
Tarantella

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TARANTELLA

Tarantella is an open-source, distributed Deep Learning framework built on top of TensorFlow, providing scalable Deep Neural Network training on CPU and GPU compute clusters.

Tarantella is easy-to-use, allows to re-use existing TensorFlow models, and does not require any knowledge of parallel computing.

WHY TARANTELLA?

Tarantella is an open-source Deep Learning framework that focuses on providing fast, scalable and efficient training of Deep Neural Networks (DNNs) on High Performance Computing (HPC) clusters.

1.1 Goals

Tarantella is designed to meet the following goals:

Tarantella...

1. ...provides strong scalability
2. ...is easy to use
3. ...follows a synchronous training scheme
4. ...integrates well with existing models
5. ...provides support for GPU and CPU systems

Tarantella provides close to linear speed-up for the training of common Deep Learning architectures, thus considerably reducing the required time-to-accuracy in many Deep Learning workflows. To make this capability accessible to as many users as possible, Tarantella's interface is designed such that its use does not require any expertise in HPC or parallel computing.

To allow integrating Tarantella into any TensorFlow-based Deep Learning workflow, we put special emphasis on strictly following the synchronous optimization scheme used to train DNNs. This guarantees that results obtained in serial execution can be reproduced when using distributed training (cf. however [these guidelines](#)), so that computation can be scaled up at any point in time without losing reproducibility of the results.

Furthermore, we made sure that existing TensorFlow models can be made ready for distributed training with minimal effort (follow the [Quick Start guide](#) to learn more). Tarantella supports distributed training on GPU and pure CPU clusters, independently of the hardware vendors.

1.2 Performance Results

To investigate the scalability of Tarantella distributed training with respect to the number of devices used, we performed several experiments across multiple machines and models used in the fields of computer vision and natural language processing.

We show below some of the results we obtained when training two state-of-the-art models in parallel with Tarantella on two types of machines: the [HPC-DA](#) cluster of the [Technical University of Dresden](#) is a machine designed for data science workloads, equipped with 6 GPUs per node, while [SuperMUC-NG](#) from the [Leibniz Supercomputing Centre](#)

is a typical HPC machine suitable for CPU-intensive simulations. The hardware details of the two machines used in our experiments are shown below.

Cluster	Hardware specifications per node
HPC-DA	<ul style="list-style-type: none"> • 6 x NVIDIA VOLTA V100 GPU with 32GB HBM2 • 2 x IBM Power9 CPU (22 cores @2.80 GHz) • NVLINK bandwidth 150 GB/s between GPUs and host • 2 x 100 Gbit/s Infiniband interconnect between nodes
SuperMUC-NG	<ul style="list-style-type: none"> • 2 x Intel Skylake Xeon Platinum 8174 CPU (48 cores @3.10 GHz) • 100 Gbit/s OmniPath network

First we look at the speedups that Tarantella can achieve when scaling up the number of devices for the ResNet-50 model trained with the ImageNet dataset. ResNet-50 is one of the most studied deep neural networks for computer vision tasks, featuring over *23 million* trainable parameters.

More specifically, Figure 1 illustrates the runtime per epoch on the *HPC-DA* cluster, when using up to 96 GPUs. Figure 2 showcases the same experiment performed on CPUs on the *SuperMUC-NG* machine, showing that training ResNet-50 distributedly scales on up to 256 processes. Compared to the baseline single-device runtime of the ResNet-50 model using TensorFlow 2.2, Tarantella succeeds in training the model **62x faster** on the CPU cluster and **57x faster** on the GPUs.

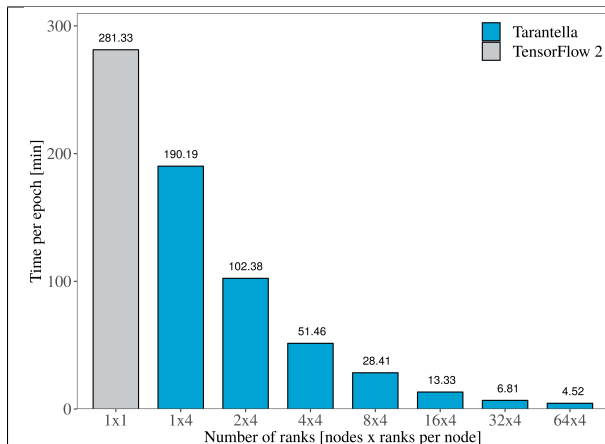


Fig. 1: Figure 1. Training Resnet-50 on CPU nodes

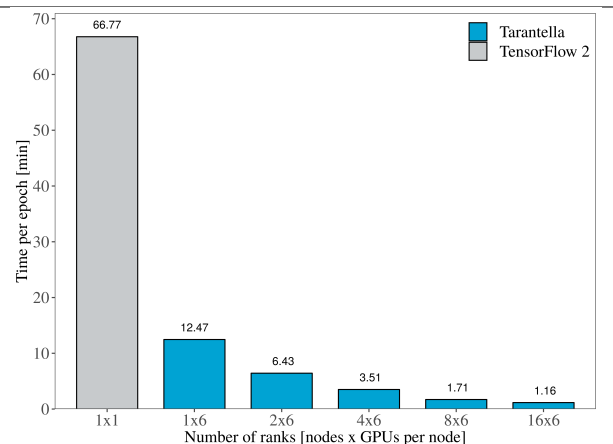


Fig. 2: Figure 2. Training Resnet-50 on GPUs

The Transformer is another widely-popular model that originated in the field of natural language processing (NLP). With more than *200 million* parameters, training the transformer (big) model heavily relies on data parallelism to achieve reasonable training times. We show that Tarantella distributed training also scales when using the Transformer for a translation task trained on the WMT14 English-German Translation dataset.

Figure 3 gives an insight of the time savings that Tarantella-based training can attain on a GPU machine such as the *HPC-DA* cluster, reaching a **34x speedup** for one epoch on 96 devices.

To find out more about training such models with Tarantella, take a look at our [tutorials](#).

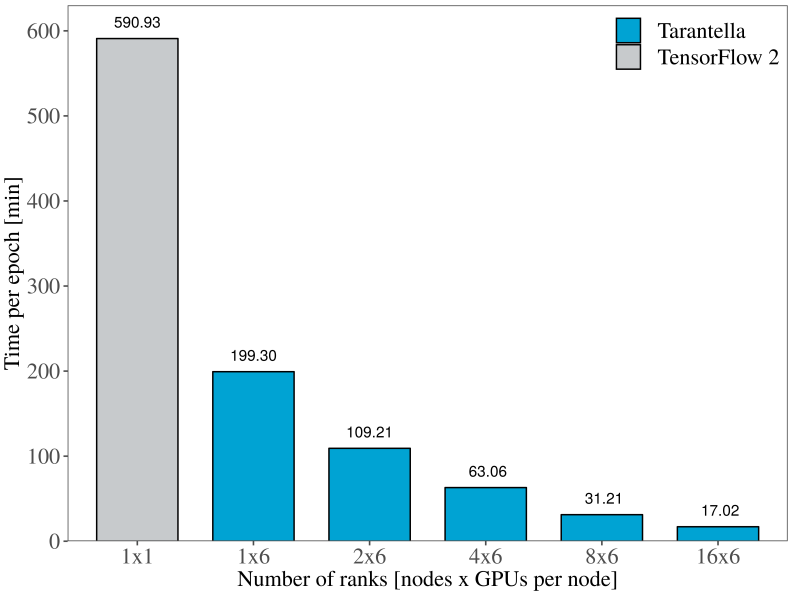


Fig. 3: Figure 3. Training the Transformer (big) on GPUs

DISTRIBUTED DATA PARALLEL TRAINING

The following section explains the parallelization strategy Tarantella uses to provide distributed training. A full understanding thereof is, however, not required to be able to use the software. Please note the *points to consider* to achieve best performance and reproducibility.

2.1 The general idea

In order to parallelize the training of DNNs, different, complementary strategies are available. The conceptually simplest and most efficient one is called *data parallelism*. This strategy is already in use when deploying batched optimizers, such as stochastic gradient descent (SGD) or ADAM. In this case, input samples are grouped together in so-called mini-batches and are processed in parallel.

2.2 Distribution of mini-batches

Tarantella extends this scheme by splitting each mini-batch into a number of micro-batches, which are then executed on different devices (e.g., GPUs). In order to do this, the DNN is replicated on each device, which then processes part of the data independently of the other devices. During the *backpropagation* pass, partial results need to be accumulated via a so-called *allreduce* collective operation.

2.3 Overlapping communication with computation

Tarantella implements this communication scheme using the *Global Address Space Programming Interface (GASPI)*. This allows in particular to overlap the communication needed to execute *allreduce* operations with the computation done in the *backpropagation* part of the DNN training. This is done by starting *allreduce* operations as soon as the required local incoming gradients are available, while continuing with *backpropagation* calculations at the same time. The final, accumulated gradients are only expected once the entire *backpropagation* is completed. This drastically mitigates the communication overhead introduced by the need to synchronize the different devices, and leads to higher scalability.

2.4 Tensor Fusion

The granularity at which Tarantella executes *allreduce* operations can be varied from one *allreduce* per layer (finest granularity) to one *allreduce* per iteration (coarsest granularity). Using coarser granularities, i.e., *fusing* gradient tensors, can lead to better bandwidth utilization, thus potentially increasing performance. *Tensor Fusion* is set up before the first iteration of training and incurs no additional communication overhead. Tarantella enables *Tensor Fusion* by default, but its granularity can be adjusted by the user, cf. [here](#).

2.5 Model initialization and loading

In order to guarantee that all devices have the same copy of the DNN when training is initiated, the model needs to be communicated from one device to all the others. This is done in Tarantella via the use of a so-called [broadcast](#) operation. This scheme applies both when the weights of a DNN are initialized randomly, or loaded from a checkpoint. As Tarantella provides this functionality automatically, the user does not have to take care of it.

DISTRIBUTED DATASETS

In order to process micro-batches independently on each device and to obtain the same results as in serial execution, the input data of each mini-batch has to be split and distributed among all devices.

Tarantella automatically takes care of this through the use of distributed datasets. The user simply provides Tarantella with a `tf.data.Dataset` that is batched with the mini-batch size. Tarantella will then automatically distribute the input data by splitting the mini-batch into individual micro-batches. Splitting is done at the level of samples (as opposed to e.g., files) to ensure *reproducibility* of serial results.

To guarantee reproducibility, it is also important that shuffling of samples is done in the same way on all devices. Tarantella does this using either the seed provided by the user, or a specific default seed. Please refer to the [Quick Start](#) for more details.

POINTS TO CONSIDER

4.1 Global versus local batch size

As explained above, when using data parallelism, there exists a *mini-batch size* (in the following also called global batch size or simply batch size) as well as a *micro-batch size* (also called local batch size). The former represents the number of samples that is averaged over in the loss function of the optimizer, and is equivalent to the (mini-)batch size used in non-distributed training. The latter is the number of samples that is processed locally by each of the devices per iteration.

Note: In Tarantella, the user always specifies the **global batch size**.

Using a strictly synchronous optimization scheme, and by carefully handling the data distribution, **Tarantella guarantees the reproducibility of DNN training results independently of the number of devices used**, as long as all hyperparameters (such as global batch size and learning rate) are kept constant.¹

However, to achieve best performance for certain DNN operators (*Conv2D*, *Dense*, etc.) it is often advisable to *keep the local batch size constant*, and scale the global batch size with the number of devices used. This, in turn, will force you to adjust other hyperparameters, such as the learning rate, in order to converge to a comparable test accuracy, as observed for instance in [Shallue].

In practice, the use of a learning rate schedule with initial *warm up* and a *linear learning rate scaling* [Goyal], as it is described [here](#), often suffices.

Tip: For best performance, scale the batch size with the number of devices used, and [adapt the learning rate schedule](#).

4.2 Batch normalization layers

The issue of global versus local batch size particularly affects the layers that calculate (and learn) statistics over entire batches. A well-known example of this type of layer is [batch normalization](#).

Caution: Tarantella always calculates batch statistics over **local batches**.

As a consequence, the training results for DNNs with batch-normalization layers **will not be identical when changing the number of devices, even if the global batch size stays the same**. At the moment, this can be circumvented by using normalization layers that do *not* average over entire batches, such as instance normalization [Ulyanov].

¹ This is strictly true, only when all randomness in TensorFlow is seeded or switched off, as explained in the [advanced topics](#)

Averaging over *local* batches instead of global batches should in practice have only minor influence on the quality of the final test accuracy. Note however, the extreme case of very small *local* batch sizes.

Caution: Avoid using `BatchNormalization` layers when the global batch size divided by the number of devices used is *smaller than 16*. A warning is issued when this occurs.

In such cases, the local batches that are used to collect statistics are too small to obtain meaningful results. This will likely reduce the benefits of batch normalization, cf. for instance [Yang] and [Uppal]. In this case, please consider increasing the global batch size, or reducing the number of devices used.

4.3 Managing individual devices

Although Tarantella’s user interface abstracts away most of the details of parallel programming, it is sometimes useful to be able to control Python code execution at device level. This can be achieved using the `GASPI` concept of a `rank`. Details on how to do this can be found in the *advanced topics*.

References

INSTALLATION

Tarantella needs to be built [from source](#). Since Tarantella is built on top of [TensorFlow](#), you will require a recent version of it. Additionally, you will need an installation of the open-source communication libraries [GaspiCxx](#) and [GPI-2](#), which Tarantella uses to implement distributed training.

Lastly, you will need [pybind11](#), which is required for Python and C++ inter-communication.

In the following we will look at the required steps in detail.

5.1 Installing dependencies

5.1.1 Compiler and build system

Tarantella can be built using a recent [gcc](#) compiler with support for C++17 (starting with `gcc 7.4.0`). You will also need the build tool [CMake](#) (from version 3.12).

5.1.2 Installing TensorFlow

First you will need to install TensorFlow. Supported versions start at `Tensorflow 2.4`, and they can be installed in a conda environment using `pip`, as recommended on the [TensorFlow website](#).

In order to do that, first install [conda](#) on your system. Then, create and activate an environment for Tarantella:

```
conda create -n tarantella
conda activate tarantella
```

Now, you can install the latest supported TensorFlow version with:

```
conda install python=3.9
pip install --upgrade tensorflow==2.9.*
```

Tarantella requires at least Python 3.7. Make sure the selected version also matches the [TensorFlow requirements](#).

5.1.3 Installing pybind11

The next dependency you will need to install is `pybind11`, which is available through pip and conda. We recommend installing `pybind11` via conda:

```
conda install pybind11 -c conda-forge
```

5.1.4 Installing GPI-2

Next, you will need to download, compile and install the GPI-2 library. GPI-2 is an API for high-performance, asynchronous communication for large scale applications, based on the [GASPI \(Global Address Space Programming Interface\)](#) standard.

The currently supported versions start with 1.5, and they need to be built with position independent flags (`-fPIC`). To download the required version, clone the [GPI-2 git repository](#) and checkout the latest tag:

```
git clone https://github.com/cc-hpc-itwm/GPI-2.git
cd GPI-2
git fetch --tags
git checkout -b v1.5.1 v1.5.1
```

Now, use `autotools` to configure and compile the code:

```
./autogen.sh
export GPI2_INSTALLATION_PATH=/your/gpi2/installation/path
CFLAGS="-fPIC" CPPFLAGS="-fPIC" ./configure --with-ethernet --prefix=${GPI2_INSTALLATION_
↪PATH}
make -j$(nproc)
```

where `${GPI2_INSTALLATION_PATH}` needs to be replaced with the path where you want to install GPI-2. Note the `--with-ethernet` option, which will use standard TCP sockets for communication. This is the correct option for laptops and workstations.

In case you want to use Infiniband, replace the above option with `--with-infiniband`. Now you are ready to install GPI-2 with:

```
make install
export PATH=${GPI2_INSTALLATION_PATH}/bin:$PATH
```

where the last two commands make the library visible to your system. If required, GPI-2 can be removed from the target directory by using `make uninstall`.

5.1.5 Installing GaspiCxx

`GaspiCxx` is a C++ abstraction layer built on top of the GPI-2 library, designed to provide easy-to-use point-to-point and collective communication primitives. Tarantella's communication layer is based on `GaspiCxx` and its `PyGPI` API for Python. Currently we support `GaspiCxx` version v1.2.0.

To install `GaspiCxx` and `PyGPI`, first download the latest release from the [git repository](#):

```
git clone https://github.com/cc-hpc-itwm/GaspiCxx.git
cd GaspiCxx
git fetch --tags
git checkout -b v1.2.0 v1.2.0
```

GaspiCxx requires an already installed version of GPI-2, which should be detected at configuration time (as long as `${GPI2_INSTALLATION_PATH}/bin` is added to the current `${PATH}` as shown [above](#)).

Compile and install the library as follows, making sure the previously created conda environment is activated:

```
conda activate tarantella

mkdir build && cd build
export GASPICXX_INSTALLATION_PATH=/your/gaspicxx/installation/path
cmake -DBUILD_PYTHON_BINDINGS=ON \
      -DBUILD_SHARED_LIBS=ON \
      -DCMAKE_INSTALL_PREFIX=${GASPICXX_INSTALLATION_PATH} ../
make -j$(nproc) install
```

where `${GASPICXX_INSTALLATION_PATH}` needs to be set to the path where you want to install the library.

5.2 SSH key-based authentication

In order to use Tarantella on a cluster, make sure you can ssh between nodes without password. For details, refer to the [FAQ section](#). In particular, to test Tarantella on your local machine, make sure you can ssh to `localhost` without password.

5.3 Building Tarantella from source

With all dependencies installed, we can now download, configure and compile Tarantella. To download the source code, simply clone the [GitHub repository](#):

```
git clone https://github.com/cc-hpc-itwm/tarantella.git
cd tarantella
git checkout tags/v0.9.0 -b v0.9.0
```

Next, we need to configure the build system using CMake. For a standard out-of-source build, we create a separate build folder and run `cmake` in it:

```
conda activate tarantella

cd tarantella
mkdir build && cd build
export TARANTELLA_INSTALLATION_PATH=/your/installation/path
cmake -DCMAKE_INSTALL_PREFIX=${TARANTELLA_INSTALLATION_PATH} \
      -DCMAKE_PREFIX_PATH=${GASPICXX_INSTALLATION_PATH} ../
```

This will configure your installation to use the previously installed GPI-2 and GaspiCxx libraries. To install Tarantella on a cluster equipped with Infiniband capabilities, make sure that GPI-2 is installed with Infiniband support as shown [here](#).

Now, we can compile and install Tarantella to `TARANTELLA_INSTALLATION_PATH`:

```
make -j$(nproc) install
export PATH=${TARANTELLA_INSTALLATION_PATH}/bin:${PATH}
```

5.4 [Optional] Building and running tests

In order to build Tarantella with tests, you will also need to install [Boost](#) (for C++ tests), and [pytest](#) (for Python tests). Additionally, the [PyYAML](#) and [NetworkX](#) libraries are required by some tests.

To install boost with the required *devel*-packages, under Ubuntu you can use

```
sudo apt install libboost-all-dev
```

while in Fedora you can use

```
sudo dnf install boost boost-devel
```

The other dependencies can be installed in the existing conda environment:

```
pip install -U pytest
pip install PyYAML==3.13
conda install networkx
```

After having installed these libraries, make sure to configure Tarantella with testing switched on:

```
cd tarantella
mkdir build && cd build
export LD_LIBRARY_PATH=`pwd`:${LD_LIBRARY_PATH}
export LD_LIBRARY_PATH=${GPI2_INSTALLATION_PATH}/lib64:${LD_LIBRARY_PATH}
export LD_LIBRARY_PATH=${GASPICXX_INSTALLATION_PATH}/lib:${LD_LIBRARY_PATH}

export PYTHONPATH=`pwd`:${PYTHONPATH}
export PYTHONPATH=${GASPICXX_INSTALLATION_PATH}/lib:${PYTHONPATH}

cmake -DENABLE_TESTING=ON ../
```

Now you can compile Tarantella and run its tests in the build directory:

```
make -j$(nproc)
ctest
```

5.5 [Optional] Building documentation

If you would like to build the documentation locally, run the following `cmake` command

```
cmake -DCMAKE_INSTALL_PREFIX=${TARANTELLA_INSTALLATION_PATH} -DBUILD_DOCS=ON ..
```

before compiling. This requires you to have [Sphinx](#) installed:

```
pip install -U sphinx
```

QUICK START

This section explains how to get started using Tarantella to distributedly train an existing TensorFlow model. First, we will examine what changes have to be made to your code, before executing it on the command line with `tarantella`. Finally, we will present the features Tarantella currently supports and what important points need to be taken into account when using the framework.

6.1 Code example: LeNet-5 on MNIST

After having *built and installed* Tarantella we are ready to add distributed training support to an existing TensorFlow model. We will first illustrate all the necessary steps, using the well-known example of **LeNet-5** on the **MNIST** dataset. Although this is not necessarily a good use case to take full advantage of Tarantella's capabilities, it will allow you to simply copy-paste the code snippets and try them out, even on your laptop.

Let's get started!

```
1 import tensorflow as tf
2 from tensorflow import keras
3
4 # Initialize Tarantella (before doing anything else)
5 import tarantella as tnt
6
7 # Skip function implementations for brevity
8 [...]
9
10 args = parse_args()
11
12 # Create Tarantella model from a `keras.Model`
13 model = tnt.Model(lenet5_model_generator())
14
15 # Compile Tarantella model (as with Keras)
16 model.compile(optimizer = keras.optimizers.SGD(learning_rate=args.learning_rate),
17               loss = keras.losses.SparseCategoricalCrossentropy(),
18               metrics = [keras.metrics.SparseCategoricalAccuracy()])
19
20 # Load MNIST dataset (as with Keras)
21 shuffle_seed = 42
22 (x_train, y_train), (x_val, y_val), (x_test, y_test) = \
23     mnist_as_np_arrays(args.train_size, args.val_size, args.test_size)
24
25 train_dataset = tf.data.Dataset.from_tensor_slices((x_train, y_train))
```

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```

26 train_dataset = train_dataset.shuffle(len(x_train), shuffle_seed)
27 train_dataset = train_dataset.batch(args.batch_size)
28 train_dataset = train_dataset.prefetch(tf.data.experimental.AUTOTUNE)
29
30 test_dataset = tf.data.Dataset.from_tensor_slices((x_test, y_test))
31 test_dataset = test_dataset.batch(args.batch_size)
32
33 # Train Tarantella model (as with Keras)
34 model.fit(train_dataset,
35           epochs = args.number_epochs,
36           verbose = 1)
37
38 # Evaluate Tarantella model (as with Keras)
39 model.evaluate(test_dataset, verbose = 1)

```

As you can see from the marked lines in the code snippet, you only need to add *two lines of code* to train LeNet-5 distributedly using Tarantella! Let us go through the code in some more detail, in order to understand what is going on.

First we need to import the Tarantella library:

```
import tarantella as tnt
```

Importing the Tarantella package will initialize the library and set up the communication infrastructure. Note that this should be done before executing any other code.

Next, we need to wrap the `keras.Model` object, generated by `lenet5_model_generator()`, into a `tnt.Model` object:

```
model = tnt.Model(lenet5_model_generator())
```

That's it!

All the necessary steps to distribute training and datasets will now be automatically handled by Tarantella. In particular, we still run `model.compile` on the new `model` to generate a compute graph, just as we would have done with a typical Keras model.

Next, we load the MNIST data for training and testing, and create `tf.data.Dataset` s from it. Note that we batch the dataset for training. This will guarantee that Tarantella is able to distribute the data later on in the correct way. Also note that the `batch_size` used here, is the same as for the original model, that is the *global* batch size. For details concerning local and global batch sizes have a look [here](#).

Now we are able to train our model using `model.fit`, in the same familiar way used by the standard Keras interface. Note, however, that Tarantella is taking care of the proper distribution of the `train_dataset` in the background. All the possibilities of how to feed datasets to Tarantella are explained in more detail below. Lastly, we can evaluate the final accuracy of our model on the `test_dataset` using `model.evaluate`.

To test and run Tarantella in the next section, you can find a full version of the above example [here](#).

6.2 Executing your model with tarantella

Next, let's execute our model distributedly using `tarantella` on the command line. Make sure to add the path to your installed *GaspiCxx* library to `LD_LIBRARY_PATH`:

```
export LD_LIBRARY_PATH=${GASPICXX_INSTALLATION_PATH}/lib:${LD_LIBRARY_PATH}
```

The simplest way to run the model is by passing its Python script to `tarantella`:

```
tarantella -- model.py
```

This will execute our model distributedly on a single node, using all the available GPUs. In case no GPUs can be found, `tarantella` will be executed in serial mode on the CPU, and a `WARNING` message will be issued. In case there are available GPUs, but we want to execute `tarantella` on CPUs nonetheless, we can add the `--no-gpu` option.

```
tarantella --no-gpu -- model.py
```

We can also set command line parameters for the python script `model.py`, which have to succeed the name of the script:

```
tarantella --no-gpu -- model.py --batch_size=64 --learning_rate=0.01
```

On a single node, we can also explicitly specify the number of TensorFlow instances we want to use. This is done with the `-n` option:

```
tarantella -n 4 -- model.py --batch_size=64
```

Here, `tarantella` will try to execute distributedly on 4 GPUs. If there are not enough GPUs available, `tarantella` will print a `WARNING` and run 4 instances of TensorFlow on the CPU instead. If there are no GPUs installed or the `--no-gpu` option is used, `tarantella` will not print a `WARNING`.

Next, let's run `tarantella` on multiple nodes. In order to do this, we need to provide `tarantella` with a hostfile that contains the hostnames of the nodes that we want to use:

```
$ cat hostfile
name_of_node_1
name_of_node_2
```

With this hostfile we can run `tarantella` on multiple nodes:

```
tarantella --hostfile hostfile -- model.py
```

In this case, `tarantella` uses *all* GPUs it can find. If no GPUs are available, `tarantella` will start *one* TensorFlow instance per node on the CPUs, and will issue a `WARNING` message. Again, this can be disabled by explicitly using the `--no-gpu` option.

As before, you can specify the number of GPUs/CPUs used per node explicitly with the option `--n-per-node <number>`:

```
tarantella --hostfile hostfile --n-per-node 4 --no-gpu -- model.py --batch_size=64
```

In this example, `tarantella` would execute 4 instances of TensorFlow on the CPUs of each node specified in hostfile.

Caution: tarantella requires all the names in the `hostfile` be **unique**, and all nodes be **homogeneous** (same number and type of CPUs and GPUs).

In addition, tarantella can be run with different levels of logging output. The log-levels that are available are INFO, WARNING, DEBUG and ERROR, and can be set with `--log-level`:

```
tarantella --hostfile hostfile --log-level INFO -- model.py
```

By default, tarantella will log on the *master rank* only. This can be changed by using the `--log-on-all-devices` option, which will print log messages for each *rank* individually.

Similarly, by default tarantella will print outputs from functions like `fit`, `evaluate` and `predict`, as well as callbacks only on the master rank. Sometimes, it might be useful to print outputs from all devices (e.g., for debugging), which can be switched on with the `--output-on-all-devices` option.

tarantella relies on GPI-2's tools for starting processes on multiple nodes (i.e., `gaspi_run`). To properly configure an execution, it will take care of exporting relevant environment variables (such as `PYTHONPATH`) for each process, and of generating an execution script from the user inputs. Details of this process can be monitored using the `--dry-run` option.

To add your own environment variables, add `-x ENV_VAR_NAME=VALUE` to your tarantella command. This option will ensure the environment variable `ENV_VAR_NAME` is exported on all ranks before executing the code. An example is shown below:

```
tarantella --hostfile hostfile -x DATASET=/scratch/data TF_CPP_MIN_LOG_LEVEL=1 -- model.  
↪py
```

Both `DATASET` and `TF_CPP_MIN_LOG_LEVEL` will be exported as environment variables before executing `model.py`, in the same order they were specified to the command line.

Additionally, you can overwrite the *Tensor Fusion* threshold tarantella uses with `--fusion-threshold FUSION_THRESHOLD_KB` (cf. [here](#) and [here](#)), and set any number of environment variables, most notably `TNT_TENSORBOARD_ON_ALL_DEVICES`, as explained [here](#).

To terminate a running tarantella instance, execute another tarantella command that specifies the `--cleanup` option in addition to the name of the program you want to interrupt.

```
tarantella --hostfile hostfile --cleanup -- model.py
```

The above command will stop the `model.py` execution on all the nodes provided in `hostfile`. You can also enable the `--force` flag to immediately terminate unresponsive processes.

Note: Any running tarantella execution can be terminated by using `Ctrl+c`, regardless of whether it was started on a single node or on multiple hosts.

6.3 Save and load Tarantella models

Storing and loading your trained `tnt.Model` is very simple.

Tarantella supports all the different ways in which you can load and store a `keras.Model` (for a guide look for instance [here](#)). In particular, you can:

- save the whole model (including the architecture, the weights and the state of the optimizer)
- save the model's architecture/configuration only
- save the model's weights only

6.3.1 Whole-model saving and loading

Saving the entire model including the architecture, weights and optimizer can be done via

```
model = ... # get `tnt.Model`
model.save('path/to/location')
```

Alternatively, you could use `tnt.models.save_model('path/to/location')`, which works on both `keras.Model`s and `tnt.Model`s.

You can then load your model back using

```
import tarantella as tnt
model = tnt.models.load_model('path/to/location')
```

which will return an instance of `tnt.Model`.

If the saved model was previously compiled, `load_model` will also return a compiled model. Alternatively, you can deliberately load the model in an uncompiled state by passing the `compile = False` flag to `load_model`.

6.3.2 Architecture saving and loading

If you only want to save the configuration (that is the architecture) of your model (in memory), you can use one of the following functions:

- `tnt.Model.get_config`
- `tnt.Model.to_json`
- `tnt.Model.to_yaml` [supported up to TF 2.6]

The architecture without its original weights and optimizer can then be restored using:

- `tnt.models.model_from_config` / `tnt.Model.from_config`
- `tnt.models.model_from_json`
- `tnt.models.model_from_yaml` [supported up to TF 2.6]

respectively. Here is an example:

```
import tarantella as tnt
model = ... # get `tnt.Model`
config = model.get_config()
new_model = tnt.models.model_from_config(config)
```

The same can be achieved through cloning:

```
import tarantella as tnt
model = ... # get `tnt.Model`
new_model = tnt.models.clone_model(model)
```

6.3.3 Weights saving and loading

Storing and loading the weights of a model to/from memory can be done using the functions `tnt.Model.get_weights` and `tnt.Model.set_weights`, respectively. Saving and loading weights to/from disk is done using the functions `tnt.Model.save_weights` and `tnt.Model.load_weights`, respectively.

Here is an example how this can be used to restore a model:

```
import tarantella as tnt
model = ... # get `tnt.Model`
config = model.get_config()
weights = model.get_weights()

# initialize a new model with original model's weights
new_model = tnt.models.model_from_config(config)
new_model.set_weights(weights)
```

6.3.4 Checkpointing via callbacks

Apart from saving and loading models manually, Tarantella also supports checkpointing via Keras' `ModelCheckpoint` callback, as it is described for instance [here](#).

```
import tensorflow as tf
import tarantella as tnt

model = ... # get `tnt.Model`

checkpoint_path = 'path/to/checkpoint/location'
model_checkpoint_callback = tf.keras.callbacks.ModelCheckpoint(
    filepath=checkpoint_path, monitor='val_acc', verbose=1, save_best_only=False,
    save_weights_only=False, mode='auto', save_freq='epoch', options=None)

model.fit(train_dataset,
          validation_data = val_dataset,
          epochs = 2,
          callbacks = [model_checkpoint_callback])
```

Note: All saving to the filesystem (including `tnt.Model.save` and `tnt.Model.save_weights`) by Tarantella will only be done on the master rank.

This is the default and will yield correct behavior when you are using a distributed filesystem. If you wish to explicitly save on all devices you can pass `tnt_save_all_devices = True` to `tnt.Model.save`, `tnt.Model.save_weights` and `tnt.models.save_model`.

6.4 Using distributed datasets

This section explains how to use Tarantella's distributed datasets.

The recommended way in which to provide your dataset to Tarantella is by passing a *batched* `tf.data.Dataset` to `tnt.Model.fit`. In order to do this, create a `Dataset` and apply the `batch` transformation using the (global) batch size to it. However, do not provide a value to `batch_size` in `tnt.Model.fit`, which would lead to double batching, and thus modified shapes for the input data.

Tarantella can distribute any `tf.data.Dataset`, regardless of the number and type of transformations that have been applied to it.

Note: When using the `dataset.shuffle` transformation without a `seed`, Tarantella will use a fixed default `seed`.

This guarantees that the input data is shuffled the same way on all devices, when no `seed` is given, which is necessary for consistency. However, when a random `seed` is provided by the user, Tarantella will use that one instead.

Tarantella also supports `batched` and `unbatched Dataset`s in `tnt.Model.fit` when setting the `tnt_micro_batch_size` argument. This can be useful to maximize performance in multi-node executions, as explained [here](#). Keep in mind however, that Tarantella still expects the `Dataset` to be `batched` with the global batch size, and that the micro-batch size has to be consistent with the global batch size.¹ This is why it is recommended to use an `unbatched Dataset` when setting `tnt_micro_batch_size` explicitly.

Tarantella does not support any other way to feed data to `fit` at the moment. In particular, Numpy arrays, TensorFlow tensors and generators are not supported.

Tarantella's automatic data distribution can be switched off by passing `tnt_distribute_dataset = False` in `tnt.Model.fit`, in which case Tarantella will issue an `INFO` message. If a validation dataset is passed to `tnt.Model.fit`, it should also be `batched` with the global batch size. You can similarly switch off its automatic micro-batching mechanism by setting `tnt_distribute_validation_dataset = False`.

6.5 Callbacks

Tarantella fully supports all pre-defined [Keras callbacks](#):

- `tf.keras.callbacks.CSVLogger`
- `tf.keras.callbacks.EarlyStopping`
- `tf.keras.callbacks.History`
- `tf.keras.callbacks.LearningRateScheduler`
- `tf.keras.callbacks.ModelCheckpoint`
- `tf.keras.callbacks.ProgbarLogger`
- `tf.keras.callbacks.ReduceLROnPlateau`
- `tf.keras.callbacks.RemoteMonitor`
- `tf.keras.callbacks.TensorBoard`
- `tf.keras.callbacks.TerminateOnNaN`

¹ That is, the global batch size must equal the micro batch size times the number of devices used.

All of these callbacks are implemented in such a way that the device-local, micro-batch information is accumulated over all devices. This leads to the same callback behavior as in a serial execution (using the full batch). That is, users do not need to make any modifications to their code when using Keras callbacks with Tarantella.

However, when using the [TensorBoard](#) callback, by default, Tarantella will only collect device-local information *on one device*. If you want to collect the local information on all devices use the environment variable `TNT_TENSORBOARD_ON_ALL_DEVICES`:

```
TNT_TENSORBOARD_ON_ALL_DEVICES=true tarantella -- model.py
```

Note: The explicit addition of `BaseLogger` callbacks is not supported in Tarantella.

6.5.1 Custom Callbacks

Any custom Keras callback can be used in a distributed fashion with Tarantella. To this end, define your own custom Keras callback as explained in the [Writing Custom Callbacks](#) guide.

Next, all you need to do is wrap the `keras_callback` into a `tnt.keras.callbacks.Callback` object and simply add it to the list of callbacks provided in the model training or inference methods:

```
class CustomCallback(keras.callbacks.Callback):
    def on_train_begin(self, logs = None):
        keys = list(logs.keys())
        print("Starting training; got log keys: {}".format(keys))
        ...

keras_callback = CustomCallback()

tnt_callback = tnt.keras.callbacks.Callback(keras_callback,
                                           aggregate_logs = True,
                                           run_on_all_ranks = True)

model.fit(train_dataset,
          epochs = 2,
          callbacks = [tnt_callback])
```

The execution of a `tnt.keras.callbacks.Callback` can be configured through the following parameters:

- `run_on_all_ranks` - defines whether the callback will be run on all devices or just the master rank (defaults to `True`). While most callbacks need to collect data from all the used devices, there are cases when this behavior is not desirable (e.g., a profiling callback might only need to measure timings on the master rank).
- `aggregate_logs` - specifies whether the logs need to be aggregated from all devices (defaults to `True`). For instance, loss values have to be aggregated across all micro-batches to provide the relevant batch-level information. Conversely, logs counting the number of iterations do not require aggregation, as the iteration counter is identical on all participating devices.

The `keras.callbacks.Callback` object can also be directly passed (without the wrapper) to the list of callbacks provided to the `model.fit` function. In this case, the `tnt.keras.callbacks.Callback` object is automatically created with the default parameter values.

6.5.2 Lambda Callbacks

A `LambdaCallback` allows users to create simple custom callbacks using a lambda function. To use this feature in Tarantella, create a Keras lambda callback as explained in the [TensorFlow guide](#).

Then, wrap the callback object into a `tnt.keras.callbacks.Callback` as shown in the previous section.

```
# Print the batch number at the beginning of every batch.
batch_print_callback = LambdaCallback(on_batch_begin = lambda batch, logs: print(batch))

# Run the callback on the master rank only
tnt_print_callback = tnt.keras.callbacks.Callback(batch_print_callback,
                                                  aggregate_logs = False,
                                                  run_on_all_ranks = False)
```

6.5.3 Rank-Local Callbacks

There are cases when user-defined callbacks do not require distributed processing, such as callbacks that print information or measure runtimes. To configure a callback to run only on the *master rank*, wrap it as a `tnt.keras.callbacks.Callback` and set the constructor parameters as follows:

```
class MyCustomCallback(keras.callbacks.Callback):
    ...

keras_callback = MyCustomCallback()
tnt_callback = tnt.keras.callbacks.Callback(keras_callback,
                                           aggregate_logs = False,
                                           run_on_all_ranks = False)
```

Note that callbacks running on a single rank will only have access to local data corresponding to that rank. For instance, even though the models are identical on all ranks, a logging callback that displays metrics will only be aware of locally collected metrics, that is, metrics generated based on the micro-batches that the rank has processed.

6.6 Important points

There is a number of points you should be aware of when using Tarantella.

Note: Tarantella does not support custom training loops.

Instead of using custom training loops, please use `Model.fit(...)`.

Note: Tarantella supports all [TensorFlow optimizers](#) with the exception of `tf.keras.optimizers.Ftrl`.

Since the `Ftrl` optimizer does not use batches, it is not supported in Tarantella.

TUTORIALS

This section delves into more advanced usage of Tarantella with the help of state-of-the-art models for two widely-used applications in Deep Learning:

- Image classification: ResNet-50
- Machine translation: Transformer

The image classification model architectures are imported through the `tf.keras.applications` module, available in recent TensorFlow releases.

The Transformer model presented in this tutorial is adapted from the [TensorFlow Model Garden](#). While the model implementations and hyperparameters are unchanged to preserve compatibility with the TensorFlow official models, we provide simplified training schemes that allow for a seamless transition from basic serial training to distributed data parallelism using Tarantella.

7.1 Prerequisites

The model can be downloaded from the [Tnt Models repository](#).

```
cd /your/models/path
git clone https://github.com/cc-hpc-itwm/tarantella_models

cd tarantella_models/src
export TNT_MODELS_PATH=`pwd`
```

This tutorial assumes the following dependencies are installed:

- TensorFlow 2.9.1
- Tarantella 0.9.0

For a step-by-step installation, follow the [Installation](#) guide.

7.2 ResNet-50

Deep Residual Networks (ResNets) represented a breakthrough in the field of computer vision, enabling deeper and more complex deep convolutional networks. Introduced in [He], ResNet-50 has become a standard model for image classification tasks, and has been shown to scale to very large number of nodes in data parallel training [Goyal].

7.2.1 Run Resnet-50 with Tarantella

Before running the model, we need to add it to the existing PYTHONPATH.

```
export PYTHONPATH=${TNT_MODELS_PATH}:${PYTHONPATH}
```

Furthermore, the ImageNet dataset needs to be installed and available on all the nodes that we want to use for training. TensorFlow provides convenience scripts to download datasets, in their `datasets` package that is installed as a dependency for the TensorFlow Model Garden. Install ImageNet to your local machine as described [here](#).

```
export TNT_DATASETS_PATH=/path/to/downloaded/datasets

python -m tensorflow_datasets.scripts.download_and_prepare \
--datasets=imagenet2012 --data_dir=${TNT_DATASETS_PATH}
```

Let's assume we have access to two nodes (saved in `hostfile`) equipped with 4 GPUs each. We can now simply run the ResNet-50 as follows:

```
tarantella --hostfile ./hostfile --devices-per-node 4 \
-- ${TNT_MODELS_PATH}/models/image_classification/train_imagenet_main.py --data_dir=$
↪ ${TNT_DATASETS_PATH} \
--model_
↪ arch=resnet50 \
--strategy=data_
↪ \
--batch_
↪ size=512 \
--train_
↪ epochs=90 \
--epochs_
↪ between_evals=10
```

The above command will train a ResNet-50 models on the 8 devices available in parallel for 90 epochs, as suggested in [Goyal] to achieve convergence. The `--val_freq` parameter specifies the frequency of evaluations of the *validation dataset* performed in between training epochs.

Note the `--batch_size` parameter, which specifies the global batch size used in training.

7.2.2 Implementation overview

We will now look closer into the implementation of the ResNet-50 training scheme. The main training steps reside in the `models/image_classification/train_imagenet_main.py` file.

The most important step in enabling data parallelism with Tarantella is to wrap the Keras model into a Tarantella model that uses data parallelism for speeding up training.

This is summarized below for the *ResNet50* model:

```
model = tf.keras.applications.resnet50.ResNet50(include_top=True, weights=None,
↪ classes=1000,
                                     input_shape=(224, 224, 3), input_
↪ tensor=None,
                                     pooling=None, classifier_activation=
↪ 'softmax')
...
if args.distribute == ParallelMethods.TNT:
    model = tnt.Model(model,
                      parallel_strategy = tnt.ParallelStrategy.DATA)
```

Next, the training procedure can simply be written down as it would be for a standard, TensorFlow-only model. No further changes are required to train the model in a distributed manner.

In particular, the ImageNet dataset is loaded and preprocessed as follows:

```
train_input_dataset = load_dataset(dataset_type='train',
                                   data_dir=args.data_dir, num_samples = args.train_num_
↪ samples,
                                   batch_size=args.batch_size, dtype=tf.float32,
                                   drop_remainder=args.drop_remainder,
                                   shuffle_seed=args.shuffle_seed)
```

The `load_dataset` function reads the input files in `data_dir`, loads the training samples, and processes them into TensorFlow datasets.

The user only needs to pass the global `batch_size` value, and the Tarantella framework will ensure that the dataset is properly distributed among devices, such that:

- each device will process an independent set of samples
- each device will group the samples into micro batches, where the micro-batch size will be roughly equal to `batch_size / num_devices`. If the batch size is not a multiple of the number of ranks, the remainder samples will be equally distributed among the participating ranks, such that some ranks will use a micro-batch of `(batch_size / num_devices) + 1`.
- each device will apply the same set of transformations to its input samples as specified in the `load_dataset` function.

The advantage of using the *automatic dataset distribution* mechanism of Tarantella is that users can reason about their I/O pipeline without taking care of the details about how to distribute it.

Before starting the training, the model is compiled using a standard Keras optimizer and loss.

```
model.compile('optimizer' : tf.keras.optimizers.SGD(learning_rate=lr_schedule,
↪ momentum=0.9),
              'loss' : tf.keras.losses.SparseCategoricalCrossentropy(),
              'metrics' : [tf.keras.metrics.SparseCategoricalAccuracy()])
```

We provide flags to enable the most commonly used Keras callbacks, such as the TensorBoard profiler, which can simply be passed to the fit function of the Tarantella model.

```
callbacks.append(tf.keras.callbacks.TensorBoard(log_dir = flags_obj.model_dir,
                                                profile_batch = 2))
```

If model checkpointing is required, it can be enabled through the ModelCheckpoint callback as usual (cf. [checkpointing models with Tarantella](#)).

```
callbacks.append(tf.keras.callbacks.ModelCheckpoint(ckpt_full_path, save_weights_
↪only=True))
```

There is no need for any further changes to proceed with distributed training:

```
history = model.fit(train_dataset,
                    validation_data = val_dataset,
                    validation_freq=args.val_freq,
                    epochs=args.train_epochs,
                    callbacks=callbacks,
                    verbose=args.verbose)
```

7.2.3 Advanced topics

Scaling the batch size

Increasing the batch size provides a simple means to achieve significant training time speed-ups, as it leads to perfect scaling with respect to the steps required to achieve the target accuracy (up to some dataset- and model- dependent critical size, after which further increasing the batch size only leads to diminishing returns) [Shallue].

This observation, together with the fact that small local batch sizes decrease the efficiency of DNN operators, represent the basis for a standard technique in data parallelism: *using a fixed micro batch size and scaling the global batch size with the number of devices*.

Tarantella provides multiple mechanisms to set the batch size, as presented in the [Quick Start guide](#).

In the case of ResNet-50, we specify the global batch size as a command line parameter, and let the framework divide the dataset into microbatches.

Scaling the learning rate

To be able to reach the same target accuracy when scaling the global batch size up, other hyperparameters need to be carefully tuned [Shallue]. In particular, adjusting the learning rate is essential for achieving convergence at large batch sizes. [Goyal] proposes to *scale the learning rate up linearly with the batch size* (and thus with the number of devices).

The scaled-up learning rate is set up at the beginning of training, after which the learning rate evolves over the training steps based on a so-called *learning rate schedule*.

In our ResNet-50 example, we use a `ExpDecayWithWarmupSchedule`.

Another type of schedule that we have implemented is the `PiecewiseConstantDecayWithWarmup` schedule, which is similar to the schedule introduced by [Goyal].

In both schedules, when training starts, the learning rate is initialized to a large value that allows to explore more of the search space. The learning rate will then decay the closer the algorithm gets to convergence.

The initial learning rate in the `ExpDecayWithWarmupSchedule` is scaled linearly with the number of devices used as follows:

```
initial_learning_rate = base_learning_rate * num_ranks
```

Learning rate warm-up

Whereas scaling up the learning rate with the batch size is necessary, a large learning rate might degrade the stability of the optimization algorithm, especially in early training. A technique to mitigate this limitation is to *warm-up* the learning rate during the first epochs, particularly when using large batches [Goyal].

In our ResNet-50 example, the *ExpDecayWithWarmupSchedule* schedule starts with a small value for the learning rate, which then increases at every step (i.e., iteration), for a number of initial *warmup_steps*.

The *warmup_steps* value defaults to the number of iterations of the first five epochs, matching the schedule proposed by [Goyal]. After the *warmup_steps* are done, the learning rate value should reach the *scaled initial learning rate* introduced above.

```
def warmup():
    # Learning rate increases linearly per step.
    multiplier = self.warmup_rate * (step / self.warmup_steps)
    return tf.multiply(self.initial_learning_rate, multiplier)
```

7.3 Transformers

The Transformer is a Deep Neural Network widely used in the field of natural language processing (NLP), in particular for tasks such as machine translation. It was first proposed by [Vaswani].

7.3.1 Prerequisites

In the following we will assume that TensorFlow was installed in a conda environment called *tarantella*.

The Transformer model architecture can be obtained from the TensorFlow official Model Garden:

```
conda activate tarantella
pip install tf-models-official==2.9.1
```

7.3.2 Run the Transformer with Tarantella

The Transformer training scheme can be found [here](#), and has to be added to the existing PYTHONPATH:

```
export PYTHONPATH=${TNT_MODELS_PATH}/models/transformer:${PYTHONPATH}
```

We will follow the training procedure presented in [Vaswani], where the authors show results for training the *big* variant of the Transformer model on a machine translation dataset called WMT14.

To install the dataset, we will use the Tensorflow datasets package, which should have been already installed in your conda environment as a dependency for the TensorFlow Model Garden, and download the English-German dataset to match the results by [Vaswani]. Detailed instructions on how to obtain the dataset are provided in the TensorFlow documentation.

Now we can start training. Once again, let's assume we have access to two nodes (specified in *hostfile*) equipped with 4 GPUs each.

```
export WMT14_PATH=/path/to/the/installed/dataset

tarantella --hostfile ./hostfile --devices-per-node 4 \
-- ${TNT_MODELS_PATH}/models/transformer/transformer_tnt.py \
    --data_dir=${WMT14_PATH} \
    --vocab_file=${WMT14_PATH}/vocab.ende.32768 \
    --bleu_ref=${WMT14_PATH}/newstest2014.de \
    --bleu_source=${WMT14_PATH}/newstest2014.en \
    --param_set=big \
    --train_epochs=30 \
    --epochs_between_evals=30 \
    --batch_size=32736
```

The above command will select the big model implementation and train it on the 8 specified devices in a distributed fashion. To reach the target accuracy, [Vaswani] specifies that the model needs to be trained for 30 epochs.

The Transformer requires access to a vocabulary file, which contains all the tokens derived from the dataset. This is provided as the `vocab_file` parameter and is part of the pre-processed dataset.

After training, one round of evaluation is conducted using the `newstest2014` dataset to translate English sentences into German. The frequency of evaluation rounds can be changed by updating the `epochs_between_evals` parameter.

7.3.3 Implementation overview

The Transformer model itself is implemented and imported from the [TensorFlow Model Garden](#). The training procedure and dataset loading and pre-processing do not require extensive changes to work with Tarantella. However, we provide a simplified version to highlight the usage of Tarantella with Keras training loops.

Thus, the Keras transformer model is created in `TransformerTntTask` class. Two different versions of the model are used, one for training (wrapped into a Tarantella model), and one for inference (serial Keras model).

```
self.train_model = create_model(internal_model, self.params, is_train = True)
# Enable distributed training
self.train_model = tnt.Model(self.train_model)

# The inference model is wrapped as a different Keras model that does not use labels
self.predict_model = create_model(internal_model, self.params, is_train = False)
```

To illustrate alternatives in the use of Tarantella, we distribute the data manually here, `data_pipeline.py` file, as explained in the [manually-distributed datasets](#) section. Alternatively, automatic dataset distribution could be used, as explained in the [Quick Start](#).

To be able to manually split the dataset across ranks, we need access to **rank IDs** and the **total number of ranks**, which are then passed to the `IO` pipeline.

The [Advanced Topics](#) section explains the API Tarantella exposes to access ranks.

```
train_ds = data_pipeline.train_input_fn(self.params,
                                       shuffle_seed = 42,
                                       num_ranks = tnt.get_size(),
                                       rank = tnt.get_rank())
```

Here, the `data_pipeline.train_input_fn` reads in the dataset and applies a series of transformations to convert it into a batched set of sentences.

Next, the user can also create callbacks, which can then be simply passed on to the training function.

```
callbacks.append(tf.keras.callbacks.TensorBoard(log_dir=self.flags_obj.model_dir))
```

Finally, we can call `model.fit` to start distributed training on all devices:

```
history = model.fit(train_ds,
                    tnt_distribute_dataset = False,
                    epochs=self.params["train_epochs"],
                    callbacks=callbacks,
                    verbose=1)
```

In the following sections we will show how we modify the `fit` loop to allow for a customized evaluation of the trained model.

7.3.4 Important points

Customized behavior based on rank

Although all ranks participating in data parallel training use identical replicas of the same model and make progress in sync, there are cases when certain tasks should be executed on a specific rank (or group or ranks). To this end, Tarantella provides a number of functions to identify the rank ID and allow users to add customized behavior based on rank, as described in this [section](#).

In the case of the Transformer model, we want to use the rank information to perform several tasks:

- print logging messages

```
if tnt.is_master_rank():
    logging.info("Start train")
```

- distribute datasets manually among participating devices
- execute other models, such as a modified, serial version of the Tarantella model for *inference*
- enable certain callbacks only on one rank (e.g., profiling callbacks)

```
if self.flags_obj.enable_time_history:
    time_callback = keras_utils.TimeHistory(self.params["batch_size"],
                                            self.params["num_sentences"],
                                            logdir = None)
    tnt_time_callback = tnt.keras.callbacks.Callback(time_callback,
                                                    aggregate_logs = False,
                                                    run_on_all_ranks = False)
    callbacks.append(tnt_time_callback)
```

Such callbacks only collect local data corresponding to the specific rank where they are executed. In this example, the *TimeHistory* callback will measure timings only on the `master_rank`. While iteration and epoch runtimes should be the same on all ranks (as all ranks train in sync), other metrics such as accuracy will only be computed based on the local data available to the rank.

A callback that should be executed on a single rank has to be wrapped within a `tnt.keras.callbacks.Callback`, to explicitly disable distributed execution (as described in the [callbacks guide](#)).

Using manually-distributed datasets

Typically, it is the task of the framework to automatically handle batched datasets, such that each rank only processes its share of the data, as explained in the [Quick Start guide](#).

However, there are complex scenarios when the user might prefer to manually build the dataset slices corresponding to each rank. Tarantella allows the user to disable the automatic distribution mechanism by passing `tnt_distribute_dataset = False` to the `model.fit` function.

This is how it is done in the case of the Transformer:

```
history = self.train_model.fit(train_ds,
                               callbacks = callbacks,
                               tnt_distribute_dataset = False,
                               initial_epoch = epoch,
                               epochs = epoch + min(self.params["epochs_between_evals"],
                                                    self.params["train_epochs"]-epoch),
                               verbose = 2)
```

Also note the use of `initial_epoch` and `epochs`. This combination of parameters is necessary to allow evaluation rounds in between training epochs, when a validation dataset cannot be simply passed to `model.fit`. In particular, our transformer implementation features a different model for inference, as described [below](#).

Now that automatic distribution is disabled, let us take a look at how to split the dataset manually among devices. The input data processing is implemented in [data_pipeline.py](#).

In the case of the Transformer model, the global `batch_size` stands for the total number of input tokens processed in a single iteration. However, as the training is performed in (fixed-sized) sentences, our global `batch_size` used for training will be in fact the number of sentences comprised in such a batch.

Furthermore, we need to divide the number of sentences across ranks, such that each rank can work on a separated shard of `micro_batch_size` sentences. Finally, the dataset itself needs to be batched using the `micro_batch_size` and each device instructed to select its own shard:

```
number_batch_sentences = batch_size // max_length

micro_batch_size = number_batch_sentences // num_ranks

# Batch the sentences and select only the shard (subset)
# corresponding to the current rank
dataset = dataset.padded_batch(micro_batch_size,
                               ([max_length], [max_length]),
                               drop_remainder=True)
dataset = dataset.shard(num_ranks, rank)
```

Mixing Keras and Tarantella models

An essential aspect of the Transformer model is that it operates on slightly different model versions during training and inference. While in training the model works on encoded tokens, inference requires translation to and from plain text. Thus, the model needs to use modified input and output layers for each of these tasks.

To illustrate the way a Tarantella model can work alongside a typical Keras model, we only execute the training phase on the Transformer within a (distributed) Tarantella model.

Take a look at the [model creation function](#). It builds two different Keras models depending on whether training is enabled or not, both of them based on the same *internal model* (i.e., using the same learned weights).

Now, when initializing our Transformer task, we only wrap one of the models as a `tnt.Model`:

```
# Transformer model used both as Tarantella model (in training) and as a serial
# model for inference
internal_model = transformer.Transformer(self.params, name="transformer_v2")

# The train model includes an additional logits layer and a customized loss
self.train_model = create_model(internal_model, self.params, is_train = True)
# Enable distributed training
self.train_model = tnt.Model(self.train_model)

# The inference model is wrapped as a different Keras model that does not use labels
self.predict_model = create_model(internal_model, self.params, is_train = False)
```

Training can now proceed as usual, by only calling the `fit` method on our `train_model`. We can however design our training loop to stop every `epochs_between_evals` epochs, evaluate the training accuracy using the serial `predict_model`, and then continue from where it left off.

```
for epoch in range(0, self.params["train_epochs"], self.params["epochs_between_evals"]):
    # as our dataset is distributed manually, disable the automatic Tarantella distribution
    history = self.train_model.fit(train_ds,
                                   callbacks = callbacks,
                                   tnt_distribute_dataset = False,
                                   initial_epoch = epoch,
                                   epochs = epoch + min(self.params["epochs_between_evals",
→"],
                                                         self.params["train_epochs"]-epoch),
                                   verbose = 2)

    if tnt.is_master_rank():
        eval_stats = self.eval()
```

The `self.eval()` method performs the translation on the test dataset using the standard Keras `predict_model`.

```
def eval(self):
    ...
    uncased_score, cased_score = transformer_main.evaluate_and_log_bleu(
        self.predict_model,
        self.params,
        self.flags_obj.bleu_source,
        self.flags_obj.bleu_ref,
        self.flags_obj.vocab_file)
```

A validation dataset can be provided in the form of a pair of input files specified at the command line as `bleu_source` and `bleu_ref`. If the validation dataset exists, the evaluation method will compute and log the corresponding BLEU scores (both case-sensitive and case-insensitive) serially.

ADVANCED TOPICS

This guide covers a number of advanced topics, such as performance, reproducibility and user customization.

8.1 GASPI ranks

To distribute the DNN training, Tarantella starts multiple processes on different devices. These processes will be assigned different IDs by the GPI-2 communication library, in order to organize communication and synchronization between the different devices. These IDs are called *ranks*. Usually, Tarantella abstracts away the concept of *ranks*, in such a way that Tarantella's user interface is essentially the same as Keras' user interface.

However, sometimes it is useful, to execute a specific part of code only on one or a subgroup of all ranks. In particular, one sometimes wants to execute a code block on the device that started `tarantella`, the so-called *master rank*.

To access ranks, Tarantella provides the following functions

- `tnt.get_rank()`
- `tnt.get_size()`
- `tnt.get_master_rank()`
- `tnt.is_master_rank()`

`tnt.get_rank()` returns the ID of the local rank. `tnt.get_size()` returns the total number of ranks. `tnt.get_master_rank()` and `tnt.is_master_rank()` return the ID of the master rank and a boolean for whether the local rank is the master rank or not, respectively.

Here is a simple example, where using the master rank can be useful to print notifications only once to `stdout`:

```
if tnt.is_master_rank():  
    print("Printing from the master rank")
```

More usage examples can be found in the *Tutorials* section.

8.2 Using local batch sizes

As it has been stated in the [points to consider](#), when using Tarantella the user always specifies the *global* batch size. This has the advantage that the optimization process during the training of a DNN, and in particular the loss function do not depend on the number of devices used during execution.

However, when the number of devices becomes very large, the (device-local) micro-batch size might become so small, that DNN kernel implementations are less efficient, resulting in overall performance degradation. This is why it is in practice often advisable to scale the global batch size with the number of nodes. This will often lead to linear speedups in terms of the time to accuracy when increasing the number of devices used, at least up to some *critical batch size*, cf. [Shallue] and [McCandlish]. Changing the batch size of the optimizer will however also imply the need to adapt the learning rate schedule.

For details, cf. for instance the [ResNet-50 tutorial](#).

If you decide to scale the batch size with the number of nodes, Tarantella provides two different ways to achieve this easily. The first option is to multiply the local batch size (for instance passed via a command-line parameter) with the number of devices used, batch your dataset with it, and call `fit` on it:

```
micro_batch_size = args.micro_batch_size
batch_size = tnt.get_size() * micro_batch_size
train_dataset = train_dataset.batch(batch_size)
tnt_model.fit(train_dataset)
```

As a second option you can also pass the local batch size directly to the `tnt_micro_batch_size` parameter in `fit`, and leave your dataset unbatched:

```
micro_batch_size = args.micro_batch_size
tnt_model.fit(train_dataset,
              tnt_micro_batch_size = micro_batch_size)
```

This parameter is also available in `evaluate` and `predict`. In addition, `fit` also supports setting the validation set micro batch size in a similar way with `tnt_validation_micro_batch_size`. For more information, please also read [using distributed datasets](#).

8.3 Setting tensor fusion threshold

Tarantella automatically uses [Tensor Fusion](#) with a default threshold of 32kB. This threshold specifies the minimal size of local buffers in *allreduce* communication operations used to accumulate partial gradients during *backpropagation*.

Note that the threshold value implies a trade-off between the potential to utilize network bandwidth, and the overlap of computation and communication during *backpropagation*. The larger the threshold, the more bandwidth-bound the *allreduce* algorithm will get, but the less potential there will be to overlap its execution with kernel computations. Also note that the ideal threshold value will generally depend on the number of nodes used.

To change the default value, you can pass a threshold value in kB to tarantella:

```
tarantella --hostfile hostfile --fusion-threshold=<FUSION_THRESHOLD_KB> -- model.py
```

8.4 Performance aspects

To increase execution performance on CPUs, it is often desirable to bind processes to physical cores or groups of cores in order to improve data locality and reduce context switching.

Tarantella provides two command-line flags to enable rank pinning to physical sockets. They rely on the `numactl` utility to detect existing NUMA domains and pin processes to them.

Tarantella pinning flags allow users to:

- pin each Tarantella process deployed on a host to a separate socket (through the `--pin-to-socket` flag)
- pin memory allocation for each Tarantella process to the socket memory (through the `--pin-memory-to-socket` flag).

Using only `--pin-to-socket` will result in memory being only preferentially allocated on the socket memory, but potentially using memory from other NUMA domains when necessary.

The example below illustrates the usage of the `--pin-to-socket` and `--pin-memory-to-socket` flags to start two Tarantella ranks on each host listed in `hostfile`, each of them pinned to a different socket.

```
tarantella --hostfile hostfile --npernode 2 --pin-to-socket -- model.py
```

8.5 Python Interpreter

The `tarantella` CLI can be used as generic tool for executing code on multiple devices simultaneously. While usually the executed program is a Python file, Tarantella uses the Python interpreter it finds in the current `$PATH`. Changing the interpreter can be easily achieved by using the `--python-interpreter` flag:

```
tarantella --hostfile hostfile --npernode 2 --python-interpreter=/path/to/python -- ↵
↵ model.py
```

Additionally, the user can also execute binary files that do not require any Python support by simply passing an empty string to the `--python-interpreter` flag.

A typical use case for the interpreter is to enable the usage of other tools that can only be enabled from the command line, such as checking for memory leaks in a parallel program with `valgrind`

```
tarantella -n 2 --python-interpreter="valgrind --leak-check=yes \
                                     --track-origins=yes --tool=memcheck \
                                     python" \
                                     -- model.py
```

8.6 Reproducibility

Reproducibility is a very important prerequisite to obtain meaningful results in scientific computing and research. Unfortunately, using stochastic algorithms, pseudo random generators and having to deal with the pitfalls of floating-point arithmetics, it is particularly difficult to achieve reproducibility in Deep Learning research.

In order to be able to reproduce results obtained with TensorFlow, when running in a multi-node/multi-device setting with Tarantella, one needs to meet at least the following requirements:

- set the random seed with `tf.random.set_seed(seed)`
- set the environment variable `os.environ['TF_DETERMINISTIC_OPS'] = '1'`

- set the environment variable `os.environ['TF_CUDNN_DETERMINISTIC'] = '1'`
- set the random seed when using layers such as `keras.layers.Dropout`
- set the shuffle seeds when using `tf.data.Dataset` with `shuffle(seed=seed)` and `list_files(seed=seed)`
- set the deterministic parameter to `True` in `Dataset` transformations such as `interleave` and `map`

Additionally, Python-specific random generators might need to be seeded, in particular:

- `random.seed(seed)`
- `numpy.random.seed(seed)`
- `os.environ['PYTHONHASHSEED'] = str(seed)`

For more details, take a look at a more in-depth study of [non-determinism sources in TensorFlow](#).

FREQUENTLY ASKED QUESTIONS (FAQ)

This is a list of frequently asked questions about Tarantella. Please feel free to *suggest new ones*!

Question

How can I ssh to localhost without password?

In order to run Tarantella programs, you will need to be able to ssh to localhost without password. In order to do that generate ssh keys first:

```
cd ~/.ssh  
ssh-keygen
```

Make sure not to overwrite existing keys. When asked for a passphrase, Enter passphrase (empty for no passphrase):, simply leave empty and return with enter. Also take specific care to set correct user rights on all files in .ssh, cf. for instance [here](#). Next, append the public key to the authorized_keys file:

```
cat id_rsa.pub >> authorized_keys
```

Now, install and start an ssh server, e.g., openssh-server on Fedora. More details can be found for instance [here](#).

Question

I get an execution error `GPI library initialization incorrect environment vars` when trying to run my script. What shall I do?

Most likely you are running your program with `python my_script.py` or `./my_script.py`. Please make sure to execute your code with `tarantella -- my_script.py` instead.

Question

I get an execution error `GPI library initialization general error`. What shall I do?

This error occurs when the GPI-2 library tries to connect to a previously used socket, which is not yet released. Try to re-run your code after a short while so that the port becomes available again.

Question

The execution seems to stall. What shall I do?

Please use the `tarantella --cleanup` command to kill any processes that might be still running from a previous (aborted) call to `tarantella` as shown [here](#). Note that you can also interrupt a running `tarantella` instance (distributed on multiple nodes) by using `Ctrl+c`.

Question

When trying to build Tarantella, CMake cannot find pybind11:
Could not find a package configuration file provided by "pybind11" with any
of the following names: [...]
What shall I do?

This error occurs when pybind11 is installed using `pip`. Please use `conda` instead, as recommended in the [installation guide](#).

Question

When trying to build Tarantella, CMake does not detect the Python interpreter from the active `conda` environment.
What shall I do?

You will need to manually add the path to the `conda` environment's `bin` directory to your `PATH`. You will also need to specify the path to the python library on the command line when configuring Tarantella:

```
PATH_TO_CONDA_ENV=/path/to/conda/env
export PATH=${PATH_TO_CONDA_ENV}/bin:${PATH}
cmake -DPYTHON_EXECUTABLE=${PATH_TO_CONDA_ENV}/bin/python \
      -DPYTHON_LIBRARY=${PATH_TO_CONDA_ENV}/lib ../
```

Question

Why do I get runtime errors when I compile Tarantella using `clang`?

Currently, Tarantella can be built properly only by using `gcc`.

The `clang` compiler relies on a different standard library (`libc++` instead of `libstdc++` that is used by `gcc`).

However, the TensorFlow `pip/conda` packages for Linux are compiled using `gcc`. The `tnt_tfops` library in Tarantella is linked against TensorFlow, which leads to linking errors at runtime if the two libraries expect a different standard library implementation.

Question

I get *undefined symbol* errors in the `libtnt-tfops.so` library at runtime. What can I do?

Such errors might be due to a TensorFlow version mismatch between Tarantella and the loaded `Conda` environment. Make sure to use the same `Conda` environment that was active when compiling Tarantella.

Question

Why does loading a Tarantella or Keras model from YAML fail?

Make sure to have the *PyYAML* Python package installed in your environment, using version *3.13* or below. Newer versions of *PyYAML* do not work with TensorFlow model loading.

```
pip install PyYAML==3.13
```

Question

Can I install Tarantella on MacOS?

Tarantella is only supported on Linux systems, as its GPI-2 dependency is built on top of a Linux kernel API called *epoll*.

BUG REPORTS

To report a bug please open an [issue on GitHub](#).

When opening an issue, please make sure you include as much information as possible about the issue. Please consider providing at least the following points:

- What version of Tarantella you are using
- What linux distribution you are using (e.g., Linux Ubuntu 20.04)
- What kind of system you are experiencing the issue on (type and number of nodes, network interconnect, etc.)
- What did you expect to see and what have you seen instead
- What exact steps are needed to reproduce the issue

FEATURE REQUESTS

For contributions other than modifications to the source code, as for example suggestions of a feature or enhancement, please open an [issue on GitHub](#) with the label **Feature**.

When providing a feature request, please consider providing at least the following information:

- What is the current behavior of the software and how does the feature improve it
- Who would benefit from the feature
- Is there a relevant reference or academic paper describing the feature
- Are you willing to contribute to and/or maintain the feature

CONTRIBUTING

Thank you for considering to contribute to Tarantella.

There are many ways to contribute to Tarantella. This includes sharing DNN models distributed through Tarantella, providing suggestions on improving the documentation, as well as contributing with changes to the [Tarantella code base](#). Even by simply providing suggestions on how we can *improve Tarantella* and help spreading the word about it are great ways to contribute and make Tarantella better software.

If you want to contribute to Tarantella with changes to its code, please open a [pull request](#) on GitHub.

CONTACT

In case you have any feature request, or want to report a bug please follow [these instructions](#).

If you consider contributing to Tarantella, please follow the instructions [here](#).

If you have any further questions or comments please email to support@tarantella.org

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