# Package 'tensorflow'

May 21, 2022

```
Description Interface to 'TensorFlow' <a href="https://www.tensorflow.org/">https://www.tensorflow.org/</a>,
      an open source software library for numerical computation using data
      flow graphs. Nodes in the graph represent mathematical operations,
      while the graph edges represent the multidimensional data arrays
      (tensors) communicated between them. The flexible architecture allows
      you to deploy computation to one or more 'CPUs' or 'GPUs' in a desktop,
      server, or mobile device with a single 'API'. 'TensorFlow' was originally
      developed by researchers and engineers working on the Google Brain Team
      within Google's Machine Intelligence research organization for the
      purposes of conducting machine learning and deep neural networks research,
      but the system is general enough to be applicable in a wide variety
      of other domains as well.
License Apache License 2.0
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BugReports https://github.com/rstudio/tensorflow/issues
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```

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2 all\_dims

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all_d	lims All dims	

### Description

This function returns an object that can be used when subsetting tensors with [. If you are familiar with python,, this is equivalent to the python Ellipsis . . . , (not to be confused with . . . in R).

### Usage

all\_dims()

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#### **Examples**

```
## Not run:
# in python, if x is a numpy array or tensorflow tensor
x[..., i]
# the ellipsis means "expand to match number of dimension of x".
# to translate the above python expression to R, write:
x[all_dims(), i]
## End(Not run)
```

as\_tensor

as\_tensor

### **Description**

Coerce objects to tensorflow tensors (potentially of a specific dtype or shape). The provided default methods will call <code>tf.convert\_to\_tensor</code>. Depending on arguments supplied it may also call some combination of

- tf.dtypes.saturate\_cast
- tf.fill or tf.reshape

•

### Usage

```
as_tensor(x, dtype = NULL, ..., name = NULL)
## Default S3 method:
as_tensor(x, dtype = NULL, ..., shape = NULL, name = NULL)
## S3 method for class 'double'
as_tensor(x, dtype = NULL, ..., name = NULL)
```

#### **Arguments**

X	object to convert
dtype	NULL, a tensorflow dtype (tf\$int32), or something coercible to one (e.g. a string "int32")
,	ignored
name	NULL or a string. Useful for debugging in graph mode, ignored while in eager mode.
shape	an integer vector, tensor, or tf.TensorShape. Can contain up to 1 unspecified dimension, encoded as a -1 or NA. This will reshape x using row-major (C-style) semantics. It will prefer reshaping using non-graph operations if possible, but will otherwise invoke tf\$reshape(). If x is a scalar and the requested shape

requested shape (it will dispatch to tf\$fill()).

is fully defined or a tensor, the value of x will be recycled to fill a tensor of the

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#### Value

a tensorflow tensor

### **Examples**

```
## Not run:
as_tensor(42, "int32")
as_tensor(as_tensor(42))
## End(Not run)
```

evaluate

Evaluate a Model

### **Description**

Evaluate a model object. See implementations in the keras and tfestimators packages.

#### Usage

```
evaluate(object, ...)
```

### **Arguments**

object An evaluatable R object.

... Optional arguments passed on to implementing methods.

### **Implementations**

- keras
- tfestimators

 $install\_tensorflow$ 

Install TensorFlow and its dependencies

### **Description**

install\_tensorflow() installs just the tensorflow python package and it's direct dependencies. For a more complete installation that includes additional optional dependencies, use keras::install\_keras().

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#### Usage

```
install_tensorflow(
  method = c("auto", "virtualenv", "conda"),
  conda = "auto",
  version = "default",
  envname = NULL,
  extra_packages = NULL,
  restart_session = TRUE,
  conda_python_version = NULL,
  ...,
  pip_ignore_installed = TRUE,
  python_version = conda_python_version
)
```

#### **Arguments**

method

Installation method. By default, "auto" automatically finds a method that will work in the local environment. Change the default to force a specific installation method. Note that the "virtualenv" method is not available on Windows.

conda

The path to a conda executable. Use "auto" to allow reticulate to automatically find an appropriate conda binary. See **Finding Conda** and conda\_binary() for more details.

version

TensorFlow version to install. Valid values include:

- "default" installs 2.9
- "release" installs the latest release version of tensorflow (which may be incompatible with the current version of the R package)
- A version specification like "2.4" or "2.4.0". Note that if the patch version is not supplied, the latest patch release is installed (e.g., "2.4" today installs version "2.4.2")
- nightly for the latest available nightly build.
- To any specification, you can append "-cpu" to install the cpu version only of the package (e.g., "2.4-cpu")
- The full URL or path to a installer binary or python \*.whl file.

envname

The name, or full path, of the environment in which Python packages are to be installed. When NULL (the default), the active environment as set by the RETICULATE\_PYTHON\_ENV variable will be used; if that is unset, then the r-reticulate environment will be used.

extra\_packages Additional Python packages to install along with TensorFlow. restart\_session

Restart R session after installing (note this will only occur within RStudio).

other arguments passed to reticulate::conda\_install() or reticulate::virtualenv\_install(), depending on the method used.

pip\_ignore\_installed

Whether pip should ignore installed python packages and reinstall all already installed python packages. This defaults to TRUE, to ensure that TensorFlow dependencies like NumPy are compatible with the prebuilt TensorFlow binaries.

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```
python_version, conda_python_version
```

Pass a string like "3.8" to request that conda install a specific Python version. This is ignored when attempting to install in a Python virtual environment. Note that the Python version must be compatible with the requested Tensorflow version, documented here: https://www.tensorflow.org/install/pip#system-requirements

#### **Details**

You may be prompted to download and install miniconda if reticulate did not find a non-system installation of python. Miniconda is the recommended installation method for most users, as it ensures that the R python installation is isolated from other python installations. All python packages will by default be installed into a self-contained conda or venv environment named "r-reticulate". Note that "conda" is the only supported method on M1 Mac.

If you initially declined the miniconda installation prompt, you can later manually install miniconda by running reticulate::install\_miniconda().

#### **Custom Installation**

install\_tensorflow() or keras::install\_keras() isn't required to use tensorflow with the package. If you manually configure a python environment with the required dependencies, you can tell R to use it by pointing reticulate at it, commonly by setting an environment variable:

```
Sys.setenv("RETICULATE_PYTHON" = "~/path/to/python-env/bin/python")
```

#### **Apple Silicon**

Tensorflow on Apple Silicon is not officially supported by the Tensorflow maintainers. However Apple has published a custom version of Tensorflow compatible with Arm Macs. install\_tensorflow() will install the special packages tensorflow-macos and tensorflow-metal on Arm Macs. See <a href="https://developer.apple.com/metal/tensorflow-plugin/">https://developer.apple.com/metal/tensorflow-macos</a> and tensorflow-metal on Arm Macs. See <a href="https://developer.apple.com/metal/tensorflow-plugin/">https://developer.apple.com/metal/tensorflow-plugin/</a> for instructions on how to do the equivalent manually. Please note that this is an experimental build of both Python and Tensorflow, with known issues. In particular, certain operations will cause errors, but can often be remedied by pinning them to the CPU. For example:

```
x <- array(runif(64*64), c(1, 64, 64))
keras::layer_random_rotation(x, .5) # Error:
# No registered 'RngReadAndSkip' OpKernel for 'GPU' devices
# Pin the operation to the CPU to avoid the error
with(tf$device("CPU"), keras::layer_random_rotation(x, .5) ) # No Error</pre>
```

#### **Additional Packages**

If you wish to add additional PyPI packages to your Keras / TensorFlow environment you can either specify the packages in the extra\_packages argument of install\_tensorflow() or install\_keras(), or alternatively install them into an existing environment using the reticulate::py\_install() function. Note that install\_keras() includes a set of additional python packages by default, see ?keras::install\_keras for details.

#### See Also

```
keras::install_keras()
```

install\_tensorflow\_extras

(Defunct) Install additional Python packages alongside TensorFlow

#### **Description**

This function is deprecated. Use the extra\_packages argument to install\_tensorflow() or reticulate::py\_install() to install additional packages.

### Usage

```
install_tensorflow_extras(packages, conda = "auto")
```

### **Arguments**

packages Python packages to install

conda Path to conda executable (or "auto" to find conda using the PATH and other

conventional install locations). Only used when TensorFlow is installed within

a conda environment.

parse\_arguments

Parse Command Line Arguments

### Description

Parse command line arguments of the form --key=value and --key value. The values are assumed to be valid yaml and will be converted using yaml.load().

#### Usage

```
parse_arguments(arguments = NULL)
```

#### Arguments

arguments

A vector of command line arguments. When NULL (the default), the command line arguments received by the current R process are used.

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parse\_flags

Parse Configuration Flags for a TensorFlow Application

#### **Description**

Parse configuration flags for a TensorFlow application. Use this to parse and unify the configuration(s) specified through a flags.yml configuration file, alongside other arguments set through the command line.

#### Usage

```
parse_flags(
  config = Sys.getenv("R_CONFIG_ACTIVE", unset = "default"),
  file = "flags.yml",
  arguments = commandArgs(TRUE)
)
```

### Arguments

config The configuration to use. Defaults to the active configuration for the current

environment (as specified by the R\_CONFIG\_ACTIVE environment variable), or

default when unset.

file The configuration file to read.

arguments The command line arguments (as a character vector) to be parsed.

#### Value

A named R list, mapping configuration keys to values.

```
## Not run:
# examine an example configuration file provided by tensorflow
file <- system.file("examples/config/flags.yml", package = "tensorflow")
cat(readLines(file), sep = "\n")

# read the default configuration
FLAGS <- tensorflow::parse_flags("default", file = file)
str(FLAGS)

# read the alternate configuration: note that
# the default configuration is inherited, but
# we override the 'string' configuration here
FLAGS <- tensorflow::parse_flags("alternate", file = file)
str(FLAGS)

# override configuration values using command
# line arguments (normally, these would be</pre>
```

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```
# passed in through the command line invocation
# used to start the process)
FLAGS <- tensorflow::parse_flags(
   "alternate",
   file = file,
   arguments = c("--foo=1")
)
str(FLAGS)
## End(Not run)</pre>
```

set\_random\_seed

Set random seed for TensorFlow

### Description

Sets all random seeds needed to make TensorFlow code reproducible.

### Usage

```
set_random_seed(seed, disable_gpu = TRUE)
```

#### **Arguments**

seed A single value, interpreted as an integer
disable\_gpu TRUE to disable GPU execution (see *Parallelism* below).

#### **Details**

This function should be used instead of use\_session\_with\_seed() if you are using TensorFlow >= 2.0, as the concept of session doesn't really make sense anymore.

This functions sets:

- The R random seed with set.seed().
- The python and Numpy seeds via (reticulate::py\_set\_seed()).
- The TensorFlow seed with (tf\$random\$set\_seed())

It also optionally disables the GPU execution as this is a potential source of non-reproducibility.

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shape

 $Create\ a\ {\tt tf.TensorShape}\ object$ 

#### **Description**

Create a tf. TensorShape object

### Usage

```
shape(..., dims = list(...))
```

#### **Arguments**

... Tensor dimensions as integers or NULL for an unknown dimensions. NA and -1 are synonyms for NULL.

dims Tensor dimensions as a vector.

#### See Also

https://www.tensorflow.org/api\_docs/python/tf/TensorShape

```
## Not run:
# --- construct ---
           # tf.TensorShape()
                                       # scalar
shape()
shape(NULL) # tf.TensorShape([None]) # 1-D array of unknown length
shape(NA) # tf.TensorShape([None]) # 1-D array of unknown length, NA is a synonym for NULL
shape(dims = NULL) # TensorShape(None)
                                          # Unknown rank, unknown size
shape(3, 4)
                   # TensorShape([3, 4]) # 2-D array (matrix) with 3 rows, 4 columns
                  # TensorShape([None, 4]) # 2-D array (matrix) with unknown rows, 4 columns
shape(NA, 4)
shape(dims = c(NA, 4)) # TensorShape([None, 4]) # same as above; bypass ... and pass dims directly
# --- inspect ---
length(shape(dims = NULL)) # NA_integer_
length(shape(1,2,3,NA))
# ---convert ---
x <- shape(dims = list(3L, 5L))</pre>
            # list(3L, 5L)
as.list(x)
as.integer(x) \# c(3L, 5L)
as.numeric(x) \# c(3, 5)
as.double(x) \# c(3, 5) \# alias for as.numeric
as_tensor(x) # tf.Tensor([3 5], shape=(2,), dtype=int32)
# convert partially undefined shapes
x <- shape(NA, 3)
```

shape 11

```
# list(NULL, 3L)
as.list(x)
as.integer(x) \# c(NA, 3L)
as_tensor(x) # tf.Tensor([-1 3], shape=(2,), dtype=int32) # unspecified dims default is -1
# as_tensor() converts undefined dimensions to -1, which is useful for
# tf functions that only accept tensors for shapes, e.g,
tf$reshape(tf$zeros(shape(8)),
           as_tensor(shape(NA, 4)))
# tf.Tensor([[0. 0. 0. 0.]
             [0. 0. 0. 0.]], shape=(2, 4), dtype=float32)
# converting fully unknown shapes raises an error
try(as.list(shape(dims = NULL))) # ValueError: as_list() is not defined on an unknown TensorShape.
# test for rank first if this a concern:
as.list_or_null <- function(x) if(is.na(length(x))) NULL else as.list(x)
as.list_or_null(shape(dims = NULL))
# --- compare ---
# Fully known shapes return TRUE if and only if each element is equal
shape(3, 4) == shape(3, 4) # TRUE
shape(3, 4) == shape(4, 4) # FALSE
# two unknown dimensions are treated as equal
shape(NA, 4) == shape(NA, 4) # TRUE
shape(NA, 4) == shape(3, 4) # FALSE
# Two unknown shapes, return TRUE
shape(dims = NULL) == shape(dims = NULL) # TRUE
# Comparing an unknown shape to a partially or fully defined shape returns FALSE
shape(dims = NULL) == shape(NULL) # FALSE
shape(dims = NULL) == shape(4)
# --- extract or replace ---
# regular R-list semantics for `[`, `[[`, `[<-`, `[[<-`</pre>
x <- shape(1, 2, 3)
x[1]
           # TensorShape([1])
x[[1]]
           # 1L
          # TensorShape([2, 3])
x[2:3]
x[-1]
           # TensorShape([2, 3])
x[1] <- 11
                 ; x # TensorShape([11, 2, 3])
x[1] \leftarrow shape(11) ; x # TensorShape([11, 2, 3])
x[1] \leftarrow list(11) ; x # TensorShape([11, 2, 3])
                      ; x # TensorShape([22, 2, 3])
x[[1]] <- 22
x[1:2] \leftarrow c(NA, 99); x # TensorShape([None, 99, 3])
x[1:2] \leftarrow shape(33, 44) ; x # TensorShape([33, 44, 3])
# --- concatenate ---
c(shape(1), shape(2, 3), shape(4, NA)) # TensorShape([1, 2, 3, 4, None])
```

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tensorboard

TensorBoard Visualization Tool

### Description

TensorBoard is a tool inspecting and understanding your TensorFlow runs and graphs.

### Usage

```
tensorboard(
  log_dir,
  action = c("start", "stop"),
  host = "127.0.0.1",
  port = "auto",
  launch_browser = getOption("tensorflow.tensorboard.browser", interactive()),
  reload_interval = 5,
  purge_orphaned_data = TRUE
)
```

### Arguments

reload\_interval

log_dir	Directories to scan for training logs. If this is a named character vector then the specified names will be used as aliases within TensorBoard.
action	Specify whether to start or stop TensorBoard (TensorBoard will be stopped automatically when the R session from which it is launched is terminated).
host	Host for serving TensorBoard
port	Port for serving TensorBoard. If "auto" is specified (the default) then an unused port will be chosen automatically.
launch_browser	Open a web browser for TensorBoard after launching. Defaults to TRUE in interactive sessions. When running under RStudio uses an RStudio window by default (pass a function e.g. utils::browseURL() to open in an external browser). Use the tensorflow.tensorboard.browser option to establish a global default behavior.

How often the backend should load more data.

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purge\_orphaned\_data

Whether to purge data that may have been orphaned due to TensorBoard restarts. Disabling purge\_orphaned\_data can be used to debug data disappearance.

#### **Details**

When TensorBoard is passed a logdir at startup, it recursively walks the directory tree rooted at logdir looking for subdirectories that contain the tensor data. Every time it encounters such a subdirectory, it loads it as a new run, and the frontend will organize the data accordingly.

The TensorBoard process will be automatically destroyed when the R session in which it is launched exits. You can pass action = "stop" to manually terminate TensorBoard.

#### Value

URL for browsing TensorBoard (invisibly).

tensorflow

TensorFlow for R

### **Description**

TensorFlow is an open source software library for numerical computation using data flow graphs. Nodes in the graph represent mathematical operations, while the graph edges represent the multidimensional data arrays (tensors) communicated between them. The flexible architecture allows you to deploy computation to one or more CPUs or GPUs in a desktop, server, or mobile device with a single API.

#### **Details**

The TensorFlow API is composed of a set of Python modules that enable constructing and executing TensorFlow graphs. The tensorflow package provides access to the complete TensorFlow API from within R.

For additional documentation on the tensorflow package see https://tensorflow.rstudio.com

tf

Main TensorFlow module

#### **Description**

Interface to main TensorFlow module. Provides access to top level classes and functions as well as sub-modules (e.g. tf\$nn, tf\$contrib\$learn, etc.).

#### Usage

tf

#### **Format**

TensorFlow module

#### **Examples**

```
## Not run:
library(tensorflow)

hello <- tf$constant('Hello, TensorFlow!')
zeros <- tf$Variable(tf$zeros(shape(1L)))

tf$print(hello)
tf$print(zeros)

## End(Not run)</pre>
```

tfe\_enable\_eager\_execution

(Deprecated) Enables, for the rest of the lifetime of this program, eager execution.

#### **Description**

This function is no longer needed since Tensorflow 2.0, when eager execution became the default.

#### Usage

```
tfe_enable_eager_execution(
  config = NULL,
  device_policy = c("explicit", "warn", "silent")
)
```

### **Arguments**

config

(Optional) A tf\$ConfigProto() protocol buffer with configuration options for the Context. Note that a lot of these options may be currently unimplemented or irrelevant when eager execution is enabled.

device\_policy

(Optional) What policy to use when trying to run an operation on a device with inputs which are not on that device. Valid values: "explicit": raises an error if the placement is not correct. "warn": copies the tensors which are not on the right device but raises a warning. "silent": silently copies the tensors. This might hide performance problems.

#### **Details**

If not called immediately on startup risks creating breakage and bugs.

After eager execution is enabled, operations are executed as they are defined and tensors hold concrete values, and can be accessed as R matrices or arrays with as.matrix(), as.array(), as.double(), etc.

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#### **Examples**

```
## Not run:
# load tensorflow and enable eager execution
library(tensorflow)
tfe_enable_eager_execution()
# create a random 10x10 matrix
x <- tf$random$normal(shape(10, 10))
# use it in R via as.matrix()
heatmap(as.matrix(x))
## End(Not run)</pre>
```

tf\_extract\_opts

Tensor extract options

#### **Description**

Tensor extract options

#### Usage

```
tf_extract_opts(
    style = getOption("tensorflow.extract.style"),
    ...,
    one_based = getOption("tensorflow.extract.one_based", TRUE),
    inclusive_stop = getOption("tensorflow.extract.inclusive_stop", TRUE),
    disallow_out_of_bounds = getOption("tensorflow.extract.dissallow_out_of_bounds",
        TRUE),
    warn_tensors_passed_asis = getOption("tensorflow.extract.warn_tensors_passed_asis",
        TRUE),
    warn_negatives_pythonic = getOption("tensorflow.extract.warn_negatives_pythonic",
        TRUE)
)
```

#### **Arguments**

```
one of NULL (the default) "R" or "python". If supplied, this overrides all other options. "python" is equivalent to all the other arguments being FALSE. "R" is equivalent to warn_tensors_passed_asis and warn_negatives_pythonic set to FALSE
... ignored

one_based TRUE or FALSE, if one-based indexing should be used
inclusive_stop TRUE or FALSE, if slices like start:stop should be inclusive of stop
```

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```
disallow_out_of_bounds
```

TRUE or FALSE, whether checks are performed on the slicing index to ensure it is within bounds.

```
warn_tensors_passed_asis
```

TRUE or FALSE, whether to emit a warning the first time a tensor is supplied to [ that tensors are passed as-is, with no R to python translation

```
warn_negatives_pythonic
```

TRUE or FALSE, whether to emit a warning the first time a negative number is supplied to [ about the non-standard (python-style) interpretation

#### Value

an object with class "tf\_extract\_opts", suitable for passing to [.tensorflow.tensor()

### Examples

```
## Not run:
x <- tf$constant(1:10)

opts <- tf_extract_opts("R")
x[1, options = opts]

# or for more fine-grained control
opts <- tf_extract_opts(
    one_based = FALSE,
    warn_tensors_passed_asis = FALSE,
    warn_negatives_pythonic = FALSE
)
x[0:2, options = opts]

## End(Not run)</pre>
```

tf\_function

Creates a callable TensorFlow graph from an R function.

### Description

tf\_function constructs a callable that executes a TensorFlow graph created by tracing the TensorFlow operations in f. This allows the TensorFlow runtime to apply optimizations and exploit parallelism in the computation defined by f.

#### Usage

```
tf_function(f, input_signature = NULL, autograph = TRUE, ...)
```

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#### **Arguments**

f the function to be compiled

input\_signature

A possibly nested sequence of tf\$TensorSpec objects specifying the shapes and dtypes of the tensors that will be supplied to this function. If NULL, a separate function is instantiated for each inferred input signature. If input\_signature

is specified, every input to f must be a tensor.

autograph TRUE or FALSE. If TRUE (the default), you can use tensors in R control flow

expressions if, while, for and break and they will be traced into the tensorflow graph. A guide to getting started and additional details can be found: here

... additional arguments passed on to tf.function (vary based on Tensorflow ver-

sion). See here for details.

#### **Details**

A guide to getting started with tf. function can be found here.

tf\_probability

TensorFlow Probability Module

### Description

TensorFlow Probability Module

### Usage

```
tf_probability()
```

#### Value

Reference to TensorFlow Probability functions and classes

```
## Not run:
library(tensorflow)
tfp <- tf_probability()
tfp$distributions$Normal(loc=0, scale=1)
## End(Not run)</pre>
```

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train\_and\_evaluate

(Deprecated) Simultaneously Train and Evaluate a Model

### Description

Train and evaluate a model object. See implementation in the tfestimators package.

### Usage

```
train_and_evaluate(object, ...)
```

### **Arguments**

object

An R object.

. . .

Optional arguments passed on to implementing methods.

### **Implementations**

• tfestimators

use\_compat

Use Compatibility

### Description

Enables TensorFlow to run under a different API version for compatibility with previous versions. For instance, this is useful to run TensorFlow 1.x code when using TensorFlow 2.x.

#### Usage

```
use_compat(version = c("v1", "v2"))
```

### **Arguments**

version

The version to activate. Must be "v1" or "v2"

```
## Not run:
library(tensorflow)
use_compat("v1")
## End(Not run)
```

use\_session\_with\_seed

### **Description**

Set various random seeds required to ensure reproducible results. The provided seed value will establish a new random seed for R, Python, NumPy, and TensorFlow. GPU computations and CPU parallelism will also be disabled by default.

#### Usage

```
use_session_with_seed(
  seed,
  disable_gpu = TRUE,
  disable_parallel_cpu = TRUE,
  quiet = FALSE
)
```

### **Arguments**

```
seed A single value, interpreted as an integer disable_gpu TRUE to disable GPU execution (see Parallelism below). disable_parallel_cpu TRUE to disable CPU parallelism (see Parallelism below). quiet TRUE to suppress printing of messages.
```

#### **Details**

This function must be called at the very top of your script (i.e. immediately after library(tensorflow), library(keras), etc.). Any existing TensorFlow session is torn down via tf\$reset\_default\_graph().

This function takes all measures known to promote reproducible results from TensorFlow sessions, however it's possible that various individual TensorFlow features or dependent libraries escape its effects. If you encounter non-reproducible results please investigate the possible sources of the problem, contributions via pull request are very welcome!

Packages which need to be notified before and after the seed is set can register for the "tensor-flow.on\_before\_use\_session" and "tensorflow.on\_use\_session" hooks (see setHook()) for additional details on hooks).

#### Value

TensorFlow session object, invisibly

#### **Parallelism**

By default the use\_session\_with\_seed() function disables GPU and CPU parallelism, since both can result in non-deterministic execution patterns (see <a href="https://stackoverflow.com/questions/42022950/">https://stackoverflow.com/questions/42022950/</a>). You can optionally enable GPU or CPU parallelism by setting the disable\_gpu and/or disable\_parallel\_cpu parameters to FALSE.

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### **Examples**

```
## Not run:
library(tensorflow)
use_session_with_seed(42)
## End(Not run)
```

view\_savedmodel

View a Saved Model

### Description

View a serialized model from disk.

### Usage

```
view_savedmodel(model_dir)
```

### **Arguments**

model\_dir

The path to the exported model, as a string.

### Value

URL for browsing TensorBoard (invisibly).

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Subset tensors with [

### Description

Subset tensors with [

### Usage

```
## $3 method for class 'tensorflow.tensor'

x[
...,
drop = TRUE,
style = getOption("tensorflow.extract.style"),
options = tf_extract_opts(style)
]
```

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### Arguments

```
x Tensorflow tensor
... slicing specs. See examples and details.
drop whether to drop scalar dimensions
style One of "python" or "R".
options An object returned by tf_extract_opts()
```

```
## Not run:
x \leftarrow as_{tensor}(array(1:15, dim = c(3, 5)))
# by default, numerics supplied to [...] are interpreted R style
x[,1] # first column
x[1:2,] # first two rows
x[,1, drop = FALSE] # 1 column matrix
# strided steps can be specified in R syntax or python syntax
x[, seq(1, 5, by = 2)]
x[, 1:5:2]
# if you are unfamiliar with python-style strided steps, see:
# https://numpy.org/doc/stable/reference/arrays.indexing.html#basic-slicing-and-indexing
# missing arguments for python syntax are valid, but they must by backticked
# or supplied as NULL
x[, `::2`]
x[, NULL:NULL:2]
x[, `2:`]
# all_dims() expands to the shape of the tensor
# (equivalent to a python ellipsis `...`)
# (not to be confused with R dots `...`)
y \leftarrow as_{tensor}(array(1:(3^5), dim = c(3,3,3,3,3)))
all.equal(y[all_dims(), 1],
          y[, , , , 1])
# tf$newaxis are valid (equivalent to a NULL)
x[,, tf$newaxis]
x[,, NULL]
# negative numbers are always interpreted python style
# The first time a negative number is supplied to `[`, a warning is issued
# about the non-standard behavior.
x[-1,] # last row, with a warning
x[-1,] # the warning is only issued once
# specifying `style = 'python'` changes the following:
```

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```
# + zero-based indexing is used
# + slice sequences in the form of `start:stop` do not include `stop`
     in the returned value
# + out-of-bounds indices in a slice are valid
# The style argument can be supplied to individual calls of `[` or set
# as a global option
# example of zero based indexing
x[0, , style = 'python'] # first row
x[1, , style = 'python'] # second row
# example of slices with exclusive stop
options(tensorflow.extract.style = 'python')
x[, 0:1] # just the first column
x[, 0:2] # first and second column
# example of out-of-bounds index
x[, 0:10]
options(tensorflow.extract.style = NULL)
# slicing with tensors is valid too, but note, tensors are never
# translated and are always interpreted python-style.
# A warning is issued the first time a tensor is passed to `[`
x[, tf$constant(0L):tf$constant(2L)]
# just as in python, only scalar tensors are valid
# https://www.tensorflow.org/api_docs/python/tf/Tensor#__getitem__
# To silence the warnings about tensors being passed as-is and negative numbers
# being interpreted python-style, set
options(tensorflow.extract.style = 'R')
# clean up from examples
options(tensorflow.extract.style = NULL)
## End(Not run)
```

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