	-2.358e-02	8.633e-03	-2.731	0.006307	**

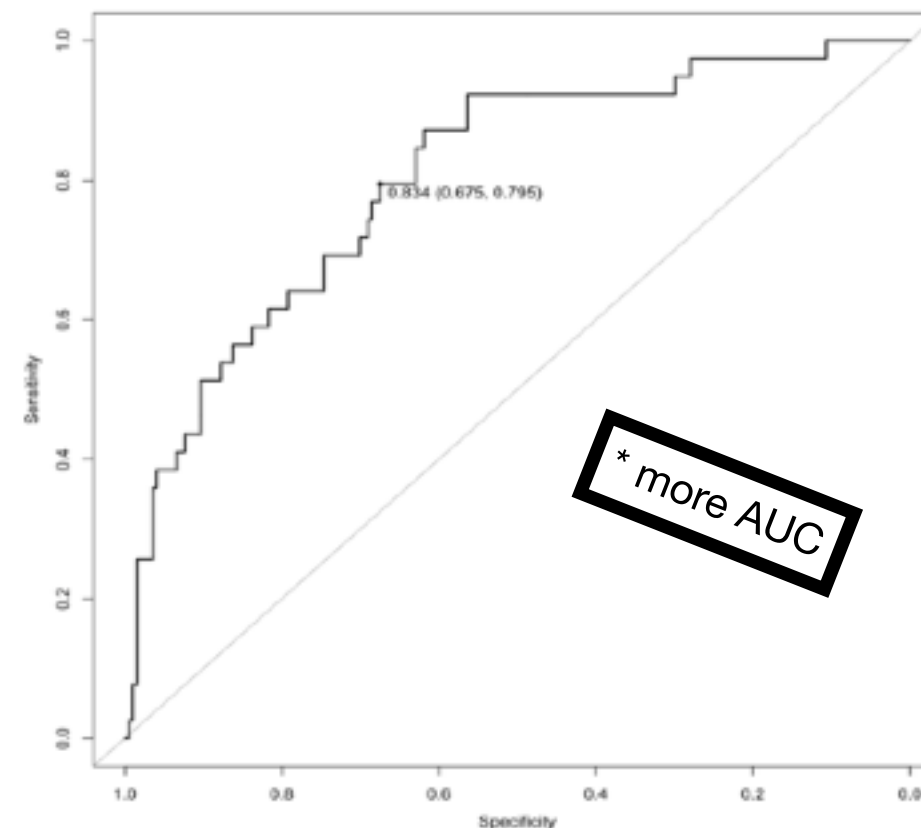
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 854.26 on 947 degrees of freedom  
Residual deviance: 703.34 on 940 degrees of freedom  
AIC: 719.34

Number of Fisher Scoring iterations: 6

\* I know significance isn't everything but hey, it's at least a little reassuring



## ... model's final form

Confusion Matrix and Statistics

	Reference	
Prediction	A	Not A
A	193	29
Not A	4	10

Accuracy : 0.8602  
95% CI : (0.8093, 0.9018)  
No Information Rate : 0.8347  
P-Value [Acc > NIR] : 0.1678

Kappa : 0.3178

McNemar's Test P-Value : 2.943e-05

Sensitivity : 0.9797  
Specificity : 0.2564  
Pos Pred Value : 0.8694  
Neg Pred Value : 0.7143  
Prevalence : 0.8347  
Detection Rate : 0.8178  
Detection Prevalence : 0.9407  
Balanced Accuracy : 0.6181

unhealthy >= 50%

Confusion Matrix and Statistics

	Reference	
Prediction	A	Not A
A	175	19
Not A	22	20

Accuracy : 0.8263  
95% CI : (0.7718, 0.8724)  
No Information Rate : 0.8347  
P-Value [Acc > NIR] : 0.6750

Kappa : 0.3891

McNemar's Test P-Value : 0.7548

Sensitivity : 0.8883  
Specificity : 0.5128  
Pos Pred Value : 0.9021  
Neg Pred Value : 0.4762  
Prevalence : 0.8347  
Detection Rate : 0.7415  
Detection Prevalence : 0.8220  
Balanced Accuracy : 0.7006

unhealthy >= 30%

Confusion Matrix and Statistics

	Reference	
Prediction	A	Not A
A	98	3
Not A	99	36

Accuracy : 0.5678  
95% CI : (0.502, 0.6319)  
No Information Rate : 0.8347  
P-Value [Acc > NIR] : 1

Kappa : 0.2116

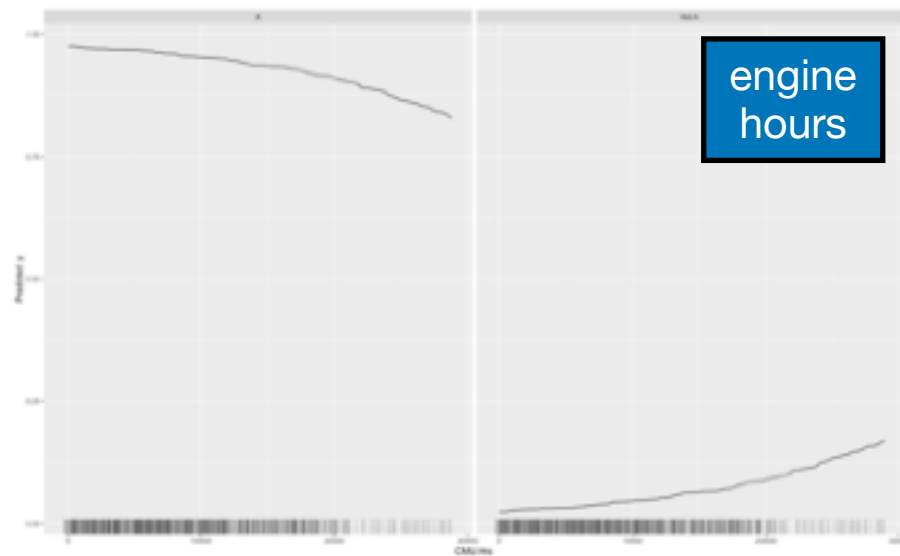
McNemar's Test P-Value : <2e-16

Sensitivity : 0.4975  
Specificity : 0.9231  
Pos Pred Value : 0.9703  
Neg Pred Value : 0.2667  
Prevalence : 0.8347  
Detection Rate : 0.4153  
Detection Prevalence : 0.4280  
Balanced Accuracy : 0.7103

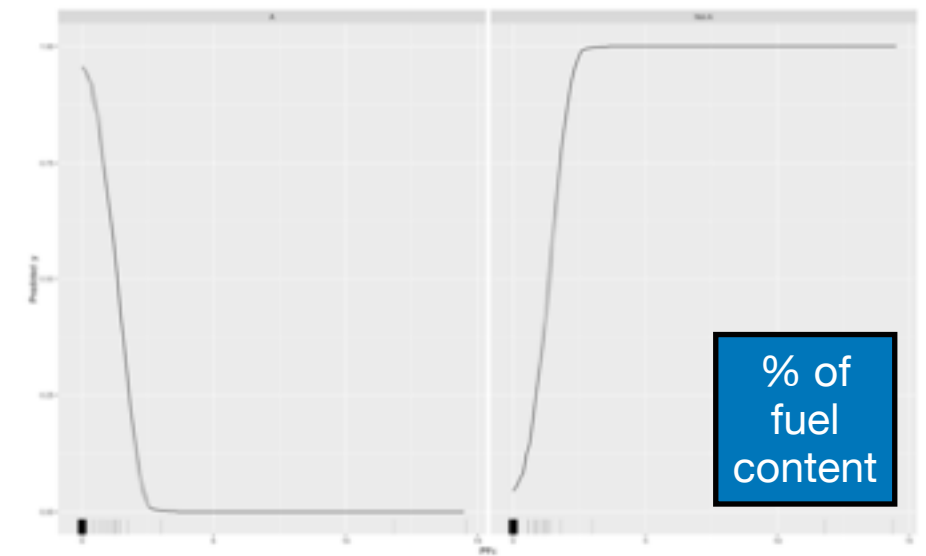
unhealthy >= 15%

\* better performance in general, especially when adjusting the probability cut-off

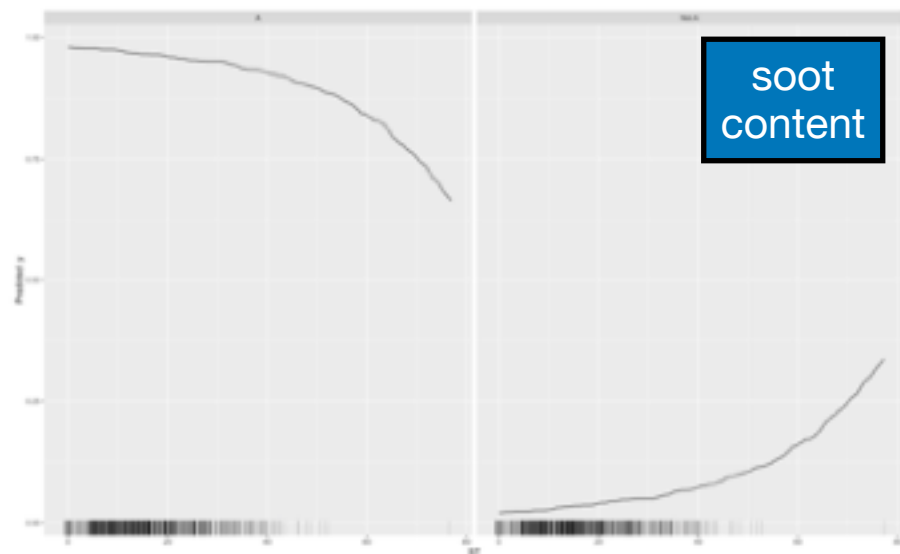
# ... simpler interpretation



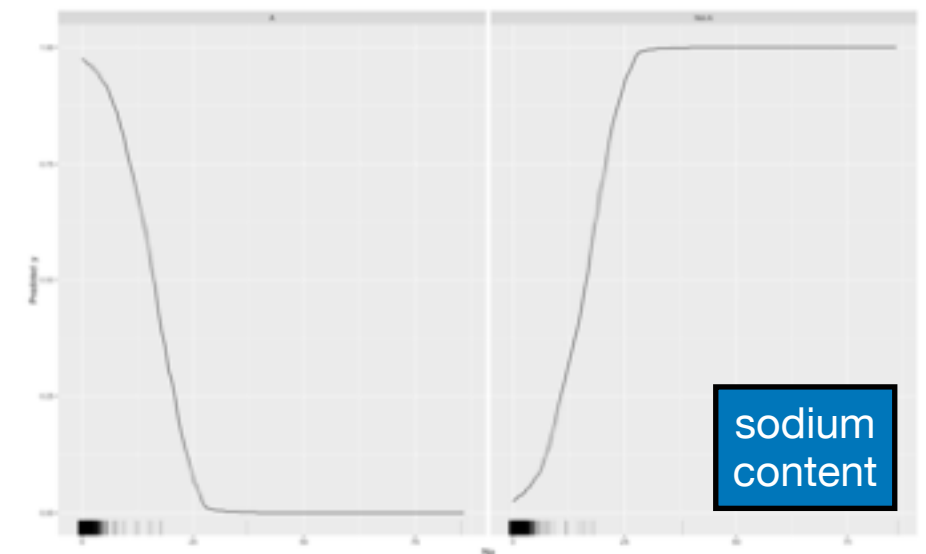
Probability of a sample being **“healthy”** is on the **left** of each plot pair!



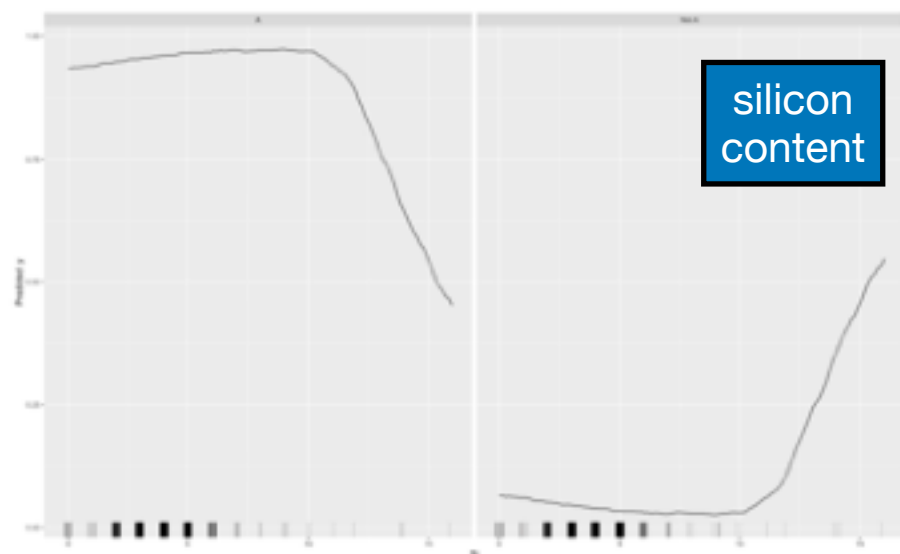
% of fuel content



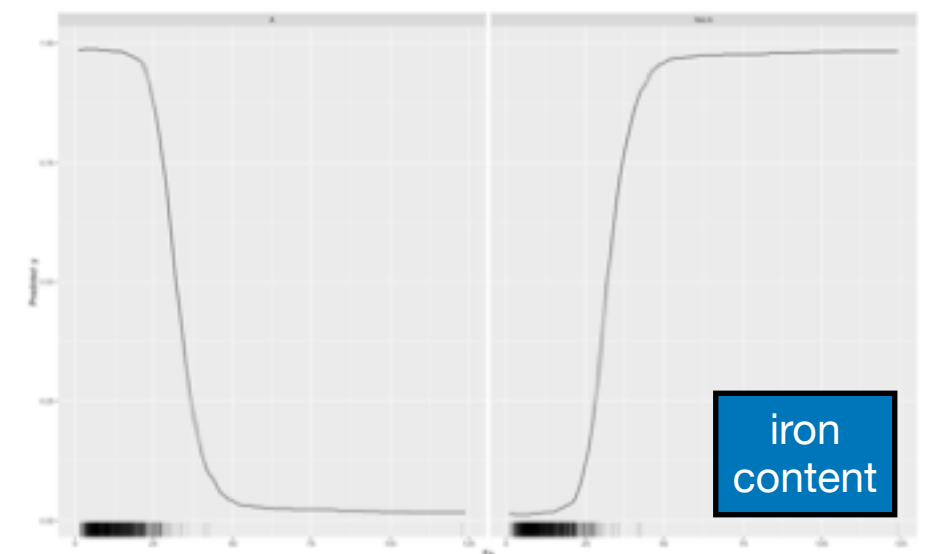
Probability of a sample being **“unhealthy”** is on the **right** of each plot pair!



sodium content



As you move from **left to right** the variable in question is increasing along the X-axis!



iron content