

# Package ‘rfcipDemand’

September 12, 2025

**Type** Package

**Title** Estimate Federal Crop Insurance Program Demand Models

**Version** 0.0.0.9000

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**Description** Tools to construct county–crop–practice–plan–unit panels from the USDA RMA Summary of Business (SOBTPU) and related sources, and to estimate FCIP demand systems with two-way cluster-robust covariance. The pipeline standardizes coverage measures, merges price and instrument variables, adds rental-rate and price-index controls, reconciles county acreage (FSA/NASS), and produces diagnostics including robust first-stage F-tests. Methods align with the empirical design in “The crop insurance demand response to premium subsidies Evidence from U.S. Agriculture” (Food Policy, 2023, 119(3)).

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**URL** <https://github.com/you/rfcipDemand>

**BugReports** <https://github.com/you/rfcipDemand/issues>

**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.3.2

**VignetteBuilder** knitr

**Depends** R (>= 4.1.0)

**Imports** rfsa, data.table, systemfit, sandwich, doBy, car, plyr, usmap, stats, utils, tidyr, nlsvr

**Remotes** github::UrbanInstitute/urbnmapr, github::dylan-turner25/rfsa, github::JanMarvin/nlsvr

**Suggests** knitr, rmarkdown, tibble, dplyr, lavaan, testthat (>= 3.0.0)

**LazyData** true

**Cite-us** If you find it useful, please consider starring the repository and citing the following studies

- Tsiboe, F. and Turner, D. (2025). “Incorporating buy-up price loss coverage into the United States farm safety net.” Applied Economic Perspectives and Policy.
- Tsiboe, F., et al. (2025). “Risk reduction impacts of crop insurance in the United States.”

Applied Economic Perspectives and Policy.  
- Gaku, S. and Tsiboe, F. (2024). Evaluation of alternative farm safety net program combination strategies. Agricultural Finance Review.

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|                |                                       |
|----------------|---------------------------------------|
| calculate_mode | <i>Calculate the Statistical Mode</i> |
|----------------|---------------------------------------|

---

Description

Returns the element that occurs most frequently in a vector.

Usage

calculate\_mode(x, na.rm = TRUE)

**Arguments**

|                    |   |
|--------------------|---|
| <code>x</code>     | A vector of any atomic type (numeric, character, factor,).  |
| <code>na.rm</code> | Logical; should missing values be ignored? Defaults to TRUE. If FALSE and <code>x</code> contains any NAs, the function returns NA. |

**Details**

Internally the function:

1. Optionally removes NAs (`na.rm = TRUE`).
2. Builds a lookup table of unique values via `unique(x)`.
3. Counts the frequency of each unique value with `tabulate(match(x, ux))`.
4. Returns the value with the maximum count.

Because it relies on base R functions, the implementation is vectorised and generally fast for typical data-frame column sizes.

**Value**

A single value giving the modal element of `x`. If two or more values are tied for the highest frequency, the first one encountered in `x` is returned.

---

```
estimate_fcip_instruments
```

*Estimate FCIP Instrumental Variables (Unloaded Rates)*

---

**Description**

Uses historical FCIP rate data to build instrumented unloaded-rate variables following:

1. Tsiboe & Turner (2023), Econometric identification of crop insurance participation *Agricultural and Resource Economics Review*, 52(3):476-497. <https://doi.org/10.1017/age.2023.13>

**Usage**

```
estimate_fcip_instruments(year, statplan)
```

**Arguments**

|                       |   |
|-----------------------|---|
| <code>year</code>     | Integer. The target crop year for which to construct instruments.   |
| <code>statplan</code> | A data.table containing FCIP rate elements, including at least:<br><b>commodity_year</b> Year of the rate observation.<br><b>state_code, county_code</b> County identifiers.<br><b>commodity_code</b> Crop identifier.<br><b>insured_area, lcr, contiguous_state_code, contiguous_county_code</b> Fields required by <code>estimate_fcip_unloaded_rate()</code> . |

## Details

1. **Task list:** Identify all unique (state, county) pairs with data in the 2-21 years before year.
2. **Unloaded-rate calculation:** For each county in task\_list, call estimate\_fcip\_unloaded\_rate() on the same 2-21 year window to get tau. Errors return NULL so processing continues.
3. **Contiguous-county smoothing:**
  - Build a lookup table of contiguous counties (using contiguous\_county).
  - For each contiguous group, compute the mean tau to get tau\_c.
4. **Merge & fill:** Left-join the raw adm and contiguous\_adm; replace any zero/NA/Inf tau with the group mean tau\_c into tau\_sob.
5. **Cleanup:** Drop helper columns (tau, tau\_c), remove invalid rows, add commodity\_year, and return the result.

## Value

A data.table with one row per county-crop for the specified year, containing:

**state\_code, county\_code, commodity\_code** Keys.

**tau\_sob** Smoothed unloaded rate (uses contiguous-county means to fill zeros/NAs).

**commodity\_year** The input year, repeated.

## See Also

Other FCIP instruments: [get\\_yu2018\\_instrument\(\)](#)

---

|                        |                               |
|------------------------|-------------------------------|
| fcip_contiguous_county |                               |
|                        | <i>fcip_contiguous_county</i> |

---

## Description

A combined dataset for fcip\_contiguous\_county

## Usage

```
data(fcip_contiguous_county)
```

## Format

A data frame with 24307 rows and 12 columns covering Inf–Inf.

## Source

USDA-RMA, Actuarial Data Master - A0123

---

fcip\_demand\_data\_dispatcher

*Build dataset to estimate Federal Crop Insurance Program (FCIP) demand (modular pipeline)*


---

## Description

End-to-end pipeline that: (1) prepares SOBTPU and coverage aggregates, (2) adds prices/instruments/rental rates/price index, (3) reconciles county acreage (FSA + NASS), and (4) finalizes bins/labels/pooling for demand estimation.

## Usage

```
fcip_demand_data_dispatcher(
  study_years = 2001:(as.numeric(format(Sys.Date(), "%Y")) - 1),
  identifiers = c("commodity_year", FCIP_INSURANCE_POOL, "insurance_plan_code",
    "unit_structure_code")
)
```

## Arguments

|             |  |
|-------------|--|
| study_years | Integer vector of commodity years to include. Defaults to 2001:(as.numeric(format(Sys.Date(), "%Y")) - 1).   |
| identifiers | Character vector of grouping keys that define the aggregation grain. Must be columns present in SOBTPU (e.g., "commodity_year", FCIP_INSURANCE_POOL, "insurance_plan_code", "unit_structure_code", and-if desired-additional keys like "commodity_code" or "practice_code"). Enrichment joins for re-codes are performed only when the required keys are included in identifiers |

## Details

Aligned with Asche, Bekkerman, & Li (2023), *Food Policy*, 119(3):102505 ([doi:10.1016/j.foodpol.2023.102505](https://doi.org/10.1016/j.foodpol.2023.102505)). Requires internet access to download release .rds assets and several in-memory lookup tables (see stage docs).

## Value

A data.table ready for FCIP demand estimation.

---

fcip\_demand\_elasticities\_lavaan

*Calibrate FCIP demand elasticities via IV-SEM (lavaan)*


---

## Description

Fits a just-identified IV-style SEM where instr\_rate instruments tilda\_rate, and tilda\_rate enters two outcome equations (tilda\_Gamma, tilda\_Theta1). Endogeneity is encoded via residual correlations between tilda\_rate and each outcome. The instrument is excluded from the outcome disturbances.

**Usage**

```
fcip_demand_elasticities_lavaan(
  data,
  estimator = c("ML", "MLR"),
  missing = c("fiml", "listwise")
)
```

**Arguments**

|           |   |
|-----------|---|
| data      | data.frame with cols: instr_rate, tilda_rate, tilda_Gamma, tilda_Theta1 |
| estimator | "ML" or "MLR" (default "MLR" = robust SE/tests)                         |
| missing   | "fiml" or "listwise" (default "fiml")                                   |

**Details**

Constraints imposed:

1.  $b1 < 0$ , 2)  $b2 < 0$ , 3)  $b1 + b2 + b1*b2 < 0$

**Value**

A data.table of parameter estimates and logical convergence flag

---

fcip\_demand\_sys\_coeff\_table

*Tidy coefficient table with cluster-robust SEs (from supplied VCOV)*

---

**Description**

Builds a clean coefficient table for a systemfit model using a **user-supplied covariance matrix** (e.g., two-way clustered from fcip\_demand\_sys\_vcov()). Estimates come from coef(fit), standard errors from diag(vcMat), then Z-scores and two-sided normal p-values are computed. The demand column is inferred from the equation prefix in the coefficient names:

- "gamma\_\*" to "gamma"
- "theta\_\*" to "theta" Otherwise the prefix itself is used.

**Usage**

```
fcip_demand_sys_coeff_table(fit, vcMat, p_digits = 5)
```

**Arguments**

|          |  |
|----------|--|
| fit      | A fitted systemfit object.   |
| vcMat    | A covariance matrix conformable with coef(fit). Row/column names are used to align; if missing, positional alignment is assumed. |
| p_digits | Integer; number of digits to keep for p-values (default 5).  |

**Value**

A data.frame with columns: demand, coef, Estimate, StdError, Zvalue, Pvalue.

---

fcip\_demand\_sys\_effect

*Delta-method "total protection response"*


---

### Description

Combines equation-specific effects into a single "total" effect for each regressor in `c(fields$endogenous, fields$included)` using `car::deltaMethod` and a supplied covariance matrix.

### Usage

```
fcip_demand_sys_effect(fit, vcMat, fields, data)
```

### Arguments

|                     |   |
|---------------------|---|
| <code>fit</code>    | A fitted <code>systemfit</code> object (the structural system).   |
| <code>vcMat</code>  | Covariance matrix conformable with <code>coef(fit)</code> (e.g., from <code>fcip_demand_sys_vcov()</code> ).                |
| <code>fields</code> | Named list carrying model fields; must include <code>outcome</code> , <code>endogenous</code> , and <code>included</code> . |
| <code>data</code>   | Estimation data used to check variable availability and build delta-method expressions.                                     |

### Value

A `data.frame` with rows `demand="total"` and columns: `demand`, `coef`, `Estimate`, `StdError`, `Zvalue`, `Pvalue`.

---

fcip\_demand\_sys\_estimate

*System estimator (modular wrapper; preserves original outputs)*


---

### Description

Runs: `per-level prep` -> `partial/tilda` -> `systemfit` -> `clustered VCOV` -> `delta-method totals` -> (optional) `restricted NLSUR` -> `diagnostics` -> `bind rows`.

### Usage

```
fcip_demand_sys_estimate(model, data)
```

### Arguments

|                    |   |
|--------------------|---|
| <code>model</code> | List with elements: <code>outcome</code> , <code>endogenous</code> , <code>included</code> , <code>excluded (opt)</code> , <code>partial (opt)</code> , <code>FE (logical)</code> , <code>disag (string colname)</code> , <code>optional restrict (logical)</code> , <code>name (string)</code> . |
| <code>data</code>  | Input <code>data.frame/data.table</code> with all referenced columns plus <code>pool</code> and <code>commodity_year</code> .   |

### Value

A `data.frame` aggregating results across all disaggregation levels.

---

|                     |   |
|---------------------|---|
| fcip_demand_sys_fit | <i>Build systemfit formulas and estimate the system</i> |
|---------------------|---|

---

### Description

Constructs the list of structural equations (g) and instrument sets (h), then runs `systemfit()` using OLS (when no excluded instruments) or 3SLS-GMM (when excluded instruments are present).

### Usage

```
fcip_demand_sys_fit(
  data,
  fields,
  tilda_included,
  tilda_endogenous,
  tilda_excluded
)
```

### Arguments

|                               |   |
|-------------------------------|---|
| <code>data</code>             | Estimation data.frame/data.table containing the <code>tilda_*</code> and <code>instr_*</code> variables referenced by the formulas. |
| <code>fields</code>           | Named list with at least outcome, included, endogenous, and optionally excluded.  |
| <code>tilda_included</code>   | Character vector of residualized included regressor names (e.g., "tilda_x1").   |
| <code>tilda_endogenous</code> | Character vector of residualized endogenous regressor names (e.g., "tilda_z1").   |
| <code>tilda_excluded</code>   | Character vector of instrument names (e.g., "instr_z1"), or NULL when no excluded instruments are used.                             |

### Value

A list with elements:

|                  |                                       |
|------------------|---------------------------------------|
| <code>fit</code> | Fitted <code>systemfit</code> object. |
| <code>g</code>   | List of structural formulas.          |
| <code>h</code>   | List of instrument formulas.          |

---

|                         |  |
|-------------------------|--|
| fcip_demand_sys_partial | <i>Residualize ("partial out") and build tilded / instrument variables</i> |
|-------------------------|--|

---

### Description

If excluded instruments exist, runs first-stage OLS for each endogenous variable  $e$ :  $e \sim 1 + \text{partial} + \text{included} + \text{excluded}$ , storing the fitted values as `instr_e`. If `partial` is non-empty, it then regresses  $\text{instr}_e \sim 1 + \text{partial}$  and replaces  $\text{instr}_e \leftarrow \text{instr}_e - \text{fitted}(\text{instr}_e \sim \text{partial})$  (i.e., removes the partial component; conceptually  $\widehat{\text{instr}}_e(\text{partial})$ ).

Outcomes, included, and endogenous variables are residualized on `partial` to create `tilda_<var>` (or copied if `partial` is empty).



**Usage**

```
fcip_demand_sys_partial(data, fields, partial_override = NULL)
```

**Arguments**

|                               |  |
|-------------------------------|--|
| <code>data</code>             | A <code>data.frame</code> / <code>data.table</code> with referenced variables.   |
| <code>fields</code>           | List with: <code>outcome</code> , <code>endogenous</code> , <code>included</code> , <code>optional excluded</code> , <code>optional partial</code> . |
| <code>partial_override</code> | Optional character vector to override <code>fields\$partial</code> .   |

**Details**

Uses defensive checks so absent columns are ignored (with a warning) rather than erroring. If `partial` is empty, residualization is a no-op and `tilda_*` simply copy the originals. Formulas are constructed via `stats::reformulate()` to avoid paste/quoting pitfalls.

**Value**

List with `data`, `tilda_included`, `tilda_endogenous`, `tilda_excluded`.

---

`fcip_demand_sys_prep`    *Prepare data for analysis*

---

**Description**

Drops incomplete/invalid rows, removes constant partials, and optionally demeanes via a fixed-effects helper.

**Usage**

```
fcip_demand_sys_prep(data, fields)
```

**Arguments**

|                     |  |
|---------------------|--|
| <code>data</code>   | Estimation dataset that already contains all columns referenced by <code>fields</code> .   |
| <code>fields</code> | Named list: <code>outcome</code> , <code>endogenous</code> , <code>included</code> , <code>excluded (opt)</code> , <code>partial (opt)</code> , <code>FE (logical)</code> , <code>disag (column name)</code> . |

**Value**

A list: `data` (prepped), `NFE` (number of FE), `partial` (possibly reduced).

---

fcip\_demand\_sys\_restricted

*Optional restricted NLSUR step (when restrict = TRUE)*


---

## Description

If enabled and feasible, estimates a nonlinear SUR with re-parameterized coefficients (negative exponents) using the optional **nlsur** package and appends "restricted\_" rows to the results. Skips gracefully if **nlsur** is not installed or the step fails.

## Usage

```
fcip_demand_sys_restricted(
  restrict,
  res,
  fit,
  data,
  outcome,
  tilda_endogenous,
  tilda_excluded,
  tilda_included
)
```

## Arguments

|                  |   |
|------------------|---|
| restrict         | Logical flag; when TRUE attempt the restricted step.                                    |
| res              | Coefficient table from the unrestricted system (used to check signs).                   |
| fit              | Fitted systemfit object from the unrestricted system.                                   |
| data             | Estimation data used to fit the restricted NLSUR model.                                 |
| outcome          | Character vector of outcome equation names (length 2 expected).                         |
| tilda_endogenous | Character vector of endogenous regressors used in the tilded system (e.g., "tilda_z1"). |
| tilda_excluded   | Character vector of excluded instruments (e.g., "instr_z1").                            |
| tilda_included   | Character vector of included regressors ("tilda_x1", ...).                              |

## Value

A `data.frame` with rows for gamma, theta, and total labeled `restricted_*`, or an empty `data.frame` if skipped.

## Note

This step uses `nlsur::nlsur()` if available; it is optional and should be listed under Suggests in DESCRIPTION.

---

|                     |                             |
|---------------------|-----------------------------|
| fcip_demand_sys_run | <i>Orchestrate analysis</i> |
|---------------------|-----------------------------|

---

## Description

Runs the full pipeline: prep -> partial/tilda creation -> systemfit -> two-way clustered VCOV -> delta-method totals -> optional restricted step -> diagnostics; then returns a tidy coefficient table with metadata.

## Usage

```
fcip_demand_sys_run(data, fields)
```

## Arguments

|        |  |
|--------|--|
| data   | Estimation dataset   |
| fields | Named list carrying model fields (see <code>fcip_demand_estimation()</code> ), including <code>disag</code> , <code>FE</code> , <code>outcome</code> , <code>endogenous</code> , <code>included</code> , <code>optional excluded</code> , <code>partial</code> , <code>restrict</code> , and <code>name</code> . |

## Value

A `data.frame` with columns `demand`, `coef`, `Estimate`, `StdError`, `Zvalue`, `Pvalue` and meta-columns `model`, `endogenous`, `FE`, `name`, `disag`

---

|                       |  |
|-----------------------|--|
| fcip_demand_sys_tests | <i>System diagnostics: two-way robust first-stage F (+ optional approx. J)</i> |
|-----------------------|--|

---

## Description

Produces diagnostics **without** re-running GMM:

- **FTest**: joint relevance of excluded instruments in each first stage, using the same two-way (pool by crop year) cluster-robust covariance via `fcip_demand_sys_vcov()` with `kind = "lm"`. Reports the **minimum** F across endogenous regressors.
- **JTest** (optional): an *approximate* over-identification test computed as the sum of per-equation Sargan statistics  $J_k \approx n_k R_k^2$  from regressions of equation residuals on that equation's instrument set. This is a quick check (not the system Hansen J).

## Usage

```
fcip_demand_sys_tests(g, h, data, fit, NFE, approx_j = FALSE)
```

**Arguments**

|                       |  |
|-----------------------|--|
| <code>g</code>        | List of system equations (the same formulas passed to <code>systemfit</code> ).  |
| <code>h</code>        | List of instrument formulas (the same formulas passed to <code>systemfit</code> ).   |
| <code>data</code>     | Estimation data.frame/data.table containing all variables in <code>g/h</code> plus clustering columns <code>pool</code> and <code>crop_yr</code> . |
| <code>fit</code>      | A fitted <code>systemfit</code> object (used for <code>N</code> and <code>residCov_*</code> extraction).   |
| <code>NFE</code>      | Integer: number of absorbed fixed effects (for reporting only).  |
| <code>approx_j</code> | Logical, compute the approximate (non-robust) Sargan J as described above. Default FALSE (returns NA for JTest).                                   |

**Value**

A data.frame with rows: `N`, `NFE`, `residCov_11`, `residCov_22`, `residCov_12`, `JTest`, `FTest`.

---

|                                   |   |
|-----------------------------------|---|
| <code>fcip_demand_sys_vcov</code> | <i>Two-way cluster-robust covariance for FCIP demand models</i> |
|-----------------------------------|---|

---

**Description**

Computes a Cameron-Gelbach-Miller two-way cluster-robust covariance matrix using inclusion-exclusion:  $V = V_{pool} + V_{year} - V_{pool\_year}$ . Works for both `systemfit` (stacked system) and `lm` (first-stage).

**Usage**

```
fcip_demand_sys_vcov(
  object,
  data,
  kind = c("systemfit", "lm"),
  pool_col = "pool",
  year_col = "commodity_year",
  NFE = 0L,
  n_partial = 0L,
  n_eq = NULL
)
```

**Arguments**

|                        |  |
|------------------------|--|
| <code>object</code>    | Fitted model: either a <code>systemfit</code> or <code>lm</code> .   |
| <code>data</code>      | Estimation data containing pool and year identifiers.  |
| <code>kind</code>      | One of <code>c("systemfit", "lm")</code> . If omitted, auto-detected.  |
| <code>pool_col</code>  | Name of the pool/cluster id column in data (default "pool").   |
| <code>year_col</code>  | Name of the year/time id column in data (default "crop_yr").   |
| <code>NFE</code>       | Integer; number of absorbed fixed effects (for df rescaling).  |
| <code>n_partial</code> | Integer; count of variables partialled out per equation.   |
| <code>n_eq</code>      | Integer; number of equations ( <code>length(object\$eq)</code> for <code>systemfit</code> , 1 for <code>lm</code> ). You can override if needed. |

**Details**

**Rescaling.** Let  $n$  be the number of observations (stacked across equations for `systemfit`). With  $k_{old}$  the number of coefficients and  $k_{new} = k_{old} + NFE + n_{partial} * n_{eq}$ , the returned matrix is scaled by  $(n - k_{old} - 1) / (n - k_{new} - 1)$ .

**Row alignment (lm).** Rows used by `lm` are inferred from `rownames(model.matrix(object))`. If they cannot be mapped back to data, the first `nobs(object)` rows are used.

**Value**

Covariance matrix aligned with `coef(object)`.

---

```
fcip_recodes_commodity_groupings
      fcip_recodes_commodity_groupings
```

---

**Description**

A combined dataset for `fcip_recodes_commodity_groupings`

**Usage**

```
data(fcip_recodes_commodity_groupings)
```

**Format**

A data frame with 3572 rows and 10 columns covering 1997-2025.

**Source**

USDA-RMA, Actuarial Data Master - A00400 and A00420 supplemented data from legacy ADM files

---

```
fcip_recodes_insurance_plan
      fcip_recodes_insurance_plan
```

---

**Description**

A combined dataset for `fcip_recodes_insurance_plan`

**Usage**

```
data(fcip_recodes_insurance_plan)
```

**Format**

A data frame with 773 rows and 10 columns covering 1989-2025.

**Source**

USDA-RMA, Actuarial Data Master - A00460 supplemented data from legacy ADM files

---

|                       |                              |
|-----------------------|------------------------------|
| fcip_recodes_practice | <i>fcip_recodes_practice</i> |
|-----------------------|------------------------------|

---

**Description**

A combined dataset for fcip\_recodes\_practice

**Usage**

```
data(fcip_recodes_practice)
```

**Format**

A data frame with 28639 rows and 8 columns covering 1997-2025.

**Source**

USDA-RMA, Actuarial Data Master - A00510 supplemented data from legacy ADM files

---

|                   |                          |
|-------------------|--------------------------|
| fcip_recodes_type | <i>fcip_recodes_type</i> |
|-------------------|--------------------------|

---

**Description**

A combined dataset for fcip\_recodes\_type

**Usage**

```
data(fcip_recodes_type)
```

**Format**

A data frame with 232709 rows and 7 columns covering 1999-2025.

**Source**

Generated internally, using `harmonize_crop_type_codes()`

---

fixed\_effect\_model\_data\_prep

*Prepare and demean data for fixed-effects models*


---

## Description

This function

1. Filters to complete cases on the specified panel, time, weight, variables, and output
2. If output is NULL, creates a dummy output column filled with 1s
3. Drops any panel with only one observation
4. Computes within-panel means for the output + each variable in varlist (`_mean_i`)
5. Computes overall sample means for the same set of variables (`_mean`)
6. Replaces each variable in varlist by `value - within_panel_mean + overall_mean`

## Usage

```
fixed_effect_model_data_prep(
  data,
  varlist,
  panel,
  time,
  wvar = NULL,
  output = NULL
)
```

## Arguments

|                      |   |
|----------------------|---|
| <code>data</code>    | A <code>data.frame</code> or <code>data.table</code> containing the data.   |
| <code>varlist</code> | Character vector of variable names to be demeaned.  |
| <code>panel</code>   | Character vector of column name(s) defining the panel identifier.   |
| <code>time</code>    | Character scalar name of the time variable.   |
| <code>wvar</code>    | Character scalar name of a variable to keep but <i>not</i> demean (optional, default NULL).   |
| <code>output</code>  | Character scalar name of an output variable whose means are computed but not altered; if NULL, a dummy column named "output" is created (optional, default NULL). |

## Value

A list with components

- **data**: a `data.table` containing
  - the original panel, time, wvar, varlist, and output columns
  - two mean columns for each of `c(output, varlist)`: `<name>_mean_i` (within-panel) and `<name>_mean` (overall)
- **NFE**: the number of panels with more than one observation

See Also

Other Estimators panel models: [panel\\_based\\_spatial\\_smoothing\\_estimator\(\)](#)

---

|                          |  |
|--------------------------|--|
| format_fcip_demand_table | <i>Table: Crop Insurance Demand System for US Federal Crop Insurance Pools (2001/22)</i> |
|--------------------------|--|

---

Description

Build a two-column, GitHub-safe panel table summarizing a crop insurance demand system. The table is organized into panels for coverage level (Theta), insured acres (Gamma), total protection response, a covariance matrix block, and additional statistics. Coefficients are formatted as estimate (std. error) with significance stars.

Usage

```
format_fcip_demand_table(df, var_labels)
```

Arguments

|            |  |
|------------|--|
| df         | A data frame containing the results with columns: <ul style="list-style-type: none"><li>demand (chr): panel identifier; expected values include "Theta", "Gamma", and "Total".</li><li>coef (chr): raw coefficient/row labels (e.g., "tilda_rate", "residCov_11", "N").</li><li>Estimate (dbl): point estimates.</li><li>StdError (dbl): standard errors (may be NA for scalars).</li><li>Pvalue (dbl): p-values used to add significance stars.</li></ul> |
| var_labels | A named character vector mapping raw names to display labels,  |

Details

Designed for README/output knitted as github\_document; use with knitr::kable(..., format = "pipe") to avoid HTML-only features.

Value

A tibble with two columns, Variables and Estimates, where panel headers have empty Estimates to enable bolding (if rendered in HTML) and coefficients are formatted as "estimate\*\*\* (se)".



---

|                 |                                  |
|-----------------|----------------------------------|
| fsa_crop_linker | <i>Simulator Helper Datasets</i> |
|-----------------|----------------------------------|

---

**Description**

A combined dataset for fsa\_crop\_linker

**Usage**

```
data(fsa_crop_linker)
```

**Format**

A data frame with 8594 rows and 8 columns covering Inf–Inf.

**Source**

Internal innovation

---

|                       |   |
|-----------------------|---|
| get_yu2018_instrument | <i>Formulate &amp; Merge National Subsidy Rate Instrument (Yu et al., 2018)</i> |
|-----------------------|---|

---

**Description**

Downloads the historical Summary of Business RDS and computes national subsidy-rate instruments at specified coverage levels, following Yu et al. (2018).

**Usage**

```
get_yu2018_instrument(
  dt,
  delivery_systems = c("RBUP", "FBUP"),
  plan_codes = c(1:3, 90, 44, 25, 42),
  coverage_levels = c(65, 75)
)
```

**Arguments**

|                  |  |
|------------------|--|
| dt               | sobcov   |
| delivery_systems | Character vector. Delivery systems to include; default c("RBUP", "FBUP").        |
| plan_codes       | Integer vector. Insurance plan codes to include; default c(1:3, 90, 44, 25, 42). |
| coverage_levels  | Numeric vector. Percent coverage levels to keep; default c(65, 75).              |

**Value**

A data.table with columns: commodity\_year, subsidy\_rate\_65, subsidy\_rate\_75.

See Also

Other FCIP instruments: [estimate\\_fcip\\_instruments\(\)](#)

---

nass\_census\_state\_beginning\_farmer\_and\_rancher\_data  
nass\_census\_state\_beginning\_farmer\_and\_rancher\_data

---

Description

A combined dataset for nass\_census\_state\_beginning\_farmer\_and\_rancher\_data

Usage

data(nass\_census\_state\_beginning\_farmer\_and\_rancher\_data)

Format

A data frame with 255 rows and 16 columns covering Inf–Inf.

Source

USDA NASS Quick Stats

---

nass\_index\_for\_price\_recived  
nass\_index\_for\_price\_recived

---

Description

A combined dataset for nass\_index\_for\_price\_recived

Usage

data(nass\_index\_for\_price\_recived)

Format

A data frame with 35 rows and 3 columns covering 1990-2024.

Source

USDA NASS Quick Stats

---

|                                      |
|--------------------------------------|
| nass_marketing_year_avg_price        |
| <i>nass_marketing_year_avg_price</i> |

---

**Description**

A combined dataset for nass\_marketing\_year\_avg\_price

**Usage**

```
data(nass_marketing_year_avg_price)
```

**Format**

A data frame with 31139 rows and 7 columns covering 1866-2024.

**Source**

USDA NASS Quick Stats

---

|                                |
|--------------------------------|
| nass_state_rental_rates        |
| <i>nass_state_rental_rates</i> |

---

**Description**

A combined dataset for nass\_state\_rental\_rates

**Usage**

```
data(nass_state_rental_rates)
```

**Format**

A data frame with 1792 rows and 5 columns covering 1994-2025.

**Source**

Output from get\_state\_rental\_rates() function

---

```
nass_us_ag_price_index_monthly
      nass_us_ag_price_index_monthly
```

---

### Description

A combined dataset for nass\_us\_ag\_price\_index\_monthly

### Usage

```
data(nass_us_ag_price_index_monthly)
```

### Format

A data frame with 2255 rows and 8 columns covering Inf–Inf.

### Source

USDA NASS: [https://www.nass.usda.gov/Charts\\_and\\_Maps/graphics/data](https://www.nass.usda.gov/Charts_and_Maps/graphics/data)

---

```
panel_based_spatial_smoothing_estimator
      Panel-based spatial smoothing estimator
```

---

### Description

This function

1. Constructs spatially-varying treatment interactions (one variable per spatial unit)
2. Applies within-panel/time fixed-effects demeaning to both outcome and interactions
3. Fits an OLS model by hand (`lm.fit`) to recover one coefficient per spatial unit

### Usage

```
panel_based_spatial_smoothing_estimator(
  data,
  output,
  treatment,
  time,
  panel,
  spatialvar
)
```

**Arguments**

|            |  |
|------------|--|
| data       | A data.table or data.frame containing panel data.                                    |
| output     | Name of the outcome variable (character scalar).                                     |
| treatment  | Name of the treatment variable whose spatial effects we estimate (character scalar). |
| time       | Name of the time variable (character scalar).  |
| panel      | Name(s) of the panel identifier variable(s) (character vector).                      |
| spatialvar | Name of the spatial grouping variable (e.g. county FIPS; character scalar).          |

**Details**

Internally, we

1. Build `treatment_code = I(spatialvar==code) * treatment` for each spatial unit code.
2. Call `fixed_effect_model_data_prep()` to demean the outcome and all `treatment_code` variables.
3. Assemble the design matrix  $X = [\text{output\_mean\_i}, \text{treatment\_}]$  and response  $y$ .
4. Solve  $\hat{\beta} = (\tilde{X}'\tilde{X})^{-1}\tilde{X}'\tilde{y}$  via `lm.fit`.
5. Return a row per spatial unit with its coefficient.

**Value**

A data.table with columns:

- `estimate`: the estimated spatial-unit coefficient
- `county_fips`: the spatial unit identifier (5-digit FIPS)
- `state_code`, `county_code`: parsed FIPS components

**See Also**

Other Estimators panel models: [fixed\\_effect\\_model\\_data\\_prep\(\)](#)

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