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## Description

`stir` reports point estimates and confidence intervals for the incidence-rate ratio (IRR) and incidence-rate difference (IRD). Stratified IRRs may be standardized to produce standardized mortality ratios.

`stir` can be used with single- or multiple-record and single- or multiple-failure `st` data.

## Quick start

IRR and IRD with confidence intervals for exposure indicator exposed using `stset` data

```
stir exposed
```

Same as above, but display exact  $p$ -values calculated without the mid- $p$  adjustment

```
stir exposed, exact
```

Crude and Mantel–Haenszel combined IRRs with test of homogeneity for strata defined by `svar`

```
stir exposed, strata(svar)
```

Same as above, and standardize the IRRs by weighting variable `wvar`

```
stir exposed, strata(svar) standard(wvar)
```

Same as above, but standardize using time at risk for the unexposed group as weights

```
stir exposed, strata(svar) estandard
```

## Menu

Statistics > Survival analysis > Summary statistics, tests, and tables > Report incidence-rate comparison

# Syntax

```
stir exposedvar [ if ] [ in ] [ , options ]
```

options	Description
Options	
strata( <i>varname</i> )	stratify on <i>varname</i>
estandard	combine external weights with within-stratum statistics
istandard	combine internal weights with within-stratum statistics
standard( <i>varname</i> )	combine user-specified weights with within-stratum statistics
pool	display pooled estimate
nocrude	do not display crude estimate
nohom	do not display homogeneity test
ird	calculate standardized IRD
midp	display <i>p</i> -values calculated using mid- <i>p</i> adjustment (unstratified only); the default
exact	display exact <i>p</i> -values without mid- <i>p</i> adjustment (unstratified only)
level(#)	set confidence level; default is level(95)
noshow	do not show st setting information

You must stset your data before using stir; see [ST] stset.  
by and collect are allowed; see [U] 11.1.10 Prefix commands.  
fweights and iweights may be specified using stset; see [ST] stset. stir may not be used with pweighted data.

# Options

Options

strata(*varname*) specifies that the calculation be stratified on *varname*, which may be a numeric or string variable. Within-stratum statistics are shown and then combined with Mantel–Haenszel weights. Also see the by() option in [R] Epitab.

estandard, istandard, standard(*varname*), pool, nocrude, nohom, and ird are relevant only if strata() is specified; see [R] Epitab.

midp and exact are relevant only if strata() is not specified; see [R] Epitab.

level(#) is relevant in all cases; see [R] Epitab.

noshow is relevant in all cases; it prevents stir from showing the key st variables. This option is seldom used because most people type stset , show or stset , noshow to set whether they want to see these variables mentioned at the top of the output of every st command; see [ST] stset.

# Remarks and examples

stir examines the incidence rate and time at risk.

```
. use https://www.stata-press.com/data/r19/page2
. stir group
    Failure _d: dead
    Analysis time _t: time
Incidence-rate comparison
Exposed:  group = 2
Unexposed: group = 1
```

	group		Total
	Exposed	Unexposed	
Failures	19	17	36
Time	5023	4095	9118
Incidence rate	.0037826	.0041514	.0039482
	Point estimate		[95% conf. interval]
Inc. rate diff.	-.0003688		-.002974 .0022364
Inc. rate ratio	.9111616		.4484366 1.866047 (exact)
Prev. frac. ex.	.0888384		-.8660469 .5515634 (exact)
Prev. frac. pop	.04894		

```
Mid-p-values for tests of incidence-rate difference:
Adj Pr(Exposed failures <= 19) = 0.3900 (lower one-sided)
Adj Pr(Exposed failures >= 19) = 0.6100 (upper one-sided)
Two-sided p-value = 0.7799
```

Specifying the exact option displays *p*-values for the tests of IRD calculated without using the mid-*p* adjustment. The noshow option suppresses the display of st variables.

```
. stir group, exact noshow
Incidence-rate comparison
Exposed:  group = 2
Unexposed: group = 1
```

	group		Total
	Exposed	Unexposed	
Failures	19	17	36
Time	5023	4095	9118
Incidence rate	.0037826	.0041514	.0039482
	Point estimate		[95% conf. interval]
Inc. rate diff.	-.0003688		-.002974 .0022364
Inc. rate ratio	.9111616		.4484366 1.866047 (exact)
Prev. frac. ex.	.0888384		-.8660469 .5515634 (exact)
Prev. frac. pop	.04894		

```
Exact p-values for tests of incidence-rate difference:
Pr(Exposed failures <= 19) = 0.4536 (lower one-sided)
Pr(Exposed failures >= 19) = 0.6737 (upper one-sided)
Two-sided p-value = 0.9071
```

See [\[R\] EpiTab](#) for details about the exact option and other stir options.

## Video example

How to calculate incidence rates and incidence-rate ratios

## Stored results

`stir` (without `strata()`) stores the following in `r()`:

Scalars

<code>r(ird)</code>	IRD
<code>r(lb_ird)</code>	lower CI bound for IRD
<code>r(ub_ird)</code>	upper CI bound for IRD
<code>r(irr)</code>	IRR
<code>r(lb_irr)</code>	lower CI bound for IRR
<code>r(ub_irr)</code>	upper CI bound for IRR
<code>r(afe)</code>	attributable fraction among the exposed
<code>r(lb_afe)</code>	lower CI bound for attributable fraction among the exposed
<code>r(ub_afe)</code>	upper CI bound for attributable fraction among the exposed
<code>r(afp)</code>	attributable fraction for the population
<code>r(p_lower_midp)</code>	lower one-sided $p$ -value with mid- $p$ adjustment
<code>r(p_upper_midp)</code>	upper one-sided $p$ -value with mid- $p$ adjustment
<code>r(p_twosided_midp)</code>	two-sided $p$ -value with mid- $p$ adjustment
<code>r(p_lower_exact)</code>	lower one-sided exact $p$ -value
<code>r(p_upper_exact)</code>	upper one-sided exact $p$ -value
<code>r(p_twosided_exact)</code>	two-sided exact $p$ -value

`stir`, `strata()` stores the following in `r()`:

Scalars

<code>r(irr)</code>	Mantel–Haenszel IRR, if option <code>ird</code> is not specified
<code>r(lb_irr)</code>	lower CI bound for Mantel–Haenszel IRR
<code>r(ub_irr)</code>	upper CI bound for Mantel–Haenszel IRR
<code>r(ird)</code>	Mantel–Haenszel IRD, if option <code>ird</code> is specified
<code>r(lb_ird)</code>	lower CI bound for Mantel–Haenszel IRD
<code>r(ub_ird)</code>	upper CI bound for Mantel–Haenszel IRD
<code>r(crude)</code>	crude IRR or, if option <code>ird</code> is specified, crude IRD
<code>r(lb_crude)</code>	lower CI bound for the crude IRR or IRD
<code>r(ub_crude)</code>	upper CI bound for the crude IRR or IRD
<code>r(pooled)</code>	pooled IRR or, if option <code>ird</code> is specified, pooled IRD
<code>r(lb_pooled)</code>	lower CI bound for pooled IRR or IRD
<code>r(ub_pooled)</code>	upper CI bound for pooled IRR or IRD
<code>r(df)</code>	degrees of freedom for homogeneity $\chi^2$ test
<code>r(chi2_mh)</code>	Mantel–Haenszel homogeneity $\chi^2$
<code>r(chi2_p)</code>	pooled homogeneity $\chi^2$ , if option <code>poo1</code> is specified

## Methods and formulas

`stir` simply accumulates numbers of failures and time at risk by exposed and unexposed (by `strata`, if necessary) and passes the calculation to `ir`; see [R] [EpiTab](#).

## Reference

Dupont, W. D. 2009. *Statistical Modeling for Biomedical Researchers: A Simple Introduction to the Analysis of Complex Data*. 2nd ed. Cambridge: Cambridge University Press.

## Also see

[ST] **stset** — Declare data to be survival-time data

[ST] **stsum** — Summarize survival-time data

[R] **EpiTab** — Tables for epidemiologists

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