

# Package ‘tfdatasets’

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**Type** Package

**Title** Interface to 'TensorFlow' Datasets

**Version** 2.4.0

**Description** Interface to 'TensorFlow' Datasets, a high-level library for building complex input pipelines from simple, re-usable pieces. See <<https://www.tensorflow.org/guide>> for additional details.

**License** Apache License 2.0

**URL** <https://github.com/rstudio/tfdatasets>

**BugReports** <https://github.com/rstudio/tfdatasets/issues>

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## R topics documented:

all_nominal . . . . .	3
all_numeric . . . . .	4
dataset_batch . . . . .	4
dataset_cache . . . . .	5
dataset_collect . . . . .	5
dataset_concatenate . . . . .	6
dataset_decode_delim . . . . .	7
dataset_filter . . . . .	7
dataset_flat_map . . . . .	8
dataset_interleave . . . . .	9
dataset_map . . . . .	10
dataset_map_and_batch . . . . .	11
dataset_padded_batch . . . . .	12
dataset_prefetch . . . . .	13
dataset_prefetch_to_device . . . . .	14
dataset_prepare . . . . .	15
dataset_reduce . . . . .	16
dataset_repeat . . . . .	17
dataset_shard . . . . .	17
dataset_shuffle . . . . .	18
dataset_shuffle_and_repeat . . . . .	19
dataset_skip . . . . .	19
dataset_take . . . . .	20
dataset_use_spec . . . . .	21
dataset_window . . . . .	22
delim_record_spec . . . . .	22
dense_features . . . . .	24
feature_spec . . . . .	24
file_list_dataset . . . . .	25
fit.FeatureSpec . . . . .	26
fixed_length_record_dataset . . . . .	27
has_type . . . . .	28
hearts . . . . .	29
input_fn.tf_dataset . . . . .	30
iterator_get_next . . . . .	31
iterator_initializer . . . . .	31
iterator_make_initializer . . . . .	32
iterator_string_handle . . . . .	32
layer_input_from_dataset . . . . .	33
make-iterator . . . . .	34
make_csv_dataset . . . . .	35
next_batch . . . . .	37

output_types . . . . .	39
range_dataset . . . . .	39
read_files . . . . .	40
sample_from_datasets . . . . .	41
scaler . . . . .	41
scaler_min_max . . . . .	42
scaler_standard . . . . .	42
selectors . . . . .	43
sparse_tensor_slices_dataset . . . . .	43
sql_record_spec . . . . .	44
steps . . . . .	45
step_bucketized_column . . . . .	45
step_categorical_column_with_hash_bucket . . . . .	46
step_categorical_column_with_identity . . . . .	48
step_categorical_column_with_vocabulary_file . . . . .	49
step_categorical_column_with_vocabulary_list . . . . .	50
step_crossed_column . . . . .	52
step_embedding_column . . . . .	53
step_indicator_column . . . . .	55
step_numeric_column . . . . .	56
step_remove_column . . . . .	57
step_shared_embeddings_column . . . . .	58
tensors_dataset . . . . .	60
tensor_slices_dataset . . . . .	60
text_line_dataset . . . . .	61
tfrecord_dataset . . . . .	61
until_out_of_range . . . . .	62
with_dataset . . . . .	63
zip_datasets . . . . .	64
<b>Index</b>	<b>65</b>

---

all_nominal	<i>Find all nominal variables.</i>
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---

### Description

Currently we only consider "string" type as nominal.

### Usage

```
all_nominal()
```

### See Also

Other Selectors: [all\\_numeric\(\)](#), [has\\_type\(\)](#)

---

`all_numeric`                      *Specify all numeric variables.*

---

### Description

Find all the variables with the following types: "float16", "float32", "float64", "int16", "int32", "int64", "half", "double".

### Usage

```
all_numeric()
```

### See Also

Other Selectors: [all\\_nominal\(\)](#), [has\\_type\(\)](#)

---

`dataset_batch`                      *Combines consecutive elements of this dataset into batches.*

---

### Description

Combines consecutive elements of this dataset into batches.

### Usage

```
dataset_batch(dataset, batch_size, drop_remainder = FALSE)
```

### Arguments

<code>dataset</code>	A dataset
<code>batch_size</code>	An integer, representing the number of consecutive elements of this dataset to combine in a single batch.
<code>drop_remainder</code>	Ensure that batches have a fixed size by omitting any final smaller batch if it's present. Note that this is required for use with the Keras tensor inputs to fit/evaluate/etc.

### Value

A dataset

### See Also

Other dataset methods: [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

dataset_cache	<i>Caches the elements in this dataset.</i>
---------------	---

---

**Description**

Caches the elements in this dataset.

**Usage**

```
dataset_cache(dataset, filename = NULL)
```

**Arguments**

dataset	A dataset
filename	String with the name of a directory on the filesystem to use for caching tensors in this Dataset. If a filename is not provided, the dataset will be cached in memory.

**Value**

A dataset

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

dataset_collect	<i>Collects a dataset</i>
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---

**Description**

Iterates through the dataset collecting every element into a list. It's useful for looking at the full result of the dataset. Note: You may run out of memory if your dataset is too big.

**Usage**

```
dataset_collect(dataset, iter_max = Inf)
```

**Arguments**

dataset	A dataset
iter_max	Maximum number of iterations. Inf until the end of the dataset

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

dataset\_concatenate     *Creates a dataset by concatenating given dataset with this dataset.*

---

**Description**

Creates a dataset by concatenating given dataset with this dataset.

**Usage**

```
dataset_concatenate(dataset, other)
```

**Arguments**

dataset	A dataset
other	Dataset to be concatenated

**Value**

A dataset

**Note**

Input dataset and dataset to be concatenated should have same nested structures and output types.

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

dataset\_decode\_delim    *Transform a dataset with delimited text lines into a dataset with named columns*

---

### Description

Transform a dataset with delimited text lines into a dataset with named columns

### Usage

```
dataset_decode_delim(dataset, record_spec, parallel_records = NULL)
```

### Arguments

dataset                Dataset containing delimited text lines (e.g. a CSV)

record\_spec            Specification of column names and types (see [delim\\_record\\_spec\(\)](#)).

parallel\_records  
(Optional) An integer, representing the number of records to decode in parallel.  
If not specified, records will be processed sequentially.

### See Also

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

dataset\_filter            *Filter a dataset by a predicate*

---

### Description

Filter a dataset by a predicate

### Usage

```
dataset_filter(dataset, predicate)
```

### Arguments

dataset                A dataset

predicate              A function mapping a nested structure of tensors (having shapes and types defined by [output\\_shapes\(\)](#) and [output\\_types\(\)](#)) to a scalar tf\$bool tensor.

**Details**

Note that the functions used inside the predicate must be tensor operations (e.g. `tf$not_equal`, `tf$less`, etc.). R generic methods for relational operators (e.g. `<`, `>`, `<=`, etc.) and logical operators (e.g. `!`, `&`, `|`, etc.) are provided so you can use shorthand syntax for most common comparisons (this is illustrated by the example below).

**Value**

A dataset composed of records that matched the predicate.

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

**Examples**

```
## Not run:

dataset <- text_line_dataset("mtcars.csv", record_spec = mtcars_spec) %>%
  dataset_filter(function(record) {
    record$mpg >= 20
  })

dataset <- text_line_dataset("mtcars.csv", record_spec = mtcars_spec) %>%
  dataset_filter(function(record) {
    record$mpg >= 20 & record$cyl >= 6L
  })

## End(Not run)
```

---

<code>dataset_flat_map</code>	<i>Maps map_func across this dataset and flattens the result.</i>
-------------------------------	---

---

**Description**

Maps `map_func` across this dataset and flattens the result.

**Usage**

```
dataset_flat_map(dataset, map_func)
```



**Arguments**

dataset	A dataset
map_func	A function mapping a nested structure of tensors (having shapes and types defined by <code>output_shapes()</code> and <code>output_types()</code> ) to a dataset.

**Value**

A dataset

---

dataset\_interleave      *Maps map\_func across this dataset, and interleaves the results*

---

**Description**

Maps map\_func across this dataset, and interleaves the results

**Usage**

```
dataset_interleave(dataset, map_func, cycle_length, block_length = 1)
```

**Arguments**

dataset	A dataset
map_func	A function mapping a nested structure of tensors (having shapes and types defined by <code>output_shapes()</code> and <code>output_types()</code> ) to a dataset.
cycle_length	The number of elements from this dataset that will be processed concurrently.
block_length	The number of consecutive elements to produce from each input element before cycling to another input element.

**Details**

The `cycle_length` and `block_length` arguments control the order in which elements are produced. `cycle_length` controls the number of input elements that are processed concurrently. In general, this transformation will apply `map_func` to `cycle_length` input elements, open iterators on the returned dataset objects, and cycle through them producing `block_length` consecutive elements from each iterator, and consuming the next input element each time it reaches the end of an iterator.

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

**Examples**

```
## Not run:

dataset <- tensor_slices_dataset(c(1,2,3,4,5)) %>%
  dataset_interleave(cycle_length = 2, block_length = 4, function(x) {
    tensors_dataset(x) %>%
      dataset_repeat(6)
  })

# resulting dataset (newlines indicate "block" boundaries):
c(1, 1, 1, 1,
  2, 2, 2, 2,
  1, 1,
  2, 2,
  3, 3, 3, 3,
  4, 4, 4, 4,
  3, 3,
  4, 4,
  5, 5, 5, 5,
  5, 5,
)

## End(Not run)
```

---

dataset\_map

*Map a function across a dataset.*


---

**Description**

Map a function across a dataset.

**Usage**

```
dataset_map(dataset, map_func, num_parallel_calls = NULL)
```

**Arguments**

dataset	A dataset
map_func	A function mapping a nested structure of tensors (having shapes and types defined by <code>output_shapes()</code> and <code>output_types()</code> ) to another nested structure of tensors. It also supports purrr style lambda functions powered by <code>rlang::as_function()</code> .
num_parallel_calls	(Optional) An integer, representing the number of elements to process in parallel. If not specified, elements will be processed sequentially.

**Value**

A dataset

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

`dataset_map_and_batch` *Fused implementation of `dataset_map()` and `dataset_batch()`*

---

**Description**

Maps ‘map\_func’ across `batch_size` consecutive elements of this dataset and then combines them into a batch. Functionally, it is equivalent to map followed by batch. However, by fusing the two transformations together, the implementation can be more efficient.

**Usage**

```
dataset_map_and_batch(
  dataset,
  map_func,
  batch_size,
  num_parallel_batches = NULL,
  drop_remainder = FALSE,
  num_parallel_calls = NULL
)
```

**Arguments**

<code>dataset</code>	A dataset
<code>map_func</code>	A function mapping a nested structure of tensors (having shapes and types defined by <a href="#">output_shapes()</a> and <a href="#">output_types()</a> ) to another nested structure of tensors. It also supports purrr style lambda functions powered by <a href="#">rlang::as_function()</a> .
<code>batch_size</code>	An integer, representing the number of consecutive elements of this dataset to combine in a single batch.
<code>num_parallel_batches</code>	(Optional) An integer, representing the number of batches to create in parallel. On one hand, higher values can help mitigate the effect of stragglers. On the other hand, higher values can increase contention if CPU is scarce.

- drop\_remainder Ensure that batches have a fixed size by omitting any final smaller batch if it's present. Note that this is required for use with the Keras tensor inputs to fit/evaluate/etc.
- num\_parallel\_calls  
(Optional) An integer, representing the number of elements to process in parallel. If not specified, elements will be processed sequentially.

### See Also

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

dataset\_padded\_batch *Combines consecutive elements of this dataset into padded batches*

---

### Description

This method combines multiple consecutive elements of this dataset, which might have different shapes, into a single element. The tensors in the resulting element have an additional outer dimension, and are padded to the respective shape in padded\_shapes.

### Usage

```
dataset_padded_batch(
    dataset,
    batch_size,
    padded_shapes,
    padding_values = NULL,
    drop_remainder = FALSE
)
```

### Arguments

- dataset A dataset
- batch\_size An integer, representing the number of consecutive elements of this dataset to combine in a single batch.
- padded\_shapes A nested structure of `tf$TensorShape` or integer vector tensor-like objects representing the shape to which the respective component of each input element should be padded prior to batching. Any unknown dimensions (e.g. `tf$Dimension(NULL)` in a `tf$TensorShape` or `-1` in a tensor-like object) will be padded to the maximum size of that dimension in each batch.
- padding\_values (Optional) A nested structure of scalar-shaped `tf$Tensor`, representing the padding values to use for the respective components. Defaults are 0 for numeric types and the empty string for string types.

`drop_remainder` Ensure that batches have a fixed size by omitting any final smaller batch if it's present. Note that this is required for use with the Keras tensor inputs to `fit/evaluate/etc.`

### Value

A dataset

### See Also

Other dataset methods: `dataset_batch()`, `dataset_cache()`, `dataset_collect()`, `dataset_concatenate()`, `dataset_decode_delim()`, `dataset_filter()`, `dataset_interleave()`, `dataset_map_and_batch()`, `dataset_map()`, `dataset_prefetch_to_device()`, `dataset_prefetch()`, `dataset_reduce()`, `dataset_repeat()`, `dataset_shuffle_and_repeat()`, `dataset_shuffle()`, `dataset_skip()`, `dataset_take()`, `dataset_window()`

---

<code>dataset_prefetch</code>	<i>Creates a Dataset that prefetches elements from this dataset.</i>
-------------------------------	--

---

### Description

Creates a Dataset that prefetches elements from this dataset.

### Usage

```
dataset_prefetch(dataset, buffer_size)
```

### Arguments

<code>dataset</code>	A dataset
<code>buffer_size</code>	An integer, representing the maximum number elements that will be buffered when prefetching.

### Value

A dataset

### See Also

Other dataset methods: `dataset_batch()`, `dataset_cache()`, `dataset_collect()`, `dataset_concatenate()`, `dataset_decode_delim()`, `dataset_filter()`, `dataset_interleave()`, `dataset_map_and_batch()`, `dataset_map()`, `dataset_padded_batch()`, `dataset_prefetch_to_device()`, `dataset_reduce()`, `dataset_repeat()`, `dataset_shuffle_and_repeat()`, `dataset_shuffle()`, `dataset_skip()`, `dataset_take()`, `dataset_window()`

---

`dataset_prefetch_to_device`*A transformation that prefetches dataset values to the given device*

---

**Description**

A transformation that prefetches dataset values to the given device

**Usage**

```
dataset_prefetch_to_device(dataset, device, buffer_size = NULL)
```

**Arguments**

<code>dataset</code>	A dataset
<code>device</code>	A string. The name of a device to which elements will be prefetched (e.g. <code>"/gpu:0"</code> ).
<code>buffer_size</code>	(Optional.) The number of elements to buffer on device. Defaults to an automatically chosen value.

**Value**

A dataset

**Note**

Although the transformation creates a dataset, the transformation must be the final dataset in the input pipeline.

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

dataset_prepare	<i>Prepare a dataset for analysis</i>
-----------------	---------------------------------------

---

### Description

Transform a dataset with named columns into a list with features (x) and response (y) elements.

### Usage

```
dataset_prepare(
    dataset,
    x,
    y = NULL,
    named = TRUE,
    named_features = FALSE,
    parallel_records = NULL,
    batch_size = NULL,
    num_parallel_batches = NULL,
    drop_remainder = FALSE
)
```

### Arguments

dataset	A dataset
x	Features to include. When <code>named_features</code> is <code>FALSE</code> all features will be stacked into a single tensor so must have an identical data type.
y	(Optional). Response variable.
named	<code>TRUE</code> to name the dataset elements "x" and "y", <code>FALSE</code> to not name the dataset elements.
named_features	<code>TRUE</code> to yield features as a named list; <code>FALSE</code> to stack features into a single array. Note that in the case of <code>FALSE</code> (the default) all features will be stacked into a single 2D tensor so need to have the same underlying data type.
parallel_records	(Optional) An integer, representing the number of records to decode in parallel. If not specified, records will be processed sequentially.
batch_size	(Optional). Batch size if you would like to fuse the <code>dataset_prepare()</code> operation together with a <code>dataset_batch()</code> (fusing generally improves overall training performance).
num_parallel_batches	(Optional) An integer, representing the number of batches to create in parallel. On one hand, higher values can help mitigate the effect of stragglers. On the other hand, higher values can increase contention if CPU is scarce.
drop_remainder	Ensure that batches have a fixed size by omitting any final smaller batch if it's present. Note that this is required for use with the Keras tensor inputs to <code>fit/evaluate/etc.</code>

**Value**

A dataset. The dataset will have a structure of either:

- When `named_features` is `TRUE`: `list(x = list(feature_name = feature_values, ...), y = response_values)`
- When `named_features` is `FALSE`: `list(x = features_array, y = response_values)`, where `features_array` is a Rank 2 array of `(batch_size, num_features)`.

Note that the `y` element will be omitted when `y` is `NULL`.

**See Also**

[input\\_fn\(\)](#) for use with **tfestimators**.

---

dataset_reduce	<i>Reduces the input dataset to a single element.</i>
----------------	---

---

**Description**

The transformation calls `reduce_func` successively on every element of the input dataset until the dataset is exhausted, aggregating information in its internal state. The `initial_state` argument is used for the initial state and the final state is returned as the result.

**Usage**

```
dataset_reduce(dataset, initial_state, reduce_func)
```

**Arguments**

<code>dataset</code>	A dataset
<code>initial_state</code>	An element representing the initial state of the transformation.
<code>reduce_func</code>	A function that maps <code>(old_state, input_element)</code> to <code>new_state</code> . It must take two arguments and return a new element. The structure of <code>new_state</code> must match the structure of <code>initial_state</code> .

**Value**

A dataset element.

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)



---

dataset_repeat	<i>Repeats a dataset count times.</i>
----------------	---------------------------------------

---

**Description**

Repeats a dataset count times.

**Usage**

```
dataset_repeat(dataset, count = NULL)
```

**Arguments**

dataset	A dataset
count	(Optional.) An integer value representing the number of times the elements of this dataset should be repeated. The default behavior (if count is NULL or -1) is for the elements to be repeated indefinitely.

**Value**

A dataset

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

dataset_shard	<i>Creates a dataset that includes only 1 / num_shards of this dataset.</i>
---------------	---

---

**Description**

This dataset operator is very useful when running distributed training, as it allows each worker to read a unique subset.

**Usage**

```
dataset_shard(dataset, num_shards, index)
```

**Arguments**

dataset	A dataset
num_shards	A integer representing the number of shards operating in parallel.
index	A integer, representing the worker index.

**Value**

A dataset

---

dataset_shuffle	<i>Randomly shuffles the elements of this dataset.</i>
-----------------	--

---

**Description**

Randomly shuffles the elements of this dataset.

**Usage**

```
dataset_shuffle(  
    dataset,  
    buffer_size,  
    seed = NULL,  
    reshuffle_each_iteration = NULL  
)
```

**Arguments**

dataset	A dataset
buffer_size	An integer, representing the number of elements from this dataset from which the new dataset will sample.
seed	(Optional) An integer, representing the random seed that will be used to create the distribution.
reshuffle_each_iteration	(Optional) A boolean, which if true indicates that the dataset should be pseudo-randomly reshuffled each time it is iterated over. (Defaults to TRUE). Not used if TF version < 1.15

**Value**

A dataset

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

`dataset_shuffle_and_repeat`*Shuffles and repeats a dataset returning a new permutation for each epoch.*

---

**Description**

Shuffles and repeats a dataset returning a new permutation for each epoch.

**Usage**

```
dataset_shuffle_and_repeat(dataset, buffer_size, count = NULL, seed = NULL)
```

**Arguments**

<code>dataset</code>	A dataset
<code>buffer_size</code>	An integer, representing the number of elements from this dataset from which the new dataset will sample.
<code>count</code>	(Optional.) An integer value representing the number of times the elements of this dataset should be repeated. The default behavior (if <code>count</code> is <code>NULL</code> or <code>-1</code> ) is for the elements to be repeated indefinitely.
<code>seed</code>	(Optional) An integer, representing the random seed that will be used to create the distribution.

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

`dataset_skip`*Creates a dataset that skips count elements from this dataset*

---

**Description**

Creates a dataset that skips count elements from this dataset

**Usage**

```
dataset_skip(dataset, count)
```

**Arguments**

dataset	A dataset
count	An integer, representing the number of elements of this dataset that should be skipped to form the new dataset. If count is greater than the size of this dataset, the new dataset will contain no elements. If count is -1, skips the entire dataset.

**Value**

A dataset

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_take\(\)](#), [dataset\\_window\(\)](#)

---

dataset_take	<i>Creates a dataset with at most count elements from this dataset</i>
--------------	--

---

**Description**

Creates a dataset with at most count elements from this dataset

**Usage**

```
dataset_take(dataset, count)
```

**Arguments**

dataset	A dataset
count	Integer representing the number of elements of this dataset that should be taken to form the new dataset. If count is -1, or if count is greater than the size of this dataset, the new dataset will contain all elements of this dataset.

**Value**

A dataset

**See Also**

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_window\(\)](#)

---

dataset_use_spec	<i>Transform the dataset using the provided spec.</i>
------------------	---

---

### Description

Prepares the dataset to be used directly in a model. The transformed dataset is prepared to return tuples (x,y) that can be used directly in Keras.

### Usage

```
dataset_use_spec(dataset, spec)
```

### Arguments

dataset	A TensorFlow dataset.
spec	A feature specification created with <a href="#">feature_spec()</a> .

### Value

A TensorFlow dataset.

### See Also

- [feature\\_spec\(\)](#) to initialize the feature specification.
- [fit.FeatureSpec\(\)](#) to create a tensorflow dataset prepared to modeling.
- [steps](#) to a list of all implemented steps.

Other Feature Spec Functions: [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_identity\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_file\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

### Examples

```
## Not run:
library(tfdatasets)
data(hearts)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ age) %>%
  step_numeric_column(age)

spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)
```

---

dataset_window	<i>Combines input elements into a dataset of windows.</i>
----------------	---

---

### Description

Combines input elements into a dataset of windows.

### Usage

```
dataset_window(dataset, size, shift = NULL, stride = 1, drop_remainder = FALSE)
```

### Arguments

dataset	A dataset
size	representing the number of elements of the input dataset to combine into a window.
shift	representing the forward shift of the sliding window in each iteration. Defaults to size.
stride	representing the stride of the input elements in the sliding window.
drop_remainder	representing whether a window should be dropped in case its size is smaller than window_size.

### See Also

Other dataset methods: [dataset\\_batch\(\)](#), [dataset\\_cache\(\)](#), [dataset\\_collect\(\)](#), [dataset\\_concatenate\(\)](#), [dataset\\_decode\\_delim\(\)](#), [dataset\\_filter\(\)](#), [dataset\\_interleave\(\)](#), [dataset\\_map\\_and\\_batch\(\)](#), [dataset\\_map\(\)](#), [dataset\\_padded\\_batch\(\)](#), [dataset\\_prefetch\\_to\\_device\(\)](#), [dataset\\_prefetch\(\)](#), [dataset\\_reduce\(\)](#), [dataset\\_repeat\(\)](#), [dataset\\_shuffle\\_and\\_repeat\(\)](#), [dataset\\_shuffle\(\)](#), [dataset\\_skip\(\)](#), [dataset\\_take\(\)](#)

---

delim_record_spec	<i>Specification for reading a record from a text file with delimited values</i>
-------------------	--

---

### Description

Specification for reading a record from a text file with delimited values

**Usage**

```
delim_record_spec(
  example_file,
  delim = ",",
  skip = 0,
  names = NULL,
  types = NULL,
  defaults = NULL
)
```

```
csv_record_spec(
  example_file,
  skip = 0,
  names = NULL,
  types = NULL,
  defaults = NULL
)
```

```
tsv_record_spec(
  example_file,
  skip = 0,
  names = NULL,
  types = NULL,
  defaults = NULL
)
```

**Arguments**

- |                           |   |
|---------------------------|---|
| <code>example_file</code> | File that provides an example of the records to be read. If you don't explicitly specify names and types (or defaults) then this file will be read to generate default values.  |
| <code>delim</code>        | Character delimiter to separate fields in a record (defaults to ",")  |
| <code>skip</code>         | Number of lines to skip before reading data. Note that if names is explicitly provided and there are column names within the file then skip should be set to 1 to ensure that the column names are bypassed.  |
| <code>names</code>        | Character vector with column names (or NULL to automatically detect the column names from the first row of <code>example_file</code> ).<br>If names is a character vector, the values will be used as the names of the columns, and the first row of the input will be read into the first row of the dataset. Note that if the underlying text file also includes column names in its first row, this row should be skipped explicitly with <code>skip = 1</code> .<br>If NULL, the first row of the <code>example_file</code> will be used as the column names, and will be skipped when reading the dataset. |
| <code>types</code>        | Column types. If NULL and defaults is specified then types will be imputed from the defaults. Otherwise, all column types will be imputed from the first 1000 rows of the <code>example_file</code> . This is convenient (and fast), but not robust. If the imputation fails, you'll need to supply the correct types yourself.   |

defaults	Types can be explicitly specified in a character vector as "integer", "double", and "character" (e.g. <code>col_types = c("double", "double", "integer")</code> ). Alternatively, you can use a compact string representation where each character represents one column: c = character, i = integer, d = double (e.g. <code>types = ddi</code> ). List of default values which are used when data is missing from a record (e.g. <code>list(0, 0, 0L)</code> ). If NULL then defaults will be automatically provided based on types ( <code>0</code> for numeric columns and <code>""</code> for character columns).
----------	---

---

dense_features	<i>Dense Features</i>
----------------	-----------------------

---

### Description

Retrieves the Dense Features from a spec.

### Usage

```
dense_features(spec)
```

### Arguments

spec            A feature specification created with [feature\\_spec\(\)](#).

### Value

A list of feature columns.

---

feature_spec	<i>Creates a feature specification.</i>
--------------	---

---

### Description

Used to create initialize a feature columns specification.

### Usage

```
feature_spec(dataset, x, y = NULL)
```

### Arguments

dataset        A TensorFlow dataset.  
x              Features to include can use [tidyselect::select\\_helpers\(\)](#) or a formula.  
y              (Optional) The response variable. Can also be specified using a formula in the x argument.



**Details**

After creating the `feature_spec` object you can add steps using the step functions.

**Value**

a `FeatureSpec` object.

**See Also**

- `fit.FeatureSpec()` to fit the `FeatureSpec`
- `dataset_use_spec()` to create a tensorflow dataset prepared to modeling.
- `steps` to a list of all implemented steps.

Other Feature Spec Functions: `dataset_use_spec()`, `fit.FeatureSpec()`, `step_bucketized_column()`, `step_categorical_column_with_hash_bucket()`, `step_categorical_column_with_identity()`, `step_categorical_column_with_vocabulary_file()`, `step_categorical_column_with_vocabulary_list()`, `step_crossed_column()`, `step_embedding_column()`, `step_indicator_column()`, `step_numeric_column()`, `step_remove_column()`, `step_shared_embeddings_column()`, `steps`

**Examples**

```
## Not run:
library(tfdatasets)
data(hearts)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ .)

# select using `tidyselect` helpers
spec <- feature_spec(hearts, x = c(thal, age), y = target)

## End(Not run)
```

---

file_list_dataset	<i>A dataset of all files matching a pattern</i>
-------------------	--

---

**Description**

A dataset of all files matching a pattern

**Usage**

```
file_list_dataset(file_pattern, shuffle = NULL, seed = NULL)
```

**Arguments**

file_pattern	A string, representing the filename pattern that will be matched.
shuffle	(Optional) If TRUE, the file names will be shuffled randomly. Defaults to TRUE.
seed	(Optional) An integer, representing the random seed that will be used to create the distribution.

**Details**

For example, if we had the following files on our filesystem: - /path/to/dir/a.txt - /path/to/dir/b.csv - /path/to/dir/c.csv

If we pass "/path/to/dir/\*.csv" as the file\_pattern, the dataset would produce: - /path/to/dir/b.csv - /path/to/dir/c.csv

**Value**

A dataset of string corresponding to file names

**Note**

The shuffle and seed arguments only apply for TensorFlow >= v1.8

---

fit.FeatureSpec	<i>Fits a feature specification.</i>
-----------------	--------------------------------------

---

**Description**

This function will fit the specification. Depending on the steps added to the specification it will compute for example, the levels of categorical features, normalization constants, etc.

**Usage**

```
## S3 method for class 'FeatureSpec'
fit(object, dataset = NULL, ...)
```

**Arguments**

object	A feature specification created with <code>feature_spec()</code> .
dataset	(Optional) A TensorFlow dataset. If NULL it will use the dataset provided when initializing the feature_spec.
...	(unused)

**Value**

a fitted FeatureSpec object.

**See Also**

- [feature\\_spec\(\)](#) to initialize the feature specification.
- [dataset\\_use\\_spec\(\)](#) to create a tensorflow dataset prepared to modeling.
- [steps](#) to a list of all implemented steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_identity\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_file\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

**Examples**

```
## Not run:
library(tfdatasets)
data(hearts)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ age) %>%
  step_numeric_column(age)

spec_fit <- fit(spec)
spec_fit

## End(Not run)
```

---

fixed\_length\_record\_dataset

*A dataset of fixed-length records from one or more binary files.*

---

**Description**

A dataset of fixed-length records from one or more binary files.

**Usage**

```
fixed_length_record_dataset(
  filenames,
  record_bytes,
  header_bytes = NULL,
  footer_bytes = NULL,
  buffer_size = NULL
)
```

**Arguments**

filenames	A string tensor containing one or more filenames.
record_bytes	An integer representing the number of bytes in each record.
header_bytes	(Optional) An integer scalar representing the number of bytes to skip at the start of a file.
footer_bytes	(Optional) A integer scalar representing the number of bytes to ignore at the end of a file.
buffer_size	(Optional) A integer scalar representing the number of bytes to buffer when reading.

**Value**

A dataset

---

has_type	<i>Identify the type of the variable.</i>
----------	---

---

**Description**

Can only be used inside the [steps](#) specifications to find variables by type.

**Usage**

```
has_type(match = "float32")
```

**Arguments**

match	A list of types to match.
-------	---------------------------

**See Also**

Other Selectors: [all\\_nominal\(\)](#), [all\\_numeric\(\)](#)

---

hearts

*Heart Disease Data Set*

---

### Description

Heart disease (angiographic disease status) dataset.

### Usage

hearts

### Format

A data frame with 303 rows and 14 variables:

**age** age in years

**sex** sex (1 = male; 0 = female)

**cp** chest pain type: Value 1: typical angina, Value 2: atypical angina, Value 3: non-anginal pain, Value 4: asymptomatic

**trestbps** resting blood pressure (in mm Hg on admission to the hospital)

**chol** serum cholestorl in mg/dl

**fbs** (fasting blood sugar > 120 mg/dl) (1 = true; 0 = false)

**restecg** resting electrocardiographic results: Value 0: normal, Value 1: having ST-T wave abnormality (T wave inversions and/or ST elevation or depression of > 0.05 mV), Value 2: showing probable or definite left ventricular hypertrophy by Estes' criteria

**thalach** maximum heart rate achieved

**exang** exercise induced angina (1 = yes; 0 = no)

**oldpeak** ST depression induced by exercise relative to rest

**slope** the slope of the peak exercise ST segment: Value 1: upsloping, Value 2: flat, Value 3: downsloping

**ca** number of major vessels (0-3) colored by flourosopy

**thal** 3 = normal; 6 = fixed defect; 7 = reversable defect

**target** diagnosis of heart disease angiographic

### Source

<https://archive.ics.uci.edu/ml/datasets/heart+Disease>

## References

The authors of the databases have requested that any publications resulting from the use of the data include the names of the principal investigator responsible for the data collection at each institution. They would be:

1. Hungarian Institute of Cardiology. Budapest: Andras Janosi, M.D.
2. University Hospital, Zurich, Switzerland: William Steinbrunn, M.D.
3. University Hospital, Basel, Switzerland: Matthias Pfisterer, M.D.
4. V.A. Medical Center, Long Beach and Cleveland Clinic Foundation: Robert Detrano, M.D., Ph.D.

---

input\_fn.tf\_dataset     *Construct a tfestimators input function from a dataset*

---

## Description

Construct a tfestimators input function from a dataset

## Usage

```
input_fn.tf_dataset(dataset, features, response = NULL)
```

## Arguments

dataset	A dataset
features	The names of feature variables to be used.
response	The name of the response variable.

## Details

Creating an input\_fn from a dataset requires that the dataset consist of a set of named output tensors (e.g. like the dataset produced by the [tfrecord\\_dataset\(\)](#) or [text\\_line\\_dataset\(\)](#) function).

## Value

An input\_fn suitable for use with tfestimators [train](#), [evaluate](#), and [predict](#) methods

---

iterator_get_next	<i>Get next element from iterator</i>
-------------------	---------------------------------------

---

**Description**

Returns a nested list of tensors that when evaluated will yield the next element(s) in the dataset.

**Usage**

```
iterator_get_next(iterator, name = NULL)
```

**Arguments**

iterator	An iterator
name	(Optional) A name for the created operation.

**Value**

A nested list of tensors

**See Also**

Other iterator functions: [iterator\\_initializer\(\)](#), [iterator\\_make\\_initializer\(\)](#), [iterator\\_string\\_handle\(\)](#), [make-iterator](#)

---

iterator_initializer	<i>An operation that should be run to initialize this iterator.</i>
----------------------	---

---

**Description**

An operation that should be run to initialize this iterator.

**Usage**

```
iterator_initializer(iterator)
```

**Arguments**

iterator	An iterator
----------	-------------

**See Also**

Other iterator functions: [iterator\\_get\\_next\(\)](#), [iterator\\_make\\_initializer\(\)](#), [iterator\\_string\\_handle\(\)](#), [make-iterator](#)

---

```
iterator_make_initializer
```

*Create an operation that can be run to initialize this iterator*

---

### Description

Create an operation that can be run to initialize this iterator

### Usage

```
iterator_make_initializer(iterator, dataset, name = NULL)
```

### Arguments

iterator	An iterator
dataset	A dataset
name	(Optional) A name for the created operation.

### Value

A `tf$Operation` that can be run to initialize this iterator on the given dataset.

### See Also

Other iterator functions: [iterator\\_get\\_next\(\)](#), [iterator\\_initializer\(\)](#), [iterator\\_string\\_handle\(\)](#), [make-iterator](#)

---

```
iterator_string_handle
```

*String-valued tensor that represents this iterator*

---

### Description

String-valued tensor that represents this iterator

### Usage

```
iterator_string_handle(iterator, name = NULL)
```

### Arguments

iterator	An iterator
name	(Optional) A name for the created operation.



**Value**

Scalar tensor of type string

**See Also**

Other iterator functions: [iterator\\_get\\_next\(\)](#), [iterator\\_initializer\(\)](#), [iterator\\_make\\_initializer\(\)](#), [make-iterator](#)

---

layer\_input\_from\_dataset

*Creates a list of inputs from a dataset*

---

**Description**

Create a list of Keras input layers that can be used together with [keras::layer\\_dense\\_features\(\)](#).

**Usage**

```
layer_input_from_dataset(dataset)
```

**Arguments**

dataset            a TensorFlow dataset or a data.frame

**Value**

a list of Keras input layers

**Examples**

```
## Not run:
library(tfdatasets)
data(hearts)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ age + slope) %>%
  step_numeric_column(age, slope) %>%
  step_bucketized_column(age, boundaries = c(10, 20, 30))

spec <- fit(spec)
dataset <- hearts %>% dataset_use_spec(spec)

input <- layer_input_from_dataset(dataset)

## End(Not run)
```

---

 make-iterator

*Creates an iterator for enumerating the elements of this dataset.*


---

### Description

Creates an iterator for enumerating the elements of this dataset.

### Usage

```
make_iterator_one_shot(dataset)

make_iterator_initializable(dataset, shared_name = NULL)

make_iterator_from_structure(
  output_types,
  output_shapes = NULL,
  shared_name = NULL
)

make_iterator_from_string_handle(
  string_handle,
  output_types,
  output_shapes = NULL
)
```

### Arguments

dataset	A dataset
shared_name	(Optional) If non-empty, the returned iterator will be shared under the given name across multiple sessions that share the same devices (e.g. when using a remote server).
output_types	A nested structure of <code>tf\$DType</code> objects corresponding to each component of an element of this iterator.
output_shapes	(Optional) A nested structure of <code>tf\$TensorShape</code> objects corresponding to each component of an element of this dataset. If omitted, each component will have an unconstrained shape.
string_handle	A scalar tensor of type string that evaluates to a handle produced by the <code>iterator_string_handle()</code> method.

### Value

An Iterator over the elements of this dataset.

### Initialization

For `make_iterator_one_shot()`, the returned iterator will be initialized automatically. A "one-shot" iterator does not currently support re-initialization.

For `make_iterator_initializable()`, the returned iterator will be in an uninitialized state, and you must run the object returned from `iterator_initializer()` before using it.

For `make_iterator_from_structure()`, the returned iterator is not bound to a particular dataset, and it has no initializer. To initialize the iterator, run the operation returned by `iterator_make_initializer()`.

### See Also

Other iterator functions: `iterator_get_next()`, `iterator_initializer()`, `iterator_make_initializer()`, `iterator_string_handle()`

---

make_csv_dataset	<i>Reads CSV files into a batched dataset</i>
------------------	---

---

### Description

Reads CSV files into a dataset, where each element is a (features, labels) list that corresponds to a batch of CSV rows. The features dictionary maps feature column names to tensors containing the corresponding feature data, and labels is a tensor containing the batch's label data.

### Usage

```
make_csv_dataset(  
    file_pattern,  
    batch_size,  
    column_names = NULL,  
    column_defaults = NULL,  
    label_name = NULL,  
    select_columns = NULL,  
    field_delim = ",",  
    use_quote_delim = TRUE,  
    na_value = "",  
    header = TRUE,  
    num_epochs = NULL,  
    shuffle = TRUE,  
    shuffle_buffer_size = 10000,  
    shuffle_seed = NULL,  
    prefetch_buffer_size = 1,  
    num_parallel_reads = 1,  
    num_parallel_parser_calls = 2,  
    sloppy = FALSE,  
    num_rows_for_inference = 100  
)
```

**Arguments**

file_pattern	List of files or glob patterns of file paths containing CSV records.
batch_size	An integer representing the number of records to combine in a single batch.
column_names	An optional list of strings that corresponds to the CSV columns, in order. One per column of the input record. If this is not provided, infers the column names from the first row of the records. These names will be the keys of the features dict of each dataset element.
column_defaults	A optional list of default values for the CSV fields. One item per selected column of the input record. Each item in the list is either a valid CSV dtype (integer, numeric, or string), or a tensor with one of the aforementioned types. The tensor can either be a scalar default value (if the column is optional), or an empty tensor (if the column is required). If a dtype is provided instead of a tensor, the column is also treated as required. If this list is not provided, tries to infer types based on reading the first num_rows_for_inference rows of files specified, and assumes all columns are optional, defaulting to 0 for numeric values and "" for string values. If both this and select_columns are specified, these must have the same lengths, and column_defaults is assumed to be sorted in order of increasing column index.
label_name	A optional string corresponding to the label column. If provided, the data for this column is returned as a separate tensor from the features dictionary, so that the dataset complies with the format expected by a TF Estimators and Keras.
select_columns	(Ignored if using TensorFlow version 1.8.) An optional list of integer indices or string column names, that specifies a subset of columns of CSV data to select. If column names are provided, these must correspond to names provided in column_names or inferred from the file header lines. When this argument is specified, only a subset of CSV columns will be parsed and returned, corresponding to the columns specified. Using this results in faster parsing and lower memory usage. If both this and column_defaults are specified, these must have the same lengths, and column_defaults is assumed to be sorted in order of increasing column index.
field_delim	An optional string. Defaults to ", ". Char delimiter to separate fields in a record.
use_quote_delim	An optional bool. Defaults to TRUE. If false, treats double quotation marks as regular characters inside of the string fields.
na_value	Additional string to recognize as NA/NaN.
header	A bool that indicates whether the first rows of provided CSV files correspond to header lines with column names, and should not be included in the data.
num_epochs	An integer specifying the number of times this dataset is repeated. If NULL, cycles through the dataset forever.
shuffle	A bool that indicates whether the input should be shuffled.
shuffle_buffer_size	Buffer size to use for shuffling. A large buffer size ensures better shuffling, but increases memory usage and startup time.
shuffle_seed	Randomization seed to use for shuffling.

prefetch_buffer_size	An int specifying the number of feature batches to prefetch for performance improvement. Recommended value is the number of batches consumed per training step.
num_parallel_reads	Number of threads used to read CSV records from files. If >1, the results will be interleaved.
num_parallel_parser_calls	(Ignored if using TensorFlow version 1.11 or later.) Number of parallel invocations of the CSV parsing function on CSV records.
sloppy	If TRUE, reading performance will be improved at the cost of non-deterministic ordering. If FALSE, the order of elements produced is deterministic prior to shuffling (elements are still randomized if shuffle=TRUE. Note that if the seed is set, then order of elements after shuffling is deterministic). Defaults to FALSE.
num_rows_for_inference	Number of rows of a file to use for type inference if record_defaults is not provided. If NULL, reads all the rows of all the files. Defaults to 100.

**Value**

A dataset, where each element is a (features, labels) list that corresponds to a batch of batch\_size CSV rows. The features dictionary maps feature column names to tensors containing the corresponding column data, and labels is a tensor containing the column data for the label column specified by label\_name.

---

next_batch	<i>Tensor(s) for retrieving the next batch from a dataset</i>
------------	---

---

**Description**

Tensor(s) for retrieving the next batch from a dataset

**Usage**

```
next_batch(dataset)
```

**Arguments**

dataset	A dataset
---------	-----------

**Details**

To access the underlying data within the dataset you iteratively evaluate the tensor(s) to read batches of data.

Note that in many cases you won't need to explicitly evaluate the tensors. Rather, you will pass the tensors to another function that will perform the evaluation (e.g. the Keras [layer\\_input\(\)](#) and [compile\(\)](#) functions).

If you do need to perform iteration manually by evaluating the tensors, there are a couple of possible approaches to controlling/detecting when iteration should end.

One approach is to create a dataset that yields batches infinitely (traversing the dataset multiple times with different batches randomly drawn). In this case you'd use another mechanism like a global step counter or detecting a learning plateau.

Another approach is to detect when all batches have been yielded from the dataset. When the tensor reaches the end of iteration a runtime error will occur. You can catch and ignore the error when it occurs by wrapping your iteration code in the `with_dataset()` function.

See the examples below for a demonstration of each of these methods of iteration.

## Value

Tensor(s) that can be evaluated to yield the next batch of training data.

## Examples

```
## Not run:

# iteration with 'infinite' dataset and explicit step counter

library(tfdatasets)
dataset <- text_line_dataset("mtcars.csv", record_spec = mtcars_spec) %>%
  dataset_prepare(x = c(mpg, disp), y = cyl) %>%
  dataset_shuffle(5000) %>%
  dataset_batch(128) %>%
  dataset_repeat() # repeat infinitely
batch <- next_batch(dataset)
steps <- 200
for (i in 1:steps) {
  # use batch$x and batch$y tensors
}

# iteration that detects and ignores end of iteration error

library(tfdatasets)
dataset <- text_line_dataset("mtcars.csv", record_spec = mtcars_spec) %>%
  dataset_prepare(x = c(mpg, disp), y = cyl) %>%
  dataset_batch(128) %>%
  dataset_repeat(10)
batch <- next_batch(dataset)
with_dataset({
  while(TRUE) {
    # use batch$x and batch$y tensors
  }
})

## End(Not run)
```

---

output_types	<i>Output types and shapes</i>
--------------	--------------------------------

---

**Description**

Output types and shapes

**Usage**

```
output_types(object)
```

```
output_shapes(object)
```

**Arguments**

object	A dataset or iterator
--------	-----------------------

**Value**

output\_types() returns the type of each component of an element of this object; output\_shapes() returns the shape of each component of an element of this object

---

range_dataset	<i>Creates a dataset of a step-separated range of values.</i>
---------------	---

---

**Description**

Creates a dataset of a step-separated range of values.

**Usage**

```
range_dataset(from = 0, to = 0, by = 1)
```

**Arguments**

from	Range start
to	Range end (exclusive)
by	Increment of the sequence

---

read_files	<i>Read a dataset from a set of files</i>
------------	---

---

### Description

Read files into a dataset, optionally processing them in parallel.

### Usage

```
read_files(  
  files,  
  reader,  
  ...,  
  parallel_files = 1,  
  parallel_interleave = 1,  
  num_shards = NULL,  
  shard_index = NULL  
)
```

### Arguments

files	List of filenames or glob pattern for files (e.g. "*.csv")
reader	Function that maps a file into a dataset (e.g. <code>text_line_dataset()</code> or <code>tfrecord_dataset()</code> ).
...	Additional arguments to pass to reader function
parallel_files	An integer, number of files to process in parallel
parallel_interleave	An integer, number of consecutive records to produce from each file before cycling to another file.
num_shards	An integer representing the number of shards operating in parallel.
shard_index	An integer, representing the worker index. Shared indexes are 0 based so for e.g. 8 shards valid indexes would be 0-7.

### Value

A dataset



---

sample\_from\_datasets    *Samples elements at random from the datasets in datasets.*

---

### Description

Samples elements at random from the datasets in datasets.

### Usage

```
sample_from_datasets(datasets, weights = NULL, seed = NULL)
```

### Arguments

datasets	A list of objects with compatible structure.
weights	(Optional.) A list of length(datasets) floating-point values where weights[[i]] represents the probability with which an element should be sampled from datasets[[i]], or a dataset object where each element is such a list. Defaults to a uniform distribution across datasets.
seed	(Optional.) An integer, representing the random seed that will be used to create the distribution.

### Value

A dataset that interleaves elements from datasets at random, according to weights if provided, otherwise with uniform probability.

---

scaler                      *List of pre-made scalers*

---

### Description

- [scaler\\_standard](#): mean and standard deviation normalizer.
- [scaler\\_min\\_max](#): min max normalizer

### See Also

[step\\_numeric\\_column](#)

---

scaler_min_max	<i>Creates an instance of a min max scaler</i>
----------------	--

---

**Description**

This scaler will learn the min and max of the numeric variable and use this to create a `normalizer_fn`.

**Usage**

```
scaler_min_max()
```

**See Also**

[scaler](#) to a complete list of normalizers

Other scaler: [scaler\\_standard\(\)](#)

---

scaler_standard	<i>Creates an instance of a standard scaler</i>
-----------------	---

---

**Description**

This scaler will learn the mean and the standard deviation and use this to create a `normalizer_fn`.

**Usage**

```
scaler_standard()
```

**See Also**

[scaler](#) to a complete list of normalizers

Other scaler: [scaler\\_min\\_max\(\)](#)

---

selectors

*Selectors*

---

### Description

List of selectors that can be used to specify variables inside steps.

### Usage

cur\_info\_env

### Format

An object of class environment of length 0.

### Selectors

- `has_type()`
- `all_numeric()`
- `all_nominal()`
- `starts_with()`
- `ends_with()`
- `one_of()`
- `matches()`
- `contains()`
- `everything()`

---

sparse\_tensor\_slices\_dataset

*Splits each rank-N tf\$SparseTensor in this dataset row-wise.*

---

### Description

Splits each rank-N tf\$SparseTensor in this dataset row-wise.

### Usage

sparse\_tensor\_slices\_dataset(sparse\_tensor)

### Arguments

sparse\_tensor A tf\$SparseTensor.

**Value**

A dataset of rank-(N-1) sparse tensors.

**See Also**

Other tensor datasets: [tensor\\_slices\\_dataset\(\)](#), [tensors\\_dataset\(\)](#)

---

sql_record_spec	<i>A dataset consisting of the results from a SQL query</i>
-----------------	---

---

**Description**

A dataset consisting of the results from a SQL query

**Usage**

```
sql_record_spec(names, types)
```

```
sql_dataset(driver_name, data_source_name, query, record_spec)
```

```
sqlite_dataset(filename, query, record_spec)
```

**Arguments**

names	Names of columns returned from the query
types	List of tf\$DType objects (e.g. tf\$int32, tf\$double, tf\$string) representing the types of the columns returned by the query.
driver_name	String containing the database type. Currently, the only supported value is 'sqlite'.
data_source_name	String containing a connection string to connect to the database.
query	String containing the SQL query to execute.
record_spec	Names and types of database columns
filename	Filename for the database

**Value**

A dataset

---

 steps
 

---

*Steps for feature columns specification.*


---

## Description

List of steps that can be used to specify columns in the `feature_spec` interface.

## Steps

- `step_numeric_column()` to define numeric columns.
- `step_categorical_column_with_vocabulary_list()` to define categorical columns.
- `step_categorical_column_with_hash_bucket()` to define categorical columns where ids are set by hashing.
- `step_categorical_column_with_identity()` to define categorical columns represented by integers in the range `[0-num_buckets)`.
- `step_categorical_column_with_vocabulary_file()` to define categorical columns when their vocabulary is available in a file.
- `step_indicator_column()` to create indicator columns from categorical columns.
- `step_embedding_column()` to create embeddings columns from categorical columns.
- `step_bucketized_column()` to create bucketized columns from numeric columns.
- `step_crossed_column()` to perform crosses of categorical columns.
- `step_shared_embeddings_column()` to share embeddings between a list of categorical columns.
- `step_remove_column()` to remove columns from the specification.

## See Also

- `selectors` for a list of selectors that can be used to specify variables.

Other Feature Spec Functions: `dataset_use_spec()`, `feature_spec()`, `fit.FeatureSpec()`, `step_bucketized_column()`, `step_categorical_column_with_hash_bucket()`, `step_categorical_column_with_identity()`, `step_categorical_column_with_vocabulary_file()`, `step_categorical_column_with_vocabulary_list()`, `step_crossed_column()`, `step_embedding_column()`, `step_indicator_column()`, `step_numeric_column()`, `step_remove_column()`, `step_shared_embeddings_column()`

---

 step\_bucketized\_column
 

---

*Creates bucketized columns*


---

## Description

Use this step to create bucketized columns from numeric columns.

**Usage**

```
step_bucketized_column(spec, ..., boundaries)
```

**Arguments**

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.
boundaries	A sorted list or tuple of floats specifying the boundaries.

**Value**

a FeatureSpec object.

**See Also**

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_identity\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_file\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

**Examples**

```
## Not run:
library(tfdatasets)
data(hearts)
file <- tempfile()
writeLines(unique(hearts$thal), file)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ age) %>%
  step_numeric_column(age) %>%
  step_bucketized_column(age, boundaries = c(10, 20, 30))
spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)
```

---

```
step_categorical_column_with_hash_bucket
```

*Creates a categorical column with hash buckets specification*

---

**Description**

Represents sparse feature where ids are set by hashing.

**Usage**

```
step_categorical_column_with_hash_bucket(
  spec,
  ...,
  hash_bucket_size,
  dtype = tf$string
)
```

**Arguments**

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.
hash_bucket_size	An int > 1. The number of buckets.
dtype	The type of features. Only string and integer types are supported.

**Value**

a FeatureSpec object.

**See Also**

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_identity\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

**Examples**

```
## Not run:
library(tfdatasets)
data(hearts)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ thal) %>%
  step_categorical_column_with_hash_bucket(thal, hash_bucket_size = 3)

spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)
```

---

step\_categorical\_column\_with\_identity  
*Create a categorical column with identity*

---

## Description

Use this when your inputs are integers in the range [0-num\_buckets).

## Usage

```
step_categorical_column_with_identity(  
  spec,  
  ...,  
  num_buckets,  
  default_value = NULL  
)
```

## Arguments

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.
num_buckets	Range of inputs and outputs is [0, num_buckets).
default_value	If NULL, this column's graph operations will fail for out-of-range inputs. Otherwise, this value must be in the range [0, num_buckets), and will replace inputs in that range.

## Value

a FeatureSpec object.

## See Also

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

## Examples

```
## Not run:  
library(tfdatasets)  
data(hearts)  
  
hearts$thal <- as.integer(as.factor(hearts$thal)) - 1L
```



```

hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ thal) %>%
  step_categorical_column_with_identity(thal, num_buckets = 5)

spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)

```

---

```
step_categorical_column_with_vocabulary_file
```

*Creates a categorical column with vocabulary file*

---

## Description

Use this function when the vocabulary of a categorical variable is written to a file.

## Usage

```

step_categorical_column_with_vocabulary_file(
  spec,
  ...,
  vocabulary_file,
  vocabulary_size = NULL,
  dtype = tf$string,
  default_value = NULL,
  num_oov_buckets = 0L
)

```

## Arguments

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.
vocabulary_file	The vocabulary file name.
vocabulary_size	Number of the elements in the vocabulary. This must be no greater than length of vocabulary_file, if less than length, later values are ignored. If None, it is set to the length of vocabulary_file.
dtype	The type of features. Only string and integer types are supported.
default_value	The integer ID value to return for out-of-vocabulary feature values, defaults to -1. This can not be specified with a positive num_oov_buckets.

num\_oov\_buckets

Non-negative integer, the number of out-of-vocabulary buckets. All out-of-vocabulary inputs will be assigned IDs in the range [vocabulary\_size, vocabulary\_size+num\_oov\_buckets) based on a hash of the input value. A positive num\_oov\_buckets can not be specified with default\_value.

### Value

a FeatureSpec object.

### See Also

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_id\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

### Examples

```
## Not run:
library(tfdatasets)
data(hearts)
file <- tempfile()
writeLines(unique(hearts$thal), file)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ thal) %>%
  step_categorical_column_with_vocabulary_file(thal, vocabulary_file = file)

spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)
```

---

step\_categorical\_column\_with\_vocabulary\_list

*Creates a categorical column specification*

---

### Description

Creates a categorical column specification

**Usage**

```
step_categorical_column_with_vocabulary_list(
  spec,
  ...,
  vocabulary_list = NULL,
  dtype = NULL,
  default_value = -1L,
  num_oov_buckets = 0L
)
```

**Arguments**

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.
vocabulary_list	An ordered iterable defining the vocabulary. Each feature is mapped to the index of its value (if present) in vocabulary_list. Must be castable to dtype. If NULL the vocabulary will be defined as all unique values in the dataset provided when fitting the specification.
dtype	The type of features. Only string and integer types are supported. If NULL, it will be inferred from vocabulary_list.
default_value	The integer ID value to return for out-of-vocabulary feature values, defaults to -1. This can not be specified with a positive num_oov_buckets.
num_oov_buckets	Non-negative integer, the number of out-of-vocabulary buckets. All out-of-vocabulary inputs will be assigned IDs in the range [length(vocabulary_list), length(vocabulary_list)+num_oov_buckets) based on a hash of the input value. A positive num_oov_buckets can not be specified with default_value.

**Value**

a FeatureSpec object.

**See Also**

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_id\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_file\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

**Examples**

```
## Not run:
library(tfdatasets)
data(hearts)
```

```

hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ thal) %>%
  step_categorical_column_with_vocabulary_list(thal)

spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)

```

---

step\_crossed\_column    *Creates crosses of categorical columns*

---

### Description

Use this step to create crosses between categorical columns.

### Usage

```
step_crossed_column(spec, ..., hash_bucket_size, hash_key = NULL)
```

### Arguments

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.
hash_bucket_size	An int > 1. The number of buckets.
hash_key	(optional) Specify the hash_key that will be used by the FingerprintCat64 function to combine the crosses fingerprints on SparseCrossOp.

### Value

a FeatureSpec object.

### See Also

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_id](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_file\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

**Examples**

```
## Not run:
library(tfdatasets)
data(hearts)
file <- tempfile()
writeLines(unique(hearts$th1), file)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ age) %>%
  step_numeric_column(age) %>%
  step_bucketized_column(age, boundaries = c(10, 20, 30))
spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)
```

---

step\_embedding\_column *Creates embeddings columns*

---

**Description**

Use this step to create embeddings columns from categorical columns.

**Usage**

```
step_embedding_column(
  spec,
  ...,
  dimension = function(x) { as.integer(x^0.25) },
  combiner = "mean",
  initializer = NULL,
  ckpt_to_load_from = NULL,
  tensor_name_in_ckpt = NULL,
  max_norm = NULL,
  trainable = TRUE
)
```

**Arguments**

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.
dimension	An integer specifying dimension of the embedding, must be > 0. Can also be a function of the size of the vocabulary.

combiner	A string specifying how to reduce if there are multiple entries in a single row. Currently 'mean', 'sqrtn' and 'sum' are supported, with 'mean' the default. 'sqrtn' often achieves good accuracy, in particular with bag-of-words columns. Each of this can be thought as example level normalizations on the column. For more information, see <code>tf.embedding_lookup_sparse</code> .
initializer	A variable initializer function to be used in embedding variable initialization. If not specified, defaults to <code>tf.truncated_normal_initializer</code> with mean $0.0$ and standard deviation $1/\sqrt{\text{dimension}}$ .
ckpt_to_load_from	String representing checkpoint name/pattern from which to restore column weights. Required if <code>tensor_name_in_ckpt</code> is not NULL.
tensor_name_in_ckpt	Name of the Tensor in <code>ckpt_to_load_from</code> from which to restore the column weights. Required if <code>ckpt_to_load_from</code> is not NULL.
max_norm	If not NULL, embedding values are l2-normalized to this value.
trainable	Whether or not the embedding is trainable. Default is TRUE.

**Value**

a FeatureSpec object.

**See Also**

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_id\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_file\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

**Examples**

```
## Not run:
library(tfdatasets)
data(hearts)
file <- tempfile()
writeLines(unique(hearts$thal), file)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ thal) %>%
  step_categorical_column_with_vocabulary_list(thal) %>%
  step_embedding_column(thal, dimension = 3)
spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)
```

---

step\_indicator\_column *Creates Indicator Columns*

---

## Description

Use this step to create indicator columns from categorical columns.

## Usage

```
step_indicator_column(spec, ...)
```

## Arguments

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.

## Value

a FeatureSpec object.

## See Also

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_id\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_file\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

## Examples

```
## Not run:
library(tfdatasets)
data(hearts)
file <- tempfile()
writeLines(unique(hearts$thal), file)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ thal) %>%
  step_categorical_column_with_vocabulary_list(thal) %>%
  step_indicator_column(thal)
spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)
```

---

step\_numeric\_column *Creates a numeric column specification*

---

### Description

step\_numeric\_column creates a numeric column specification. It can also be used to normalize numeric columns.

### Usage

```
step_numeric_column(
  spec,
  ...,
  shape = 1L,
  default_value = NULL,
  dtype = tf$float32,
  normalizer_fn = NULL
)
```

### Arguments

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.
shape	An iterable of integers specifies the shape of the Tensor. An integer can be given which means a single dimension Tensor with given width. The Tensor representing the column will have the shape of batch_size + shape.
default_value	A single value compatible with dtype or an iterable of values compatible with dtype which the column takes on during <code>tf.parse_example</code> if data is missing. A default value of NULL will cause <code>tf.parse_example</code> to fail if an example does not contain this column. If a single value is provided, the same value will be applied as the default value for every item. If an iterable of values is provided, the shape of the default_value should be equal to the given shape.
dtype	defines the type of values. Default value is <code>tf\$float32</code> . Must be a non-quantized, real integer or floating point type.
normalizer_fn	If not NULL, a function that can be used to normalize the value of the tensor after default_value is applied for parsing. Normalizer function takes the input Tensor as its argument, and returns the output Tensor. (e.g. <code>function(x) (x - 3.0) / 4.2</code> ). Please note that even though the most common use case of this function is normalization, it can be used for any kind of Tensorflow transformations. You can also a pre-made <a href="#">scaler</a> , in this case a function will be created after <code>fit.FeatureSpec</code> is called on the feature specification.

### Value

a FeatureSpec object.



## See Also

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_id\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_file\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

## Examples

```
## Not run:
library(tfdatasets)
data(hearts)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ age) %>%
  step_numeric_column(age, normalizer_fn = standard_scaler())

spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)
```

---

step_remove_column	<i>Creates a step that can remove columns</i>
--------------------	---

---

## Description

Removes features of the feature specification.

## Usage

```
step_remove_column(spec, ...)
```

## Arguments

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.

## Value

a FeatureSpec object.

**See Also**

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_id\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_file\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_shared\\_embeddings\\_column\(\)](#), [steps](#)

**Examples**

```
## Not run:
library(tfdatasets)
data(hearts)
hearts <- tensor_slices_dataset(hearts) %>% dataset_batch(32)

# use the formula interface
spec <- feature_spec(hearts, target ~ age) %>%
  step_numeric_column(age, normalizer_fn = scaler_standard()) %>%
  step_bucketized_column(age, boundaries = c(20, 50)) %>%
  step_remove_column(age)

spec_fit <- fit(spec)
final_dataset <- hearts %>% dataset_use_spec(spec_fit)

## End(Not run)
```

---

step\_shared\_embeddings\_column

*Creates shared embeddings for categorical columns*

---

**Description**

This is similar to [step\\_embedding\\_column](#), except that it produces a list of embedding columns that share the same embedding weights.

**Usage**

```
step_shared_embeddings_column(
  spec,
  ...,
  dimension,
  combiner = "mean",
  initializer = NULL,
  shared_embedding_collection_name = NULL,
  ckpt_to_load_from = NULL,
  tensor_name_in_ckpt = NULL,
  max_norm = NULL,
```

```

    trainable = TRUE
)

```

### Arguments

spec	A feature specification created with <a href="#">feature_spec()</a> .
...	Comma separated list of variable names to apply the step. <a href="#">selectors</a> can also be used.
dimension	An integer specifying dimension of the embedding, must be > 0. Can also be a function of the size of the vocabulary.
combiner	A string specifying how to reduce if there are multiple entries in a single row. Currently 'mean', 'sqrtn' and 'sum' are supported, with 'mean' the default. 'sqrtn' often achieves good accuracy, in particular with bag-of-words columns. Each of this can be thought as example level normalizations on the column. For more information, see <code>tf.embedding_lookup_sparse</code> .
initializer	A variable initializer function to be used in embedding variable initialization. If not specified, defaults to <code>tf.truncated_normal_initializer</code> with mean 0.0 and standard deviation $1/\sqrt{\text{dimension}}$ .
shared_embedding_collection_name	Optional collective name of these columns. If not given, a reasonable name will be chosen based on the names of <code>categorical_columns</code> .
ckpt_to_load_from	String representing checkpoint name/pattern from which to restore column weights. Required if <code>tensor_name_in_ckpt</code> is not NULL.
tensor_name_in_ckpt	Name of the Tensor in <code>ckpt_to_load_from</code> from which to restore the column weights. Required if <code>ckpt_to_load_from</code> is not NULL.
max_norm	If not NULL, embedding values are l2-normalized to this value.
trainable	Whether or not the embedding is trainable. Default is TRUE.

### Value

a FeatureSpec object.

### Note

Does not work in the eager mode.

### See Also

[steps](#) for a complete list of allowed steps.

Other Feature Spec Functions: [dataset\\_use\\_spec\(\)](#), [feature\\_spec\(\)](#), [fit.FeatureSpec\(\)](#), [step\\_bucketized\\_column\(\)](#), [step\\_categorical\\_column\\_with\\_hash\\_bucket\(\)](#), [step\\_categorical\\_column\\_with\\_id](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_file\(\)](#), [step\\_categorical\\_column\\_with\\_vocabulary\\_list\(\)](#), [step\\_crossed\\_column\(\)](#), [step\\_embedding\\_column\(\)](#), [step\\_indicator\\_column\(\)](#), [step\\_numeric\\_column\(\)](#), [step\\_remove\\_column\(\)](#), [steps](#)

---

tensors_dataset	<i>Creates a dataset with a single element, comprising the given tensors.</i>
-----------------	---

---

**Description**

Creates a dataset with a single element, comprising the given tensors.

**Usage**

```
tensors_dataset(tensors)
```

**Arguments**

tensors          A nested structure of tensors.

**Value**

A dataset.

**See Also**

Other tensor datasets: [sparse\\_tensor\\_slices\\_dataset\(\)](#), [tensor\\_slices\\_dataset\(\)](#)

---

tensor_slices_dataset	<i>Creates a dataset whose elements are slices of the given tensors.</i>
-----------------------	--

---

**Description**

Creates a dataset whose elements are slices of the given tensors.

**Usage**

```
tensor_slices_dataset(tensors)
```

**Arguments**

tensors          A nested structure of tensors, each having the same size in the 0th dimension.

**Value**

A dataset.

**See Also**

Other tensor datasets: [sparse\\_tensor\\_slices\\_dataset\(\)](#), [tensors\\_dataset\(\)](#)

---

text\_line\_dataset      *A dataset comprising lines from one or more text files.*

---

### Description

A dataset comprising lines from one or more text files.

### Usage

```
text_line_dataset(
  filenames,
  compression_type = NULL,
  record_spec = NULL,
  parallel_records = NULL
)
```

### Arguments

filenames      String(s) specifying one or more filenames

compression\_type      A string, one of: NULL (no compression), "ZLIB", or "GZIP".

record\_spec      (Optional) Specification used to decode delimited text lines into records (see [delim\\_record\\_spec\(\)](#)).

parallel\_records      (Optional) An integer, representing the number of records to decode in parallel. If not specified, records will be processed sequentially.

### Value

A dataset

---

tfrecord\_dataset      *A dataset comprising records from one or more TFRecord files.*

---

### Description

A dataset comprising records from one or more TFRecord files.

### Usage

```
tfrecord_dataset(
  filenames,
  compression_type = NULL,
  buffer_size = NULL,
  num_parallel_reads = NULL
)
```

**Arguments**

filenames       String(s) specifying one or more filenames

compression\_type       A string, one of: NULL (no compression), "ZLIB", or "GZIP".

buffer\_size       An integer representing the number of bytes in the read buffer. (0 means no buffering).

num\_parallel\_reads       An integer representing the number of files to read in parallel. Defaults to reading files sequentially.

**Details**

If the dataset encodes a set of TFExample instances, then they can be decoded into named records using the `dataset_map()` function (see example below).

**Examples**

```
## Not run:

# Creates a dataset that reads all of the examples from two files, and extracts
# the image and label features.
filenames <- c("/var/data/file1.tfrecord", "/var/data/file2.tfrecord")
dataset <- tfrecord_dataset(filenames) %>%
  dataset_map(function(example_proto) {
    features <- list(
      image = tf$FixedLenFeature(shape(), tf$string, default_value = ""),
      label = tf$FixedLenFeature(shape(), tf$int32, default_value = 0L)
    )
    tf$parse_single_example(example_proto, features)
  })

## End(Not run)
```

---

until_out_of_range	<i>Execute code that traverses a dataset until an out of range condition occurs</i>
--------------------	---

---

**Description**

Execute code that traverses a dataset until an out of range condition occurs

**Usage**

```
until_out_of_range(expr)

out_of_range_handler(e)
```

**Arguments**

expr	Expression to execute (will be executed multiple times until the condition occurs)
e	Error object

**Details**

When a dataset iterator reaches the end, an out of range runtime error will occur. This function will catch and ignore the error when it occurs.

**Examples**

```
## Not run:
library(tfdatasets)
dataset <- text_line_dataset("mtcars.csv", record_spec = mtcars_spec) %>%
  dataset_batch(128) %>%
  dataset_repeat(10) %>%
  dataset_prepare(x = c(mpg, disp), y = cyl)

iter <- make_iterator_one_shot(dataset)
next_batch <- iterator_get_next(iter)

until_out_of_range({
  batch <- sess$run(next_batch)
  # use batch$x and batch$y tensors
})

## End(Not run)
```

---

with\_dataset

*Execute code that traverses a dataset*


---

**Description**

Execute code that traverses a dataset

**Usage**

```
with_dataset(expr)
```

**Arguments**

expr	Expression to execute
------	-----------------------

**Details**

When a dataset iterator reaches the end, an out of range runtime error will occur. You can catch and ignore the error when it occurs by wrapping your iteration code in a call to `with_dataset()` (see the example below for an illustration).

## Examples

```
## Not run:
library(tfdatasets)
dataset <- text_line_dataset("mtcars.csv", record_spec = mtcars_spec) %>%
  dataset_prepare(x = c(mpg, disp), y = cyl) %>%
  dataset_batch(128) %>%
  dataset_repeat(10)

iter <- make_iterator_one_shot(dataset)
next_batch <- iterator_get_next(iter)

with_dataset({
  while(TRUE) {
    batch <- sess$run(next_batch)
    # use batch$x and batch$y tensors
  }
})

## End(Not run)
```

---

zip\_datasets

*Creates a dataset by zipping together the given datasets.*

---

## Description

Merges datasets together into pairs or tuples that contain an element from each dataset.

## Usage

```
zip_datasets(...)
```

## Arguments

... Datasets to zip (or a single argument with a list or list of lists of datasets).

## Value

A dataset



# Index

- \* **Dataset methods**
    - dataset\_shard, 17
  - \* **Feature Spec Functions**
    - dataset\_use\_spec, 21
    - feature\_spec, 24
    - fit.FeatureSpec, 26
    - step\_bucketized\_column, 45
    - step\_categorical\_column\_with\_hash\_bucket, 46
    - step\_categorical\_column\_with\_identity, 48
    - step\_categorical\_column\_with\_vocabulary\_file, 49
    - step\_categorical\_column\_with\_vocabulary\_list, 50
    - step\_crossed\_column, 52
    - step\_embedding\_column, 53
    - step\_indicator\_column, 55
    - step\_numeric\_column, 56
    - step\_remove\_column, 57
    - step\_shared\_embeddings\_column, 58
    - steps, 45
  - \* **Selectors**
    - all\_nominal, 3
    - all\_numeric, 4
    - has\_type, 28
  - \* **dataset methods**
    - dataset\_batch, 4
    - dataset\_cache, 5
    - dataset\_collect, 5
    - dataset\_concatenate, 6
    - dataset\_decode\_delim, 7
    - dataset\_filter, 7
    - dataset\_interleave, 9
    - dataset\_map, 10
    - dataset\_map\_and\_batch, 11
    - dataset\_padded\_batch, 12
    - dataset\_prefetch, 13
    - dataset\_prefetch\_to\_device, 14
    - dataset\_reduce, 16
    - dataset\_repeat, 17
    - dataset\_shuffle, 18
    - dataset\_shuffle\_and\_repeat, 19
    - dataset\_skip, 19
    - dataset\_take, 20
    - dataset\_window, 22
  - \* **datasets**
    - hearts, 29
    - selectors, 43
  - \* **iterator functions**
    - iterator\_get\_next, 31
    - iterator\_initializer, 31
    - iterator\_make\_initializer, 32
    - iterator\_string\_handle, 32
    - make-iterator, 34
  - \* **reading datasets**
    - until\_out\_of\_range, 62
  - \* **scaler**
    - scaler\_min\_max, 42
    - scaler\_standard, 42
  - \* **tensor datasets**
    - sparse\_tensor\_slices\_dataset, 43
    - tensor\_slices\_dataset, 60
    - tensors\_dataset, 60
  - \* **text datasets**
    - text\_line\_dataset, 61
- 
- all\_nominal, 3, 4, 28
  - all\_nominal(), 43
  - all\_numeric, 3, 4, 28
  - all\_numeric(), 43
  - compile(), 37
  - contains(), 43
  - csv\_record\_spec (delim\_record\_spec), 22
  - cur\_info\_env (selectors), 43
  - dataset\_batch, 4, 5–9, 11–14, 16–20, 22
  - dataset\_cache, 4, 5, 6–9, 11–14, 16–20, 22

- dataset\_collect, [4](#), [5](#), [5](#), [6–9](#), [11–14](#), [16–20](#), [22](#)
- dataset\_concatenate, [4–6](#), [6](#), [7–9](#), [11–14](#), [16–20](#), [22](#)
- dataset\_decode\_delim, [4–6](#), [7](#), [8](#), [9](#), [11–14](#), [16–20](#), [22](#)
- dataset\_filter, [4–7](#), [7](#), [9](#), [11–14](#), [16–20](#), [22](#)
- dataset\_flat\_map, [8](#)
- dataset\_interleave, [4–8](#), [9](#), [11–14](#), [16–20](#), [22](#)
- dataset\_map, [4–9](#), [10](#), [12–14](#), [16–20](#), [22](#)
- dataset\_map(), [62](#)
- dataset\_map\_and\_batch, [4–9](#), [11](#), [11](#), [13](#), [14](#), [16–20](#), [22](#)
- dataset\_padded\_batch, [4–9](#), [11](#), [12](#), [12](#), [13](#), [14](#), [16–20](#), [22](#)
- dataset\_prefetch, [4–9](#), [11–13](#), [13](#), [14](#), [16–20](#), [22](#)
- dataset\_prefetch\_to\_device, [4–9](#), [11–13](#), [14](#), [16–20](#), [22](#)
- dataset\_prepare, [15](#)
- dataset\_reduce, [4–9](#), [11–14](#), [16](#), [17–20](#), [22](#)
- dataset\_repeat, [4–9](#), [11–14](#), [16](#), [17](#), [18–20](#), [22](#)
- dataset\_shard, [17](#)
- dataset\_shuffle, [4–9](#), [11–14](#), [16](#), [17](#), [18](#), [19](#), [20](#), [22](#)
- dataset\_shuffle\_and\_repeat, [4–9](#), [11–14](#), [16–18](#), [19](#), [20](#), [22](#)
- dataset\_skip, [4–9](#), [11–14](#), [16–19](#), [19](#), [20](#), [22](#)
- dataset\_take, [4–9](#), [11–14](#), [16–20](#), [20](#), [22](#)
- dataset\_use\_spec, [21](#), [25](#), [27](#), [45–48](#), [50–52](#), [54](#), [55](#), [57–59](#)
- dataset\_use\_spec(), [25](#), [27](#)
- dataset\_window, [4–9](#), [11–14](#), [16–20](#), [22](#)
- delim\_record\_spec, [22](#)
- delim\_record\_spec(), [7](#), [61](#)
- dense\_features, [24](#)
  
- ends\_with(), [43](#)
- evaluate, [30](#)
- everything(), [43](#)
  
- feature\_spec, [21](#), [24](#), [27](#), [45–48](#), [50–52](#), [54](#), [55](#), [57–59](#)
- feature\_spec(), [21](#), [24](#), [26](#), [27](#), [46–49](#), [51–53](#), [55–57](#), [59](#)
- file\_list\_dataset, [25](#)
  
- fit.FeatureSpec, [21](#), [25](#), [26](#), [45–48](#), [50–52](#), [54–59](#)
- fit.FeatureSpec(), [21](#), [25](#)
- fixed\_length\_record\_dataset, [27](#)
  
- has\_type, [3](#), [4](#), [28](#)
- has\_type(), [43](#)
- hearts, [29](#)
  
- input\_fn(input\_fn.tf\_dataset), [30](#)
- input\_fn(), [16](#)
- input\_fn.tf\_dataset, [30](#)
- iterator\_get\_next, [31](#), [31](#), [32](#), [33](#), [35](#)
- iterator\_initializer, [31](#), [31](#), [32](#), [33](#), [35](#)
- iterator\_initializer(), [35](#)
- iterator\_make\_initializer, [31](#), [32](#), [33](#), [35](#)
- iterator\_make\_initializer(), [35](#)
- iterator\_string\_handle, [31](#), [32](#), [32](#), [35](#)
- iterator\_string\_handle(), [34](#)
  
- keras::layer\_dense\_features(), [33](#)
  
- layer\_input(), [37](#)
- layer\_input\_from\_dataset, [33](#)
  
- make\_iterator, [34](#)
- make\_csv\_dataset, [35](#)
- make\_iterator\_from\_string\_handle  
(make\_iterator), [34](#)
- make\_iterator\_from\_structure  
(make\_iterator), [34](#)
- make\_iterator\_initializable  
(make\_iterator), [34](#)
- make\_iterator\_one\_shot (make\_iterator),  
[34](#)
- matches(), [43](#)
  
- next\_batch, [37](#)
  
- one\_of(), [43](#)
- out\_of\_range\_handler  
(until\_out\_of\_range), [62](#)
- output\_shapes(output\_types), [39](#)
- output\_shapes(), [7](#), [9–11](#)
- output\_types, [39](#)
- output\_types(), [7](#), [9–11](#)
  
- predict, [30](#)
  
- range\_dataset, [39](#)

read\_files, 40  
rlang::as\_function(), 10, 11

sample\_from\_datasets, 41  
scaler, 41, 42, 56  
scaler\_min\_max, 41, 42, 42  
scaler\_standard, 41, 42, 42  
selectors, 43, 45–49, 51–53, 55–57, 59  
sparse\_tensor\_slices\_dataset, 43, 60  
sql\_dataset (sql\_record\_spec), 44  
sql\_record\_spec, 44  
sqlite\_dataset (sql\_record\_spec), 44  
starts\_with(), 43  
step\_bucketized\_column, 21, 25, 27, 45, 45,  
47, 48, 50–52, 54, 55, 57–59  
step\_bucketized\_column(), 45  
step\_categorical\_column\_with\_hash\_bucket,  
21, 25, 27, 45, 46, 46, 48, 50–52, 54,  
55, 57–59  
step\_categorical\_column\_with\_hash\_bucket(),  
45  
step\_categorical\_column\_with\_identity,  
21, 25, 27, 45–47, 48, 50–52, 54, 55,  
57–59  
step\_categorical\_column\_with\_identity(),  
45  
step\_categorical\_column\_with\_vocabulary\_file,  
21, 25, 27, 45–48, 49, 51, 52, 54, 55,  
57–59  
step\_categorical\_column\_with\_vocabulary\_file(),  
45  
step\_categorical\_column\_with\_vocabulary\_list,  
21, 25, 27, 45–48, 50, 50, 52, 54, 55,  
57–59  
step\_categorical\_column\_with\_vocabulary\_list(),  
45  
step\_crossed\_column, 21, 25, 27, 45–48, 50,  
51, 52, 54, 55, 57–59  
step\_crossed\_column(), 45  
step\_embedding\_column, 21, 25, 27, 45–48,  
50–52, 53, 55, 57–59  
step\_embedding\_column(), 45  
step\_indicator\_column, 21, 25, 27, 45–48,  
50–52, 54, 55, 57–59  
step\_indicator\_column(), 45  
step\_numeric\_column, 21, 25, 27, 41, 45–48,  
50–52, 54, 55, 56, 58, 59  
step\_numeric\_column(), 45  
step\_remove\_column, 21, 25, 27, 45–48,  
50–52, 54, 55, 57, 57, 59  
step\_remove\_column(), 45  
step\_shared\_embeddings\_column, 21, 25,  
27, 45–48, 50–52, 54, 55, 57, 58, 58  
step\_shared\_embeddings\_column(), 45  
steps, 21, 25, 27, 28, 45, 46–48, 50–52, 54,  
55, 57–59

tensor\_slices\_dataset, 44, 60, 60  
tensors\_dataset, 44, 60, 60  
text\_line\_dataset, 61  
text\_line\_dataset(), 30, 40  
tfrecord\_dataset, 61  
tfrecord\_dataset(), 30, 40  
tidyselect::select\_helpers(), 24  
train, 30  
tsv\_record\_spec (delim\_record\_spec), 22

until\_out\_of\_range, 62

with\_dataset, 63

zip\_datasets, 64