

Practical handout for the workshop “Introduction to the analysis of large-scale data on social connections”

Prepared for the Historical Demography Scientific Research Network
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Paul Lambert and Dave Griffiths, University of Stirling, April 2012

Version 1



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Introduction

This handout accompanies the lab sessions for the workshop 'Introduction to the analysis of large-scale data on social connections' (26 April 2012, Utrecht).

For Stata and R, the step-by-step implementation instructions for each session are largely to be found within the specific 'syntax' files for the relevant sessions (.do and .R files). For Pajek, step-by-step instructions with screenshots are provided below.

Parts of this handout are extracted from a more extended handout on using data analysis packages for social science research, produced by Lambert for the DAMES Node workshop programme (see www.dames.org.uk) and for his course 'Introduction to multilevel models with applications' to the Essex Summer School in Social Science data analysis (www.staff.stir.ac.uk/paul.lambert/essex_summer_school).

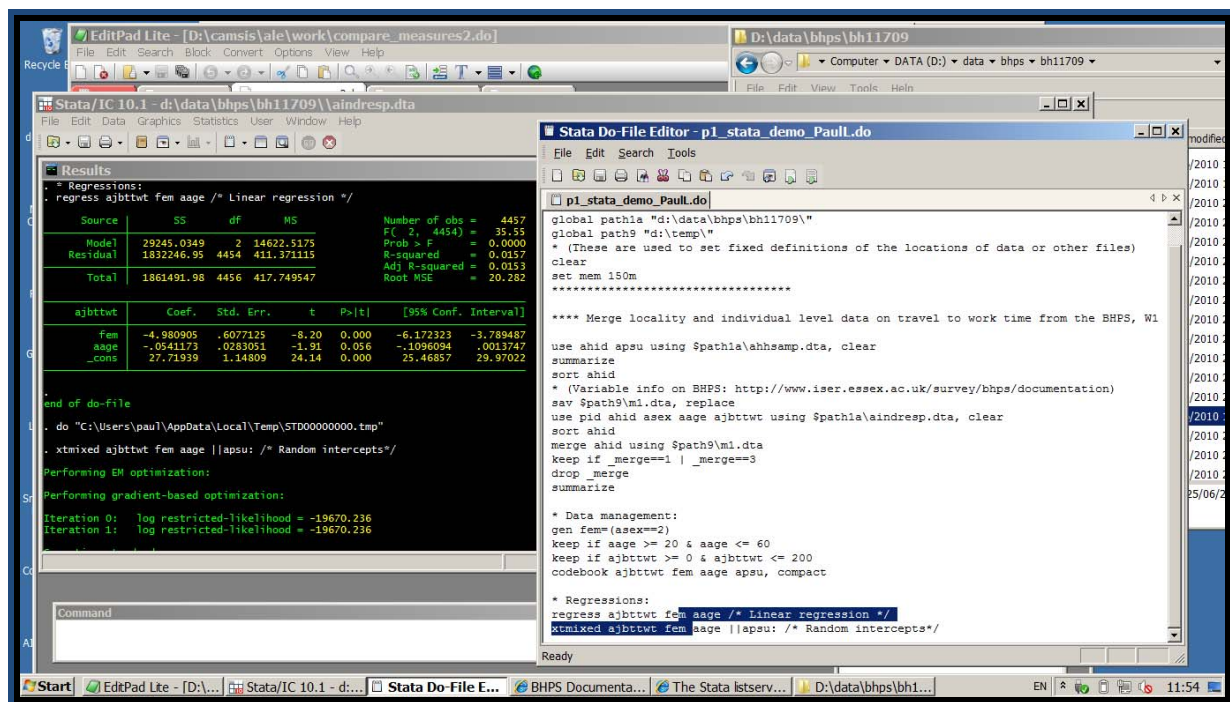
General arrangements for the practicals

Unless noted otherwise the data files used are for distribution for this lab session only and should not be transferred elsewhere. The sources of these files are ultimately available online from international data providers such as NAPP (<http://www.nappdata.org/napp/>) or the UK's ESDS (www.esds.ac.uk).

You will need to have the relevant packages installed to undertake the relevant exercises (though it should be possible to use only some of the packages as relevant). Introductory notes on the packages are included below under 'lab 1'. Some of the packages and lab exercises have online dependencies (e.g. they need to use a data file or programme extension which is available online). We have tried to note the details below, but information may not be comprehensive.

In general terms, the task in the labs is to open up the relevant syntax files, and work your way through them, digesting the examples shown (and potentially adding your own notes, adjustments or examples). You'll ordinarily need to have the analytical software open and the relevant tool for working with a syntax file (e.g. 'syntax window' or 'do file editor'). In addition it will typically also be helpful to have open some applications to remind you of where the data is stored, and perhaps a plain text editor allowing you to conveniently open up several other syntax files for purposes of comparison.

- When working with Stata a typical view of your desktop might be something like:



Description: The first two interfaces you can see in this screenshot are respectively the Stata do file editor (where I write commands and send them to be processed, such as by highlighting the relevant lines and clicking 'ctrl-d'); and the main Stata window (here Version 10) which includes the results page. Note that the syntax file open is a modified (personalised) version of the supplied illustrative syntax file – the name has been changed so that I can save the original file plus my own version of it after edits (e.g. with my own comments). Behind the scenes I've also got open an 'Editpadlite' session which I'm using to conveniently open up and compare some other syntax files that I don't particularly want in my do file editor itself; I've also got a file manager software open showing the data I'm drawing upon (in what Stata will call 'path1a'); and I've got some Internet Explorer (IE) sessions open (I'm looking up the online BHPS documentation, and the Stata listserv, where information on Stata commands is available).

Materials referred to in the sessions will include:

- data files (copies of survey and other data used);
- sample command files (pre-prepared materials which include programming commands in the language of the relevant software)
- supplementary 'macros' or 'sub-files' (further pre-prepared materials featuring programming commands in relevant languages, usually invoked as a sub-routine within the main sample command files)

An important point to make is that some of the command files will need to draw upon other files (e.g. data files) in order to run. To do this, they need to be able to reference the location of the required files. In most applications, we do this by defining 'macros' which point to specific 'paths' on your computer (see also software sections below). For the labs to work successfully, it will be necessary to ensure that the command file you are trying to run is pointing to the right paths at the right time. In general, this only requires one specification to be made at the start of the session, for instance whereby in Stata we define 'macros' for the relevant paths. Sometimes however it can be necessary to edit the full path reference of a particular file in order to be able to access it.

For example, in the text below, we show some Stata (and SPSS) commands which in both cases define a macro (called 'path3a') which gives the directory location of the data file 'aindresp.dta' or 'aindresp.sav', so that subsequent commands calling it will go directly to that path:

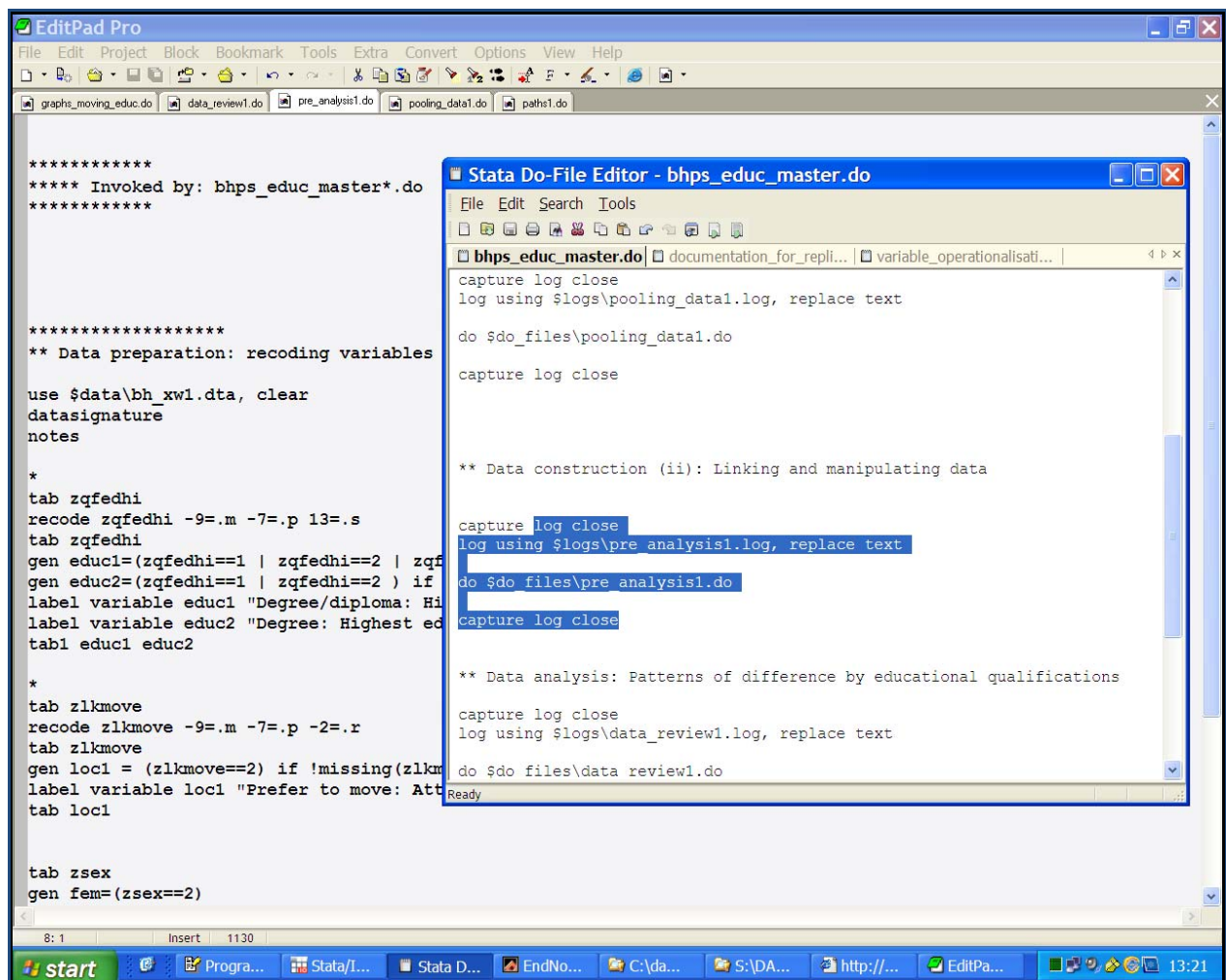
<i>Stata example</i>	<i>comparable SPSS example</i>
global path3a "d:\data\bhps\"	define !path3a () "d:\data\bhps\" !enddefine.
use pid asex using \$path3a\aindresp.dta, clear	get file=!path3a+" aindresp.sav" .
tab asex	fre var=asex.

Relevant background: Thinking about workflows

There are very good expositions of the idea of workflows in the social science data analysis process in, amongst others, Long (2009); Treiman (2009), and Kohler and Kreuter (2008). A workflow in itself is a representation of a series of tasks which contribute to a project or activity. It can be a useful exercise to conceptualise a research project as a workflow (with components such as data collection, data processing, data analysis, report writing). However, when dealing with large scale data, a really useful contribution is to organise your data and command files that are associated with a project in a consistent style that recognises that relevant contributions to the workflow structure.

What does that involve? The issue is that we want to construct a replicable trail of our data oriented research, which allows us to go all the way from opening the initial data file, to producing the publication quality graph or statistical results which are our end products. We need the replicable trail in order to adjust our analysis on the basis of minor changes at any possible stage of the process (or to be able to transfer a record of our work on to others). However because when dealing with large-scale and complex data (e.g. on social connections) the trail is long and complex (and not entirely linear), we can only do this, realistically, if we break down our activities into multiple separate components.

There are different ways to organise files for these purposes, but a popular and highly effective approach is to design a 'master' syntax command file and a series of 'sub-files' which it draws upon. In this model, the sub-files cover different parts of the research analysis. Personally, my preference is to construct both the master and sub-files in the principle software package being used, though Long (2009) notes that creating a documentation master file in a different software (e.g. MS Excel) is an effective way to record a wider range of activities which span across different software. Here's an example of a series of tasks being called upon via a Stata format 'master' file:



(This screenshot shows the Stata master file, and the sub-files which are mostly open within the EditPad editor - except for a few other files which I've opened in the do file editor. The Stata output file is not visible but is open behind the scenes).

Here's an example of a project documentation file that might be constructed in Excel:

	A	B	C	D	E	F
1						
2			File names	Location	Description	
3		Stata work:				
4	1	Command files:	bhps_educ_master.do	c:\dames\workshops\2010\4\applic_1\work\	Master file invoking sub-files	
5	2		paths1.do	c:\dames\workshops\2010\4\applic_1\work\	Sets the paths used	
6	3		pooling_data_1.do	c:\dames\workshops\2010\4\applic_1\work\	Combines BHPS data from multiple waves	
7	4		pre_analysis_1.do	c:\dames\workshops\2010\4\applic_1\work\	Recodes variables, missing data declarations	
8	5		data_review_1.do	c:\dames\workshops\2010\4\applic_1\work\	Generates summary statistics	
9	6		graphs_moving_educ_1.do	c:\dames\workshops\2010\4\applic_1\work\	Generates graphs and produces emfs	
10						
11						
12	7	Macros	casoc_iscs.do	http://www.camsis.stir.ac.uk/downloads/data/other/cas	Allows conversion of string ISCO to valid numeric	
13						
14	8	Data:	BHPS source files	c:\data\bhps\bh11709\	UKDA, SN: 5151	
15			[Files derived by above]	c:\dames\workshops\2010\4\applic_1\data\		
16						
17		MLwiN work:				
18	9	Command files:	read_bhps_data.mac	c:\dames\workshops\2010\4\applic_1\work\	Data setup, after (4) (variable names etc)	
19	10		bhps_educ_3level.mac	c:\dames\workshops\2010\4\applic_1\work\	Runs a 3-level model on outcomes	
20	11	Data:	[Files derived by above]	c:\dames\workshops\2010\4\applic_1\data\		
21						
22						

Note that the other tabs in the Excel file can be used to show things like author details, context of the analysis, and last update time. The file also notes some (though not all) dependencies within the workflow – for instance step 9 requires step 4 to have been taken (the macro reads in a plain text data file that was generated in Stata by do file pre_analysis1.do).

I must keep my files well-organised
 I must keep my files well-organised
 I must keep my files well-organised
 I must keep my files well-organised
 I must keep my files well-organised
 I must keep my files well-organised
 I must keep my files...

In summary, we can't advise you strongly enough on the value of organising your data files around a workflow conceptualisation, such as through master and sub-files. Read the opening chapters of Long (2009), or the other references mentioned above, for more on this theme. We'd encourage you to look at the workshop materials from the 'DAMES' research Node, at www.dames.org.uk, for more on this topic.

Software alternatives

Many different software packages can be used effectively for applied research using complex data on social connections. Various packages support the estimation of a wide range of statistical models including association models, and there are numerous (mostly different) packages which feature techniques for social network analysis.

In this session we focus upon three packages which bring slightly different contributions to multilevel modelling.

- **Stata** is used because it is a popular general purpose package for data management and data analysis which also includes a substantial range of analysis options for dealing with data on social connections. Stata is attractive to applied researchers for many reasons, including its good facilities for storing and summarising estimation results; its support of a wide range of advanced analytical methods which complement a multilevel analysis (e.g. clustering estimators used in Economics); and its wide range of data management functions suited to complex data. Stata is proprietary and may be purchased from: www.stata.com.
- **R** is used because it is a popular freeware that supports many forms of statistical model estimation, social network analysis examples, and has various graphical and data construction capabilities. Many of its facilities are available via extension 'libraries' which are usually installed online. R is a difficult language for social scientists to work effectively with, however, because it brings with it very high 'overheads' in its programming requirements, especially for large and complex data. R is available to install as freeware from: <http://www.r-project.org/>
- **Pajek** is used because it is a freely available and popular package for social network analysis, featuring a wide range of graphical and statistical analysis possibilities. Pajek may be downloaded and installed as freeware from: <http://pajek.imfm.si/doku.php>

We should stress that many more packages can be used effectively for the analyses used below. In addition, an exciting software development in the area being led in the UK is the construction of a generic interface for specifying and estimating complex statistical models of 'arbitrary complexity'. These cover most forms of multilevel models, as well as many other statistical modelling devices. This project is called '**e-Stat**' and is expecting to generate its first publicly available resources over the period 2010-2012 (see <http://www.cmm.bristol.ac.uk/research/NCESS-EStat/>).

Lab 1: Introduction to the analysis of social connections data

This lab introduces a few examples of datasets on social connections, and provides illustrative analyses in Stata, R and Pajek.

The work of the Stata and R exercises is done by the corresponding command files, which should (hopefully) be self-explanatory. We introduce using Stata and R below.

The Pajek exercises are described through step-by-step instructions, provided below.



Background: Introducing Stata

Stata was first developed in 1984 and was originally used mainly in academic research in economics. From approximately the mid 1990's its functionalities for social survey data analysis began to filter through to other social science disciplines, and in the last decade it has displaced SPSS as the most popular intermediate-to-advanced level statistical analysis package in most academic disciplines which use social survey data (e.g. sociology, educational research, geography).

Stata is popular for many good reasons. The list of features of Stata that lead me personally to favour this package above others are:

- It supports explicit documentation of complex processes through a concise and 'human readable' syntax language
- It supports a wide range of data management functions including many routines useful in complex survey data which are not readily performed in other packages (e.g. 'egen', 'xtdes')
- It supports a very full range of statistical modelling options, including several advanced model specifications which are not widely available elsewhere
- It has excellent graphics capabilities, supporting the specification and export of publication quality graphs (in a syntactical, replicable manner)
- It features very convenient tools for storing the results from multiple models or analyses and compiling them in summary tables or files (e.g. 'est store', 'statsby')
- It can read online data files and run command files and macros from online locations
- It supports extended add-on programming capabilities, and benefits from a large, constructive community of user-contributed extensions (see e.g. <http://www.stata.com/links/resources3.html>)

In pragmatic terms, most users of Stata are reasonably confident programmers, and getting started with the package does need a little effort in learning about data manipulation and data analysis. This is one reason why Stata is not yet widely taught in introductory social science courses, though, in the UK for example, it is increasingly used in intermediate and advanced level teaching (e.g. MSc programmes or Undergraduate social science programmes with extended statistical components).

A common problem with working with Stata is that many institutions do not have site-level access to the software, and accordingly many individual researchers don't have access to the package - Stata is generally sold as an 'n-user' package, which means that an institution buys a specified number of copies at any one time. Recently however, access to Stata for academic researchers has probably be made easier by the Stata 'GradPlan', which allows purchase of personal copies of the package for students and faculty at fairly low price – see <http://www.stata.com/order/new/edu/gradplan.html> . Stata also comes in several different forms with different upper limits on the scale of data it may handle – 'Small Stata' is not normally adequate for working with advanced survey datasets; 'Intercooled' Stata (I/C) usually has more than enough capacity to support social survey research analysis (although, working with a large scale resources you may occasionally hit upper limits, such as on the number of variables or cases, it is usually possible to find an easy work-around such as by dropping unnecessary variables); Stata SE and MP offer greater capacity regarding the size of datasets and faster processing power, but they are more expensive to purchase. To my knowledge, most academic researchers use Intercooled Stata.

In summary, many users of Stata favour the package not because it offers one particular functionality which others don't, but because it offers an integrated set of advanced functionalities covering data management and data analysis which can't easily be matched by any other software. For other texts which explain the strengths and attractions of Stata, see for example Treiman (2009).

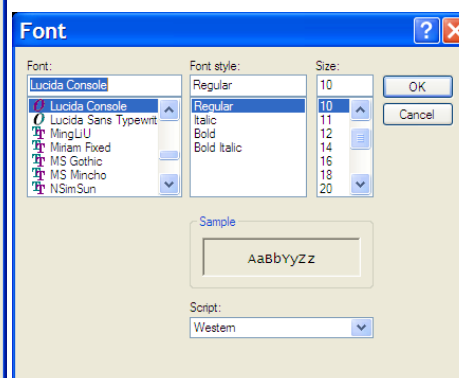
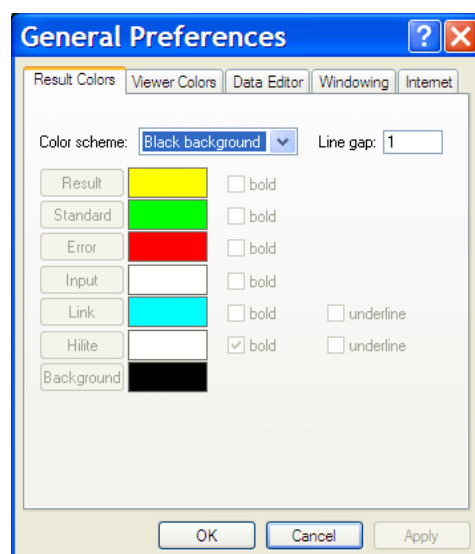
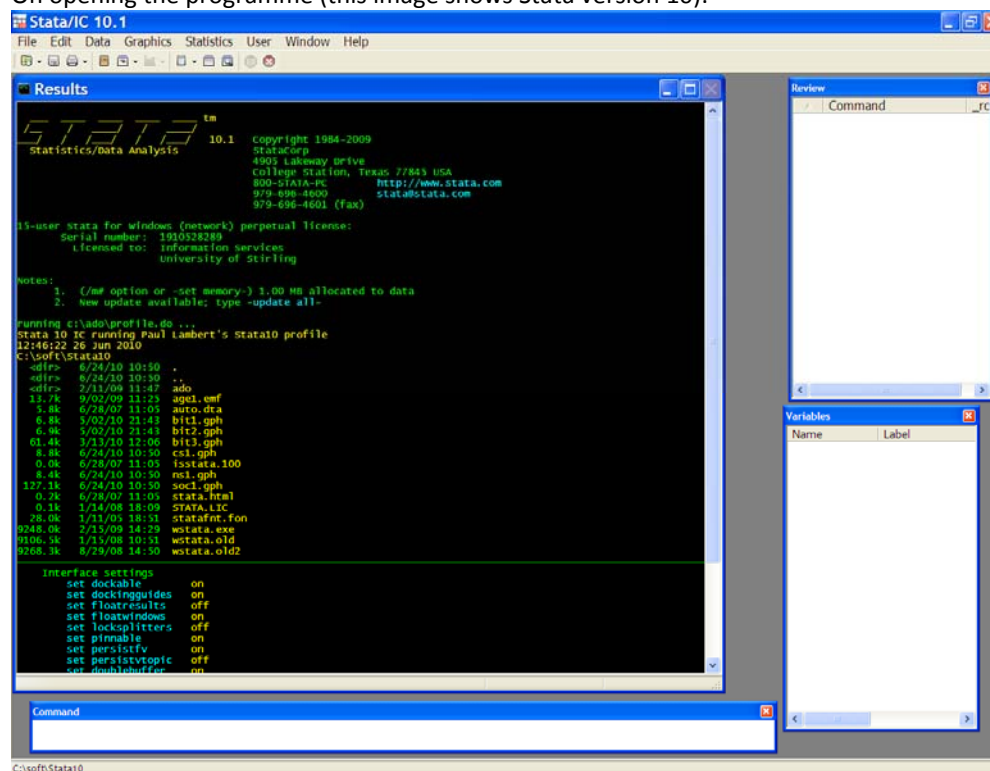
The steps below give you some relevant instructions on working with Stata for the purposes of the module (the examples are mostly from the Practical 1 Stata file). Many online resources on Stata are available, in particular we highlight:

- UCLA's ATS pages: <http://www.ats.ucla.edu/stat/stata/> (Features a wide range of materials including videos of using Stata and routines across the range of the package)
- The CMM's LEMMA online course: <http://www.cmm.bristol.ac.uk/learning-training/course.shtml> (includes detailed descriptions of running basic regression models and of specifying random effects multilevel models in Stata)
- In the first lab session we point you to an illustrative do file which serves as an introduction to Stata, available from www.longitudinal.stir.ac.uk

When you launch the package, you see the basic Stata window, here for version I/C 10.1.

You can customise its appearance (e.g. right click on the results window) – the image on the right is how I set up the windows on my machine and will be slightly different to what you see on first launching the package in the lab at Essex.

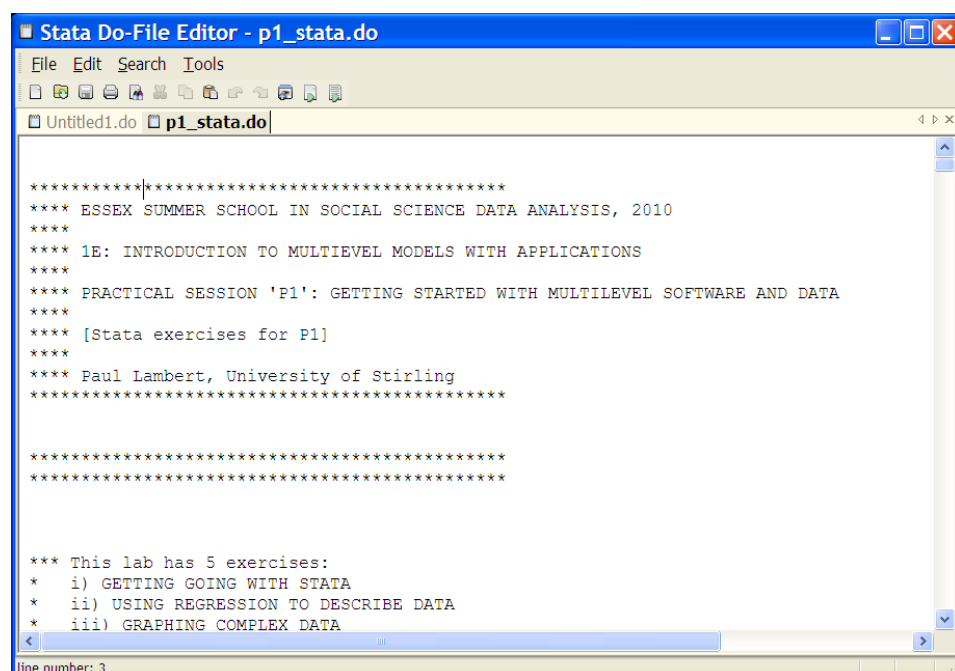
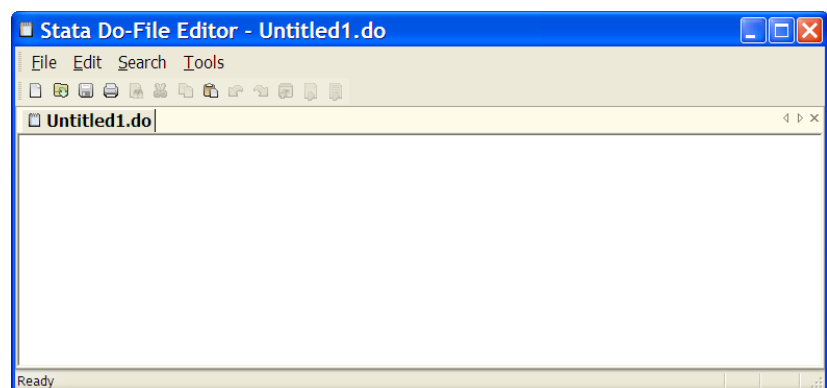
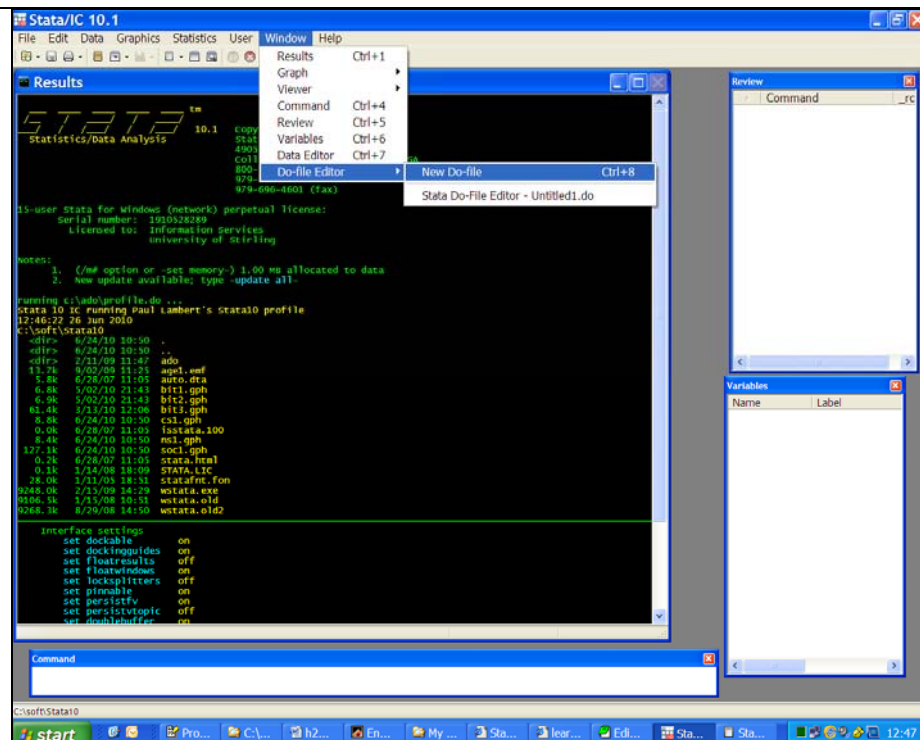
On opening the programme (this image shows Stata version 10):



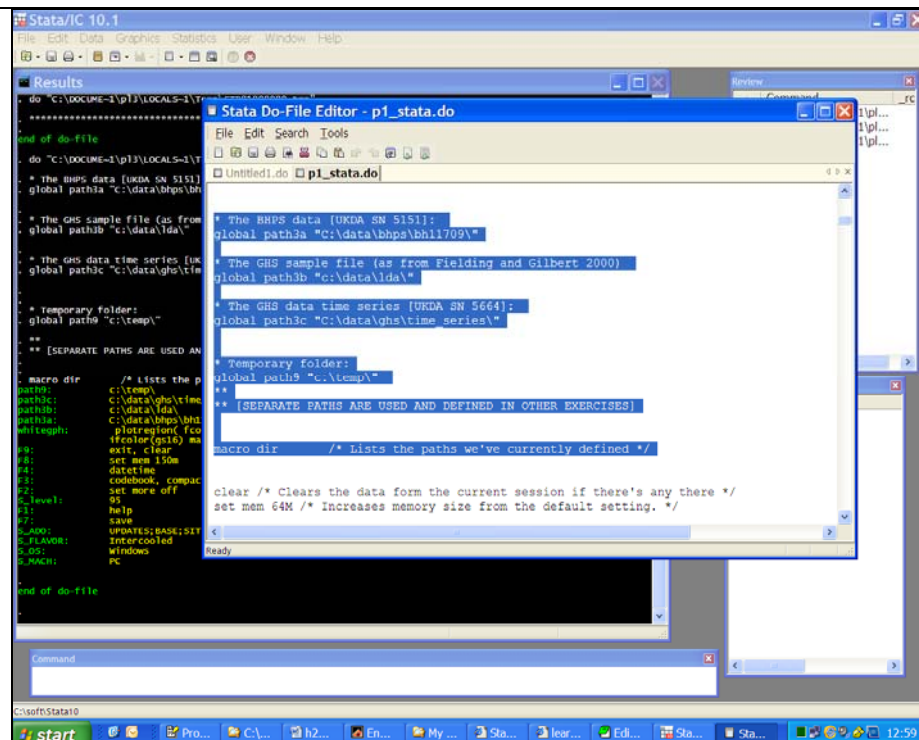
The very first thing you should do at the start of every session it so ask explicitly to open the 'do file' editor with 'ctrl+8' or via the GUI.

Note below that we can have several do files open at once.

Not shown below, but from Stata 11 onwards, it is possible to permit various formatting options in the do file editor (e.g. colour coding). It is also possible to set up Stata to run directly from a plain text editor if you wish to (search online for how to do this).



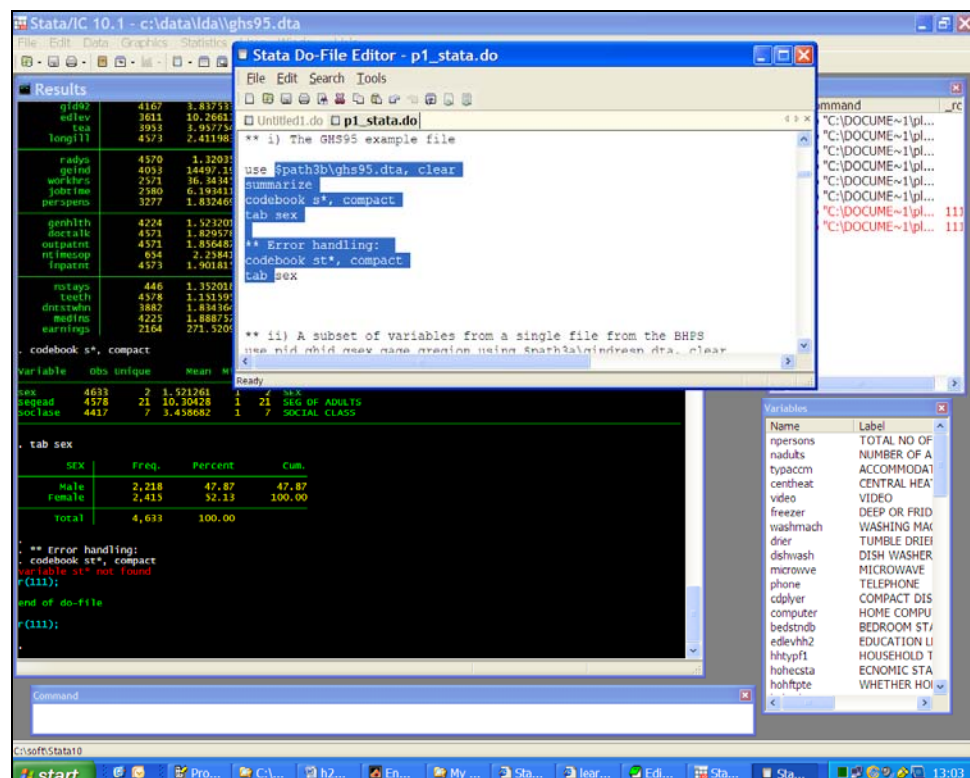
Once you've opened a 'do file' you can begin running commands by clicking on the segments of the relevant command lines and clicking 'ctrl+R'.



Important: Defining macros for paths. This particular image shows an important component of the start of every session in the module lab exercises. The lines beginning with 'global' are ways of defining permanent 'macros' for the session. The macros serve to define locations on my machine where my files (e.g. data files) are actually stored. Doing this means that in later commands (e.g. the image below) I can call on files via their macro folder locations rather than their full path – this aids transferability of work between machines/locations.

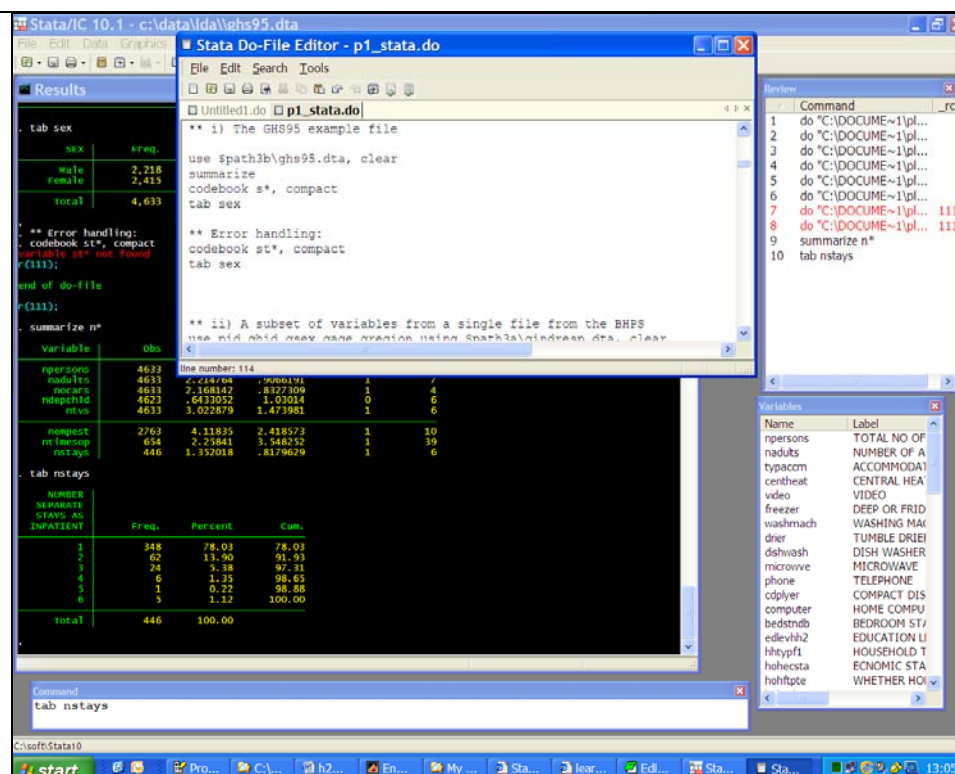
The results are shown in the results window. Error messages are by default shown in red text and lead to the termination of the sequence of commands at that point

(unlike in SPSS, which carries on, disregarding the error).



In the above, the macro which I have called 'path3b' means that when Stata reads the line:
 use \$path3b\ghs95.dta, clear
 what it reads behind the scenes is
 use c:\data\lda\ghs95.dta, clear

You can also submit commands line by line through the command interface (e.g. if you don't want to log them in the do file).



Note how the 'review' window shows lines that were entered through the command window, but it just shows some programming code for commands run through the do file editor.

Note some of the features of the Stata syntax language:

You need to 'clear' the dataspace to read in a new file, e.g.

```
use $path3b\ghs95.dta, clear
```

You can't create a new variable with the same name as an existing one – if it's there already you need to delete it first, e.g.

```
drop fem
gen fem=(sex==2)
```

The 'capture' command suppresses potential error messages so is a useful way to make commands generic

```
capture drop fem
gen fem=(sex==2)
```

Using 'by:' or 'if..' within a command can usefully restrict the specifications:

```
tab nstates if sex==1
bysort sex: tab nstates if age >= 20 & age <= 30
```

The 'numlabel' command is a useful way to show both numeric values and categorical labels compared to the default (labels only), e.g.;

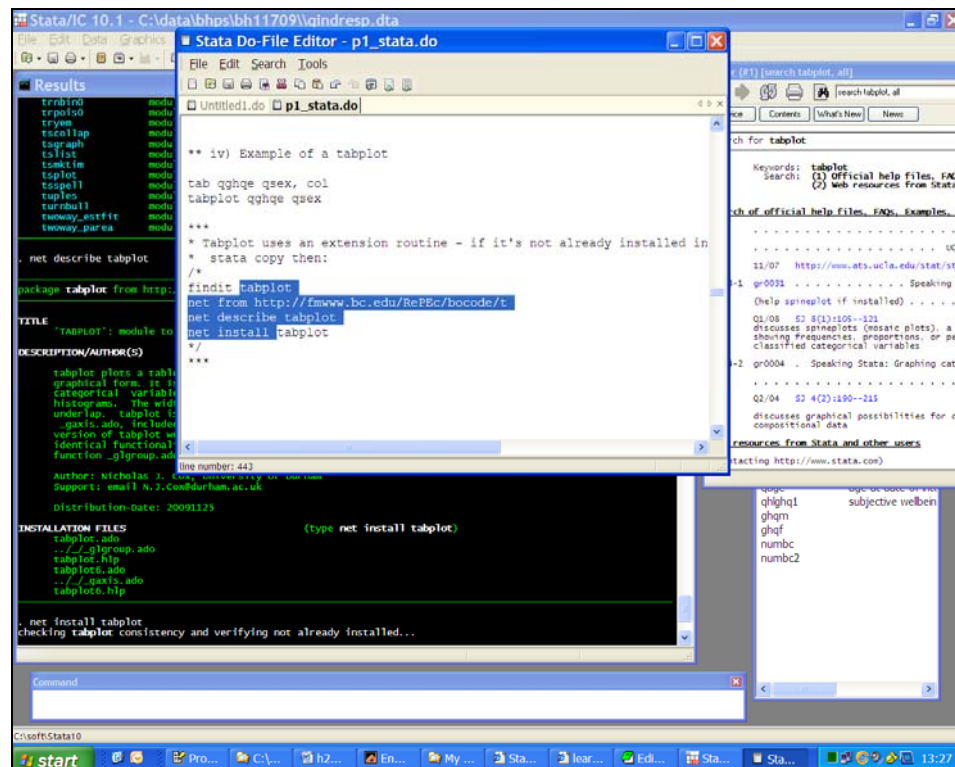
```
tab ecstaa
numlabel _all, add
tab ecstaa
```

There's no requirement for full stops at the end of lines, but a carriage return serves as the default delimiter, and so we usually use '///' to extend a command over more than one line.

```
bysort sex: tab nstates ///
if age >= 20 & age <= 60 & ecstaa==1
```

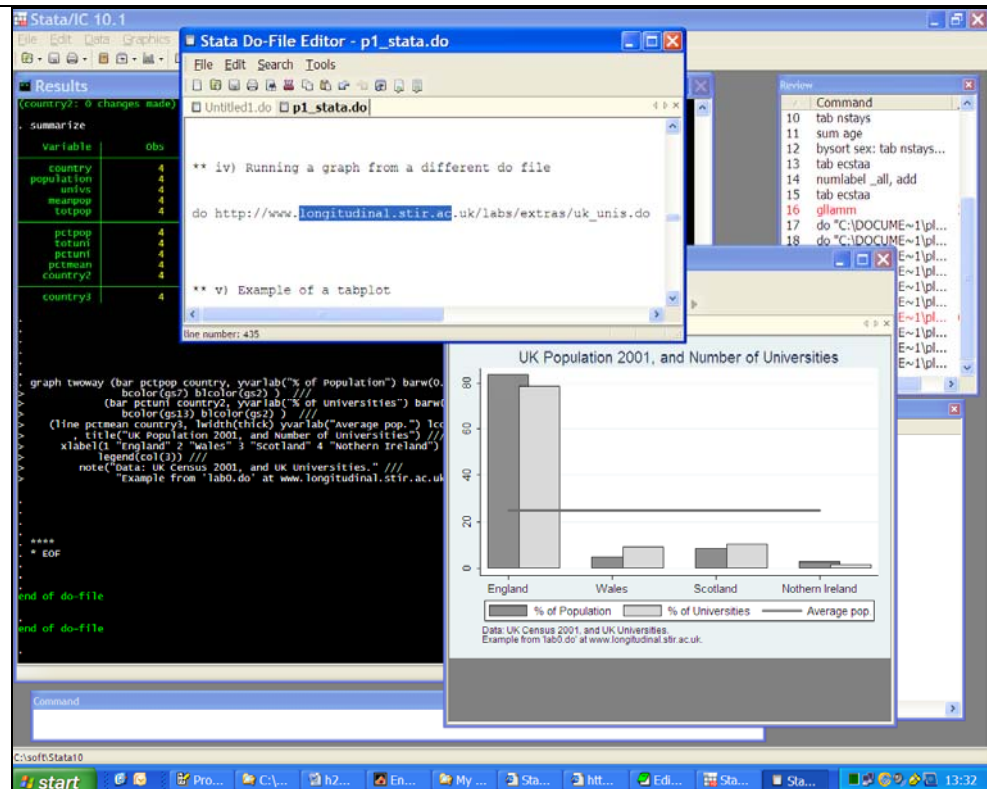

Extension routines are often written by users and made available to the wider community. To exploit them, you need to run either 'net install' or 'ssc install'

Example of finding and installing the 'tabplot' extension routine:

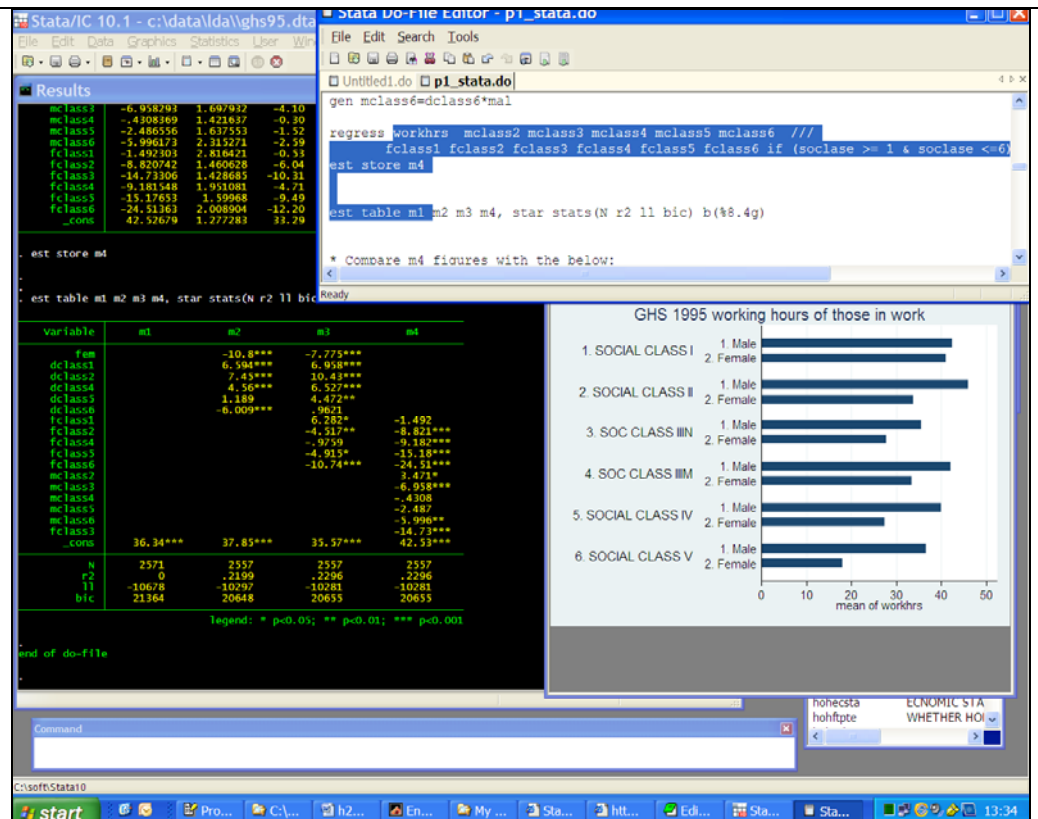


(the exact code needed may depend on which machine you are working on – you may have to define a folder for the installation that you have permission to write to)

We often run subfiles, or define macros or programmes, via calling upon other do files with the 'do' command



'est store' is a great way to collate and review multiple model results



Extension:
You can write a '.profile' file to load some starting specifications into your session

For lots more on Stata, see the links and references given above, or the DAMES Node workshops at dames.org.uk

```

** Stata profile

global F2 "set more off"
global F3 "codebook, compact"
global F4 "datetime"
global F8 "set mem 150m"
global F9 "exit, clear"

global whitegph " plotregion( fcolor(gs16) ifcolor(gs16) lcolor(gs16) ilcolor(gs16) ) graphreg
doedit

noisily display "Stata 10 IC running Paul Lambert's Stata10 profile"
noisily display "`c(current_time)' `c(current_date)'"

noisily pwd
noisily dir
set more off
set scrollbufsize 320000
noisily query interface

version 7.0
capture program drop datetime
program define datetime
disp "DateTime: $$_DATE $$_TIME"
end

```



Background: Introducing R

R is a freeware which is a popular tool amongst statisticians and a small community of social science researchers with advanced programming skills. It is an 'object oriented' programming language which supports a vast range of statistical analysis routines, and many data management tasks, through its 'base' or extension commands. Being 'object oriented' is important and means the package appears to behave in a rather different way to the other packages described above. The other packages essentially have one principal quantitative dataset in memory at any one time, plus they store metadata on the matrix and typically some other statistical results in the form of other scalars and matrices. In the other packages, commands are automatically applied to the variables of the principal dataset. In R, however, different quantitative datasets ('data frames'), matrices, vectors, scalars and metadata, are all stored as different 'objects', potentially alongside each other. R therefore works by first defining objects, then second performing operations on one or many objects, however defined.

Some researchers are very enthusiastic about R, the common reasons being that it is free and that it often supports exciting statistical models or functions which aren't available in other packages. However, my perspective is that R isn't an efficient package for a social survey researcher interested in applied research, as the programming demands to exploit it are very high, and, because it isn't widely used in applied research, it hasn't yet developed robust and helpful routines, working interfaces, or documentation standards, to address popular social science data-oriented requirements.

An important concept in R is the 'extension library', which is how 'shortcut' programmes to undertake many routines are supplied. In fact, you will rarely use R without exploiting extension libraries. The idea here is that R has a 'base' set of commands and support, and that many user-contributed programmes have been written in that base language. Those extensions typically provide shortcut routes to useful outcome analyses. A few extension libraries in R are specifically designed to support random effects multilevel model estimation – e.g. the lme package (Bates, 2005; Pinhero & Bates, 2000).

R is installed as freeware and since it is frequently updated it is wise to regularly revisit the distribution site and re-download

The screenshot shows the R Project for Statistical Computing website in a Mozilla Firefox browser window. The browser's address bar shows the URL <http://www.r-project.org/>. The website features the R logo, a navigation menu with links like 'About R', 'What is R?', 'Contributors', 'Screenshots', 'What's new?', 'Download, Packages', 'CRAN', 'R Project', 'Foundation', 'Members & Donors', 'Mailing Lists', 'Bug Tracking', 'Developer Page', 'Conferences', and 'Search'. The main content area displays a 'Getting Started' section with the following text:

Getting Started:

- R is a free software environment for statistical computing and graphics. It compiles and runs on a wide variety of UNIX platforms, Windows and MacOS. To [download R](#), please choose your preferred [CRAN mirror](#).
- If you have questions about R like how to download and install the software, or what the license terms are, please read our [answers to frequently asked questions](#) before you send an email.

The page also includes a sidebar with a search bar and a list of links. The main content area features a large scatter plot titled 'PCA 5 vars' and a smaller plot titled 'Clustering 4 groups'.

When you open R, you will see something like this 'R console'

```

R R Console
File Edit Misc Packages Windows Help

R version 2.10.1 (2009-12-14)
Copyright (C) 2009 The R Foundation for Statistical Computing
ISBN 3-900051-07-0

R is free software and comes with ABSOLUTELY NO WARRANTY.
You are welcome to redistribute it under certain conditions.
Type 'license()' or 'licence()' for distribution details.

Natural language support but running in an English locale

R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

[Previously saved workspace restored]

Paul Lambert, R for Windows using customised .Rprofile 2
Started at Mon Jun 28 00:42:05 2010
> |

```

(on my machine I use a '.rprofile' settings file so my starting display is marginally different to the default)

The first few lines show me defining a new object (a short vector) and listing the objects in current memory.

```

R R Console
File Edit Misc Packages Windows Help

R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

[Previously saved workspace restored]

Paul Lambert, R for Windows using customised .Rprofile 2
Started at Mon Jun 28 00:42:05 2010
> new_object <- c(1,2,3,4,56,67,78.3,10006)
> new_object
[1] 1.0 2.0 3.0 4.0 56.0 67.0 78.3 10006.0
> next_object <- new_object*3
> next_object
[1] 3.0 6.0 9.0 12.0 168.0 201.0 234.9 30018.0
> ls()
[1] "brdata" "data" "fit.1" "income1" "keep" "kept.cases" "matched.cases"
[8] "n.year" "nes.year" "new_object" "next_object" "plotyear" "region.codes" "rvote"
[15] "used_data" "used_data2" "vote1"
> |

```

R's basic help functions point to webpages.

R: Fitting Linear Models - Windows Internet Explorer provided by Uni...

File Edit View Favorites Tools Help

Address <http://127.0.0.1:16241/library/stats/html/lm.html> Go

lm {stats} R Documentation

Fitting Linear Models

Description

lm is used to fit linear models. It can be used to carry out regression, single stratum analysis of variance and analysis of covariance (although [aov](#) may provide a more convenient interface for these).

Usage

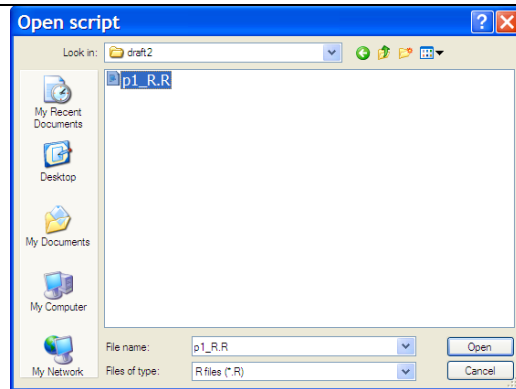
```
lm(formula, data, subset, weights, na.action,
  method = "qr", model = TRUE, x = FALSE, y = FALSE, qr = TRUE,
  singular.ok = TRUE, contrasts = NULL, offset, ...)
```

Arguments

formula	an object of class " formula " (or one that can be coerced to that class): a symbolic description of the model to be fitted. The details of model specification are given under 'Details'.
data	an optional data frame, list or environment (or object coercible by as.data.frame to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment (formula), typically the environment from which lm is called.

The general help pages mostly have generic information, and are not in general provided with worked examples. Many R users get their help from other online sources, e.g. <http://www.statmethods.net/>

In general, with R, the first thing you should do is ask to open a new or existing script and work from that. Scripts in R work in a similar way to a syntax file in Stata or SPSS – highlight a line or lines, and press 'ctrl+R'.



```

R C:\research\workshops\essex_summer_school\intro_multilevel\labs\syntax\...
File Edit Packages Help

#####
#### ESSEX SUMMER SCHOOL IN SOCIAL SCIENCE DATA ANALYSIS, 2010
####
#### 1E: INTRODUCTION TO MULTILEVEL MODELS WITH APPLICATIONS
####
#### PRACTICAL SESSION 'P1': GETTING STARTED WITH MULTILEVEL SOFTWARE AND DATA
####
#### [R exercises for P1]
####
#### Paul Lambert, University of Stirling
#####

## Some examples of mixed models in R

install.packages("foreign")
library(foreign)
# Package for reading Stata and other format files

install.packages("arm")
library(arm)
# Package to assist displaying model results

install.packages("Matrix")
library(Matrix)
# Package to support multilevel model routines

install.packages("lme4")
library(lme4)
# Package to support multilevel model routines

options(digits=4)

#####

popular2 <- read.dta("c:/refs/books/hox2010/data/stata/popular2.dta",
                    convert.factors=F)
# Reads in the data from Hox c2

names(popular2)
attach(popular2)
hist(popular)

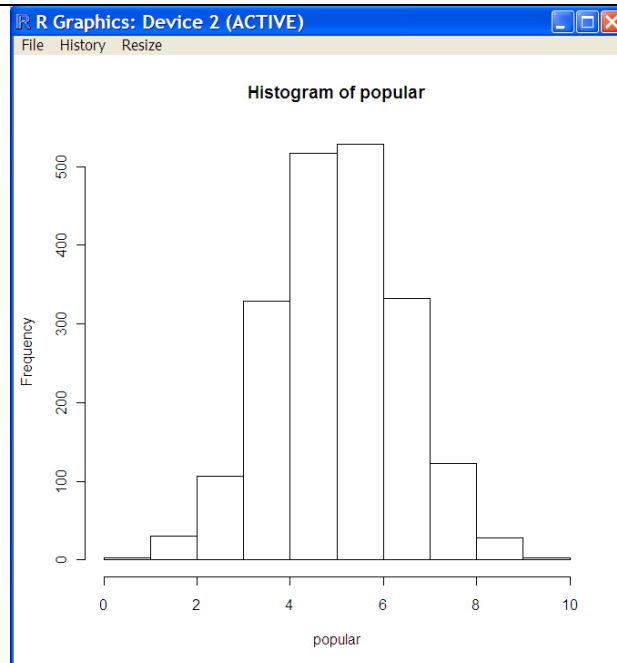
## Linear regression

lin1 <- lm(formula = popular ~ sex + extrav + texp )
summary(lin1)
display(lin1)

## Random intercepts multilevel model with 2-levels, clustering by psu:

```

After running commands, output is sent either to the main console or a separate graphics window



```
R R Console
File Edit Misc Packages Windows Help

>
> lin1 <- lm(formula = popular ~ sex + extrav + texp )
> summary(lin1)

Call:
lm(formula = popular ~ sex + extrav + texp)

Residuals:
    Min       1Q   Median       3Q      Max
-3.72043 -0.60641 -0.01323  0.59580  3.41835

Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept)  0.632307   0.123545   5.118 3.38e-07 ***
sex          1.379262   0.042308  32.601 < 2e-16 ***
extrav       0.475434   0.018126  26.229 < 2e-16 ***
texp         0.088869   0.003488  25.480 < 2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.9358 on 1996 degrees of freedom
Multiple R-squared:  0.5426,    Adjusted R-squared:  0.5419
F-statistic: 789.1 on 3 and 1996 DF,  p-value: < 2.2e-16

> display(lin1)
lm(formula = popular ~ sex + extrav + texp)
      coef.est coef.se
(Intercept)  0.63    0.12
sex          1.38    0.04
extrav       0.48    0.02
texp         0.09    0.00
---
n = 2000, k = 4
residual sd = 0.94, R-Squared = 0.54
>
>
> ## Random intercepts multilevel model with 2-levels, clustering by psu:
>
> mlml <- lmer(formula = popular ~ + (1 | class ) )
> summary(mlml)
Linear mixed model fit by REML
Formula: popular ~ + (1 | class)
      AIC   BIC logLik deviance REMLdev
 6337 6353  -3165    6327    6331
Random effects:
Groups   Name              Variance Std.Dev.
class    (Intercept)  0.7021   0.83791
Residual              1.2218   1.10535
Number of obs: 2000, groups: class, 100

Fixed effects:
            Estimate Std. Error t value
(Intercept)  5.07786   0.08739  58.11
> display(mlml)
```



Lab 1: Pajek exercises

Pajek is a social network analysis software package which has been developed by Vladimir Batagelj and Andrej Mrvar from the University of Ljubljana (hence the name, Slovenian for 'spider'). Some argue the software is not as advanced as competing generalist software such as UCINET or the Social Network package within R, but it has the following benefits:

- It is simple and free to install (for non-commercial use)
- It has easy methods for importing data
- It is simple to use and covers most common network commands
- It is more robust than other packages for dealing with very large datasets.

There are several limitations with Pajek. Unlike R there is a requirement to use drop-down menus meaning it is not possible to run syntax files (although all processes can be saved). It cannot perform some of the emerging analyses such as random graph models, forcing users to use SIENA, PNET or other specialist package. However, Pajek performs the basic elements of network analysis in a very user-friendly manner, which makes it the ideal package for people unfamiliar with network methods (and, therefore, less likely to require the more advanced methods central to other packages). It retains sufficient sophistication to be utilised by many experienced researchers. Most other SNA packages (for instance, UCINET, Siena and PNET) have strong links to Pajek and enable data to be readily imported.

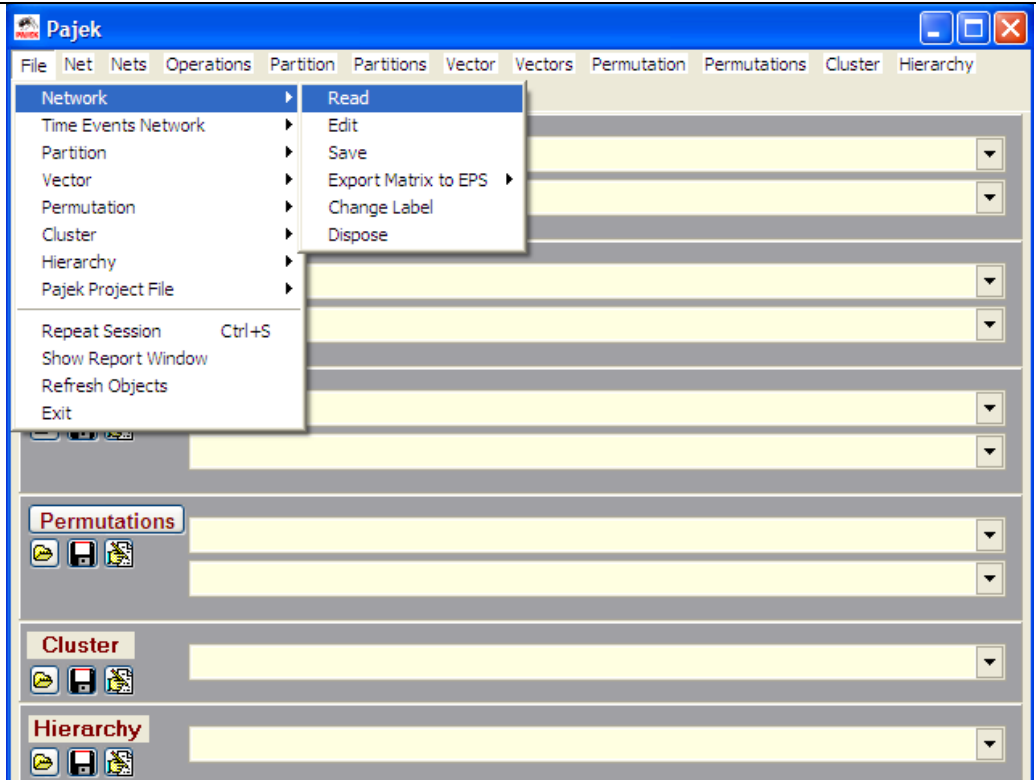
Pajek also benefits from having a comprehensive book providing good examples of how to use the software (de Nooy, W., Mrvar, A., & Batagelj, V. (2012) *Exploratory Social Network Analysis with Pajek*. Cambridge: Cambridge University Press. 2nd edition). This book is an excellent introduction to both Pajek and SNA more generally, providing an overview of each method described and working through examples which convey not only how to perform such analysis but also spells out the benefits of each technique.

The manual, however, is less helpful if you're unsure of how to use Pajek. It provides detailed information on Pajek but in a manner which assumes prior understanding of the operation.

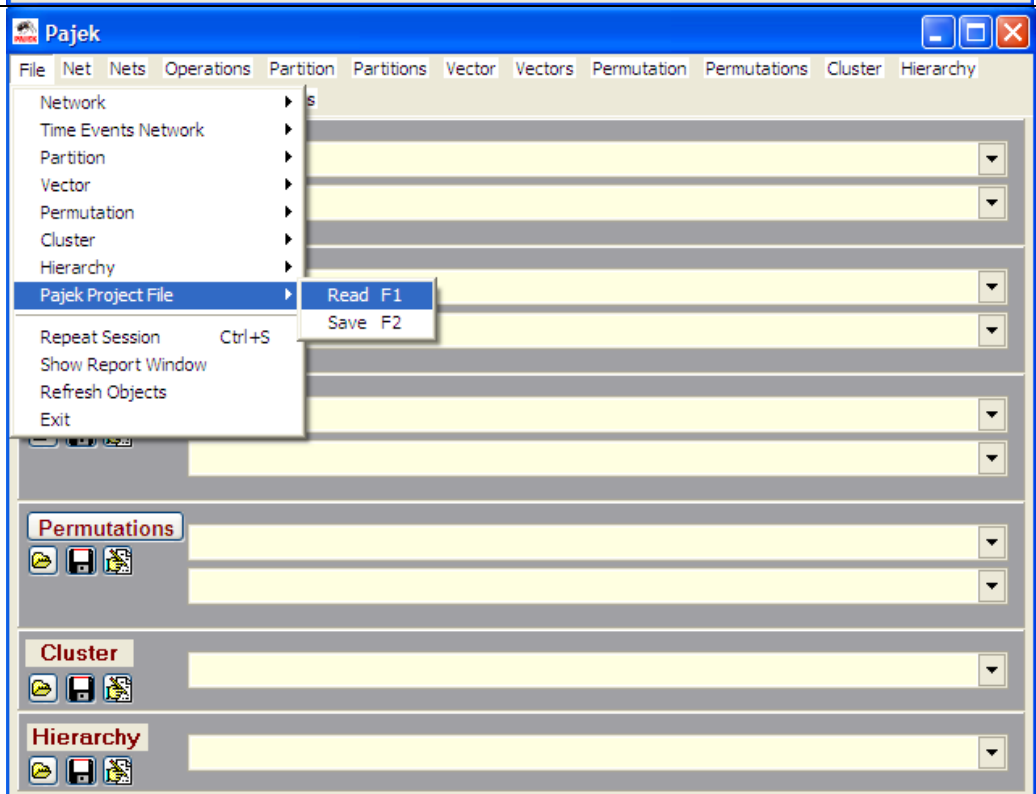
Therefore, it provides many useful resources for experienced users (such as the default colours for vertices and labels for triad censuses), providing in-depth knowledge of the finer points of the package, but the manual is more helpful for advancing your familiarity with the software.

Pajek frequently updates the software (usually fixing tiny glitches, adding new procedures or speeding up processes) so given the ease in installing it's often worth checking you're using the most up-to-date version before starting a piece of work. There is also a dedicated e-mail list which provides rapid answers to complex questions. A new development is the Pajek-XXL programme, which replicates Pajek but operates much faster on huge datasets (tens of thousands of nodes).

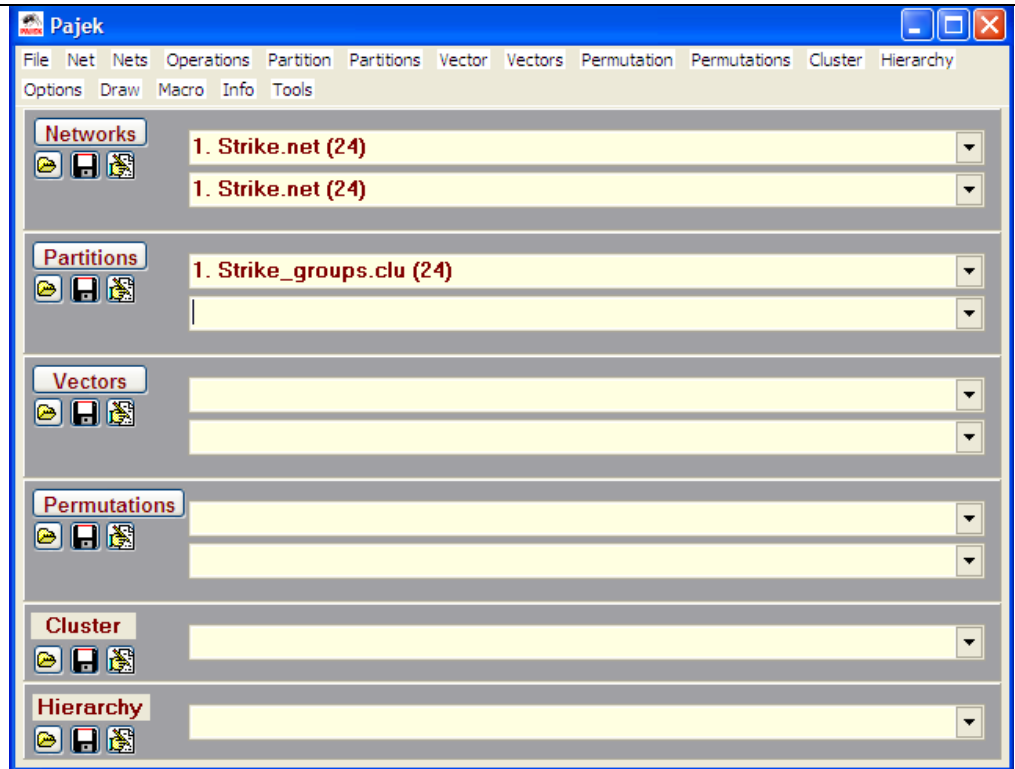
When you first open Pajek you need to import your data. You can import in individual networks one at a time using either the drop down menus, or the 'file open' button. You can then open any saved partitions, vectors and so on in the same way.



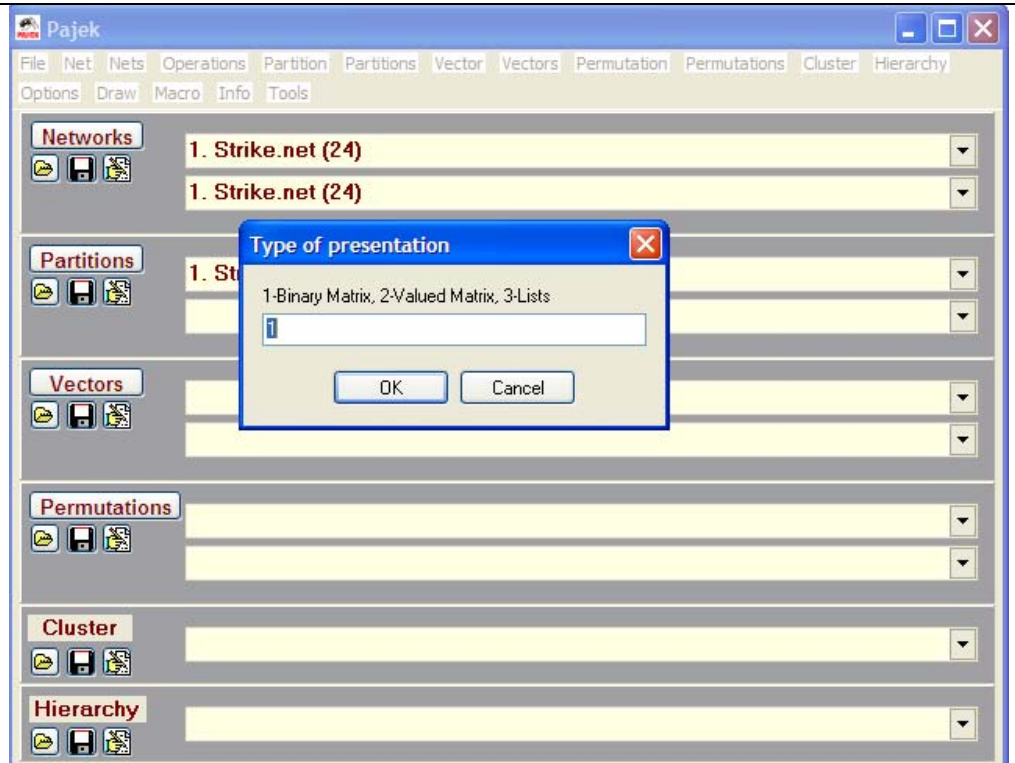
Alternatively, you could open a 'Pajek Project File' which imports all saved data relevant to the piece of work.



This provides the data useful for the research. We can simply add any new networks, partitions etc. to this file by simply opening additional individual data as before. However, opening a new 'Pajek Project File' will remove all the data from the package. Multiple files can be open at once in Pajek. The labels for each file have the structure for a number for the item in the drop list, the file name (or method of construction) and finally the number of cases in brackets.



We can view the matrix which is providing all the information Pajek needs to operate. To do this, click on the actual name on the network in the yellow drop-drop section of the networks tab (the line marked "1. Strike.net (24)" in this example). This produces a dialogue box enabling us to see whether a binary matrix (# marks a link), a valued list (showing the numeric value of the link) or a list (a list of the ties which are formed).

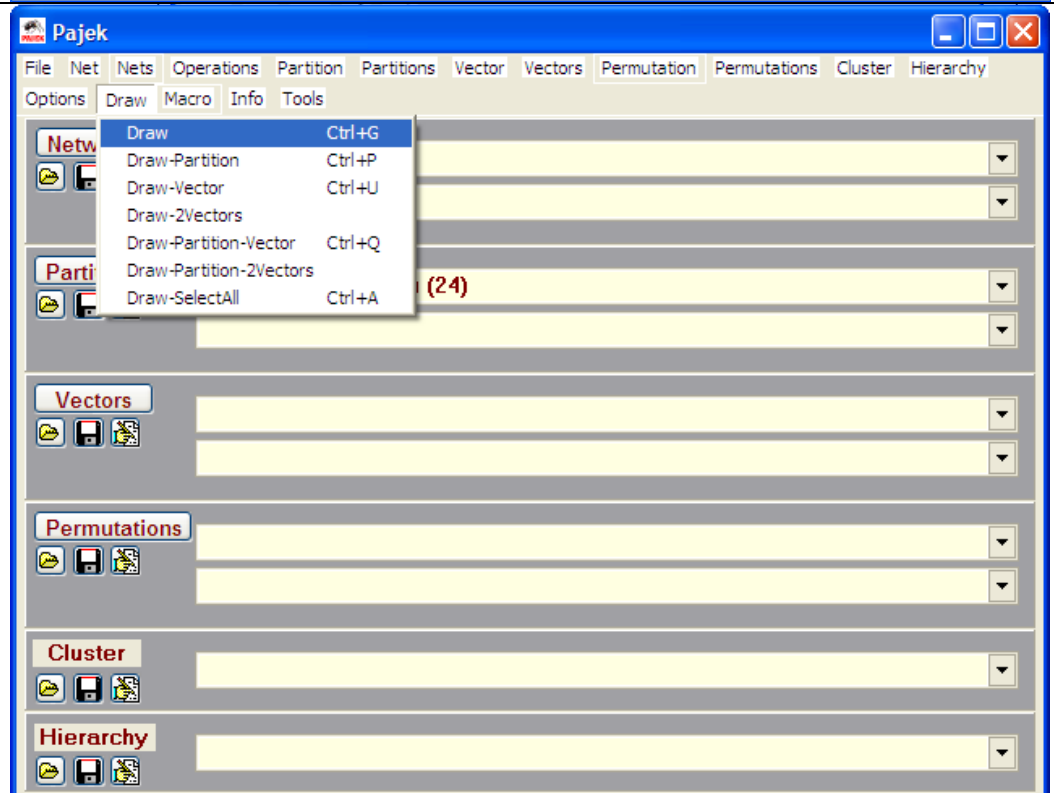


As this is non-numeric data, we have selected a binary matrix (1). This shows the presence of links between actors. The actors in the rows (which are labelled) send a link to the actors in the columns (which are in the same order). In this case, the links must be reciprocated (i.e., if A speaks to B, B speaks to A). This is not always the case (i.e., if A likes B, that doesn't necessarily mean B likes A).

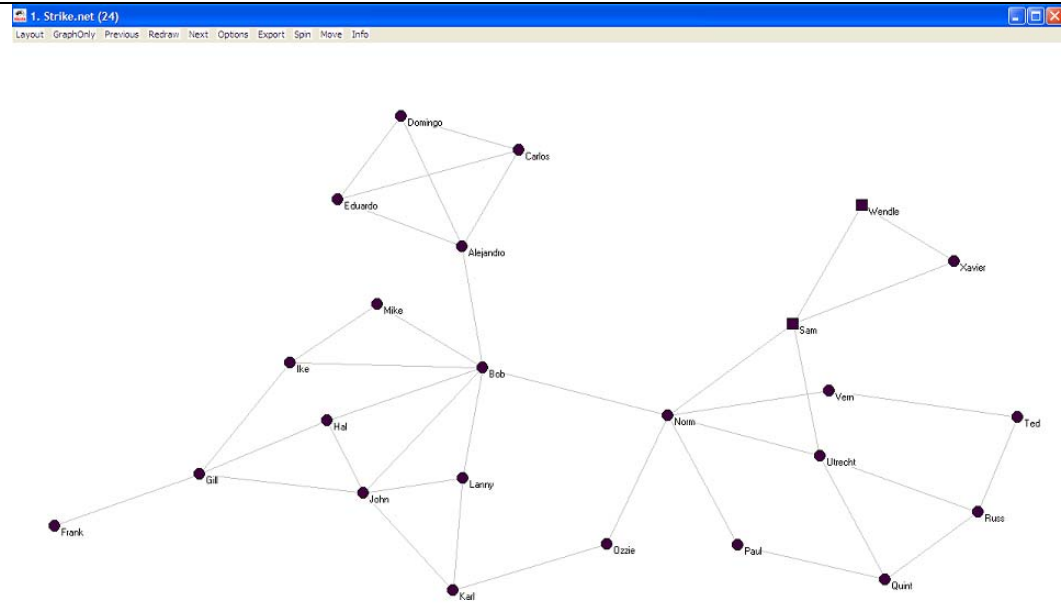
Viewing Network as a Matrix --- 1. Strike.net (24)

	1	2	3	4	5	6	7	8	9	0	1	2	3	4	5	6	7	8	9	0	1	2	3	4	Label
1.	.	#	Frank
2.	#	.	#	#	Gill
3.	.	#	Ike
4.	.	.	#	Mike
5.	.	#	.	#	Hal
6.	.	#	.	.	#	#	John
7.	#	#	Karl
8.	#	Lanny
9.	.	.	#	#	#	Bob
10.	#	#	Alejandro
11.	#	Carlos
12.	#	#	Eduardo
13.	#	#	Domingo
14.	#	#	Norm
15.	#	#	Ozzie
16.	#	Vern
17.	#	Paul
18.	#	#	Quint
19.	#	Utrecht
20.	#	Russ
21.	#	Ted
22.	#	Sam
23.	#	Xavier
24.	#	Wendle

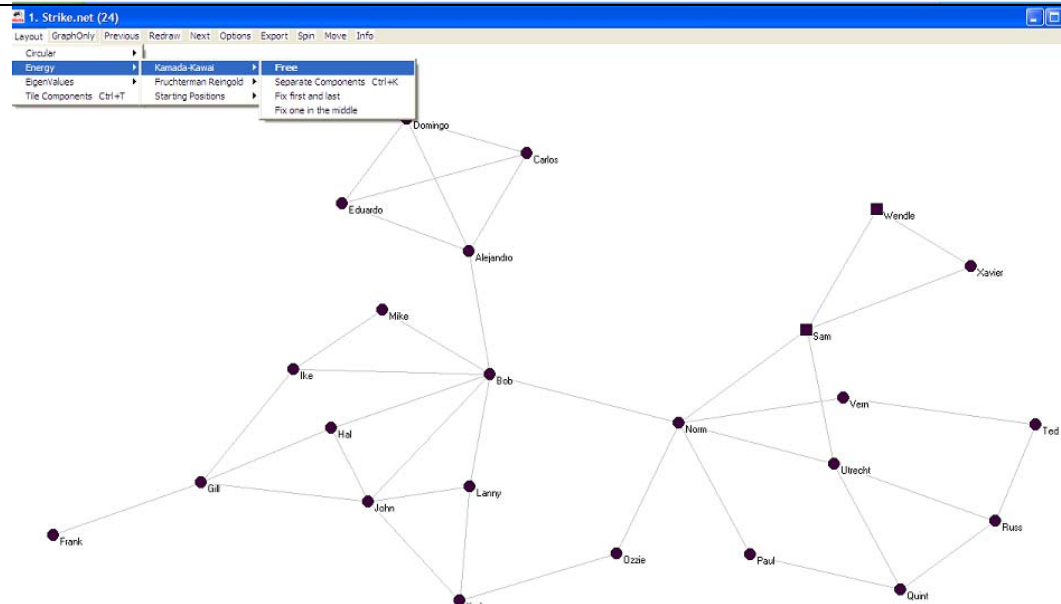
You can also visual the data as a network. Firstly, we will look at the basic structure of the network. This can be done through either using the 'DRAW' drop-down menu, or CTRL+G.



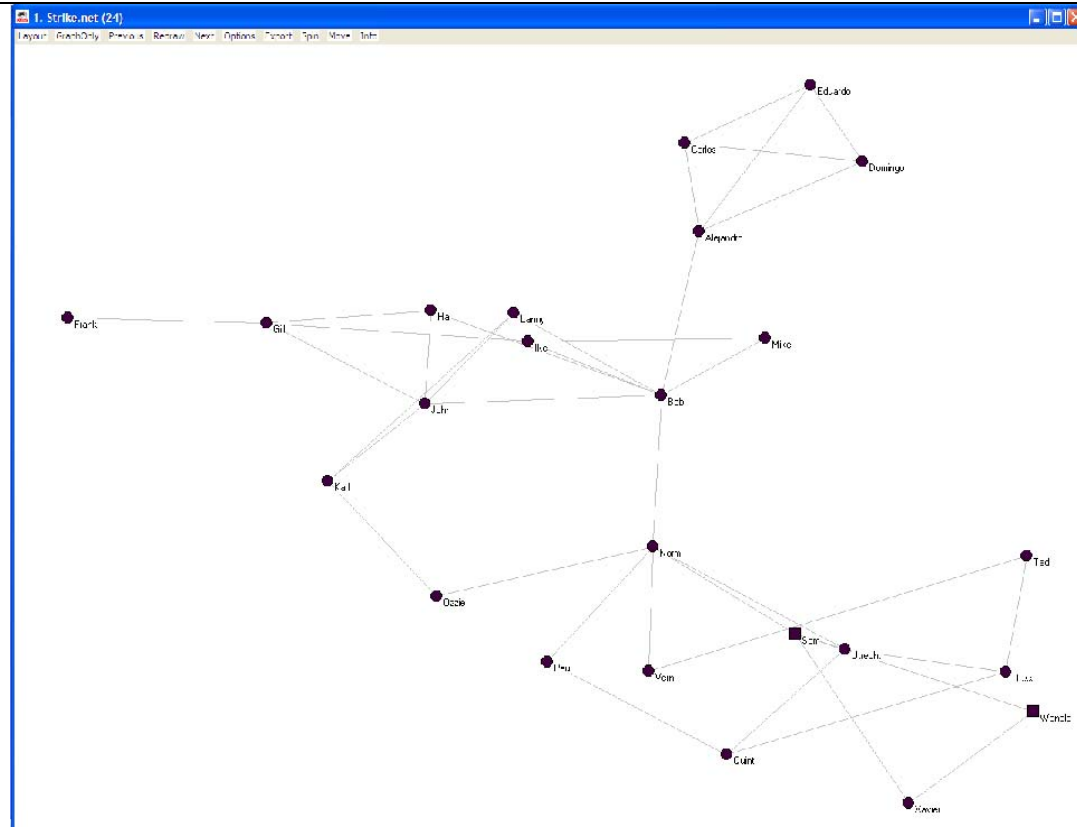
This produces a basic visualisation (in this case, who speaks with whom). Pajek places the nodes in what it believes is the most appropriate position. This is based upon a calculation which can be different each time. Sometimes the graph is displayed nicely the first time, as occurred for me in this example. The graph will be positioned slightly differently each time you generate it.



Using the layout drop-down menu, you can use 'energy', then 'Kamada-Kawai', then 'separate components' (or CTRL+K) to rearrange the network using the same criteria as originally if you want a slightly different visualisation. The links will remain the same, but the nodes will be positioned slightly differently as it tries again to produce the most appropriate layout.



This is an example of how the network can look if repositioned using the Kamada-Kawai equation. Substantially there is no difference between the networks, but the first was spaced more nicely and made the links easier to read. Sometimes the layout is not optimal, therefore it is always useful to press CTRL+K a few times to see a few representations of the data.



This data thus far has not distinguished between actors.. We might have some characteristics of the actors we wish to group them by. This project file contains a partition. Clicking on the name of the partition (highlighted on the Pajek window), brings up a list of the partitions and labels. We can see, therefore, these individuals are split into three groups.

Viewing Partition --- 1. Strike_groups.clu (...)

File	
1.	2 - Frank
2.	2 - Gill
3.	2 - Ike
4.	2 - Mike
5.	2 - Hal
6.	2 - John
7.	2 - Karl
8.	2 - Lanny
9.	2 - Bob
10.	1 - Alejandro
11.	1 - Carlos
12.	1 - Eduardo
13.	1 - Domingo
14.	3 - Norm
15.	3 - Ozzie
16.	3 - Vern
17.	3 - Paul
18.	3 - Quint
19.	3 - Utrech
20.	3 - Russ
21.	3 - Ted
22.	3 - Sam
23.	3 - Xavie
24.	3 - Wendl

Pajek

File Net Nets Operations Partition Partitions Vector Vectors Permutation Permutations Cluster Hierarchy Options Draw Macro Info Tools

Networks

1. Strike.net (24)

1. Strike.net (24)

Partitions

1. Strike_groups.clu (24)

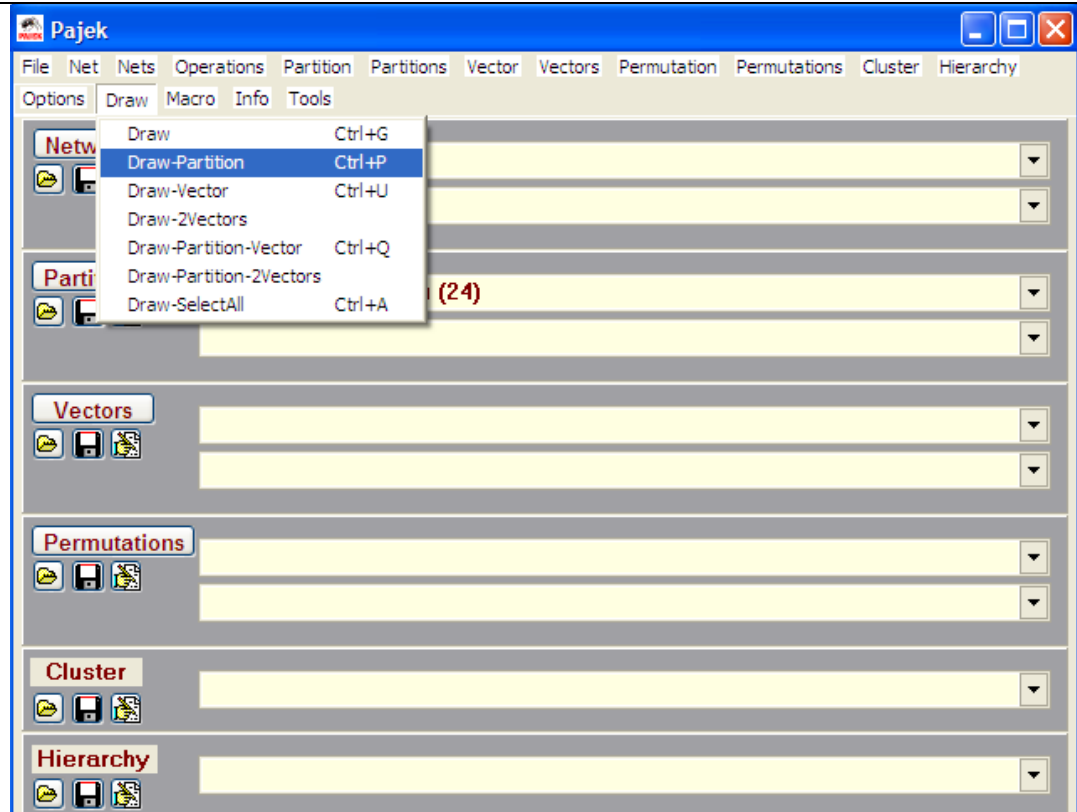
Vectors

Permutations

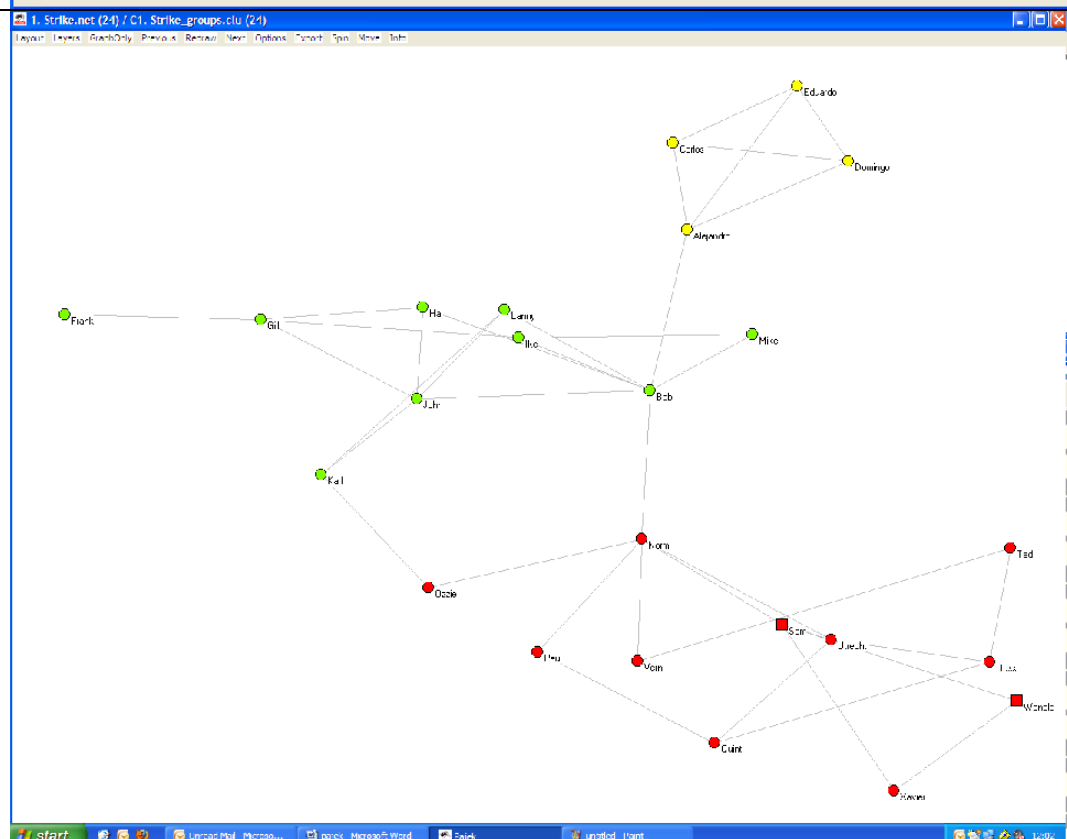
Cluster

Hierarchy

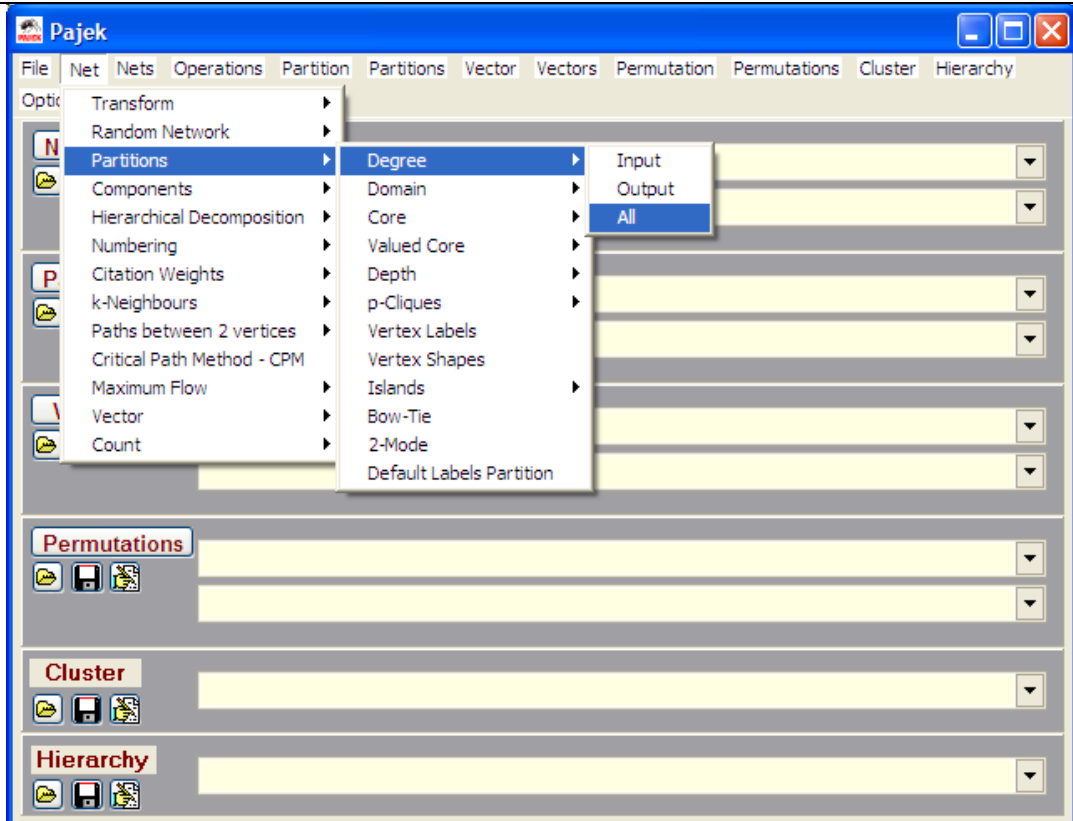
Therefore, we may wish to know how well the grouping within this partition corresponds with the network we've created. We can again go to the draw menu, but select 'draw-partition' (or CTRL+P). This shades the nodes dependent on their partition.



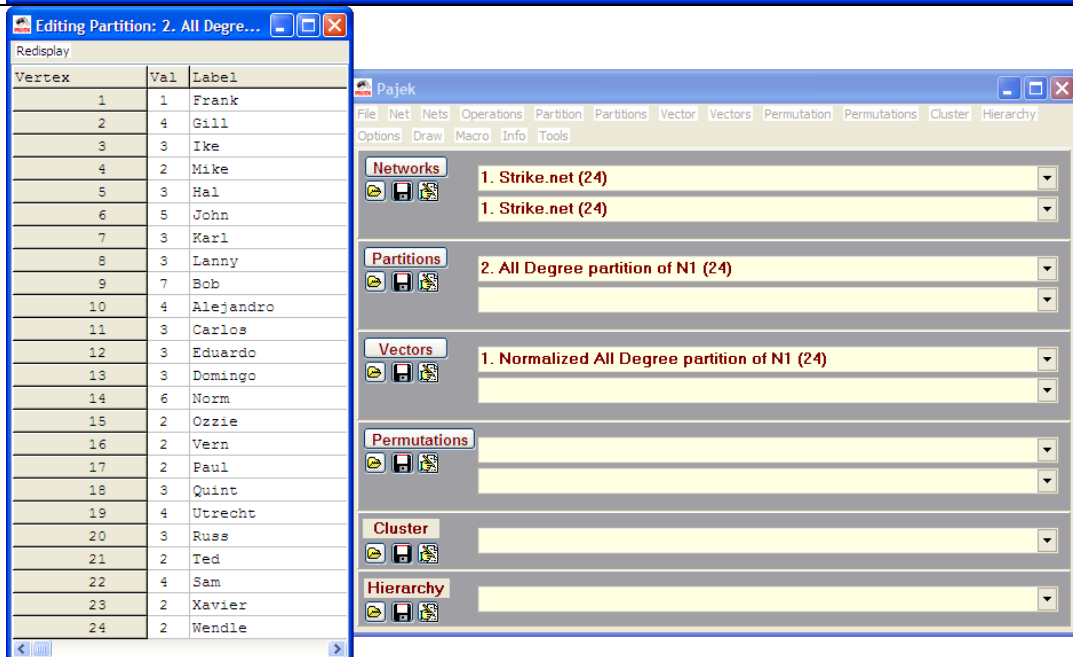
This graph shows we can explain very well the network positions by the characteristics used in creating the partition. In this example, the upper grouping are Spanish speakers, the middle grouping are younger English speakers and the lower group older English speakers. The hypothesis that age and language influences communication networks is clearly supported here.



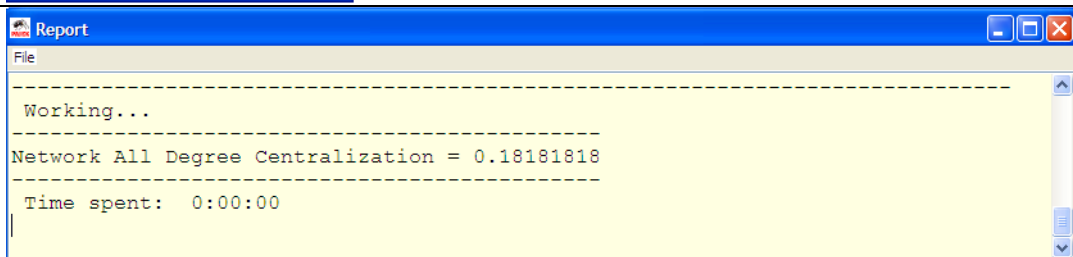
Visualisation is not all that Pajek can do. We might want to gather some statistics about our network. We can see who the most central actors are. We can do this by creating a new partition of the degree for each actor. *(Note: we have the option of input (incoming ties), output (outgoing ties) or all – as ties must be reciprocated within this network there are no differences, but if they were not reciprocated by design the choice would be important.)*



This creates a new partition. Clicking on the name of the new partition (2. All Degree partition of N1 (24)) enables us to see the number of ties each actor has. If we are interested in degree centrality (the number of people each individual speaks to in this case) we can get the information here.

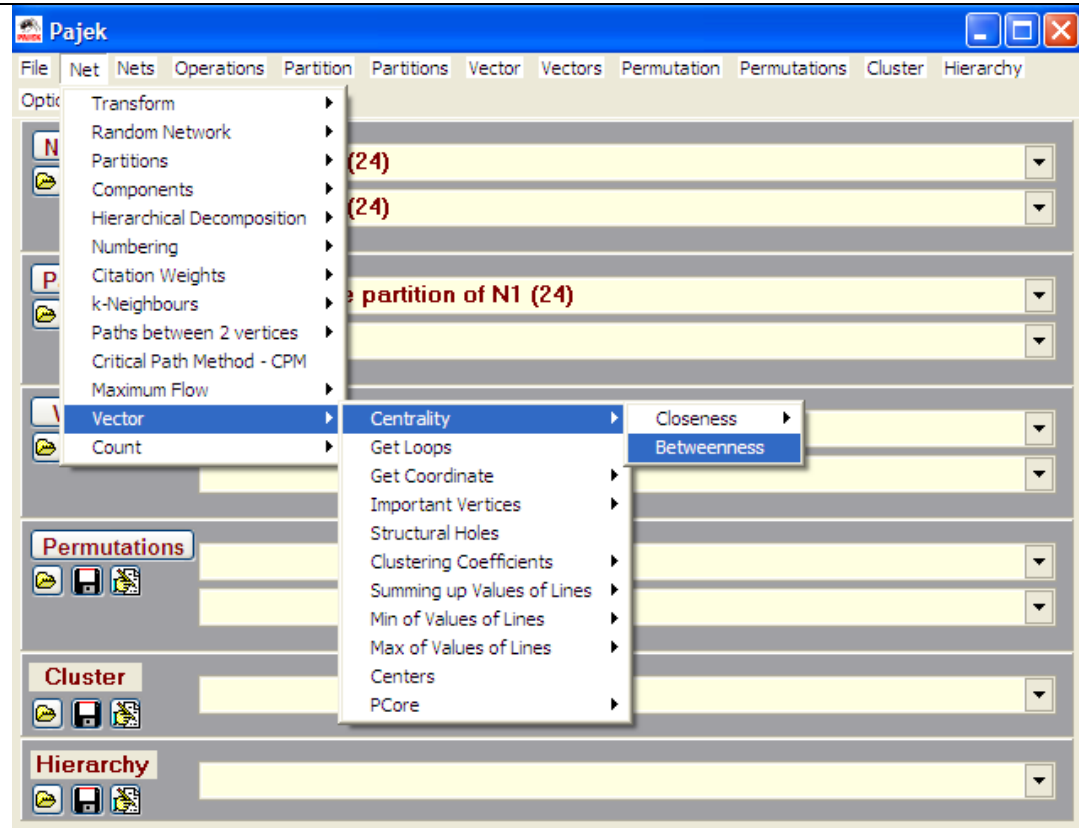


Also created is a report window. This explains what we have done, but also gives us the degree centralization of the network (i.e., the percentage of possible ties which were created). In this case it is 0.18, showing that

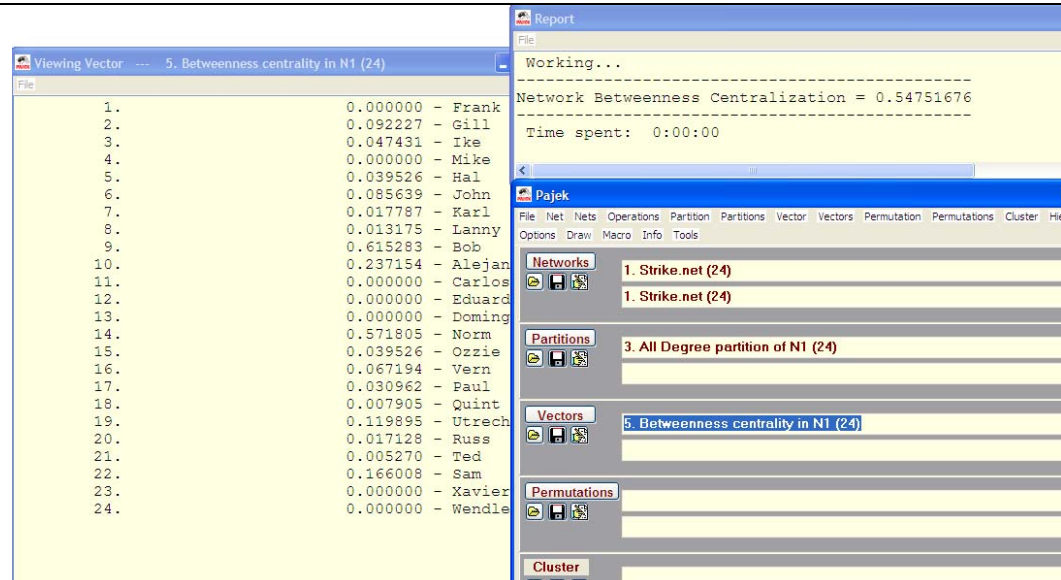


18% of all possible connections (i.e., where everyone speaks to everyone else) have been formed.

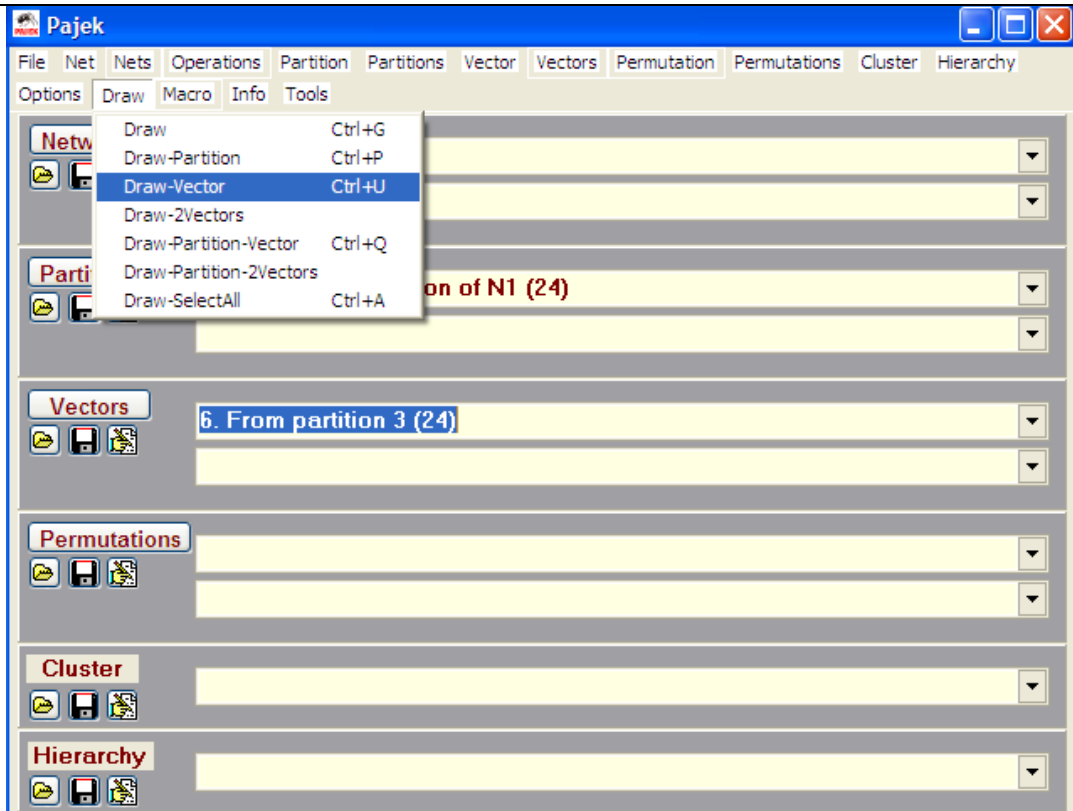
We might be interested in other forms of centrality. Betweenness centrality measures how often an actor is part of a path between two other actors (i.e., where A and B can only communicate through C, whether directly or through others). The higher the betweenness centrality, the increased opportunity for actors to influence and control information which flows around a network. This can be found from the 'net', 'centrality', 'betweenness' drop down menu.



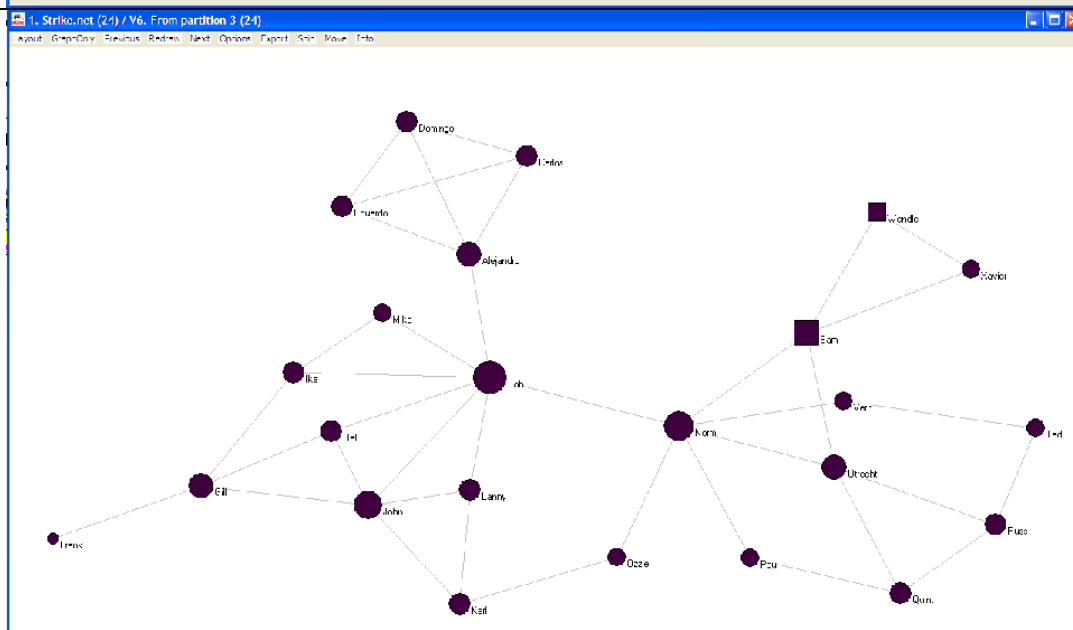
Again, the report window gives us a score for the overall network (.548) which can be compared to other similar networks. Clicking on the highlighted vector enables us to see the scores for each individual. Frank, in the first row, has a score of 0 as he never connects people. Bob (9) has the highest value of 0.61.



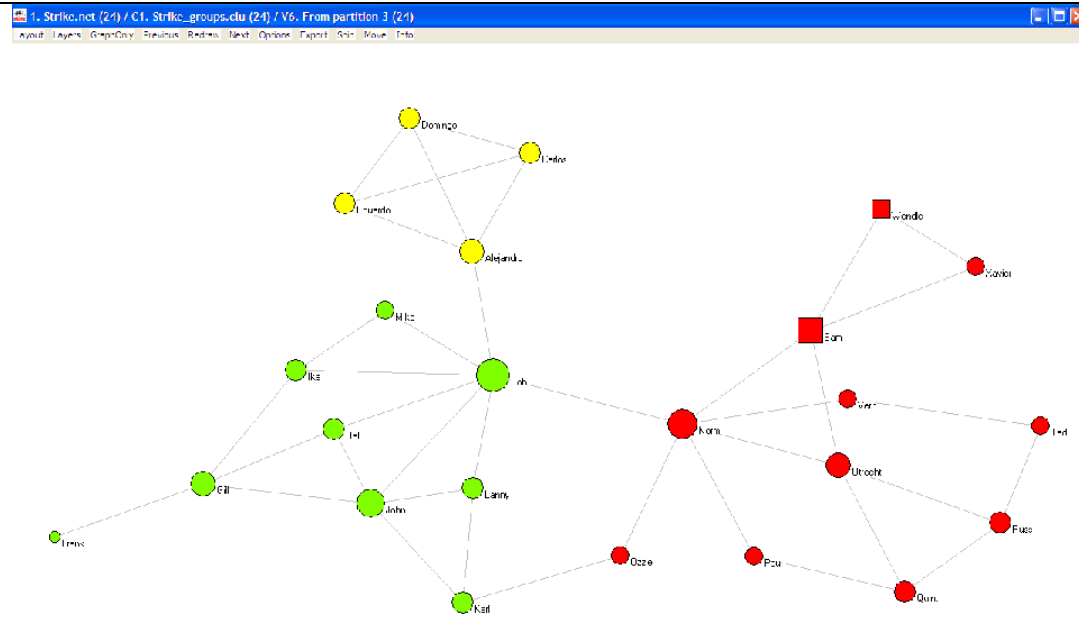
We might want to visualise this to understand the importance of betweenness in terms of the network. We can do this as we did with the partitions, but selecting 'Draw-Vector' (or CTRL+U).



Draw-Vector sizes the node by the scale of the vector. This demonstrates that, for instance, Gill has a reasonable level of influence despite being on the left of the diagram away from the more central actors. This is because they alone can communicate to Frank and manipulate his opinions to the group.

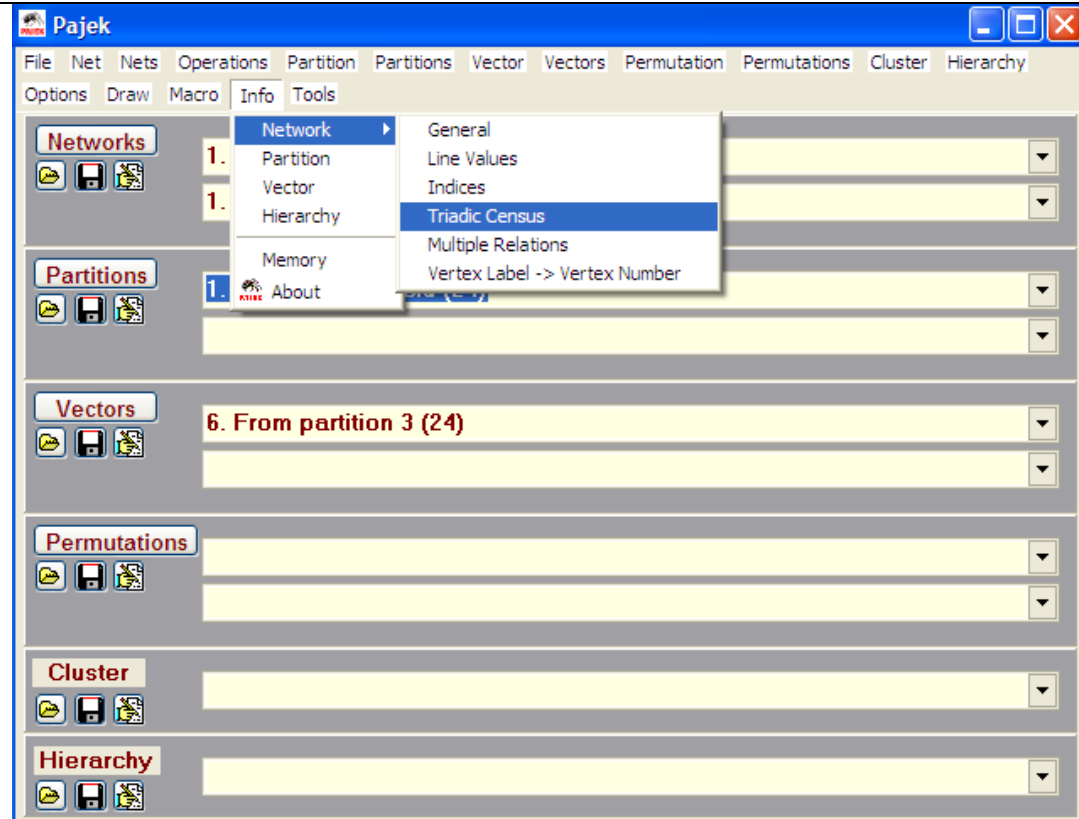


We can mix together different ways of visualising the data. We could look at both partition (in this instance the three groups) and a vector (in this case betweenness) to see how the groups differ. Whilst the upper group of Spanish speakers appears marginalised, using betweenness values we can see they have as much potential influence as the other groups.



We can look at other elements of the network. We might be interested in a triadic census of the data. This can be performed using the info, network, triadic census drop down menu.

A triadic census takes every possible combination of three actors and looks at the structures between them.



This produces a report of the number of each triad we have observed and the expected number given the number of actors and links. The triads are also labelled according to what they show. A chi-square test is conducted, which in this case shows a difference from what would be expected. There is a large percentage of triads which all connect to each other, showing there is balance within the network (i.e., my friends' friend is my friend).

Report

Triadic Census 1. Strike.net (24)

Working...

Type	Number of triads (ni)	Expected (ei)	(ni-ei)/ei	Model
3 - 102	664	63.64	9.43	Balance
16 - 300	12	0.01	869.41	Balance
1 - 003	1280	832.18	0.54	Clusterability
4 - 021D	0	63.64	-1.00	Ranked Clusters
5 - 021U	0	63.64	-1.00	Ranked Clusters
9 - 030T	0	20.32	-1.00	Ranked Clusters
12 - 120D	0	1.62	-1.00	Ranked Clusters
13 - 120U	0	1.62	-1.00	Ranked Clusters
2 - 012	0	797.22	-1.00	Transitivity
14 - 120C	0	3.24	-1.00	Hierarchical Clusters
15 - 210	0	0.52	-1.00	Hierarchical Clusters
6 - 021C	0	127.29	-1.00	Forbidden
7 - 111D	0	20.32	-1.00	Forbidden
8 - 111U	0	20.32	-1.00	Forbidden
10 - 030C	0	6.77	-1.00	Forbidden
11 - 201	68	1.62	40.91	Forbidden

Chi-Square: 20167.3828***
6 cells (37.50%) have expected frequencies less than 5.
The minimum expected cell frequency is 0.01.

Viewing Vector ... 8. Expected Triadic Census of N1 (16)

1.	832.180928
2.	797.215342
3.	63.643242
4.	63.643242
5.	63.643242
6.	127.286483
7.	20.323052
8.	20.323052
9.	20.323052
10.	6.774351

Pajek

File Net Nets Operations Partition Partitions Vector Vectors Permutation Permutations Cluster Hierarchy

Options Draw Macro Info Tools

Networks

- 1. Strike.net (24)
- 1. Strike.net (24)

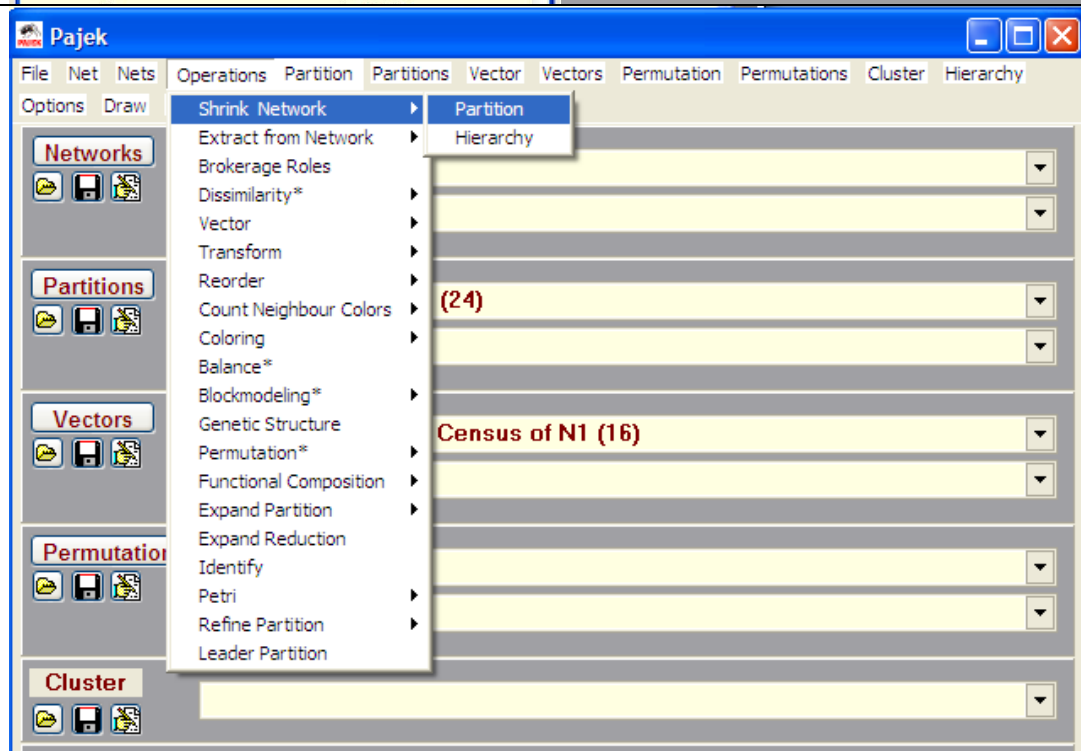
Partitions

- 1. Strike_groups.clu (24)

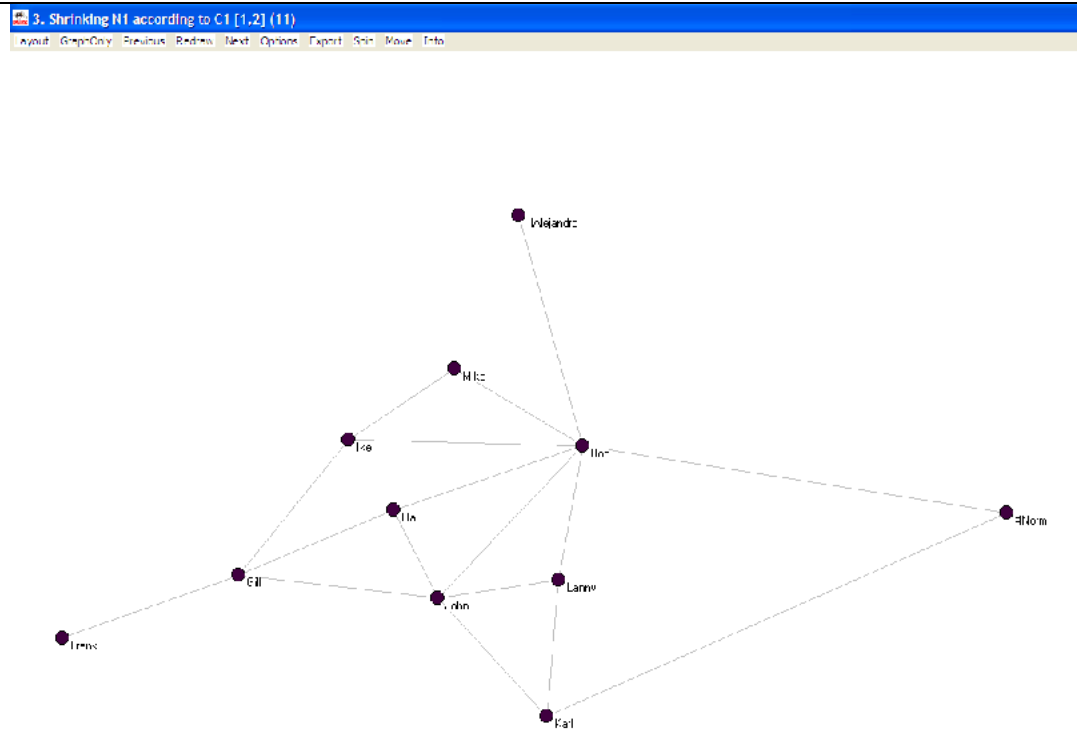
Vectors

- 8. Expected Triadic Census of N1 (16)

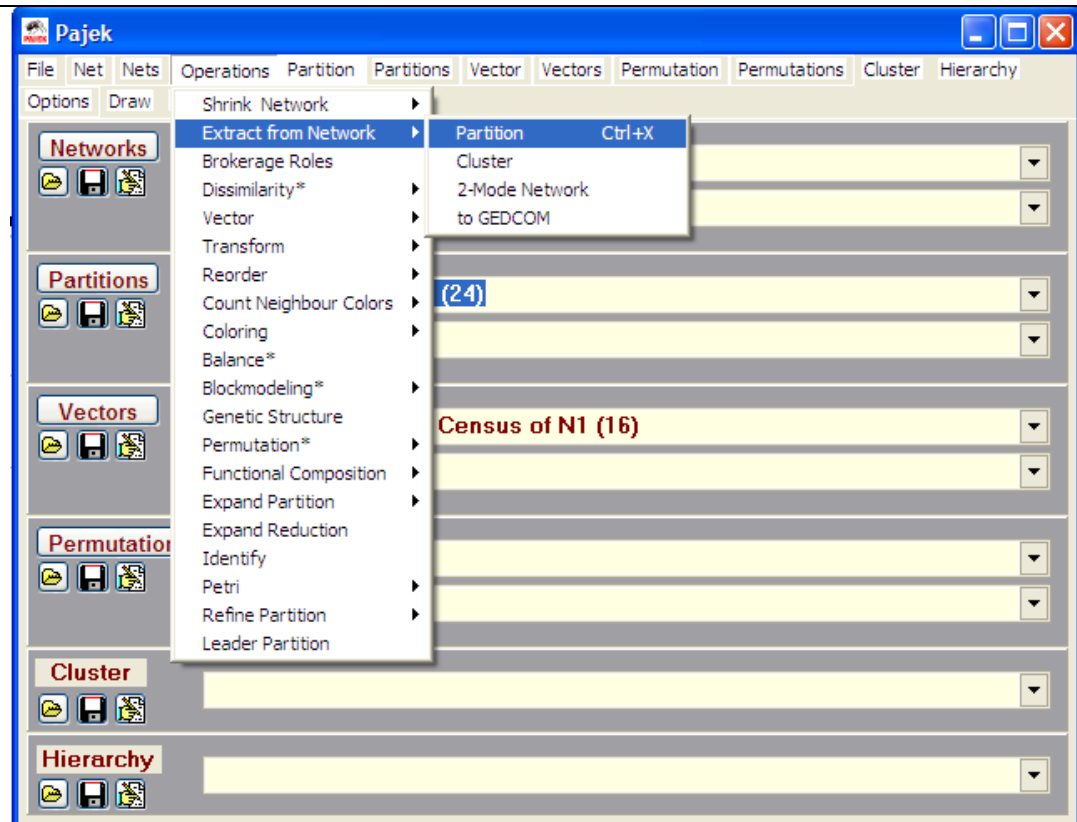
We can also perform analysis grouping together particular clusters. For instance, we can shrink the network into partition. This produces a dialogue box asking for the minimum number of ties to connect to another cluster (default of 1), before asking which partition(s) should not be shrunk. Once selected, it reduces all other clusters to a single actor (assuming the analysed cluster hold sufficient number of links to its members).



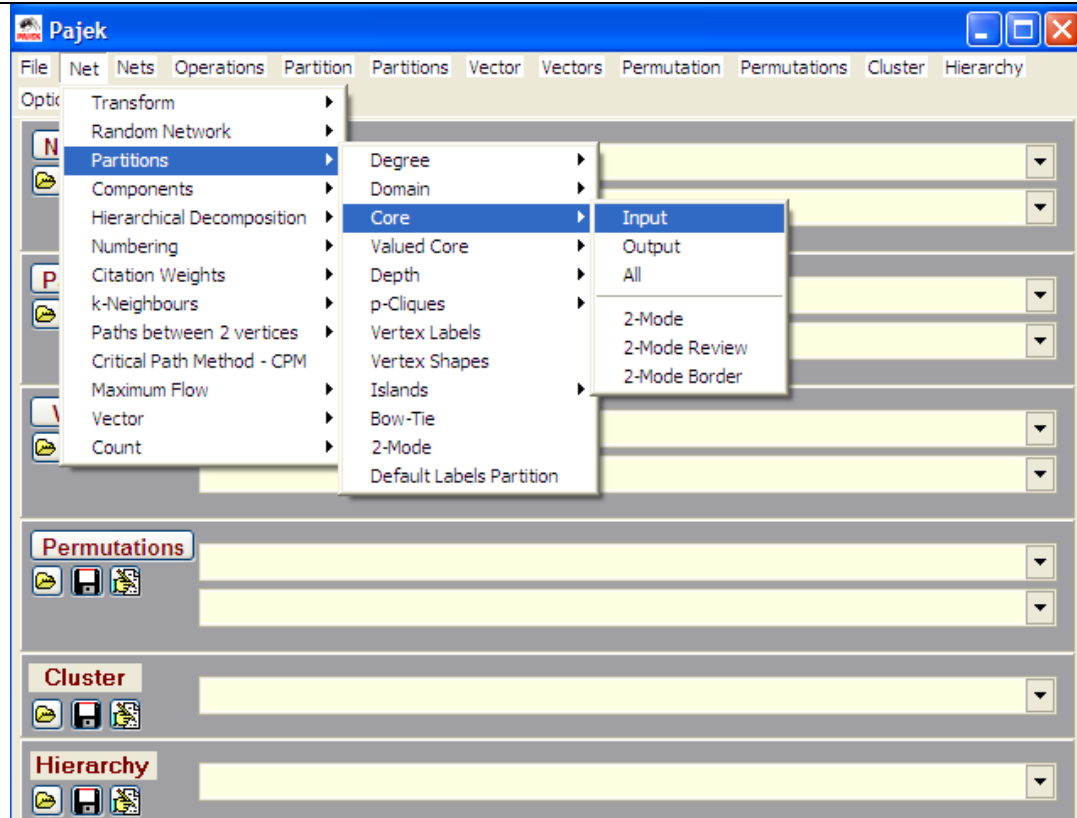
Here the network is shown, but only for the younger English cohort. The older English cohort have been shrunk into one group, and the Spanish cohort into another (labelled by #). We can now rerun any of our analysis (such as centrality methods) viewing the other clusters as sub-groups rather than their individual actors.



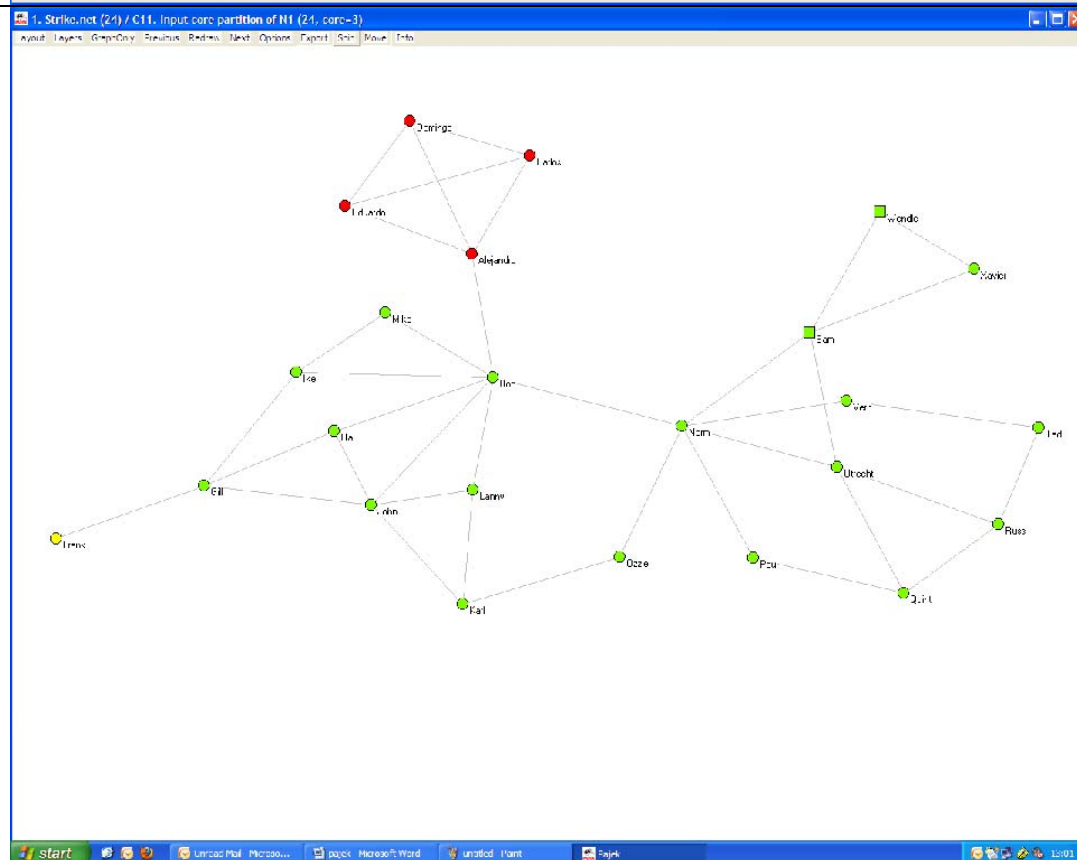
Alternatively, we can just decide to extract a particular partition from the network (by drop down menu or CTRL+X) which enables us to perform analysis irrespective of the wider network the actors are part of. (Note: this is particular useful if you have, for instance, international trade networks and you wish to look at Europe in isolation, or European countries by the other European countries and continents they trade with).



We could also look at the whole network and decide to look at the cores. This produces sub-graphs whereby each actor is connected to X actors who all have at least Y links. The core for each actor is the largest possible Y that they can be part of a core of.



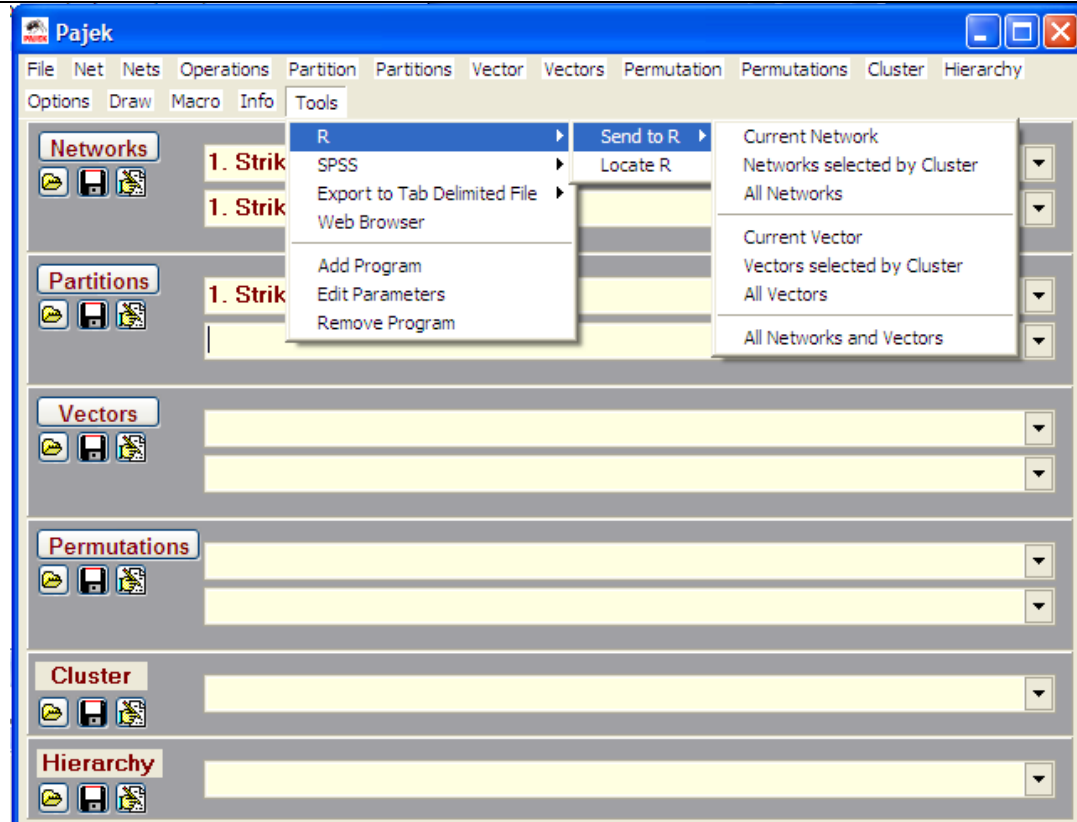
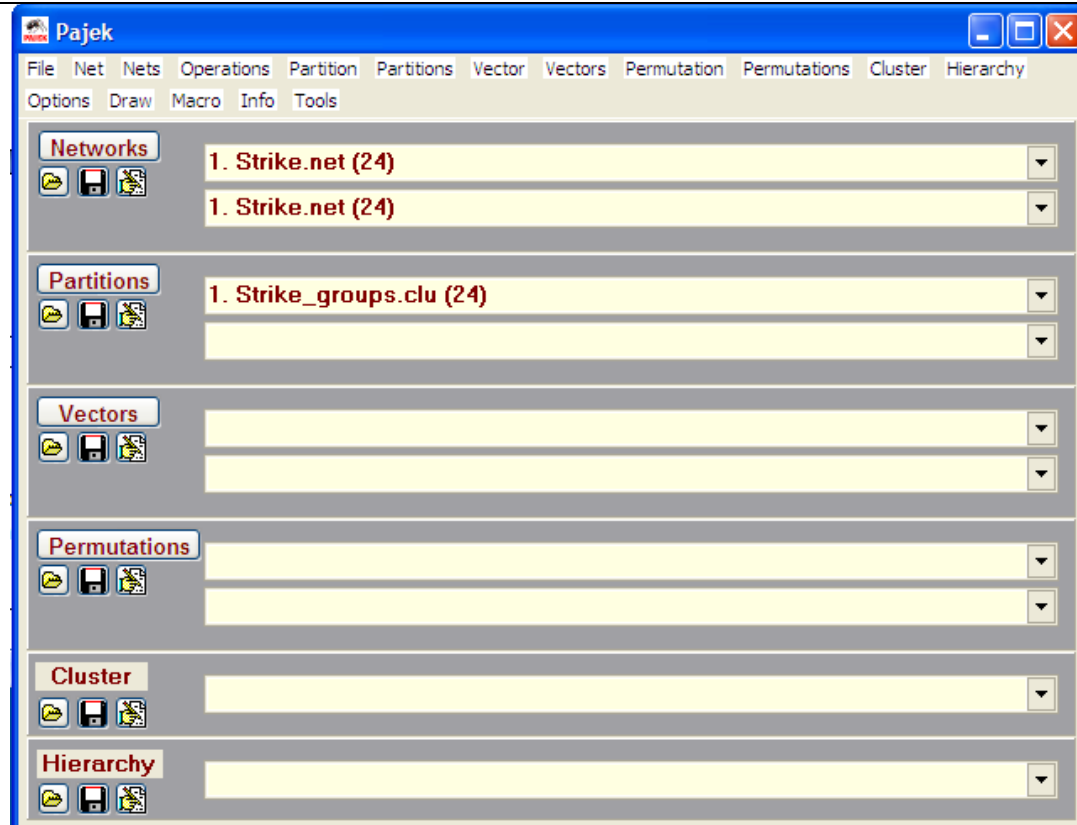
Therefore, we see that Frank can only be incorporated in a core measured by one link. The English speakers, Frank aside, all form part of the two core. The three Spanish speakers form a three-core. Whilst Karl has three connections, Ozzy does not. Therefore, removing Ozzy from the 3-core removes Karl, which in turn removes Lanny. Losing Mike loses Ike, which removes Gill and quickly the English structure falls apart. Therefore, this is ideal for spotting well-constructed elements of networks and identify cores (even if of non-central actors).



Saving data in Pajek involves using the file buttons under each tab (i.e., networks, partitions etc.), or the 'file' drop down menu. Each file you've used needs to be saved separately. You can move between them using the right-hand arrows. Alternatively, use 'file, Pajek project file, save' to save everything in one large file.

Note: Pajek keeps every file you've opened in its list which can become large if working on multiple tasks. Therefore, it is often beneficial to close and reopen the window to avoid confusion.

Pajek also has the options to export data directly to R and SPSS.



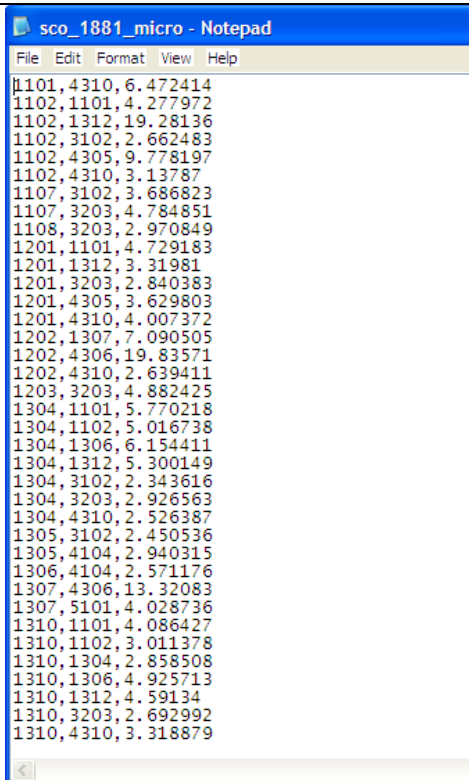
Preparing data for Pajek

The Pajek website (<http://pajek.imfm.si>) offers a range of useful datasets for exploring network theory. It also offers the Excel2Pajek and Text2Pajek tools for formatting data. Converting a dataset into Pajek format is simple.

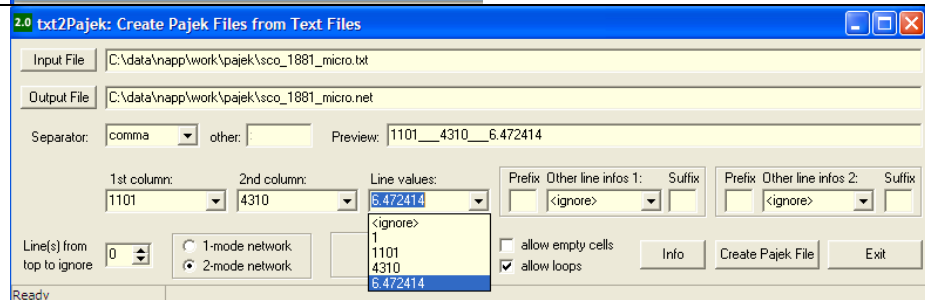
Firstly, produce a dataset which has two columns of nodes, for instance, male and female occupations within married couples. A value for strength of line can also be saved. This can be produced in any format.

	hocc	wocc	val _mi n
1.	1101	4310	5. 770497
2.	1102	1101	3. 565027
3.	1102	1312	17. 26048
4.	1102	3102	2. 306733
5.	1102	4305	8. 565514
6.	1102	4310	2. 706895
7.	1107	3102	3. 024689
8.	1107	3203	4. 27487
9.	1108	3203	2. 468718
10.	1201	1101	4. 309639
11.	1201	1312	2. 850363
12.	1201	3203	2. 701401
13.	1201	4305	3. 21621
14.	1201	4310	3. 734819
15.	1202	1307	5. 940358
16.	1202	4306	19. 30663
17.	1202	4310	2. 422515
18.	1203	3203	4. 090449
19.	1304	1101	5. 060044
20.	1304	1102	4. 12995

To use Text2Pajek, save the list as a text file. In this example it's a comma-separated version. Just the rows of data are required, with no additional information.

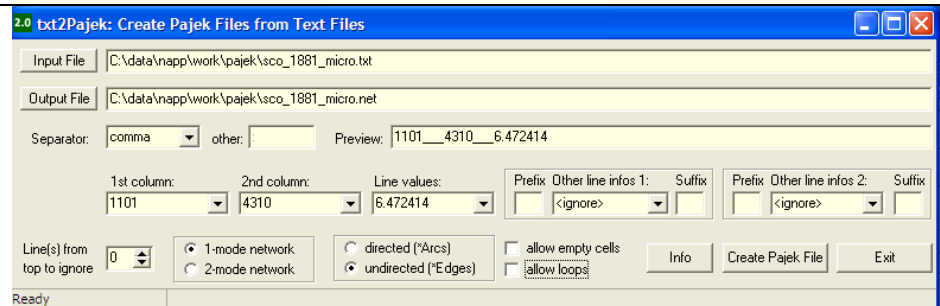


We can then open the txt2Pajek software. Firstly, we use the 'input file' button to select the file. It then defaults to save the file in the same folder with the same name but

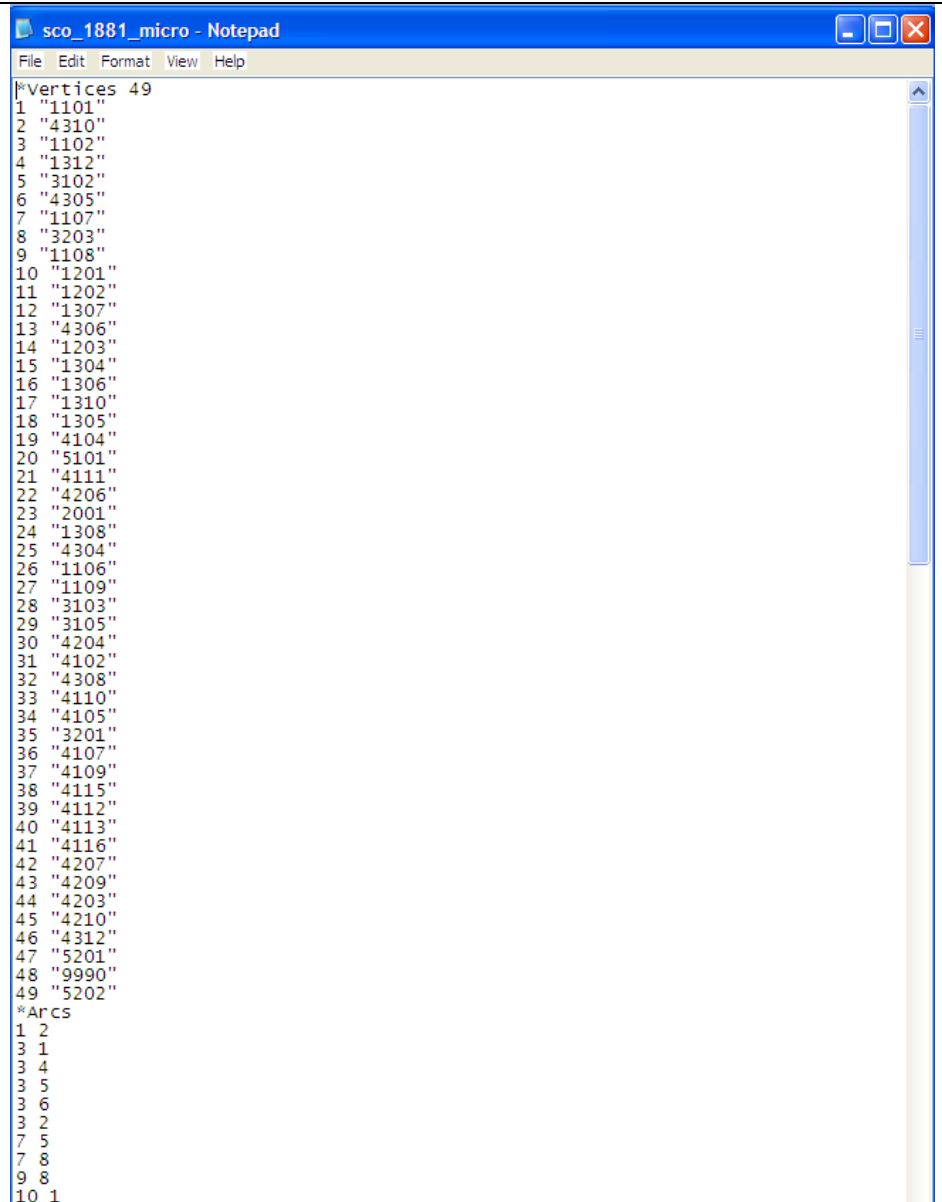


different suffix (clicking on the output name allows you to change it). We can specify the separator (comma) and select the 1st and 2nd columns from the available columns. We can include or ignore line values.

We can specify if it is a one-mode (the same actors in both columns, such as jobs) or two-mode (different actors in each column, such as employees-employers) and whether the network is directed (i.e., if ties are assumed to always be replicated). We then click on 'Create Pajek File'.



This produces a .net file which we need for Pajek. It starts off by specifying how many vertices exist, then giving a number to each label (as we saw above with the matrix). It also produces a list of the arcs/edges (links, whether directed or undirected), just showing the two numbers which are connected. In this example the labels are the occupational numbers. The data could be exported with the labels instead, which might in some cases be beneficial if they are to be shown in the graphs (obviously, with limitations on the size of what will be readable).



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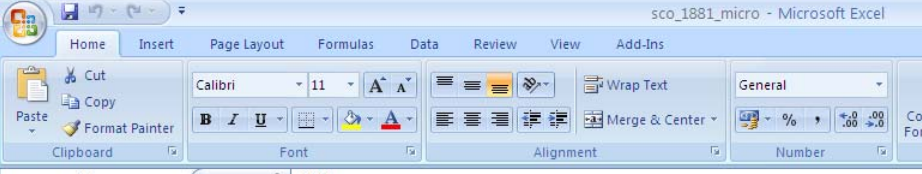
sco_1881_micro - Notepad
File Edit Format View Help

*Vertices 49
1 "1101"
2 "4310"
3 "1102"
4 "1312"
5 "3102"
6 "4305"
7 "1107"
8 "3203"
9 "1108"
10 "1201"
11 "1202"
12 "1307"
13 "4306"
14 "1203"
15 "1304"
16 "1306"
17 "1310"
18 "1305"
19 "4104"
20 "5101"
21 "4111"
22 "4206"
23 "2001"
24 "1308"
25 "4304"
26 "1106"
27 "1109"
28 "3103"
29 "3105"
30 "4204"
31 "4102"
32 "4308"
33 "4110"
34 "4105"
35 "3201"
36 "4107"
37 "4109"
38 "4115"
39 "4112"
40 "4113"
41 "4116"
42 "4207"
43 "4209"
44 "4203"
45 "4210"
46 "4312"
47 "5201"
48 "9990"
49 "5202"

*Arcs
1 2
3 1
3 4
3 5
3 6
3 2
7 5
7 8
9 8
10 1

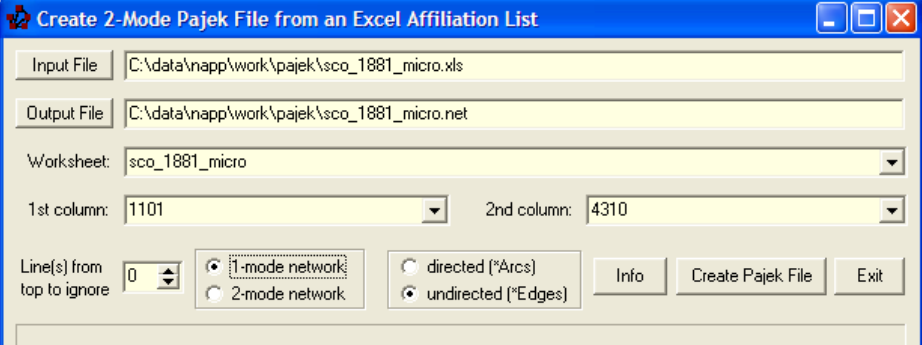
```

Using the Excel2Pajek tool is just as simple. Start off with the data saved as an Excel file. Again, how two columns showing the linkages which form part of the network. You have the option to have a third variable for the strength of the line. It does not matter if there is additional information stored in the Excel file, as you will select the columns you wish. Therefore, it would be possible to have both the occupational codes and labels in the same file, which you could export as two different networks (if required).



	A	B	C	D	E	F	G	H	I	J	K	L
1	1101	4310	6.472414									
2	1102	1101	4.277972									
3	1102	1312	19.28136									
4	1102	3102	2.662483									
5	1102	4305	9.778197									
6	1102	4310	3.13787									
7	1107	3102	3.686823									
8	1107	3203	4.784851									
9	1108	3203	2.970849									
10	1201	1101	4.729183									
11	1201	1312	3.31981									
12	1201	3203	2.840383									
13	1201	4305	3.629803									
14	1201	4310	4.007372									
15	1202	1307	7.090505									
16	1202	4306	19.83571									
17	1202	4310	2.639411									
18	1203	3203	4.882425									
19	1304	1101	5.770218									
20	1304	1102	5.016738									
21	1304	1306	6.154411									
22	1304	1312	5.300149									
23	1304	3102	2.343616									
24	1304	3203	2.926563									
25	1304	4310	2.526387									
26	1305	3102	2.450536									
27	1305	4104	2.940315									
28	1306	4104	2.571176									
29	1307	4306	13.32083									
30	1307	5101	4.028736									
31	1310	1101	4.086427									
32	1310	1102	3.011378									
33	1310	1304	2.858508									

Again, we can select the input file and it defaults in the same way. We can decide which worksheet we want to use, and which columns are important. We can save as a 1- or 2-mode network, and decide to ignore the top line (if it is merely column labels). Click on 'Create Pajek file' and the file will be ready to be opened in Pajek.



Create 2-Mode Pajek File from an Excel Affiliation List

Input File: C:\data\ncapp\work\pajek\sco_1881_micro.xls

Output File: C:\data\ncapp\work\pajek\sco_1881_micro.net

Worksheet: sco_1881_micro

1st column: 1101 2nd column: 4310

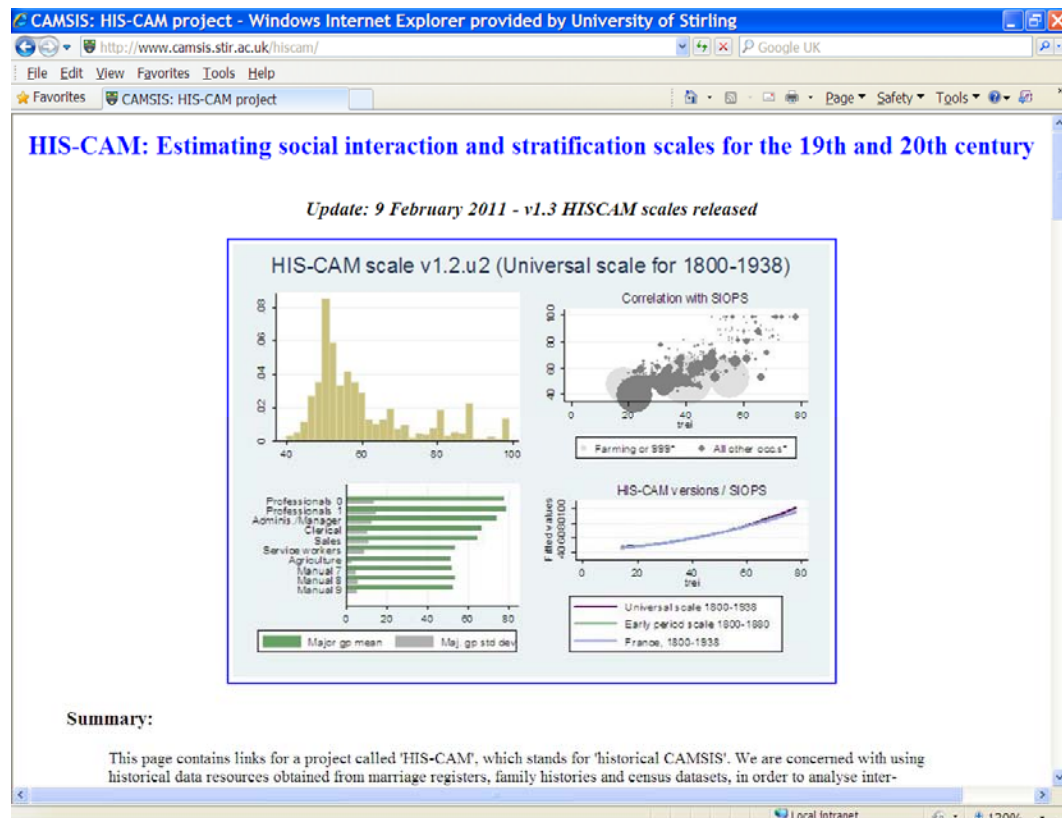
Line(s) from top to ignore: 0

☒ 1-mode network
 ☐ directed (*Arcs)
 ☒ undirected (*Edges)
 ☐ 2-mode network

Info Create Pajek File Exit

Lab 2: Historical data on occupations and its analysis using SNA and SID approaches

This lab features some examples focussed upon using large scale occupational data on social connections. Examples include operationalising key measures using occupational records, and worked examples of SID and SNA techniques. The Stata and R exercises can be found in the accompanying command files, whilst the Pajek instructions are given below (the Pajek exercise begins with a brief use of Stata, though that can be skipped if necessary).



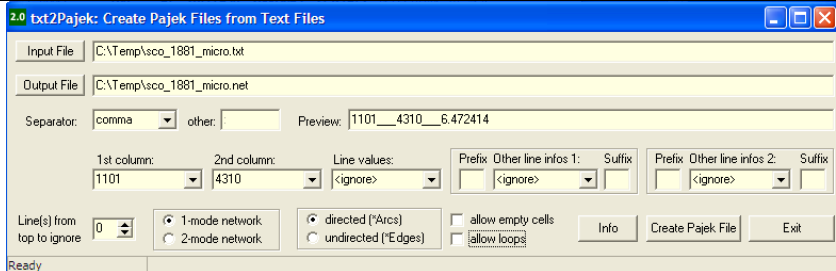
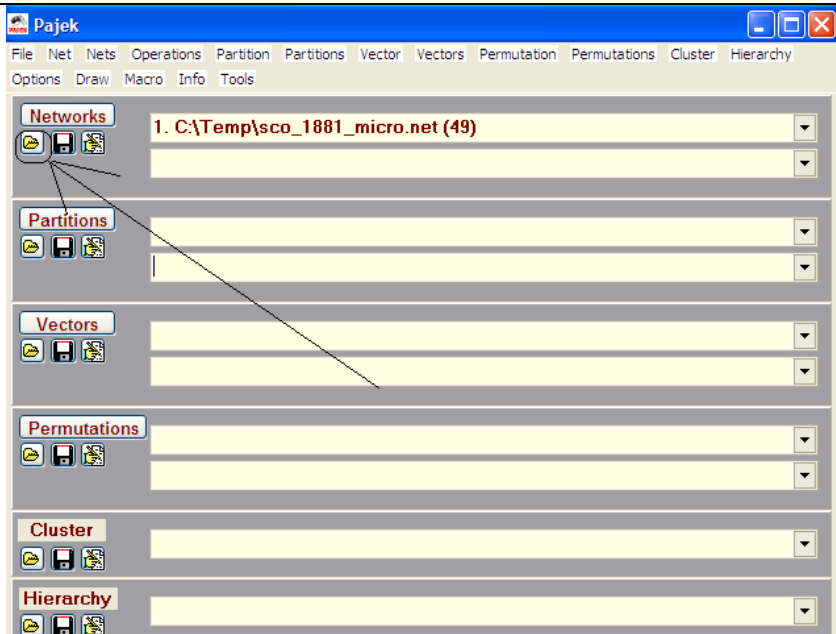
Screenshot of the HIS-CAM website, a useful resource for data on occupations

Lab 2: Pajek exercises

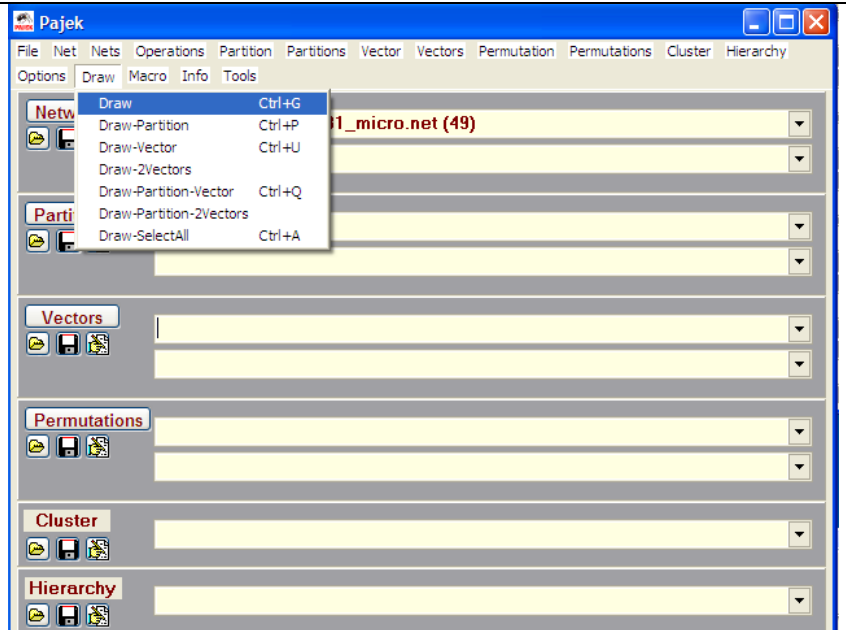
Using SNA to analyse occupational stratification in the past

In this lab we explore ways we can use social network analysis to understand more about occupational stratification. Research in this area always benefits by consideration of the occupational structure and the national context of vocations. Research using social connections, similarly, can benefit from understanding underlying patterns of social interactions. This analysis involves processes from both Stata and Pajek. Stata is utilised to generate the data which can be utilised by network analysis, which is processed within Pajek.

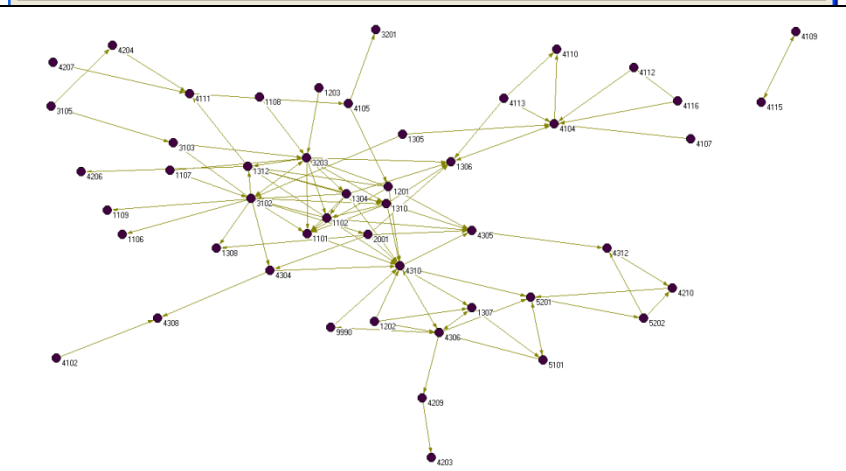
Our first step is to find a dataset consisting of pairs of occupations. For this lab, a dataset has been created consisting of within household combinations of microclasses. Only male ties with an age-gap of 16 years, aged between 16 and 75, are included. Combinations within the same microclass have been excluded.	<pre>. use \$path9\scot81.dta, clear</pre> <pre>. codebook, compact</pre> <table><thead><tr><th>Variable</th><th>Obs</th><th>Unique</th><th>Mean</th><th>Min</th><th>Max</th><th>Label</th></tr></thead><tbody><tr><td>serial</td><td>258740</td><td>136853</td><td>403449.1</td><td>2</td><td>784262</td><td>Household index number</td></tr><tr><td>pernum1</td><td>258740</td><td>67</td><td>3.802242</td><td>1</td><td>87</td><td>Person index within ...</td></tr><tr><td>age1</td><td>258740</td><td>48</td><td>52.48311</td><td>28</td><td>75</td><td>Age</td></tr><tr><td>sex1</td><td>258740</td><td>1</td><td>1</td><td>1</td><td>1</td><td>Sex</td></tr><tr><td>occgb1</td><td>258740</td><td>384</td><td>210.265</td><td>1</td><td>413</td><td>Occupation, Britain ...</td></tr><tr><td>hocc</td><td>258740</td><td>64</td><td>4164.652</td><td>1101</td><td>9990</td><td></td></tr><tr><td>pernum</td><td>258740</td><td>76</td><td>6.289681</td><td>1</td><td>110</td><td>Person index within ...</td></tr><tr><td>age</td><td>258740</td><td>48</td><td>21.85543</td><td>12</td><td>59</td><td>Age</td></tr><tr><td>sex</td><td>258740</td><td>1</td><td>1</td><td>1</td><td>1</td><td>Sex</td></tr><tr><td>occgb</td><td>258740</td><td>382</td><td>199.2149</td><td>1</td><td>413</td><td>Occupation, Britain ...</td></tr><tr><td>wocc</td><td>258740</td><td>64</td><td>4061.158</td><td>1101</td><td>9990</td><td></td></tr><tr><td>pershh</td><td>258740</td><td>47</td><td>7.602937</td><td>2</td><td>49</td><td></td></tr><tr><td>pairshh</td><td>258740</td><td>47</td><td>62.56172</td><td>1</td><td>1176</td><td></td></tr><tr><td>aged</td><td>258740</td><td>48</td><td>30.62768</td><td>16</td><td>63</td><td></td></tr></tbody></table>	Variable	Obs	Unique	Mean	Min	Max	Label	serial	258740	136853	403449.1	2	784262	Household index number	pernum1	258740	67	3.802242	1	87	Person index within ...	age1	258740	48	52.48311	28	75	Age	sex1	258740	1	1	1	1	Sex	occgb1	258740	384	210.265	1	413	Occupation, Britain ...	hocc	258740	64	4164.652	1101	9990		pernum	258740	76	6.289681	1	110	Person index within ...	age	258740	48	21.85543	12	59	Age	sex	258740	1	1	1	1	Sex	occgb	258740	382	199.2149	1	413	Occupation, Britain ...	wocc	258740	64	4061.158	1101	9990		pershh	258740	47	7.602937	2	49		pairshh	258740	47	62.56172	1	1176		aged	258740	48	30.62768	16	63																													
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Next we run a syntax file which automatically generates some information on pairs of occupations.	<pre>do http://www.camsis.stir.ac.uk/sonocs/do/pajek.do</pre>																																																																																																																																					
This provides 16 variables. hocc is the older cohort occupation and wocc the younger cohort. freq is the number of connections for each combination. ewocc is the expected number of ties if the data was random. val_min is the value of over-representation (taking standard errors into account). These are the most important variables.	<table><thead><tr><th>Variable</th><th>Obs</th><th>Unique</th><th>Mean</th><th>Min</th><th>Max</th><th>Label</th></tr></thead><tbody><tr><td>hocc</td><td>3131</td><td>64</td><td>3458.181</td><td>1101</td><td>9990</td><td></td></tr><tr><td>wocc</td><td>3131</td><td>64</td><td>3401.974</td><td>1101</td><td>9990</td><td></td></tr><tr><td>freq</td><td>3131</td><td>395</td><td>82.63813</td><td>1</td><td>25594</td><td>(count) freq</td></tr><tr><td>tot</td><td>3131</td><td>1</td><td>258740</td><td>258740</td><td>258740</td><td>total number in ...</td></tr><tr><td>nhocc</td><td>3131</td><td>64</td><td>4849.383</td><td>4</td><td>39522</td><td>total number of ...</td></tr><tr><td>nwocc</td><td>3131</td><td>64</td><td>4828.742</td><td>5</td><td>33491</td><td>total number of ...</td></tr><tr><td>phocc</td><td>3131</td><td>64</td><td>.0187423</td><td>.0000155</td><td>.1527479</td><td>percentage of me...</td></tr><tr><td>pwocc</td><td>3131</td><td>64</td><td>.0186625</td><td>.0000193</td><td>.1294388</td><td>percentage of wo...</td></tr><tr><td>ewocc</td><td>3131</td><td>3129</td><td>79.17888</td><td>.0091598</td><td>5115.681</td><td>expected number ...</td></tr><tr><td>prop</td><td>3131</td><td>395</td><td>.0003194</td><td>3.86e-06</td><td>.0989178</td><td></td></tr><tr><td>staner</td><td>3131</td><td>395</td><td>.0000229</td><td>3.86e-06</td><td>.0005869</td><td>Standard error f...</td></tr><tr><td>pro_obs</td><td>3131</td><td>395</td><td>.0003194</td><td>3.86e-06</td><td>.0989178</td><td>Observed proport...</td></tr><tr><td>pro_exp</td><td>3131</td><td>3129</td><td>.000306</td><td>3.54e-08</td><td>.0197715</td><td>Expected proport...</td></tr><tr><td>pro_min</td><td>3131</td><td>395</td><td>.0002965</td><td>7.28e-12</td><td>.0983309</td><td>Lower confidence...</td></tr><tr><td>pro_max</td><td>3131</td><td>395</td><td>.0003423</td><td>7.73e-06</td><td>.0995048</td><td>Higher confidence...</td></tr><tr><td>value</td><td>3131</td><td>3116</td><td>1.592005</td><td>.0257491</td><td>109.173</td><td>Observed value o...</td></tr><tr><td>val_min</td><td>3131</td><td>3131</td><td>.9458267</td><td>4.85e-08</td><td>24.38851</td><td>Value of represe...</td></tr><tr><td>val_max</td><td>3131</td><td>3131</td><td>2.238184</td><td>.0503272</td><td>218.3458</td><td>Value of represe...</td></tr></tbody></table>	Variable	Obs	Unique	Mean	Min	Max	Label	hocc	3131	64	3458.181	1101	9990		wocc	3131	64	3401.974	1101	9990		freq	3131	395	82.63813	1	25594	(count) freq	tot	3131	1	258740	258740	258740	total number in ...	nhocc	3131	64	4849.383	4	39522	total number of ...	nwocc	3131	64	4828.742	5	33491	total number of ...	phocc	3131	64	.0187423	.0000155	.1527479	percentage of me...	pwocc	3131	64	.0186625	.0000193	.1294388	percentage of wo...	ewocc	3131	3129	79.17888	.0091598	5115.681	expected number ...	prop	3131	395	.0003194	3.86e-06	.0989178		staner	3131	395	.0000229	3.86e-06	.0005869	Standard error f...	pro_obs	3131	395	.0003194	3.86e-06	.0989178	Observed proport...	pro_exp	3131	3129	.000306	3.54e-08	.0197715	Expected proport...	pro_min	3131	395	.0002965	7.28e-12	.0983309	Lower confidence...	pro_max	3131	395	.0003423	7.73e-06	.0995048	Higher confidence...	value	3131	3116	1.592005	.0257491	109.173	Observed value o...	val_min	3131	3131	.9458267	4.85e-08	24.38851	Value of represe...	val_max	3131	3131	2.238184	.0503272	218.3458	Value of represe...
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Next we need some criteria for which ties are needed. We are interested in cases occurring at least once in every 10,000 cases (therefore, frequency of at least 28) and which occur at least twice as often as we would expect.	<pre>. keep if freq>=27</pre> <p>(1942 observations deleted)</p> <pre>. keep if val_min>=2</pre> <p>(1081 observations deleted)</p>																																																																																																																																					

<p>This produces 108 combinations which are both frequently constructed and occur more commonly than expected. As the val_min shows, these occur up to 20 times more than we would anticipate. There are 41 different microclasses for the older cohort and 37 for the younger cohort.</p>	<table><thead><tr><th>Variable</th><th>Obs</th><th>Unique</th><th>Mean</th><th>Min</th><th>Max</th><th>Label</th></tr></thead><tbody><tr><td>hocc</td><td>108</td><td>41</td><td>2992.852</td><td>1101</td><td>9990</td><td></td></tr><tr><td>wocc</td><td>108</td><td>37</td><td>3223.398</td><td>1101</td><td>9990</td><td></td></tr><tr><td>freq</td><td>108</td><td>85</td><td>466.1852</td><td>29</td><td>25594</td><td>(count) freq</td></tr><tr><td>tot</td><td>108</td><td>1</td><td>258740</td><td>258740</td><td>258740</td><td>total number in s...</td></tr><tr><td>nhocc</td><td>108</td><td>41</td><td>4435.796</td><td>146</td><td>39522</td><td>total number of m...</td></tr><tr><td>nwocc</td><td>108</td><td>37</td><td>4578.713</td><td>281</td><td>33491</td><td>total number of f...</td></tr><tr><td>phocc</td><td>108</td><td>41</td><td>.0171438</td><td>.0005643</td><td>.1527479</td><td>percentage of men...</td></tr><tr><td>pwocc</td><td>108</td><td>37</td><td>.0176962</td><td>.001086</td><td>.1294388</td><td>percentage of wom...</td></tr><tr><td>ewocc</td><td>108</td><td>108</td><td>114.8665</td><td>3.142614</td><td>5115.681</td><td>expected number o...</td></tr><tr><td>prop</td><td>108</td><td>85</td><td>.0018018</td><td>.0001121</td><td>.0989178</td><td></td></tr><tr><td>staner</td><td>108</td><td>85</td><td>.0000512</td><td>.0000208</td><td>.0005869</td><td>Standard error fo...</td></tr><tr><td>pro_obs</td><td>108</td><td>85</td><td>.0018018</td><td>.0001121</td><td>.0989178</td><td>Observed proporti...</td></tr><tr><td>pro_exp</td><td>108</td><td>108</td><td>.0004439</td><td>.0000121</td><td>.0197715</td><td>Expected proporti...</td></tr><tr><td>pro_min</td><td>108</td><td>85</td><td>.0017506</td><td>.0000913</td><td>.0983309</td><td>Lower confidence ...</td></tr><tr><td>pro_max</td><td>108</td><td>85</td><td>.0018529</td><td>.0001329</td><td>.0995048</td><td>Higher confidence...</td></tr><tr><td>value</td><td>108</td><td>108</td><td>4.659531</td><td>2.122135</td><td>19.83571</td><td>Observed value of...</td></tr><tr><td>val_min</td><td>108</td><td>108</td><td>4.150986</td><td>2.011253</td><td>19.30663</td><td>Value of represen...</td></tr><tr><td>val_max</td><td>108</td><td>108</td><td>5.168075</td><td>2.22231</td><td>21.30224</td><td>Value of represen...</td></tr></tbody></table>	Variable	Obs	Unique	Mean	Min	Max	Label	hocc	108	41	2992.852	1101	9990		wocc	108	37	3223.398	1101	9990		freq	108	85	466.1852	29	25594	(count) freq	tot	108	1	258740	258740	258740	total number in s...	nhocc	108	41	4435.796	146	39522	total number of m...	nwocc	108	37	4578.713	281	33491	total number of f...	phocc	108	41	.0171438	.0005643	.1527479	percentage of men...	pwocc	108	37	.0176962	.001086	.1294388	percentage of wom...	ewocc	108	108	114.8665	3.142614	5115.681	expected number o...	prop	108	85	.0018018	.0001121	.0989178		staner	108	85	.0000512	.0000208	.0005869	Standard error fo...	pro_obs	108	85	.0018018	.0001121	.0989178	Observed proporti...	pro_exp	108	108	.0004439	.0000121	.0197715	Expected proporti...	pro_min	108	85	.0017506	.0000913	.0983309	Lower confidence ...	pro_max	108	85	.0018529	.0001329	.0995048	Higher confidence...	value	108	108	4.659531	2.122135	19.83571	Observed value of...	val_min	108	108	4.150986	2.011253	19.30663	Value of represen...	val_max	108	108	5.168075	2.22231	21.30224	Value of represen...
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<p>We can then export the data as a comma-separated text file, showing hocc wocc and val_min</p>	<pre>outsheet hocc wocc val_min using /// "\$path9\sco_1881_micro.txt", /// comma nonames nolabel replace</pre>																																																																																																																																					
<p>We can then use txt2Pajek to convert the data into a Pajek file. We select the input file, which then defaults to saving the output file to the same folder. We specify it is comma separated and select the two microclass labels. We then assert it is a one-mode directed network before clicking on 'create Pajek file'.</p>																																																																																																																																						
<p>We can then open the file in Pajek, using either the file button below network, or the drop-down menu.</p>																																																																																																																																						

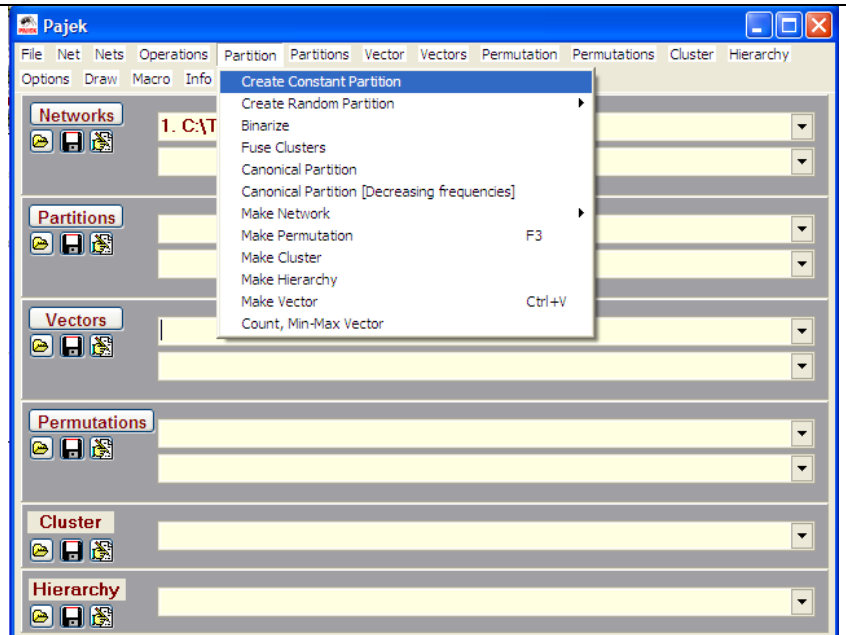
We can then select 'draw' or CTRL+G to visual the network.



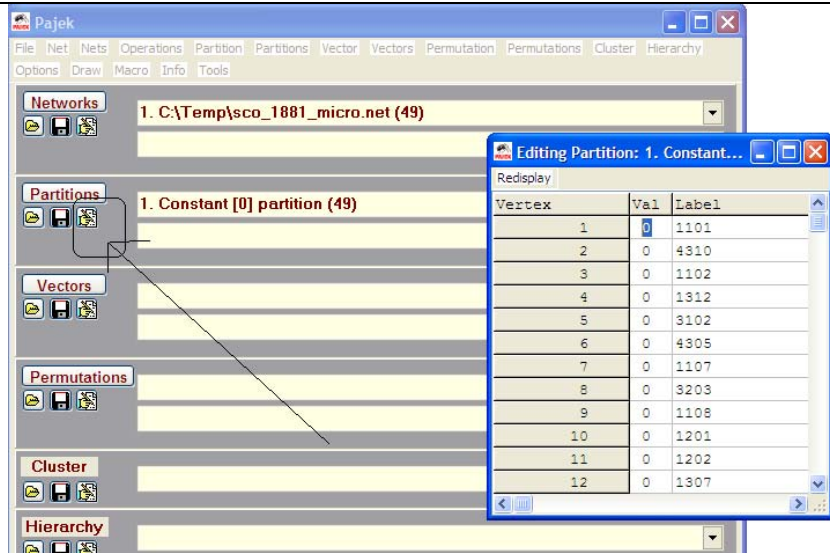
This provides a diagram of the network. The nodes are distributed where the software believes they are best placed. Sometimes it can be a little strange, so pressing CTRL+K will regenerate the network. Clicking on an individual node enables it to be moved around.



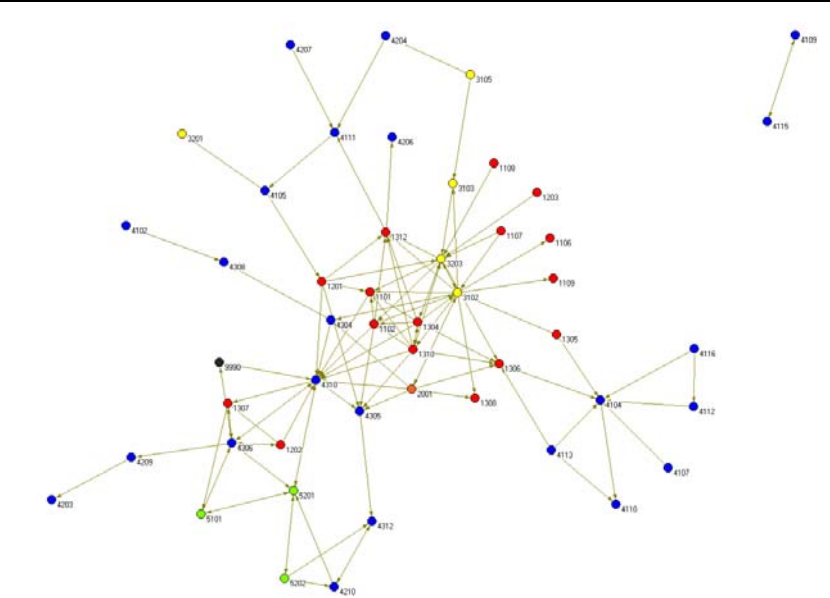
We can change the colour of the nodes to see differences more clearly. If we close the graph window we can click on 'partition', 'create constant partition' to create a blank partition. The first dialogue box regards the size of the partition (which defaults to the number of nodes, as it needs to be). The second dialogue box asks for the constant term to be used (default is 0).



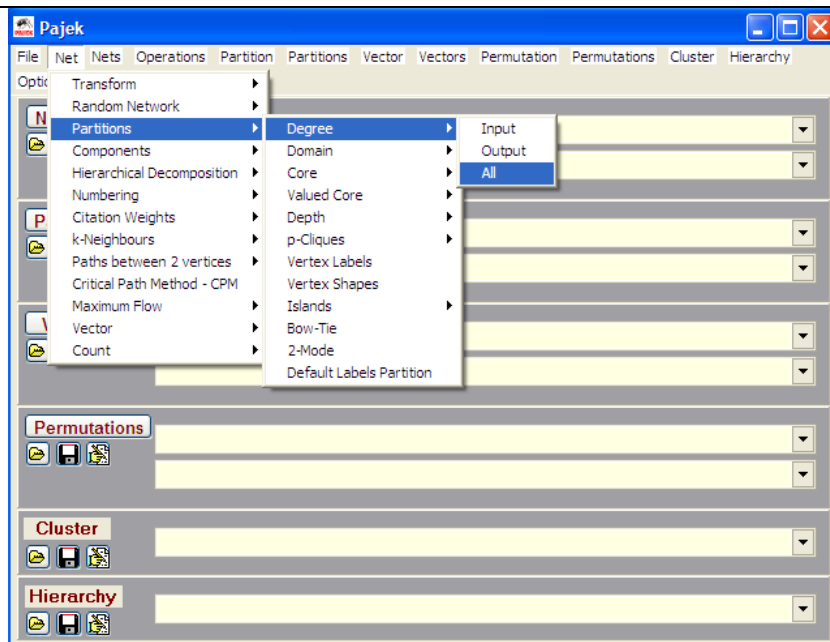
If we click on the last of the three buttons we can edit the partition. This enables us to group the nodes. In this example, we will place the first number of each label as the partition (which enables us to code the nodes by macroclass).



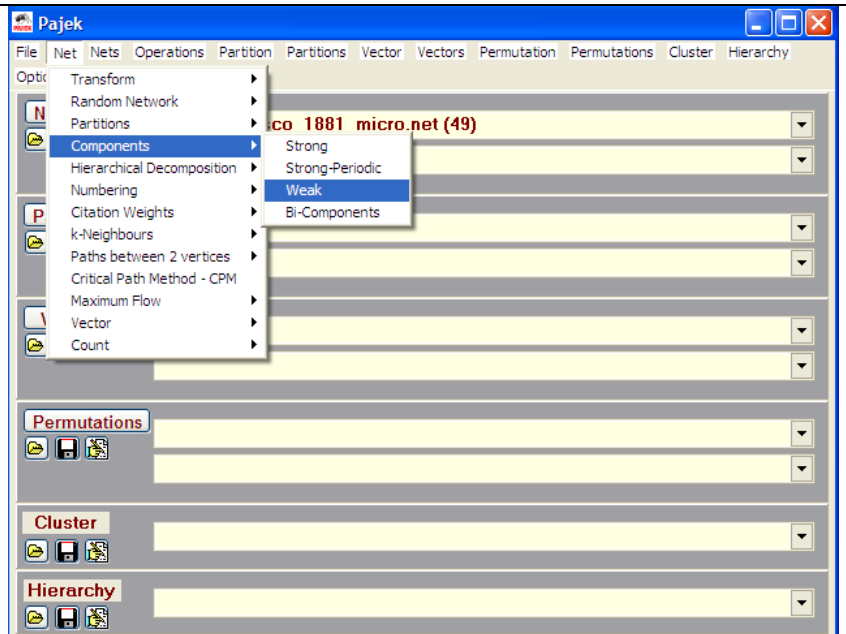
This enables us to see the differences between the macroclasses more clearly. The colours of the partitions can be altered using the 'options', 'colors', 'partition colors', 'of vertices' drop down menu within the draw window. Therefore, the colours might not be the same as displayed.



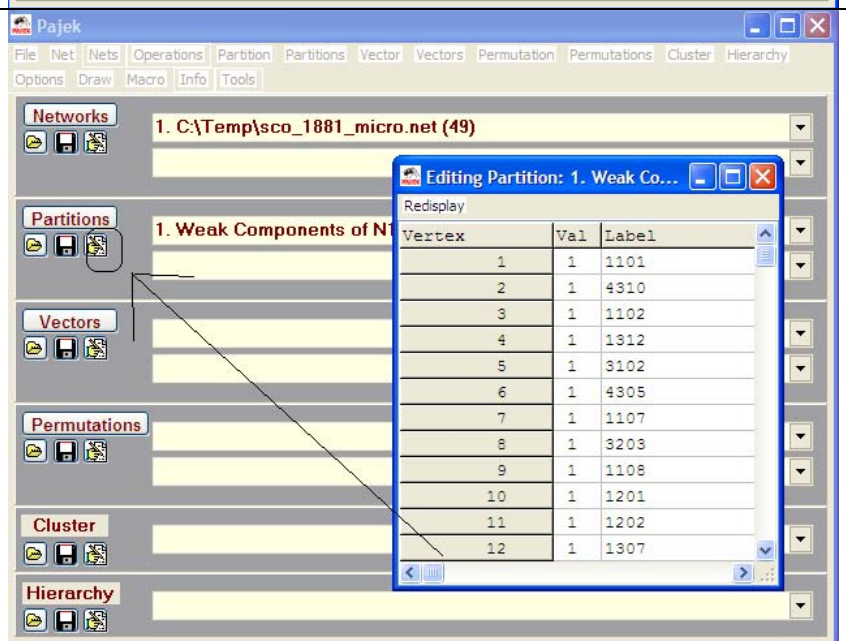
We can also find out statistics about the network. Using 'net', 'partition', 'degree', 'all' we can see the degree centrality of the network (as well as getting the degree for each occupation).



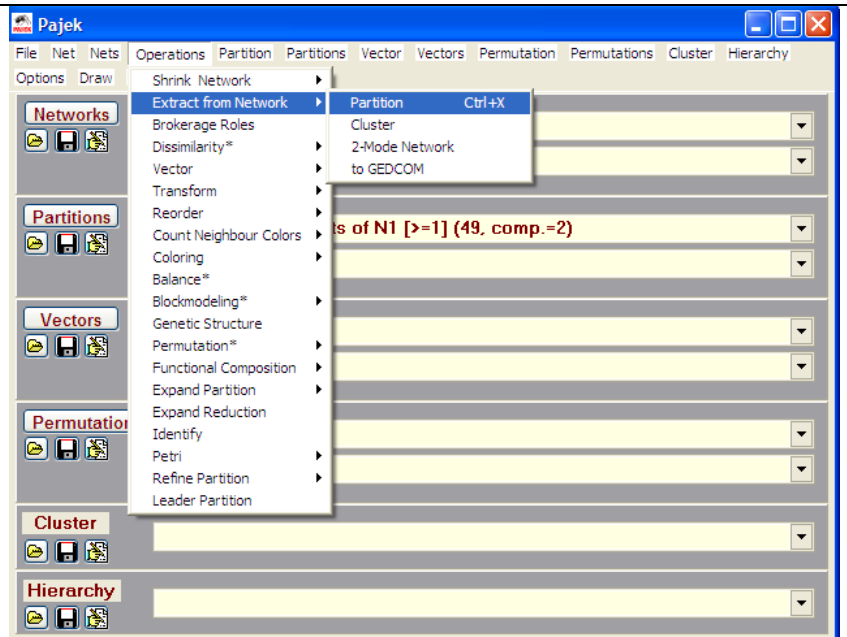
We can see how many components exist, using 'net', 'components', 'weak'. From the dialogue box we select '1' in this example (this is just about the strength of ties needed to form part of the community. We are interested here in nodes which have at least one tie to a member of the sub-population which are connected).



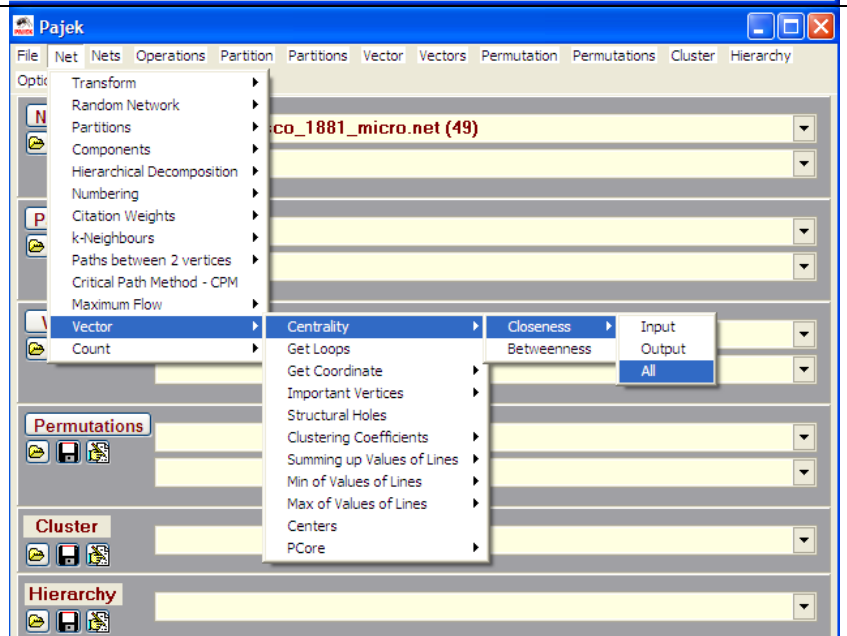
We can then reduce our analysis to just the largest component. This allows us to perform some further analysis. Firstly, we click on the partition named 'components' to see which is largest.



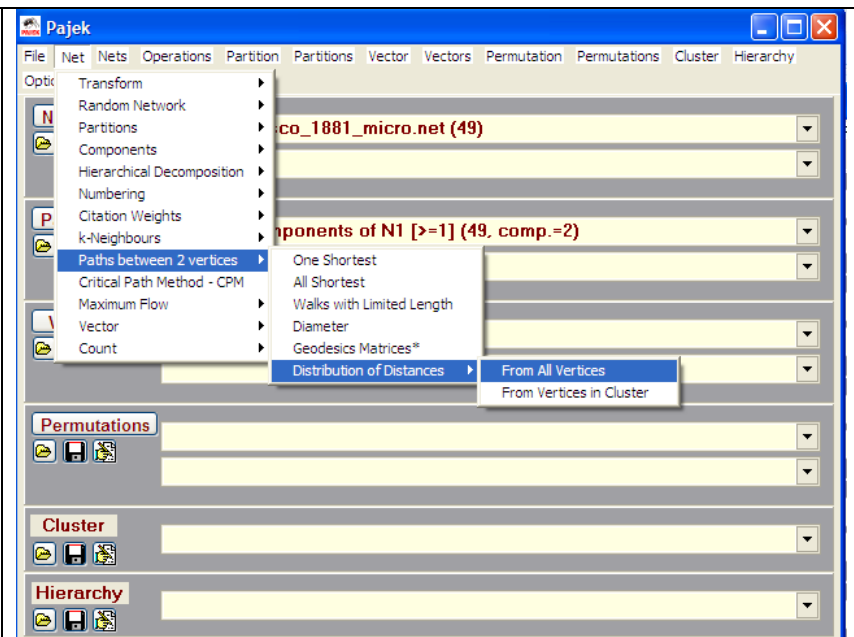
We then click on 'operations', 'extract from network', 'partition' (or CTRL+X) and decide to keep only '1'. This removes those cases which do not connect to the main component.



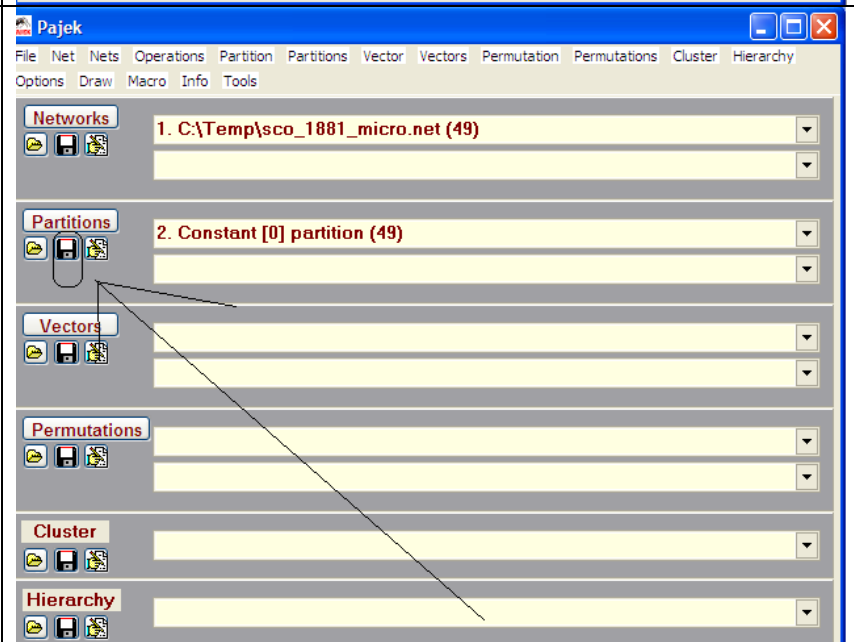
We can use 'net', 'vector', 'centrality', 'closeness', 'all' to see the closeness centrality.



We can use 'net', 'Paths between 2 vertices', 'distribution of distances', 'from all vertices' to see the average distance between the nodes.



We can then save the partition we created to ensure we have the data next time we analyse it.



We can analyse more than just microclasses. In this second example we will look at Canada 1891 data, looking at religion. Again, a dataset has been created.

```
. use $path9\canada91.dta, clear
. codebook, compact
```

Vari able	Obs	Uni que	Mean	Min	Max	Label
seri al	21699	7675	51919.54	12	76715	Househol d i ndex ...
age1	21699	48	51.6469	28	75	Age
sex1	21699	1	1	1	1	Sex
hi sco1	21699	312	60062.34	1110	98990	
mi crocl ass1	21699	71	3658.492	1101	9990	
age	21699	46	23.66455	12	57	Age
sex	21699	1	1	1	1	Sex
hi sco	21699	337	62402.54	1110	98990	
mi crocl ass	21699	70	3820.816	1101	9990	
rel i gi on1	21699	66	3595.099	1100	9997	Rel i gi on, fi rst ...
rel i gi on	21699	61	3609.406	1100	9997	Rel i gi on, fi rst ...
hi scam1	21699	68	58.69302	45.69142	99	(mean) hi scam
hi scam	21699	68	57.72059	45.69142	99	(mean) hi scam

From the religion data, we can dichotomise the microclasses by placing a 1 in front of the microclasses for Catholics.	<pre>. capture drop cath* . gen cath=religion==1100 . gen cath1=religion1==1100 . capture drop hocc . gen hocc=microclass . replace hocc=microclass+10000 if cath==1 (30927 real changes made) . capture drop wocc . gen wocc=microclass1 . replace wocc=microclass1+10000 if cath1==1 (29994 real changes made)</pre>																																																																																																																																					
This enables us to run the analysis as above.	do http://www.camsis.stir.ac.uk/sonocs/do/pajek.do																																																																																																																																					
This provides us with over 4,000 combinations, although many of these are of little relevance to us.	<table><tr><th>Variable</th><th>Obs</th><th>Unique</th><th>Mean</th><th>Min</th><th>Max</th><th>Label</th></tr><tr><td>hocc</td><td>4471</td><td>137</td><td>7575.914</td><td>1101</td><td>19990</td><td></td></tr><tr><td>wocc</td><td>4471</td><td>138</td><td>7164.323</td><td>1101</td><td>19990</td><td></td></tr><tr><td>freq</td><td>4471</td><td>77</td><td>4.66115</td><td>1</td><td>1111</td><td>(count) freq</td></tr><tr><td>tot</td><td>4471</td><td>1</td><td>20840</td><td>20840</td><td>20840</td><td>total number in ...</td></tr><tr><td>nhocc</td><td>4471</td><td>107</td><td>265.926</td><td>1</td><td>1566</td><td>total number of ...</td></tr><tr><td>nwocc</td><td>4471</td><td>102</td><td>309.9512</td><td>1</td><td>2662</td><td>total number of ...</td></tr><tr><td>phocc</td><td>4471</td><td>107</td><td>.0127604</td><td>.000048</td><td>.075144</td><td>percentage of me...</td></tr><tr><td>pwocc</td><td>4471</td><td>102</td><td>.0148729</td><td>.000048</td><td>.1277351</td><td>percentage of wo...</td></tr><tr><td>ewocc</td><td>4471</td><td>3832</td><td>3.305061</td><td>.0004319</td><td>200.0332</td><td>expected number ...</td></tr><tr><td>prop</td><td>4471</td><td>77</td><td>.0002237</td><td>.000048</td><td>.0533109</td><td></td></tr><tr><td>staner</td><td>4471</td><td>77</td><td>.0000831</td><td>.000048</td><td>.0015562</td><td>Standard error f...</td></tr><tr><td>pro_obs</td><td>4471</td><td>77</td><td>.0002237</td><td>.000048</td><td>.0533109</td><td>Observed proport...</td></tr><tr><td>pro_exp</td><td>4471</td><td>3818</td><td>.0001586</td><td>2.07e-08</td><td>.0095985</td><td>Expected proport...</td></tr><tr><td>pro_min</td><td>4471</td><td>77</td><td>.0001406</td><td>1.15e-09</td><td>.0517548</td><td>Lower confidence...</td></tr><tr><td>pro_max</td><td>4471</td><td>77</td><td>.0003067</td><td>.000096</td><td>.0548671</td><td>Higher confidenc...</td></tr><tr><td>value</td><td>4471</td><td>3968</td><td>5.717776</td><td>.0432525</td><td>2315.556</td><td>Observed value o...</td></tr><tr><td>val_min</td><td>4471</td><td>4217</td><td>1.115635</td><td>1.04e-06</td><td>81.03288</td><td>Value of represe...</td></tr><tr><td>val_max</td><td>4471</td><td>4214</td><td>10.31992</td><td>.0691135</td><td>4631.055</td><td>Value of represe...</td></tr></table>	Variable	Obs	Unique	Mean	Min	Max	Label	hocc	4471	137	7575.914	1101	19990		wocc	4471	138	7164.323	1101	19990		freq	4471	77	4.66115	1	1111	(count) freq	tot	4471	1	20840	20840	20840	total number in ...	nhocc	4471	107	265.926	1	1566	total number of ...	nwocc	4471	102	309.9512	1	2662	total number of ...	phocc	4471	107	.0127604	.000048	.075144	percentage of me...	pwocc	4471	102	.0148729	.000048	.1277351	percentage of wo...	ewocc	4471	3832	3.305061	.0004319	200.0332	expected number ...	prop	4471	77	.0002237	.000048	.0533109		staner	4471	77	.0000831	.000048	.0015562	Standard error f...	pro_obs	4471	77	.0002237	.000048	.0533109	Observed proport...	pro_exp	4471	3818	.0001586	2.07e-08	.0095985	Expected proport...	pro_min	4471	77	.0001406	1.15e-09	.0517548	Lower confidence...	pro_max	4471	77	.0003067	.000096	.0548671	Higher confidenc...	value	4471	3968	5.717776	.0432525	2315.556	Observed value o...	val_min	4471	4217	1.115635	1.04e-06	81.03288	Value of represe...	val_max	4471	4214	10.31992	.0691135	4631.055	Value of represe...
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We set limits of frequency of 5 and val_min of 2.	<pre>. keep if freq>=5 (3546 observations deleted) . keep if val_min>=2 (617 observations deleted)</pre>																																																																																																																																					
This provides us with over 300 combinations of microclass_religion linkages.	<table><tr><th>Variable</th><th>Obs</th><th>Unique</th><th>Mean</th><th>Min</th><th>Max</th><th>Label</th></tr><tr><td>hocc</td><td>308</td><td>92</td><td>8767.581</td><td>1101</td><td>15202</td><td></td></tr><tr><td>wocc</td><td>308</td><td>91</td><td>8542.578</td><td>1101</td><td>15202</td><td></td></tr><tr><td>freq</td><td>308</td><td>57</td><td>22.03571</td><td>5</td><td>1111</td><td>(count) freq</td></tr><tr><td>tot</td><td>308</td><td>1</td><td>20840</td><td>20840</td><td>20840</td><td>total number in s...</td></tr><tr><td>nhocc</td><td>308</td><td>85</td><td>273.1526</td><td>14</td><td>1566</td><td>total number of m...</td></tr><tr><td>nwocc</td><td>308</td><td>77</td><td>330.25</td><td>12</td><td>2662</td><td>total number of f...</td></tr><tr><td>phocc</td><td>308</td><td>85</td><td>.0131071</td><td>.0006718</td><td>.075144</td><td>percentage of men...</td></tr><tr><td>pwocc</td><td>308</td><td>77</td><td>.0158469</td><td>.0005758</td><td>.1277351</td><td>percentage of wom...</td></tr><tr><td>ewocc</td><td>308</td><td>304</td><td>4.442368</td><td>.0735605</td><td>200.0332</td><td>expected number o...</td></tr><tr><td>prop</td><td>308</td><td>57</td><td>.0010574</td><td>.0002399</td><td>.0533109</td><td></td></tr><tr><td>staner</td><td>308</td><td>57</td><td>.0001857</td><td>.0001073</td><td>.0015562</td><td>Standard error fo...</td></tr><tr><td>pro_obs</td><td>308</td><td>57</td><td>.0010574</td><td>.0002399</td><td>.0533109</td><td>Observed proporti...</td></tr><tr><td>pro_exp</td><td>308</td><td>304</td><td>.0002132</td><td>3.53e-06</td><td>.0095985</td><td>Expected proporti...</td></tr><tr><td>pro_min</td><td>308</td><td>57</td><td>.0008717</td><td>.0001326</td><td>.0517548</td><td>Lower confidence...</td></tr><tr><td>pro_max</td><td>308</td><td>57</td><td>.0012431</td><td>.0003472</td><td>.0548671</td><td>Higher confidence...</td></tr><tr><td>value</td><td>308</td><td>304</td><td>9.38363</td><td>2.268401</td><td>109.2243</td><td>Observed value of...</td></tr><tr><td>val_min</td><td>308</td><td>306</td><td>6.628953</td><td>2.010274</td><td>81.03288</td><td>Value of represen...</td></tr><tr><td>val_max</td><td>308</td><td>306</td><td>12.13831</td><td>2.521529</td><td>137.4158</td><td>Value of represen...</td></tr></table>	Variable	Obs	Unique	Mean	Min	Max	Label	hocc	308	92	8767.581	1101	15202		wocc	308	91	8542.578	1101	15202		freq	308	57	22.03571	5	1111	(count) freq	tot	308	1	20840	20840	20840	total number in s...	nhocc	308	85	273.1526	14	1566	total number of m...	nwocc	308	77	330.25	12	2662	total number of f...	phocc	308	85	.0131071	.0006718	.075144	percentage of men...	pwocc	308	77	.0158469	.0005758	.1277351	percentage of wom...	ewocc	308	304	4.442368	.0735605	200.0332	expected number o...	prop	308	57	.0010574	.0002399	.0533109		staner	308	57	.0001857	.0001073	.0015562	Standard error fo...	pro_obs	308	57	.0010574	.0002399	.0533109	Observed proporti...	pro_exp	308	304	.0002132	3.53e-06	.0095985	Expected proporti...	pro_min	308	57	.0008717	.0001326	.0517548	Lower confidence...	pro_max	308	57	.0012431	.0003472	.0548671	Higher confidence...	value	308	304	9.38363	2.268401	109.2243	Observed value of...	val_min	308	306	6.628953	2.010274	81.03288	Value of represen...	val_max	308	306	12.13831	2.521529	137.4158	Value of represen...
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We can then export the data as above.	<pre>outsheet hocc wocc val_min using /// "\$path9\ca_1891_micro_cath.txt", /// comma nonames nolabel replace</pre>																																																																																																																																					
We can then rerun the analysis in Pajek, as above.	<p>Follow the steps from the txt2pajek example, using the Canadian data. The practices in Pajek are the same, regardless of the type of data.</p> <p><i>(Note: When we create the partition to distinguish between Catholics and non-Catholics, if you ask it to set a constant partition with a value of 0, you can simply change the value to a '1' if it is a five-digit number (i.e., it has a one added to the front).</i></p> <p><i>If you wish, you might want to code the macroclass additionally (the first number if 4 digits, first two if 5 digits). This will create potential values of 1-5, 9, 11-15 & 9. Using 'options', 'colors', 'partition colors',</i></p>																																																																																																																																					

	<p><i>'of vertices' in the graph window, you can click on the colors and assign them a new value. This would enable a 'bluescale' system for Catholics and a 'redscale' for non-Catholics akin to the usual 'greyscale' we see of the macroclasses being represented by brightness and the religion represented by colours).</i></p>
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