# Package 'sparktf'

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Type Package
Title Interface for 'TensorFlow' 'TFRecord' Files with 'Apache Spark'
Version 0.1.0
<b>Description</b> A 'sparklyr' extension that enables reading and writing 'TensorFlow' TFRecord files via 'Apache Spark'.
<b>License</b> Apache License (>= 2.0)
Encoding UTF-8
SystemRequirements TensorFlow (https://www.tensorflow.org/)
LazyData true
<b>Depends</b> R (>= $3.1.2$ )
Imports sparklyr (>= 1.0)
RoxygenNote 6.1.0
Suggests testthat, dplyr
NeedsCompilation no
<b>Author</b> Kevin Kuo [aut, cre] ( <a href="https://orcid.org/0000-0001-7803-7901">https://orcid.org/0000-0001-7803-7901</a> )
Maintainer Kevin Kuo <kevin.kuo@rstudio.com></kevin.kuo@rstudio.com>
Repository CRAN
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spark\_read\_tfrecord Read a TFRecord File

### **Description**

Read a TFRecord file as a Spark DataFrame.

## Usage

```
spark_read_tfrecord(sc, name = NULL, path = name, schema = NULL,
  record_type = c("Example", "SequenceExample"), overwrite = TRUE)
```

## Arguments

sc	A spark conneciton.
name	The name to assign to the newly generated table or the path to the file. Note that if a path is provided for the 'name' argument then one cannot specify a name.
path	The path to the file. Needs to be accessible from the cluster. Supports the "hdfs://", "s $3a$ ://" and "file://" protocols.
schema	(Currently unsupported.) Schema of TensorFlow records. If not provided, the schema is inferred from TensorFlow records.
record_type	Input format of TensorFlow records. By default it is Example.
overwrite	Boolean; overwrite the table with the given name if it already exists?

## **Examples**

```
## Not run:
iris_tbl <- copy_to(sc, iris)
data_path <- file.path(tempdir(), "iris")
df1 <- iris_tbl %>%
ft_string_indexer_model(
    "Species", "label",
    labels = c("setosa", "versicolor", "virginica")
)

df1 %>%
spark_write_tfrecord(
    path = data_path,
        write_locality = "local"
)

spark_read_tfrecord(sc, data_path)

## End(Not run)
```

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spark\_write\_tfrecord Write a Spark DataFrame to a TFRecord file

#### **Description**

Serialize a Spark DataFrame to the TensorFlow TFRecord format for training or inference.

### Usage

```
spark_write_tfrecord(x, path, record_type = c("Example",
   "SequenceExample"), write_locality = c("distributed", "local"),
   mode = NULL)
```

#### **Arguments**

x A Spark DataFrame

path The path to the file. Needs to be accessible from the cluster. Supports the

"hdfs://", "s3a://", and "file://" protocols.

record\_type Output format of TensorFlow records. One of "Example" and "SequenceExample".

write\_locality Determines whether the TensorFlow records are written locally on the workers

or on a distributed file system. One of "distributed" and "local". See Details

for more information.

mode A character element. Specifies the behavior when data or table already exists.

Supported values include: 'error', 'append', 'overwrite' and 'ignore'. Notice

that 'overwrite' will also change the column structure.

For more details see also http://spark.apache.org/docs/latest/sql-programming-guide.

html#save-modes for your version of Spark.

#### **Details**

For write\_locality = local, each of the workers stores on the local disk a subset of the data. The subset that is stored on each worker is determined by the partitioning of the DataFrame. Each of the partitions is coalesced into a single TFRecord file and written on the node where the partition lives. This is useful in the context of distributed training, in which each of the workers gets a subset of the data to work on. When this mode is activated, the path provided to the writer is interpreted as a base path that is created on each of the worker nodes, and that will be populated with data from the DataFrame.

#### **Examples**

```
## Not run:
iris_tbl <- copy_to(sc, iris)
data_path <- file.path(tempdir(), "iris")
df1 <- iris_tbl %>%
ft_string_indexer_model(
    "Species", "label",
```

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```
labels = c("setosa", "versicolor", "virginica")
)

df1 %>%
spark_write_tfrecord(
  path = data_path,
   write_locality = "local"
)

## End(Not run)
```

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