

Package ‘psvr’

September 24, 2026

Title Percentage-Error Support Vector Regression

Version 0.1.0

Description Implements four support vector regression (SVR) models derived from a unified mathematical framework for percentage-error loss functions: epsilon-SVR minimizing the mean absolute percentage error (MAPE), its symmetric kernel extension, least-squares SVR (LS-SVR) minimizing the root mean square percentage error (RMSPE), and its symmetric counterpart. All models require strictly positive targets. The epsilon-SVR models are solved via a built-in sequential minimal optimization (SMO) algorithm (with 'osqp' available as an optional alternative backend) and the LS-SVR models via a linear system (base R). See Benavides-Herrera et al. (2026) <[doi:10.3390/math14101679](https://doi.org/10.3390/math14101679)> for the mathematical derivations.

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coef.psvr_mape	<i>Extract coefficients from a psvr_mape model</i>
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Description

Extract coefficients from a psvr_mape model

Usage

```
## S3 method for class 'psvr_mape'
coef(object, ...)
```

Arguments

object	An object of class "psvr_mape".
...	Ignored.

Value

A named list with five components:

alpha, alpha_star The length-N pre-pruning dual variables α_k and α_k^* .

beta The pruned dual differences $\beta_k = \alpha_k - \alpha_k^*$ over the support-vector indices (length n_sv); this is what predict() uses.

b Bias term.

support_data Support vector input matrix.

The LS-SVR classes return **three** components rather than five, since they have no alpha_star and no pruned beta; the absent components are not materialised as NULL. So names(coef(fit)) depends on the model family. That is deliberate: each class is family-specific, and inventing empty slots to make the two agree would add structure with nothing to inherit it from.

Renamed in 0.0.2.9011

alpha previously held the pruned β and support_data was named X_sv, which made coef(fit)\$alpha mean the length-n_sv β here but the length-N dual α on a fit from the superseded psvr(): one generic returning two different vectors under one name, silently, depending on entry point. The β -under-alpha meaning is the one 0.0.2.9004 moved away from on the object itself; this aligns coef() with it.

coef.psvr_mape_sym	<i>Extract coefficients from a psvr_mape_sym model</i>
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Description

Extract coefficients from a psvr_mape_sym model

Usage

```
## S3 method for class 'psvr_mape_sym'
coef(object, ...)
```

Arguments

object	An object of class "psvr_mape_sym".
...	Ignored.

Value

A named list with five components, identical in meaning to `coef.psvr_mape()`:

alpha, alpha_star The length-N pre-pruning dual variables α_k and α_k^* .

beta The pruned dual differences $\beta_k = \alpha_k - \alpha_k^*$ over the support-vector indices (length n_sv); this is what `predict()` uses.

b Bias term.

support_data Support vector input matrix.

Renamed in 0.0.2.9011

See `coef.psvr_mape()` — the same rename, for the same reason, applied to both MAPE classes together.

coef.psvr_rmspe	<i>Extract coefficients from a psvr_rmspe model</i>
-----------------	---

Description

Extract coefficients from a psvr_rmspe model

Usage

```
## S3 method for class 'psvr_rmspe'
coef(object, ...)
```

Arguments

object	An object of class "psvr_rmspe".
...	Ignored.

Value

A named list with components:

alpha Dual variables / Lagrange multipliers (length N).

b Bias term.

support_data Training input matrix (all N observations).

Three components, against five for the MAPE classes (`coef.psvr_mape()`): LS-SVR has no `alpha_star` and no pruned beta, and they are **not** materialised as NULL. So `names(coef(fit))` depends on the model family, which is a decision rather than an oversight – each class is family-specific, and inventing empty slots to make the two agree would add structure with nothing to inherit it from. `$alpha_star` and `$beta` yield NULL on both, so every accessor still agrees.

Renamed in 0.0.2.9011

`support_data` was named `X_sv`. LS-SVR performs no pruning — every training point contributes to $f(x)$ — so there are no support vectors to name: it was an epsilon-SVR name on an LS-SVR value. The value is unchanged.

<code>coef.psvr_rmspe_sym</code>	<i>Extract coefficients from a <code>psvr_rmspe_sym</code> model</i>
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Description

Extract coefficients from a `psvr_rmspe_sym` model

Usage

```
## S3 method for class 'psvr_rmspe_sym'
coef(object, ...)
```

Arguments

object	An object of class "psvr_rmspe_sym".
...	Ignored.

Value

A named list with components:

alpha Dual variables / Lagrange multipliers (length N).

b Bias term.

support_data Training input matrix (all N observations).

Three components, not the five the MAPE classes return: LS-SVR has no alpha_star and no pruned beta, and they are not materialised as NULL. See [coef.psvr_rmspe\(\)](#).

Renamed in 0.0.2.9011

See [coef.psvr_rmspe\(\)](#) — the same rename, for the same reason, applied to both LS-SVR classes together.

cost_psvr	<i>Cost parameter with extended range for psvr models</i>
-----------	---

Description

A dials parameter for the regularisation parameter in psvr models. The default range [-2, 10] on the log2 scale (corresponding to approximately 0.25 to 1024) is wider than `dials::cost()` to accommodate the larger regularisation values typically needed by LS-SVR models.

Usage

```
cost_psvr(range = c(-2, 10), trans = scales::log2_trans())
```

Arguments

range Numeric vector of length 2 on the log2 scale. Default `c(-2, 10)`.

trans A scales transformation object. Default `scales::log2_trans()`.

Details

For LS-SVR (m3, m4) models the static range is still frequently too narrow, because Γ enters the LS-SVR system through the y_k^2/Γ diagonal and so scales with $\text{var}(y) * N$ rather than sitting at a fixed magnitude. Prefer [cost_psvr_ls_data\(\)](#), whose range is computed from the training outcome.

Value

A quant_param dials object.

See Also

[cost_psvr_ls_data\(\)](#) for a data-driven LS-SVR variant.

Examples

```
cost_psvr()
```

cost_psvr_ls_data	<i>Data-driven cost range for LS-SVR psvr models</i>
-------------------	--

Description

Returns a `quant_param` whose search range scales with $\text{var}(y) * N$, the standard heuristic for the LS-SVR regularisation parameter Γ (Suykens et al. 2002, *Least Squares Support Vector Machines*, §3.1.3). On the log2 scale, the lower bound is -2 (i.e. $\Gamma \geq 0.25$) and the upper bound is $\log_2(\text{var}(y) * N) + \text{width_log2}$.

Usage

```
cost_psvr_ls_data(y, n = length(y), width_log2 = 4)
```

Arguments

<code>y</code>	Numeric vector of strictly positive training targets.
<code>n</code>	Sample size. Default <code>length(y)</code> .
<code>width_log2</code>	Scalar giving the half-width (in log2 units) added above $\log_2(\text{var}(y) * n)$ to set the upper bound. Default 4 (about 16× headroom). Negative values are accepted with a warning, since the resulting upper bound falls below the $\text{var}(y) * n$ heuristic and is unlikely to be useful.

Details

The default `width_log2 = 4` places the upper bound about 16 times above $\text{var}(y) * n$, which is headroom for a Bayesian or grid search to work in without pinning against the boundary. Because the bound tracks $\text{var}(y) * n$, it moves with the outcome: the static `cost_psvr()` range [-2, 10] (i.e. $\Gamma \leq 1024$) is fixed, so it falls short whenever $\text{var}(y) * n$ exceeds a few hundred — which is the usual case, not the exception.

Use this function for m3 (LS-SVR) and m4 (symmetric LS-SVR) workflows. Stick to `cost_psvr()` for m1/m2 (ϵ -SVR), where cost maps to C and typical optima lie in [10, 100].

Value

A `quant_param` dial object.

See Also

`cost_psvr()`, `psvr_option_add_cost_ls()`

Examples

```
cost_psvr_ls_data(c(10, 20, 30, 40, 50))
```

make_kernel	Create a kernel function
-------------	--------------------------

Description

Returns a closure $K(x_i, x_j)$ suitable for use with all four psvr model fitting functions. The returned function accepts numeric vectors of any sign, which is required by the symmetric models (Models 2 and 4) that evaluate $K(x_k, -x_l)$.

Usage

```
make_kernel(
  type = c("rbf", "linear", "polynomial"),
  sigma = 1,
  degree = 3L,
  coef0 = 1
)
```

Arguments

type	Kernel type: "rbf", "linear", or "polynomial".
sigma	Bandwidth for the RBF kernel, $\sigma > 0$ (default 1).
degree	Integer degree for the polynomial kernel, $\text{degree} \geq 1$ (default 3).
coef0	Constant term for the polynomial kernel (default 1).

Details

The three supported kernels are:

- **RBF:** $K(x_i, x_j) = \exp(-\|x_i - x_j\|^2 / (2 * \sigma^2))$
- **Linear:** $K(x_i, x_j) = x_i \cdot x_j$
- **Polynomial:** $K(x_i, x_j) = (x_i \cdot x_j + \text{coef0})^{\text{degree}}$

RBF and even-degree polynomial kernels satisfy Assumption 3 of the paper (kernel symmetry), making them compatible with the symmetric models. The linear kernel and odd-degree polynomial kernels do **not** satisfy Assumption 3 and should not be used with the symmetric models, i.e. with `psvr_mape()` / `psvr_rmspe()` or a `parsnip` spec whose `sym_type` is "even" or "odd".

Value

A function $K(x_i, x_j)$ where x_i and x_j are numeric vectors of the same length, returning a scalar kernel evaluation.

Examples

```
K <- make_kernel("rbf", sigma = 0.5)
K(c(1, 2), c(3, 4))

K_lin <- make_kernel("linear")
K_lin(c(1, 0), c(0, 1))

K_poly <- make_kernel("polynomial", degree = 2, coef0 = 1)
K_poly(c(1, 2), c(3, 4))
```

margin_percentage	<i>Insensitivity margin in percentage units</i>
-------------------	---

Description

A dials parameter for the epsilon tube half-width in psvr MAPE models. Unlike `dials::svm_margin()` which uses absolute units, this parameter is expressed as a percentage of each target value. The default range [1, 20] means the insensitivity tube spans 1% to 20% of each target.

Usage

```
margin_percentage(range = c(1, 20), trans = NULL)
```

Arguments

range	Numeric vector of length 2. Default <code>c(1, 20)</code> .
trans	A scales transformation object. Default <code>NULL</code> .

Value

A `quant_param` dials object.

Examples

```
margin_percentage()
margin_percentage(range = c(0.5, 10))
```

predict.psvr_mape	<i>Predict from a fitted epsilon-SVR with MAPE model</i>
-------------------	--

Description

Method dispatched on the "psvr_mape" class, which both `psvr_mape()` and the parsnip engine fit wrappers return.

Usage

```
## S3 method for class 'psvr_mape'
predict(object, newdata, ...)
```

Arguments

object	An object of class "psvr_mape", as returned by <code>psvr_mape()</code> with <code>sym_type = "none"</code> or by the parsnip engine fit wrappers (see <code>psvr-fit-wrappers</code> ; <code>sym_type = "even"</code> or <code>"odd"</code> yields "psvr_mape_sym" instead). Unwrap a parsnip fit with <code>parsonip::extract_fit_engine()</code> to obtain it.
newdata	Numeric matrix of new inputs, one observation per row ($M \times p$).
...	Ignored.

Value

Numeric vector of length M with predicted values.

predict.psvr_mape_sym	<i>Predict from a fitted symmetric epsilon-SVR with MAPE model</i>
-----------------------	--

Description

Method dispatched on the "psvr_mape_sym" class, which both `psvr_mape()` and the parsnip engine fit wrappers return. Uses the symmetric representer theorem

$$f(x) = \frac{1}{2} \sum_k \beta_k K_s(x_k, x) + b$$

with $K_s(x_k, x) = K(x_k, x) + aK(x_k, -x)$.

Usage

```
## S3 method for class 'psvr_mape_sym'
predict(object, newdata, ...)
```

Arguments

object	An object of class "psvr_mape_sym", as returned by <code>psvr_mape()</code> with <code>sym_type = "even"</code> or <code>"odd"</code> , or by the parsnip engine fit wrappers (see psvr-fit-wrappers ; <code>sym_type = "none"</code> yields "psvr_mape" instead). Unwrap a parsnip fit with <code>parsnip::extract_fit_engine()</code> to obtain it.
newdata	Numeric matrix of new inputs, one observation per row ($M \times p$).
...	Ignored.

Value

Numeric vector of length M with predicted values.

predict.psvr_rmspe	<i>Predict from a fitted LS-SVR with RMSPE model</i>
--------------------	--

Description

Method dispatched on the "psvr_rmspe" class, which both `psvr_rmspe()` and the parsnip engine fit wrappers return.

Usage

```
## S3 method for class 'psvr_rmspe'
predict(object, newdata, ...)
```

Arguments

object	An object of class "psvr_rmspe", as returned by <code>psvr_rmspe()</code> with <code>sym_type = "none"</code> or by the parsnip engine fit wrappers (see psvr-fit-wrappers ; <code>sym_type = "even"</code> or <code>"odd"</code> yields "psvr_rmspe_sym" instead). Unwrap a parsnip fit with <code>parsnip::extract_fit_engine()</code> to obtain it.
newdata	Numeric matrix of new inputs, one observation per row ($M \times p$).
...	Ignored.

Value

Numeric vector of length M with predicted values.

predict.psvr_rmspe_sym

Predict from a fitted symmetric LS-SVR with RMSPE model

Description

Method dispatched on the "psvr_rmspe_sym" class, which both [psvr_rmspe\(\)](#) and the parsnip engine fit wrappers return. Uses the symmetric representer

$$f(x) = \sum_k \alpha_k \cdot \frac{1}{2} (K(x_k, x) + aK(x_k, -x)) + b$$

Usage

```
## S3 method for class 'psvr_rmspe_sym'
predict(object, newdata, ...)
```

Arguments

object	An object of class "psvr_rmspe_sym", as returned by psvr_rmspe() with sym_type = "even" or "odd", or by the parsnip engine fit wrappers (see psvr-fit-wrappers ; sym_type = "none" yields "psvr_rmspe" instead). Unwrap a parsnip fit with parsnip::extract_fit_engine() to obtain it.
newdata	Numeric matrix of new inputs, one observation per row ($M \times p$).
...	Ignored.

Value

Numeric vector of length M with predicted values.

print.psvr_mape

Print method for psvr_mape objects

Description

Print method for psvr_mape objects

Usage

```
## S3 method for class 'psvr_mape'
print(x, ...)
```

Arguments

x	An object of class "psvr_mape".
...	Ignored.

Value

x, invisibly.

print.psvr_mape_sym	<i>Print method for psvr_mape_sym objects</i>
---------------------	---

Description

Print method for psvr_mape_sym objects

Usage

```
## S3 method for class 'psvr_mape_sym'
print(x, ...)
```

Arguments

x	An object of class "psvr_mape_sym".
...	Ignored.

Value

x, invisibly.

print.psvr_rmspe	<i>Print method for psvr_rmspe objects</i>
------------------	--

Description

Print method for psvr_rmspe objects

Usage

```
## S3 method for class 'psvr_rmspe'
print(x, ...)
```

Arguments

x	An object of class "psvr_rmspe".
...	Ignored.

Value

x, invisibly.

```
print.psvr_rmspe_sym    Print method for psvr_rmspe_sym objects
```

Description

Print method for psvr_rmspe_sym objects

Usage

```
## S3 method for class 'psvr_rmspe_sym'
print(x, ...)
```

Arguments

x An object of class "psvr_rmspe_sym".
 ... Ignored.

Value

x, invisibly.

```
psvr-fitted            Extract training fitted values from a psvr model
```

Description

Returns the length- N in-sample predictions $f(x_k)$ recorded when the model was fitted. No kernel matrix is rebuilt: the values are recovered from state the solver already holds. For the MAPE models that is a matvec against the retained Ω ; for the LS-SVR models it is the KKT stationarity identity

$$f(x_k) = y_k - (10^{-6} + y_k^2/\Gamma) \alpha_k$$

which costs $O(N)$ and holds in both preconditioner branches. The training inputs X are not retained for this purpose.

Usage

```
## S3 method for class 'psvr_mape'
fitted(object, ...)

## S3 method for class 'psvr_mape_sym'
fitted(object, ...)

## S3 method for class 'psvr_rmspe'
fitted(object, ...)

## S3 method for class 'psvr_rmspe_sym'
fitted(object, ...)
```

Arguments

object	A fitted object of class "psvr_mape", "psvr_mape_sym", "psvr_rmspe" or "psvr_rmspe_sym", from <code>psvr_mape()</code> , <code>psvr_rmspe()</code> , or a parsnip fit unwrapped with <code>parsnip::extract_fit_engine()</code> .
...	Ignored.

Details

The result equals `predict(object, X_train)` to machine precision. It is not bit-identical: the two use different summation orders (a BLAS matvec versus the column-wise reduction in `predict()`), and for the LS-SVR models the identity above is exact only up to the residual of the linear solve. Observed agreement is within $3e-12$ relative across the four models.

Value

Numeric vector of length N (the number of training observations), in training-row order.

Not reachable through parsnip

`parsnip` registers neither `residuals.model_fit` nor `fitted.model_fit`, so calling either generic on a `model_fit` dispatches to the stats default and returns NULL **silently** - no error, no warning. Reach the `psvr` object first with `parsnip::extract_fit_engine()`, then call `fitted()` or `residuals()` on that. `psvr` deliberately does not register S3 methods on `parsnip`'s class. Note also that `parsnip::augment()` recomputes predictions on whatever `new_data` it is given and reports response residuals only.

See Also

`residuals.psvr_mape()` and the other residuals methods

Examples

```
set.seed(1)
X <- matrix(runif(40, 0.5, 3), 20, 2)
y <- 2 + X[, 1]^2
fit <- psvr_rmspe(X, y, kernel = make_kernel("rbf"), gamma = 100)
head(fitted(fit))

# Through parsnip both generics return NULL on the model_fit wrapper;
# extract the engine object first.
df <- data.frame(x1 = X[, 1], x2 = X[, 2], y = y)
spec <- psvr_rmspe_rbf(cost = 10, rbf_sigma = 0.8)
pfit <- parsnip::fit(spec, y ~ x1 + x2, data = df)
fitted(pfit) # NULL
head(fitted(parsnip::extract_fit_engine(pfit))) # the fitted values
```

psvr-residuals

Extract training residuals from a psvr model

Description

Three residual types are available. They are **different quantities with different denominators**, not interchangeable scalings of one another:

Usage

```
## S3 method for class 'psvr_mape'
residuals(object, type = c("response", "percentage", "multiplicative"), ...)

## S3 method for class 'psvr_mape_sym'
residuals(object, type = c("response", "percentage", "multiplicative"), ...)

## S3 method for class 'psvr_rmspe'
residuals(object, type = c("response", "percentage", "multiplicative"), ...)

## S3 method for class 'psvr_rmspe_sym'
residuals(object, type = c("response", "percentage", "multiplicative"), ...)
```

Arguments

object	A fitted object of class "psvr_mape", "psvr_mape_sym", "psvr_rmspe" or "psvr_rmspe_sym", from <code>psvr_mape()</code> , <code>psvr_rmspe()</code> , or a parsnip fit unwrapped with <code>parsnip::extract_fit_engine()</code> .
type	One of "response" (default), "percentage", or "multiplicative". See above; the denominators differ.
...	Ignored.

Details

"response" $y - \hat{y}$. The default, following the R convention for `stats::residuals()`. Units of the response.

"percentage" $(y - \hat{y})/y$. Divides by the **observed target**. This is the per-observation contribution to the MAPE loss the epsilon-SVR models are fitted under, so it is the residual that corresponds to the estimated objective. `mean(abs(.)) * 100` is the training MAPE.

"multiplicative" $(y - \hat{y})/\hat{y}$. Divides by the **fitted value**. This is the $\hat{\eta}$ of the multiplicative-noise model $Y = f(x)(1 + \eta)$, under which $\eta = (Y - f(x))/f(x)$. Use it to inspect the assumed noise structure, e.g. checking whether $\hat{\eta}$ is homoscedastic and centred at zero.

The denominators differ and the choice matters: "percentage" divides by the observed target because that is what the MAPE loss does, "multiplicative" divides by the fitted value because that is what the noise model does. They agree only when $y = \hat{y}$, and diverge as the fit degrades. Choose by the question being asked: the loss actually minimised ("percentage"), or the noise model assumed ("multiplicative").

Value

Numeric vector of length N (the number of training observations), in training-row order.

Near-zero fitted values

"multiplicative" divides by $\hat{y}$, which an SVR does not constrain to be positive even though the targets are. Where $\hat{y}$ is at or below $\sqrt{.Machine\$double.eps} * \text{mean}(\text{abs}(y))$ - including zero and negative fitted values - the ratio is inflated, infinite, or sign-flipped.

This function does **not** drop those observations: it always returns exactly N values in training-row order, so the result stays aligned with `y_train`, `fitted()`, and the training rows. The raw value (possibly Inf or NaN) is returned and a single warning reports how many observations are affected.

For *pooled* diagnostics (a mean multiplicative error, a variance, a histogram) the affected observations must be excluded, or one near-zero fitted value dominates the summary. This follows the standard treatment of percentage errors near zero (Makridakis): screen the fitted values against a threshold and drop the observations below it before pooling, rather than altering the per-observation values. Exclude at the point of aggregation, e.g.:

```
e <- residuals(fit, type = "multiplicative")
mean(e[is.finite(e)])
```

Not reachable through parsnip

parsonip registers neither `residuals.model_fit` nor `fitted.model_fit`, so calling either generic on a `model_fit` dispatches to the stats default and returns NULL **silently** - no error, no warning. Reach the `psvr` object first with `parsonip::extract_fit_engine()`, then call `residuals()` or `fitted()` on that. `psvr` deliberately does not register S3 methods on `parsonip`'s class. Note also that `parsonip::augment()` recomputes predictions on whatever `new_data` it is given and reports response residuals only, so it is not a substitute for the "percentage" and "multiplicative" types.

See Also

`fitted.psvr_mape()` and the other fitted methods

Examples

```
set.seed(1)
X <- matrix(runif(40, 0.5, 3), 20, 2)
y <- 2 + X[, 1]^2
fit <- psvr_rmspe(X, y, kernel = make_kernel("rbf"), gamma = 100)
head(residuals(fit))
mean(abs(residuals(fit, type = "percentage"))) * 100 # training MAPE

# Through parsnip both generics return NULL on the model_fit wrapper;
# extract the engine object first.
df <- data.frame(x1 = X[, 1], x2 = X[, 2], y = y)
spec <- psvr_rmspe_rbf(cost = 10, rbf_sigma = 0.8)
pfit <- parsnip::fit(spec, y ~ x1 + x2, data = df)
residuals(pfit) # NULL
```

```
head(residuals(parsnip::extract_fit_engine(pfit))) # the residuals
```

psvr_cv

Cross-validate psvr_mape() with automatic warm-start across folds

Description

Fits a `psvr_mape()` model on each split in `splits`, carrying the converged dual variables (`alpha`, `alpha_star`) from one fold into the next as the SMO warm-start, so each solve starts from the previous fold's optimum instead of from zero. Because consecutive folds share most of their training rows, that starting point is already close to feasible; before each solve the carried vectors are projected back onto the constraint set (the equality $\sum_k \beta_k = 0$ and the per-sample box), with the residual violation absorbed by the rows that are new to this fold. The warm-start procedure is Algorithm 1 of arXiv:2605.01446 v3. Returns a tibble with one row per fold.

Usage

```
psvr_cv(
  splits,
  ...,
  X_var = NULL,
  y_var = NULL,
  warm_start = TRUE,
  verbose = FALSE
)
```

Arguments

<code>splits</code>	Either an <code>rsample::rset</code> object (e.g. from <code>rsample::vfold_cv()</code>), or a list of named lists each containing <code>analysis</code> (data frame), <code>assessment</code> (data frame), and optionally <code>row_ids</code> (integer vector of original training-row indices used for warm-start alignment across folds; defaults to positional).
<code>...</code>	Arguments forwarded to <code>psvr_mape()</code> . Must specify kernel and the MAPE hyperparameters (<code>C</code> , <code>eps</code>). <code>alpha_init</code> and <code>alpha_star_init</code> are managed internally; supplying them via <code>...</code> is an error, and so is <code>loss</code> , which is not an argument of <code>psvr_mape()</code> .
<code>X_var</code>	Character vector of predictor column names.
<code>y_var</code>	Single character giving the target column name.
<code>warm_start</code>	Logical; if <code>FALSE</code> , each fold fits cold-start (useful for benchmarking the T5 speedup).
<code>verbose</code>	Logical; if <code>TRUE</code> , report per-fold progress via <code>message()</code> (suppressible with <code>suppressMessages()</code>).

Details

This helper is **MAPE-only**, and there is no loss argument. That is a limitation of the implementation, not of the method: only `psvr_mape()` was ever wired to it. LS-SVR cross-validates perfectly well, it simply has no carryover state to exploit (each fold is a single linear-system solve), so for `psvr_rmspe()` use `tune::tune_grid()` with parallel cold-start.

Value

A tibble with one row per split and columns:

`split_id` 1-based fold index.

`fit` A list-column of `psvr_mape` objects, or `psvr_mape_sym` when `sym_type` is "even" or "odd".

`predictions` A list-column of numeric vectors (predictions on the assessment set).

`metrics` A list-column of named numeric vectors (`mape`, `rmspe`, `mse`, `r2`).

`iter_count` Integer; SMO iterations from `fit$iterations`.

`elapsed_sec` Numeric; wall-clock seconds for the fit.

`warm_started` Logical; TRUE for fold > 1 when `warm_start` = TRUE.

Examples

```
if (requireNamespace("rsample", quietly = TRUE) &&
    requireNamespace("tibble", quietly = TRUE)) {
  set.seed(2026)
  d <- data.frame(
    y = stats::rlnorm(80, sdlog = 1.0),
    x1 = stats::rnorm(80),
    x2 = stats::rnorm(80)
  )
  folds <- rsample::vfold_cv(d, v = 5)
  res <- psvr_cv(folds, X_var = c("x1", "x2"), y_var = "y",
    kernel = make_kernel("rbf", sigma = 1),
    C = 10, eps = 5)
  median(vapply(res$metrics, function(m) m[["mape"]], numeric(1)))
}
```

psvr_mape

Fit an epsilon-SVR with MAPE loss

Description

Fits the percentage-error epsilon-SVR of the paper: Model 1 when `sym_type` = "none", and the symmetric-kernel Model 2 when `sym_type` is "even" or "odd". The dual is a quadratic program with the sample-dependent box $|\beta_k| \leq 100C/y_k$ and $\sum_k \beta_k = 0$, solved by the built-in SMO loop or by `osqp`.

Usage

```
psvr_mape(
  X,
  y,
  sym_type = c("none", "even", "odd"),
  kernel,
  C,
  eps,
  solver = c("smo", "osqp"),
  tol = 0.001,
  max_iter = 100000L,
  alpha_init = NULL,
  alpha_star_init = NULL,
  warm_start_check = TRUE,
  block_k4_enabled = TRUE,
  engine = c("rcpp", "r"),
  ...,
  alpha_couple = 0.5,
  precomputed_Omega = NULL,
  precomputed_Omega_s = NULL
)
```

Arguments

<code>X</code>	Numeric matrix of training inputs, one observation per row ($N \times p$).
<code>y</code>	Numeric vector of training targets, length N . Must satisfy $y_k > 0$ for every k ; percentage-error loss is undefined otherwise, and this is checked rather than coerced.
<code>sym_type</code>	Symmetry type, one of "none" (default), "even" or "odd". Maps onto the symmetry parameter a of the paper: "none" fits Model 1 and imposes no symmetry constraint; "even" sets $a = +1$, enforcing $f(x) = f(-x)$; "odd" sets $a = -1$, enforcing $f(x) = -f(-x)$. This is the same vocabulary as the <code>sym_type</code> argument of the <code>parsnip</code> specifications, so the two public surfaces agree. The symmetric variants require a kernel satisfying Assumption 3 of the paper – see make_kernel() .
<code>kernel</code>	A kernel function created by make_kernel() .
<code>C</code>	Regularization parameter, $C > 0$. Required.
<code>eps</code>	Insensitivity tube half-width in percentage units, $\epsilon \geq 0$. Required.
<code>solver</code>	Backend for the dual quadratic program, "smo" (default) or "osqp". See the section above.
<code>tol</code>	Numerical tolerance, default 1e-3. Its meaning depends on solver. Under "smo" it is the convergence tolerance of the SMO loop, and tighter values produce more iterations. Under "osqp" the solver runs at its own fixed tolerances and <code>tol</code> is used only afterwards, as the threshold below which a dual variable counts as zero when identifying free, saturated and support-vector sets.

max_iter	Maximum SMO iterations, default 100000L. The solver emits a warning() and returns converged = FALSE if it does not converge within max_iter. Ignored for solver = "osqp".
alpha_init, alpha_star_init	Optional warm-start vectors for the SMO solver, each a finite numeric vector of length N . Projected onto $\sum_k (\alpha_k - \alpha_k^*) = 0$ intersected with the per-sample box $[0, 100C/y_k]$ before the solve. NULL (default) cold-starts. <code>psvr_cv()</code> manages these automatically across folds.
warm_start_check	Logical; if TRUE (default), validate post-projection feasibility and stop() on violation. A surviving equality residual is fatal rather than cosmetic: SMO conserves $\sum_k (\alpha_k - \alpha_k^*)$, so an infeasible start is carried through to the returned solution.
block_k4_enabled	Logical; if TRUE (default), enable the block-k=4 SMO inner loop. Each outer iteration may select a second working pair and apply a two-dimensional joint update when the descent-guaranteed decoupling criterion holds. FALSE restores the k=2 behaviour bit-identically.
engine	One of "rcpp" (default) or "r". Selects the SMO backend: the C++ core, or the R reference implementation. Both produce bit-identical results on the development toolchain; "r" is retained as the reference and will be deprecated in 0.0.4.0 and removed in 0.1.0.
...	Must be empty. Present only so that alpha_couple, precomputed_Omega and precomputed_Omega_s must be matched by their exact names rather than by position or partial matching – without it precomputed_Omega would be a partial-match prefix of precomputed_Omega_s. Passing anything here is an error, which is how a mistyped argument name is caught.
alpha_couple	Numeric in $[0, 1]$, default 0.5. Coupling penalty in the second-pair selection score gain $\times (1 - \alpha_{\text{couple}} \cdot \text{coupling})$. Exposed for empirical tuning; rarely needs adjustment. Ignored when block_k4_enabled = FALSE.
precomputed_Omega, precomputed_Omega_s	INTERNAL – used by <code>psvr_cv()</code> to share one full-dataset kernel matrix across folds. Users should not set these. precomputed_Omega applies when sym_type = "none", precomputed_Omega_s otherwise.

Details

For the least-squares / RMSPE family (Models 3 and 4) see `psvr_rmspe()`. The two are deliberately separate functions: they share no solver, no dual structure and no hyperparameter search space, so a single signature would make most of its own arguments conditional. The name `psvr()` is reserved for a future automatic-selection front end and is **not** a synonym for either.

Value

For `sym_type = "none"`, an object of class "psvr_mape": a list with components beta (the pruned dual differences $\beta = \alpha - \alpha^*$ over the support-vector indices, used by `predict()`), alpha and alpha_star (the length- N pre-pruning duals, retained for warm starts), b, X_sv, y_sv, y_train, fitted_values, kernel, C, eps, n_train, p_train, iterations, converged and block_k4.

For `sym_type = "even"` or `"odd"`, an object of class `"psvr_mape_sym"`: the same components plus a (the symmetry parameter) and `spectral` (adaptive spectral-shift diagnostics).

Methods are available for `predict()`, `print()`, `coef()`, `summary()`, `fitted()` and `residuals()`.

Choosing a solver

`solver = "smo"` (the default) uses the built-in sequential minimal optimisation loop. On some problems it does **not** converge within `max_iter`: it emits a `warning()`, `converged` is `FALSE` and `iterations` reaches the cap. This was first observed with linear and polynomial kernels, but it also occurs with the RBF kernel, so the kernel is **not** what determines it, and what does has not been established. Check `converged` on the returned fit, and refit with `solver = "osqp"` when it is `FALSE` and accuracy matters.

See Also

`psvr_rmspe()` for the LS-SVR / RMSPE family, `psvr_cv()` for cross-validation with warm-start carryover, `make_kernel()` for kernels.

Examples

```
set.seed(1)
X <- matrix(rnorm(40), 20, 2)
y <- rlnorm(20)
K <- make_kernel("rbf", sigma = 1)

fit <- psvr_mape(X, y, kernel = K, C = 10, eps = 5)
predict(fit, X[1:3, , drop = FALSE])

# Even-symmetric variant (Model 2): f(x) = f(-x).
fit_sym <- psvr_mape(X, y, sym_type = "even", kernel = K, C = 10, eps = 5)
predict(fit_sym, X[1:3, , drop = FALSE])
```

psvr_mape_specs

Parsnip model specs: epsilon-SVR with MAPE loss (Model 1)

Description

Create parsnip model specifications for `psvr_mape()` with a fixed kernel type. Kernel parameters are tunable parsnip arguments; the symmetry parameter `a` and solver tolerance are engine arguments passed via `set_engine()`.

Usage

```
psvr_mape_rbf(
  mode = "regression",
  engine = "psvr",
  cost = NULL,
```

```

    margin = NULL,
    rbf_sigma = NULL,
    sym_type = NULL
  )

psvr_mape_poly(
  mode = "regression",
  engine = "psvr",
  cost = NULL,
  margin = NULL,
  degree = NULL,
  scale_factor = NULL,
  sym_type = NULL
)

psvr_mape_linear(
  mode = "regression",
  engine = "psvr",
  cost = NULL,
  margin = NULL,
  sym_type = NULL
)

```

Arguments

mode	Only "regression" is supported.
engine	Only "psvr" is available.
cost	Regularization parameter $C > 0$. Use <code>hardhat::tune()</code> to optimize. Mapped to <code>cost_psvr()</code> , whose default range is $[-2, 10]$ on the log2 scale (about 0.25 to 1024). That range is adequate here, where typical ϵ -SVR optima lie in $[10, 100]$. The LS-SVR specs (<code>psvr_rmspe_specs</code>) map cost to Γ and need a much wider range; see there.
margin	Epsilon tube half-width $\epsilon \geq 0$ expressed as a percentage of each target value. Use <code>hardhat::tune()</code> to optimize. Mapped to <code>margin_percentage()</code> with default range $[1, 20]$ (percentage units). Named to match <code>parsnip::svm_rbf()</code> ; note the units differ from <code>dials::svm_margin()</code> , which is absolute rather than percentage.
rbf_sigma	RBF bandwidth $\sigma > 0$. Use <code>hardhat::tune()</code> to optimize. Mapped to <code>rbf_sigma_psvr()</code> , whose default range $[-3, 1]$ on the log10 scale is a fixed, conservative fall-back. The range does not finalize automatically from the training data. <code>rbf_sigma_psvr()</code> sets <code>finalize = NULL</code> , so <code>dials::finalize()</code> leaves it untouched. To centre the range on the data, pass <code>rbf_sigma_psvr_data()</code> computed on the preprocessed predictors explicitly — via <code>update()</code> on the extracted parameter set, or via <code>psvr_option_add()</code> for a workflow set. (RBF specs only.)
sym_type	Symmetry type: "none" (default) fits the non-symmetric ϵ -SVR of Model 1; "even" ($a = 1$) and "odd" ($a = -1$) fit the symmetric ϵ -SVR of Model 2. Use

	<code>hardhat::tune()</code> to optimise over the levels during CV; see <code>sym_type_param()</code> to restrict which levels are searched.
degree	Polynomial degree ≥ 1 . Use <code>hardhat::tune()</code> to optimize. (Polynomial specs only.)
scale_factor	Polynomial constant term (coef0). Use <code>hardhat::tune()</code> to optimize. (Polynomial specs only.)

Value

A parsnip model_spec object of the corresponding class.

Examples

```
library(parsnip)
spec <- psvr_rape_rbf(cost = 10, margin = 1, rbf_sigma = 1) |>
  set_engine("psvr")

spec_poly <- psvr_rape_poly(cost = 10, margin = 1, degree = 2,
  scale_factor = 1) |>
  set_engine("psvr")

spec_lin <- psvr_rape_linear(cost = 10, margin = 1) |>
  set_engine("psvr")

# Symmetric epsilon-SVR (Model 2) via the sym_type argument:
spec_sym <- psvr_rape_rbf(cost = 10, margin = 1, rbf_sigma = 1,
  sym_type = "even") |>
  set_engine("psvr")
```

psvr_option_add	<i>Apply data-driven rbf_sigma to all psvr workflows in a workflow set</i>
-----------------	--

Description

A convenience wrapper that calls `workflowsets::option_add()` for every psvr workflow in `wf_set` (those whose `wflow_id` contains "m1", "m2", "m3", or "m4"), replacing the `rbf_sigma` dial's parameter with a data-driven one built from `X` via `rbf_sigma_psvr_data()`.

Usage

```
psvr_option_add(wf_set, X, width = 10, sample_size = 500L, seed = NULL)
```

Arguments

<code>wf_set</code>	A <code>workflow_set</code> object.
<code>X</code>	A numeric matrix or data frame of preprocessed predictors.
<code>width</code>	Positive scalar. Passed to <code>rbf_sigma_psvr_data()</code> . Default 10.

`sample_size` Integer. Passed to [sigma_heuristic\(\)](#). Default 500L.

`seed` Integer seed for subsampling. Default NULL.

Value

The updated `workflow_set` (the same object with `option_add()` applied to each psvr workflow).

See Also

[rbf_sigma_psvr_data\(\)](#), [sigma_heuristic\(\)](#)

Examples

```
## Not run:
# After building wf_set and preprocessing:
train_baked <- rec |> prep() |> bake(new_data = train)
wf_set <- psvr_option_add(wf_set, train_baked |> select(-outcome))

## End(Not run)
```

`psvr_option_add_cost_ls`

Apply data-driven LS-SVR cost range to all m3/m4 workflows in a workflow set

Description

A convenience wrapper that calls [workflowsets::option_add\(\)](#) for every LS-SVR psvr workflow in `wf_set` (those whose `wflow_id` matches "m3" or "m4"), replacing the cost dials parameter with one built from `y` via [cost_psvr_ls_data\(\)](#).

Usage

```
psvr_option_add_cost_ls(wf_set, y, width_log2 = 4)
```

Arguments

`wf_set` A `workflow_set` object.

`y` Numeric vector of strictly positive training targets.

`width_log2` Passed to [cost_psvr_ls_data\(\)](#). Default 4.

Details

Workflows for m1/m2 (ϵ -SVR) are intentionally skipped — for those, cost maps to C and the static `cost_psvr()` range is usually adequate.

Note: `workflowsets::option_add()` replaces the whole `param_info` option for each matched workflow. If you also need a data-driven `rbf_sigma` (via `psvr_option_add()`), build the full `param_info` manually with `tune::extract_parameter_set_dials()` and call `workflowsets::option_add()` in one shot, or call this helper first and then `psvr_option_add()` (which only touches `rbf_sigma`) — the latter currently overwrites the former, so prefer the manual one-shot approach when both are needed.

Value

The updated `workflow_set`.

See Also

`cost_psvr_ls_data()`, `psvr_option_add()`

Examples

```
## Not run:
# After building wf_set with at least one m3 or m4 workflow:
wf_set <- psvr_option_add_cost_ls(wf_set, y = train_df$y)

## End(Not run)
```

psvr_rmspe

Fit a least-squares SVR with RMSPE loss

Description

Fits the percentage-error LS-SVR of the paper: Model 3 when `sym_type = "none"`, and the symmetric-kernel Model 4 when `sym_type` is "even" or "odd". There is no quadratic program and no sparsity: the fit is a single solve of the $(N + 1) \times (N + 1)$ augmented linear system

$$\begin{pmatrix} 0 & 1^\top \\ 1 & \Omega + Y_\Gamma \end{pmatrix} \begin{pmatrix} b \\ \alpha \end{pmatrix} = \begin{pmatrix} 0 \\ y \end{pmatrix}$$

with $Y_\Gamma = \text{diag}(y_1^2/\Gamma, \dots, y_N^2/\Gamma)$, and Ω_s replacing Ω in the symmetric case. Every training point contributes to the prediction.

Usage

```
psvr_rmspe(
  X,
  y,
  sym_type = c("none", "even", "odd"),
  kernel,
  gamma,
  precondition = "auto",
  ...
)
```

Arguments

<code>X</code>	Numeric matrix of training inputs, one observation per row ($N \times p$).
<code>y</code>	Numeric vector of training targets, length N . Must satisfy $y_k > 0$ for every k ; percentage-error loss is undefined otherwise, and this is checked rather than coerced.
<code>sym_type</code>	Symmetry type, one of "none" (default), "even" or "odd". Maps onto the symmetry parameter a of the paper: "none" fits Model 3 and imposes no symmetry constraint; "even" sets $a = +1$, enforcing $f(x) = f(-x)$; "odd" sets $a = -1$, enforcing $f(x) = -f(-x)$. This is the same vocabulary as the <code>sym_type</code> argument of the <code>parsnip</code> specifications, so the two public surfaces agree. The symmetric variants require a kernel satisfying Assumption 3 of the paper – see make_kernel() .
<code>kernel</code>	A kernel function created by make_kernel() .
<code>gamma</code>	Regularization parameter $\Gamma > 0$. Required. Larger values weight the squared percentage residuals more heavily against the norm penalty.
<code>precondition</code>	One of "auto" (default), "always", "never", or a positive numeric threshold. Controls a symmetric rescaling of the linear system by $P = \text{diag}(1/y)$. Unpreconditioned, the system carries the target-weighted diagonal y_k^2/Γ , whose entries span the square of the target range; solving $P\Omega P$ instead replaces it with the constant $1/\Gamma$, so the diagonal no longer inflates the condition number when the targets differ by orders of magnitude. The multipliers are rescaled back on the way out, leaving the solution unchanged in exact arithmetic. "auto" applies it when the target ratio $\max(y)/\min(y)$ exceeds 10; a numeric value sets that threshold explicitly. Whether it fired is reported in <code>fit\$precondition_applied</code> .
<code>...</code>	Must be empty. Passing anything here is an error, which is how a mistyped argument name is caught.

Details

For the epsilon-SVR / MAPE family (Models 1 and 2) see [psvr_mape\(\)](#). The two are deliberately separate functions: they share no solver, no dual structure and no hyperparameter search space. The name `psvr()` is reserved for a future automatic-selection front end and is **not** a synonym for either.

Value

For `sym_type = "none"`, an object of class `"psvr_rmspe"`: a list with components `alpha` (the length- N multipliers), `b`, `X_train`, `y_train`, `fitted_values`, `kernel`, `gamma`, `n_train`, `p_train` and `precondition_applied`.

For `sym_type = "even"` or `"odd"`, an object of class `"psvr_rmspe_sym"`: the same components plus `a` (the symmetry parameter).

Methods are available for `predict()`, `print()`, `coef()`, `summary()`, `fitted()` and `residuals()`. Note that `coef()` returns three components here (`alpha`, `b`, `support_data`) against five for the MAPE classes: LS-SVR has no `alpha_star` and no pruned `beta`, and the absent components are not materialised as `NULL`.

See Also

`psvr_mape()` for the epsilon-SVR / MAPE family, `make_kernel()` for kernels.

Examples

```
set.seed(1)
X <- matrix(rnorm(40), 20, 2)
y <- rlnorm(20)
K <- make_kernel("rbf", sigma = 1)

fit <- psvr_rmspe(X, y, kernel = K, gamma = 100)
predict(fit, X[1:3, , drop = FALSE])

# Even-symmetric variant (Model 4):  $f(x) = f(-x)$ .
fit_sym <- psvr_rmspe(X, y, sym_type = "even", kernel = K, gamma = 100)
predict(fit_sym, X[1:3, , drop = FALSE])
```

psvr_rmspe_specs

Parsnip model specs: LS-SVR with RMSPE loss (Model 3)

Description

Create parsnip model specifications for `psvr_rmspe()` with a fixed kernel type. `cost` maps to the regularization parameter Γ .

Usage

```
psvr_rmspe_rbf(
  mode = "regression",
  engine = "psvr",
  cost = NULL,
  rbf_sigma = NULL,
  sym_type = NULL
)
```

```

psvr_rmspe_poly(
  mode = "regression",
  engine = "psvr",
  cost = NULL,
  degree = NULL,
  scale_factor = NULL,
  sym_type = NULL
)

psvr_rmspe_linear(
  mode = "regression",
  engine = "psvr",
  cost = NULL,
  sym_type = NULL
)

```

Arguments

mode	Only "regression" is supported.
engine	Only "psvr" is available.
cost	Regularization parameter $\Gamma > 0$. Use <code>hardhat::tune()</code> to optimize. Mapped to <code>cost_psvr()</code> , whose default range $[-2, 10]$ on the log2 scale ($\Gamma \leq 1024$) is the ϵ -SVR range and is too narrow for LS-SVR . Γ enters the LS-SVR system only through the y_k^2/Γ diagonal, so the value that balances that term against the kernel scales with the outcome's variance and with the sample size — the quantity <code>cost_psvr_ls_data()</code> computes. The LS-SVR optimum is therefore routinely orders of magnitude above the static ceiling, and a grid over the default is boundary-trapped whenever it is. Pass <code>cost_psvr_ls_data()</code> built from the training outcome explicitly, via <code>update()</code> on the extracted parameter set or via <code>psvr_option_add_cost_ls()</code> for a workflow set. This cannot be automated: <code>tune</code> finalizes parameters from the molded predictors only and never passes the outcome to <code>dials::finalize()</code> , so no <code>finalize</code> function on <code>cost</code> could compute it.
rbf_sigma	RBF bandwidth $\sigma > 0$. Use <code>hardhat::tune()</code> to optimize. Mapped to <code>rbf_sigma_psvr()</code> , whose default range $[-3, 1]$ on the log10 scale is a fixed, conservative fall-back. The range does not finalize automatically from the training data. <code>rbf_sigma_psvr()</code> sets <code>finalize = NULL</code> , so <code>dials::finalize()</code> leaves it untouched. To centre the range on the data, pass <code>rbf_sigma_psvr_data()</code> computed on the preprocessed predictors explicitly — via <code>update()</code> on the extracted parameter set, or via <code>psvr_option_add()</code> for a workflow set. (RBF specs only.)
sym_type	Symmetry type: "none" (default) fits the non-symmetric LS-SVR of Model 3; "even" ($a = 1$) and "odd" ($a = -1$) fit the symmetric LS-SVR of Model 4. Use <code>hardhat::tune()</code> to optimise over the levels during CV; see <code>sym_type_param()</code> to restrict which levels are searched.
degree	Polynomial degree ≥ 1 . Use <code>hardhat::tune()</code> to optimize. (Polynomial specs only.)

`scale_factor` Polynomial constant term (`coef0`). Use `hardhat::tune()` to optimize. (Polynomial specs only.)

Value

A parsnip `model_spec` object of the corresponding class.

Engine arguments

The precondition argument of `psvr_rmspe()` is exposed as a non-tunable engine argument. Pass it via `parsnip::set_engine()`, e.g. `set_engine("psvr", precondition = "always")`. Default is "auto". See `psvr_rmspe()` for accepted values and semantics.

Examples

```
library(parsnip)
spec <- psvr_rmspe_rbf(cost = 1000, rbf_sigma = 1) |>
  set_engine("psvr")

spec_poly <- psvr_rmspe_poly(cost = 1000, degree = 2, scale_factor = 1) |>
  set_engine("psvr")

spec_lin <- psvr_rmspe_linear(cost = 1000) |>
  set_engine("psvr")

# Symmetric LS-SVR (Model 4) via the sym_type argument:
spec_sym <- psvr_rmspe_rbf(cost = 1000, rbf_sigma = 1,
  sym_type = "even") |>
  set_engine("psvr")
```

rbf_sigma_psvr	<i>RBF sigma parameter for psvr models</i>
----------------	--

Description

A dial parameter for the RBF kernel bandwidth in psvr models. The default range `[-3, 1]` on the `log10` scale is a conservative fallback. For best results, override the range using `sigma_heuristic()` computed on the preprocessed training data:

Usage

```
rbf_sigma_psvr(range = c(-3, 1), trans = scales::log10_trans())
```

Arguments

`range` Numeric vector of length 2 on the `log10` scale. Default `c(-3, 1)`.

`trans` A scales transformation object. Default `scales::log10_trans()`.

Details

```

train_baked <- rec |> prep() |> bake(new_data = train)
sigma_med   <- sigma_heuristic(train_baked |> select(-outcome))
rbf_sigma_custom <- rbf_sigma_psvr(
  range = c(log10(sigma_med / 10), log10(sigma_med * 10))
)
# Then inject via option_add():
wf_set |> option_add(
  param_info = extract_parameter_set_dials(wf) |>
    update(rbf_sigma = rbf_sigma_custom),
  id = "your_workflow_id"
)

```

Value

A quant_param dials object.

See Also

[sigma_heuristic\(\)](#)

Examples

```

rbf_sigma_psvr()

# Override with data-driven range:
X <- matrix(rnorm(200), ncol = 4)
sigma_med <- sigma_heuristic(X)
rbf_sigma_psvr(range = c(log10(sigma_med / 10), log10(sigma_med * 10)))

```

rbf_sigma_psvr_data	<i>RBF sigma parameter with data-driven range for psvr models</i>
---------------------	---

Description

A convenience wrapper that combines [sigma_heuristic\(\)](#) and [rbf_sigma_psvr\(\)](#) in a single call. It computes the median pairwise distance from X and returns a quant_param whose search range spans one order of magnitude either side of the heuristic value on the log10 scale (i.e., $[\log_{10}(\text{sigma_med} / \text{width}), \log_{10}(\text{sigma_med} * \text{width})]$).

Usage

```

rbf_sigma_psvr_data(
  X,
  width = 10,
  sample_size = 500L,
  seed = NULL,

```

```

    trans = scales::log10_trans()
  )

```

Arguments

<code>X</code>	A numeric matrix or data frame of predictors (already preprocessed — centred, scaled, etc.).
<code>width</code>	Positive scalar. Multiplier that sets the half-width of the search range around the heuristic sigma. Default 10 (one decade).
<code>sample_size</code>	Integer. Passed to <code>sigma_heuristic()</code> . Default 500L.
<code>seed</code>	Integer seed for subsampling. Passed to <code>sigma_heuristic()</code> . Default NULL.
<code>trans</code>	A scales transformation object. Default <code>scales::log10_trans()</code> .

Value

A `quant_param` dials object with a data-driven search range.

See Also

`sigma_heuristic()`, `rbf_sigma_psvr()`

Examples

```

X <- matrix(rnorm(200), ncol = 4)
rbf_sigma_psvr_data(X)
rbf_sigma_psvr_data(X, width = 5)

```

<code>sigma_heuristic</code>	<i>Median-distance heuristic for RBF kernel bandwidth</i>
------------------------------	---

Description

Returns the median pairwise Euclidean distance between rows of `X`, which is a standard data-driven starting point for the RBF kernel bandwidth (Schölkopf & Smola, 2002). Use the result to define a sensible `rbf_sigma` search range centred on this value via `dials::rbf_sigma(range = c(log10(sigma / 10), log10(sigma * 10)))`.

Usage

```
sigma_heuristic(X, sample_size = 500L, seed = NULL)
```

Arguments

<code>X</code>	A numeric matrix or data frame of predictors (already preprocessed — centred, scaled, etc.).
<code>sample_size</code>	Integer. If <code>nrow(X) > sample_size</code> , a random subsample is used to avoid $O(n^2)$ memory cost on large datasets. Default 500L.
<code>seed</code>	Integer seed for the subsample. Default NULL. The caller's RNG stream is restored on exit, so passing <code>seed</code> makes the subsample reproducible without affecting the caller's subsequent random draws.

Value

A scalar numeric: the median pairwise Euclidean distance.

Examples

```
X <- matrix(rnorm(200), ncol = 4)
sigma_heuristic(X)
```

summary.psvr_mape	<i>Summarize a fitted epsilon-SVR with MAPE loss</i>
-------------------	--

Description

Prints the kernel, the training and support-vector counts, the hyperparameters, and the SMO iteration count with its convergence status. Every training point contributes for LS-SVR but not here: the support-vector percentage is the sparsity of the fit.

Usage

```
## S3 method for class 'psvr_mape'
summary(object, ...)
```

Arguments

<code>object</code>	An object of class "psvr_mape", from <code>psvr_mape()</code> with <code>sym_type = "none"</code> , or from a <code>parsnip</code> fit unwrapped with <code>parsnip::extract_fit_engine()</code> .
<code>...</code>	Ignored.

Value

`object`, invisibly. Called for the printed summary.

See Also

`print.psvr_mape()`, `coef.psvr_mape()`

Examples

```

set.seed(1)
X <- matrix(rnorm(40), 20, 2)
y <- rlnorm(20)
fit <- psvr_mape(X, y, kernel = make_kernel("rbf", sigma = 1),
                C = 10, eps = 5)
summary(fit)

```

summary.psvr_mape_sym *Summarize a fitted symmetric epsilon-SVR with MAPE loss*

Description

As [summary.psvr_mape\(\)](#), with the symmetry parameter *a* reported.

Usage

```

## S3 method for class 'psvr_mape_sym'
summary(object, ...)

```

Arguments

object	An object of class "psvr_mape_sym", from psvr_mape() with sym_type = "even" or "odd".
...	Ignored.

Value

object, invisibly. Called for the printed summary.

See Also

[print.psvr_mape_sym\(\)](#), [coef.psvr_mape_sym\(\)](#)

Examples

```

set.seed(1)
X <- matrix(rnorm(40), 20, 2)
y <- rlnorm(20)
fit <- psvr_mape(X, y, sym_type = "even",
                kernel = make_kernel("rbf", sigma = 1), C = 10, eps = 5)
summary(fit)

```

summary.psvr_rmspe	<i>Summarize a fitted LS-SVR with RMSPE loss</i>
--------------------	--

Description

Prints the kernel, the training count, the hyperparameter, and whether the `diag(1/y)` preconditioner fired. No support-vector count is reported: LS-SVR performs no pruning, so every training point contributes to the prediction.

Usage

```
## S3 method for class 'psvr_rmspe'
summary(object, ...)
```

Arguments

object	An object of class "psvr_rmspe", from <code>psvr_rmspe()</code> with <code>sym_type = "none"</code> , or from a parsnip fit unwrapped with <code>parsonip::extract_fit_engine()</code> .
...	Ignored.

Value

object, invisibly. Called for the printed summary.

See Also

`print.psvr_rmspe()`, `coef.psvr_rmspe()`

Examples

```
set.seed(1)
X <- matrix(rnorm(40), 20, 2)
y <- rlnorm(20)
fit <- psvr_rmspe(X, y, kernel = make_kernel("rbf", sigma = 1), gamma = 100)
summary(fit)
```

summary.psvr_rmspe_sym	<i>Summarize a fitted symmetric LS-SVR with RMSPE loss</i>
------------------------	--

Description

As `summary.psvr_rmspe()`, with the symmetry parameter *a* reported.

Usage

```
## S3 method for class 'psvr_rmspe_sym'
summary(object, ...)
```

Arguments

object An object of class "psvr_rmspe_sym", from `psvr_rmspe()` with `sym_type = "even"` or `"odd"`.

... Ignored.

Value

object, invisibly. Called for the printed summary.

See Also

`print.psvr_rmspe_sym()`, `coef.psvr_rmspe_sym()`

Examples

```
set.seed(1)
X <- matrix(rnorm(40), 20, 2)
y <- rlnorm(20)
fit <- psvr_rmspe(X, y, sym_type = "even",
                  kernel = make_kernel("rbf", sigma = 1), gamma = 100)
summary(fit)
```

sym_type_param

Dials parameter for symmetry type

Description

Returns a qualitative `dials::new_qual_param()` describing the `sym_type` argument of the `psvr` model specs. `"none"` fits the non-symmetric model; `"even"` maps to $a = 1L$ (standard symmetric kernel); `"odd"` maps to $a = -1L$ (anti-symmetric kernel).

Usage

```
sym_type_param(values = c("none", "even", "odd"))
```

Arguments

values Character vector of levels to search over. Any subset of `c("none", "even", "odd")`; defaults to all three. Pass `values = c("even", "odd")` to tune over the symmetric models only, which reproduces the two-level grid offered before `psvr 0.0.2.9011`.

Value

A `qual_param` object.

Examples

```
sym_type_param()  
sym_type_param(values = c("even", "odd"))
```

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