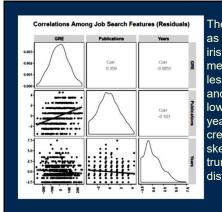
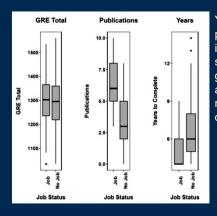
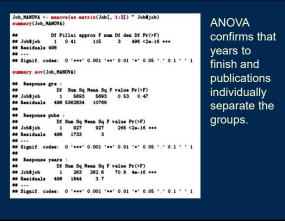
Discriminant Analysis	
Today • Another example • Classification details • Alternatives when assumptions fail	
In this hypothetical example, data from 500 graduate students seeking jobs were examined. Available for each student were three predictors: GRE(V+Q), Years to Finish the Degree, and Number of Publications. The outcome measure was categorical: "Got a job" versus "Did not get a job."	



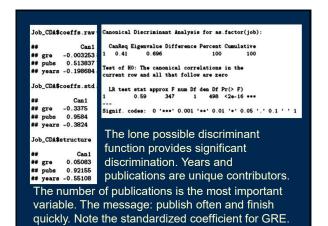
The data are not as clean as the iris data. The measures are less continuous and years has a lower bound (4 years) that creates a skewed and truncated distribution.



Years and publications individually separate the groups. GRE appears to make no difference.



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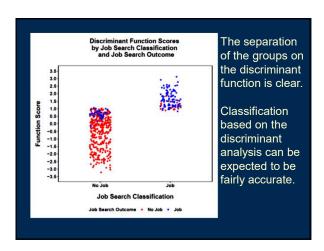
Job_CDA\$coeffs.raw

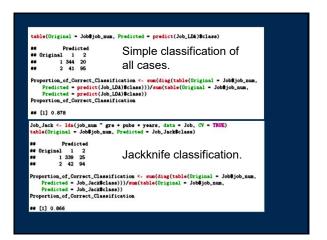
"" Cant
"" gre -0.003253
"" pubs 0.513837
"" years -0.198684

Job_CDA\$coeffs.std

"" Cant
"" we have no information about their
standard errors. Those are important
if we want to make claims about
coefficients being different from 0.

Later we will consider bootstrapping
as a means to get that information.





```
training_sample <- sample(1:500, 250)

Job_Train <- lad_job_num * gre * pube * years, data * Job_ CV * FALSE, subset* training_sample)

Job_Predict <- predict(Job_Train, newdata * Job_Ctraining_sample, ])

Job_Cross <- chief(Job_Original, Job_PredictsClass)

names(Job_Cross) <- c(*Original*, * Predictsd*)

table(Original * Job_CrossSOriginal, * Predictsd*)

table(Original * Job_CrossSOriginal, * Predictsd*)

se Predictsd

se Original * Original * Job_CrossSOriginal, * Predictsd*)

se Original * Job_CrossSOriginal, * Predictsd*)

Proportion_of_Correct_Classification <- sum(diag(table(Original * Job_CrossSOriginal, Predicted * Job_C
```

Under the assumption of multivariate normality, the probability of a particular profile of scores (X) given a particular group centroid and pooled covariance matrix can be estimated:

$$p(X|Group_g) \propto \frac{1}{\sqrt{|C_w|(2\pi)^{\frac{k}{2}}}} e^{-\frac{1}{2}(X-\overline{X_g})'(C_w^{-1})(X-\overline{X_g})}$$

The pooled within-group covariance matrix is used if the homogeneity assumption holds.

The posterior probabilities take prior probability of group membership into account:

$$p(Group_g|X) = \frac{p(Group_g)p(X|Group_g)}{\sum_{i=1}^{G} p(Group_i)p(X|Group_i)}$$

In the iris data, the prior probabilities were equal (.33) but for the current data, the prior probability of getting a job is smaller (.272) than the prior probability of not getting a job (.728; from the sample proportions). We assume the sample is representative.

The case is assigned to the group with the highest posterior probability. The posterior probabilities are particularly useful because they indicate our confidence in classification.

	LD1	1	2	
1	-0.2686	0.9077	0.092347	1
2	-2.0386	0.9963	0.003695	1
3			0.502488	
4	0.4677	0.7125	0.287474	1
5	0.1521	0.8173	0.182719	1
6	-0.9081	0.9702	0.029837	1

Is the classification better than would be expected by chance? We first need to know the correct classification that would occur by chance:

		Ехрє		
		No Job	Job	All
Actual	No Job	(q ₁ N) ² N		q₁N
ual	Job		(q ₂ N) ² N	q ₂ N
	All	q₁N	q ₂ N	N

Note that we assume the expected marginals to be the same as the actual marginals. Is the classification better than would be expected by chance? We first need to know the correct classification that would occur by chance:

		Ехрє		
		No Job	Job	All
Actual	No Job	(q ₁ N) ² N		364
ual	Job		(q ₂ N) ² N	136
	All	364	136	500

Note that we assume the expected marginals to be the same as the actual marginals.

Is the classification better than would be expected by chance? We first need to know the correct classification that would occur by chance:

		Ехре		
		No Job	Job	All
Actual	No Job	264.99 ^	99.01	364
tual	Job	99.01	36.99	136
	All	364	136	500
			(364	x 364)/500

The total number of correct classifications that would occur by chance (302, 60.4%) can be tested against the actual number of correct classifications given the discriminant analysis model (439, 87.8%, for the simple prediction of all cases).

A *t*-test can be calculated:

$$t = \frac{439 - 302}{\sqrt{500(.604)(1 - .604)}} = 12.53$$

where the denominator is the standard error of the number of correct classifications by chance (the null hypothesis). The difference between chance expected and actual classification can be tested with a chisquare as well.

$$\chi^2 = \sum_{i=1}^{C} \frac{(f_{0i} - f_{e_i})^2}{f_{e_i}}$$

$$\chi^2 = \frac{(439 - 302)^2}{302} + \frac{(61 - 198)^2}{198} = 156.94$$

Because this is a single degree of freedom test, $t^2 = \gamma^2$

Klecka's $tau(\tau)$ is sometimes calculated for classification results:

$$\tau = \frac{n_0 - \sum_{i=1}^G p_i n_i}{N - \sum_{i=1}^G p_i n_i}$$

 n_{o} is the number of correct classifications, n_{i} is the number of cases in group i, p_{i} is the proportion of the total sample expected to be in group i, G is the number of groups, and N is the total sample size.

$$\tau = \frac{439 - 302}{500 - 302} = .69$$

This is interpreted as the proportional improvement in classification over random assignment to groups. Values can range from 0 to 1. This is very similar to Cohen's kappa.

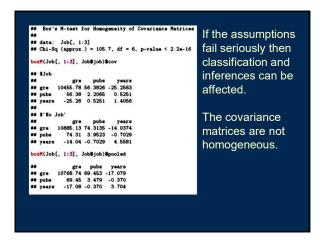
Is an obtained τ value different from 0? The sampling distribution of τ is unknown and so a convenient formula for the standard error is not available. The bootstrap procedure will help here as well.

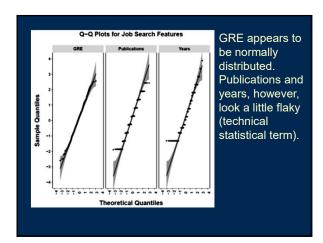
- Several additional classification indices can be useful. In the following, an "event" is "getting a job":
- Precision: "What percentage of predicted events are correct?"
- Recall: "What percentage of events were correctly predicted?"
- F1: Harmonic mean of precision and recall.
- Prevalence: "What is the proportion of actual events in the sample?"
- Detection Rate: "What proportion of the entire sample are correctly predicted events?"

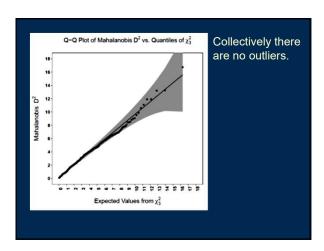
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$			Ac			
Prediction Present b a Row 2 = b+ Marginal Column 1 = d+b Column 2 = c+a N=a+b+c+c $Recall = \frac{a}{a+c} Precision = \frac{a}{a+b}$ $F1 = \frac{2*Precision*Recall}{Precision+Recall}$			Absent	Present		
Present b a Row 2 = b+. Marginal Column 1 = d+b Column 2 = c+a N=a+b+c+c $Recall = \frac{a}{a+c} Precision = \frac{a}{a+b}$ $F1 = \frac{2 * Precision * Recall}{Precision + Recall}$	Prediction			c		
$Recall = \frac{a}{a+c} \qquad Precision = \frac{a}{a+b}$ $F1 = \frac{2 * Precision * Recall}{Precision + Recall}$	Trediction					
$F1 = \frac{2 * Precision * Recall}{Precision + Recall}$		Marginal	Column 1 = d+b	Column 2 = c+a	N=a+b+c+d	
$Prevalence = \frac{a+c}{a+b+c+d} \qquad Detection \ Rate = \frac{a}{a+b+c+d}$		Precision + Recall				
		11	Precision + R	ecall		
	Preval		Precision + R		$\frac{a}{+b+c+d}$	

	Referen	ice		
Predicti	on No Job	Job		
No J	ob 339	42		
Job	25	94		
	Ac	curacy		
		95% CI	;	(0.833, 0.895
No I	nformatio	n Rate	:	0.728
P-Va	lue [Acc	> NIR]	:	8.02e-14
		Kappa	:	0.648
Mcnemar	's Test F	-Value	:	0.0506
	Sensi	tivity	:	0.691
	Speci	ficity	:	0.931
	Pos Pred	Value	:	0.790
	Neg Pred	Value	:	0.890
	Pre	cision	:	0.790
	1600	Recall	:	0.691
		F1	:	0.737
	Prev	alence	:	0.272
	Detection	n Rate	:	0.188
Detec	tion Prev	alence	:	0.238
	lanced Ac	curacu		0.911

The discriminant analysis produced predictions that were correct quite often (.79) and in particular correctly identified getting a job fairly well (.69), despite getting a job being a relatively rare event (.27).







The significance tests for normality are surely being influenced by the very large sample size.

If we have doubts about the legitimacy of significance tests or classification based on normality, we can use methods that don't rely on that assumption.

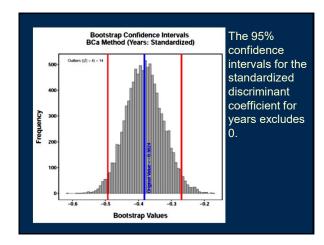
When the homogeneity of covariance matrices assumption fails, a pooled within-groups covariance matrix is not appropriate. An alternative form of analysis known as quadratic discriminant analysis can be used. This uses the separate group variance- covariance matrices in the classification process.

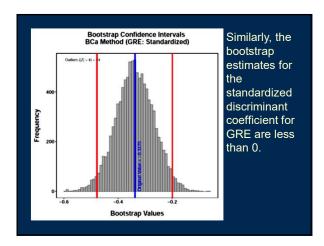
The jackknife classification is nearly identical to the original analysis.

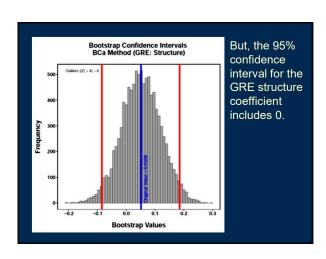
If the assumptions underlying the discriminant analysis (homogeneous covariance matrices, multivariate normality) are not viable, the bootstrapping approach can be taken.

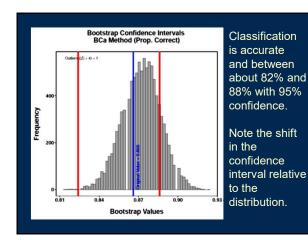
In bootstrapping, we assume that whatever population the sample came from, the sample is representative of that population. Therefore we can sample randomly from the sample, with replacement, to get multiple samples of the same size on which we can repeat the analyses.

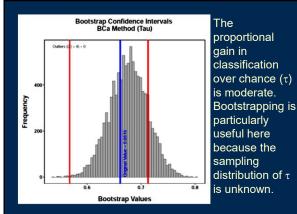
The resulting <u>empirical</u> sampling distributions can be examined to assist inferences.











particularly useful here because the sampling distribution of τ is unknown.

Bootstrapping is helpful if multivariate normality is out of the question, homogeneity of covariance matrices is not tenable, or sampling distributions are

But, other "problems" require a different approach. What if the groups are ordered in some way that we would like to incorporate into the statistical model? What if we have particular interest in comparisons among some of the groups? What if we would like to include interactions or polynomial terms with the predictors?

Next time	
Logistic regression	