### tca

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#### Abstract

tca is a C implementation of the non-parametric dynamic topic model from ?. It allows both the topic mixtures and the word-topic distributions to change over epochs (time) through two-parameter Poisson-Dirichlet processes (PDP). The software can also be configured to run more like a standard LDA model with proper parameters, though this is intended only for testing. "Document completion" testing, burstiness modelling, coherence measurements with PMI and a few other diagnostic values are supported. A multi-core version is in development but not recommended currently.

## 1 Synopsis

tca [-?] [-?Arq] DataStem RepStem

## 2 Description

tca reads the collection of files with stem DataStem that form the input set of data. When checkpointing, or at termination, the output is written to files with stem RepStem. On restart with the -r option, some of these are also read initially to restore the previous state. A log of the run is reported to stderr if the -e option is used. By default, the log goes to RepStem.log.

The programme is research software, so not all options or combinations of options work correctly. Note that in this release, all the more experimental features have been stripped, so this release contains only moderately well tested components.

The programme runs a Gibbs sampler for a model consisting of the following three parts:

theta: this is the prior on topic vector (theta) for each document. tca uses a Poisson-Dirichlet distribution with parameters  $a_{\theta}$  and  $b_{\theta}$ . The vector has dimension T (the number of topics).

**phi:** this is the prior on word vector (phi) for each topic. tca uses a Poisson-Dirichlet distribution with parameters  $a_{\phi}$  and  $b_{\phi}$ . The vector has dimension W (the number of words).

mu: this is the prior on topic vector (mu) for each epoch. tca uses a Poisson-Dirichlet distribution with parameters  $a_{\mu}$  and  $b_{\mu}$ . The vector has dimension T (the number of topics).

**burst:** this is the burstiness component which has a document specific variant of the word vector for each topic. This is not used by default.

It is possible to find-tune the model specification through the **-S** option.

The programme uses generalised second order Stirling numbers with the library extracted from *libstb* version 1.8 released at https://github.com/wbuntine/libstb. This is initialised with predefined bounds on the tables, and these can be modified with the -N option. This should be used for collections with more than 20,000 documents, but its best to run first and on error, increase the bounds.

## 3 Options

Options have a single letter followed by a possible single argument. Options are grouped under the following functions: setting of hyperparameters, controlling sampling of hyperparameters, general control, and testing and reports

### 3.1 Setting of hyperparameters

-Svar=value Set variable var to float value, where var can be one of:

am discount parameter for mu, the topic distribution.

at discount parameter for theta, topic distribution per document.

ap0 discount parameter for phi, the word distribution per topic, in the first epoch.

ap1 discount parameter for phi for the second epoch onwards.

**bm0** concentration parameter for mu in the first epoch.

bm1 concentration parameter for mu for the second epoch onwards.

bt concentration parameter for theta

**b0p** concentration parameter for on the prior of phi.

b0m concentration parameter for on the prior of mu

**bp0** concentration parameter for **phi** in the first epoch.

**bp1** concentration parameter for **phi** for the second epoch onwards.

ab discount parameter for burstiness

**bb** concentration parameter for burstiness, a constant initially but subsequent sampling will allow a different value per topic.

If the discount parameter is set to 0, the distribution will become a symmetric Dirichlet, the LDA default.

#### 3.2 Controlling sampling of hyperparameters

**-F**var Fix the variable var where it takes the value of a hyper-parameter to the **-S** option.

-gvar,int Set the extra integer parameter for sampling var to int.

-Gvar, cycles, start Sample the variable var where it takes the value of a hyper-parameter to the -S option. The sampling will be after start cycles and then repeat every cycles cycles.

#### 3.3 General control

**-bepochs** Specify the maximum number of previous epochs that table indicators can use for computation.

-ccycles Do a checkpoint every this many cycles. This saves the output statistics and the parameter file adequate to do a restart with -r option.

-Ccycles Stop after this many cycles. Default is 100.

-ddots For really big batches of data, print a "." every dots documents within a single cycle.

-e Reroute logging to the stderr.

-fformat Read input data from data formatted according to the type format. Data is expected to come from an input file with name DataStem.Suff where Suff is an appropriate suffix. These are given with Input Files below. Allowed formats are: ldac, witdit, docword, bag and lst.

- **-K**topics Set T the maximum number of topics. Default is 10.
- -NmaxN,maxM Set maximum for the Stirling number tables to count maxM and maximum table counts for  $a_{\mu}$  and  $a_{\phi}$  to maxM. Default is 1000,10000. Higher numbers are needed when running with large number of documents.
- -qthreads If compiled with threading, enables this many threads. Default is 1.
- -rtype Restart using something built previously, where type is:
  - **hca** Restart from the output file from *hca* and the beta side is held constant and not sampled. Can significantly speed up testing or querying sometimes.
  - mu Load up the mu matrix from file RepStem.mu where it is in standard sparse format (each line is "e k val" for e=epoch, k=topic index and the value). This means mu will not be estimated. Note the probabilities are expected to be normalised along the topic dimension.
  - phi Load up the phi matrix from files RepStem.phiNNN for NNN is a 3-digit representation of the topic index (so, "000", "001", ...). The file is in standard sparse format (each line is "w k val" for w=word index, k=topic index and the value). This means phi will not be estimated. Note the probabilities are expected to be normalised along the word dimension.
  - tca Restart from a previous run, reloading data and parameters.
- **-sseed** Initialise the random number seed.
- -v Up verbosity by one increment. Starts at zero and currently understands 0-3.
- ${ extstyle -V}$  load the dictionary from the DataStem.tokens file for use in reporting. It has one token per line.
- -Wmax Set to the maximum number of unique words to max

#### 3.4 Testing and reports

- -hHold,arg Do document completion testing on the test set. There are three styles of document completion implemented. When Hold is the string "doc", then every arg-th word is held out in estimating the latent variables (like theta) for the document and used instead for testing of perplexity. That is, words at document positions arg-1, 2\*arg-1, etc. When Hold is the string "dict", then every arg-th word in the dictionary is held out in estimating and used for testing. So if a word has dictionary index arg-1, 2\*arg-1, etc., it is held out. When Hold is the string "fract", then the fract proportion at the tail of the document is held out. The initial proportion is used in estimating.
- -lDiag, cycles, start Do a run-time estimation of the diagnostic Diag starting after the start cycle and then taking the estimate every cycles cycle. Diagnostics are:
  - mu Estimate the topic probability vector across epochs. Stored in RepStem.mu.
  - phi Estimate the word probability vector for each topic. Stored in the RepStem.phiNNN files where NNN indicates topic number ("000", "001", etc.).
  - **prog** How often to do the standard diagnostic reports (default is every 5-th cycle).
- **-L**Diag,cycles,start Do a diagnostic estimate Diag after all Gibbs sampling is complete. Sampling of the estimate starts after the start cycle and goes for a total of cycles cycles (including the starting ones). Diagnostics are:
  - like Estimate likelihood/perplexity on the test set using the standard (biased) document likelihood.

-oscore Scoring rule to pick top words for printing. Methods are 'count', 'idf', and 'cost'.

-p Report topic coherency in the log file, and save results in the parameter file. This requires a DataStem.pmi or DataStem.pmi.gz file exist in the right format. This can be created with the *mkmat.pl* and *cooc2pmi.pl* scripts in the scripts directory of the release. The format is a simple sparse matrix form with lines of the form "N M PMI" for word indices (offset by 0) N and M and PMI value.

-tsize Specify size of training set. It takes the first size entries in the data set. Default is all the set minus the test data.

-Tsize Specify size of test set. It takes the last size entries as the test set and must agree with whats in the DataStem.epoch file.

-V load the dictionary from the DataStem.tokens file for use in reporting. It has one token per line.

## 4 Input Files

The following files provide details about the dataset. The filenames are constructed by adding a suffix to the data stem. The data (document+word) format itself can be one of four different formats and is specified with the **-f** option.

DataStem.docs A list of documents including their titles

DataStem.epoch epoch information file. The first number indicates the total number of epochs in the dataset, followed by the number of entries in each epoch in subsequent lines.

DataStem.ldac Standard LdaC format. Word indices to the dictionary are offset by 1.

DataStem.srcpar summary information on the total number of documents and features of the dataset.

DataStem.stops List of stop words

DataStem.titles A list of titles for all entries

DataStem.tokens tokens/words in the dictionary, one per line.

 ${\tt DataStem.tpc} \ \ {\rm Meta} \ \ {\rm data} \ \ {\rm of} \ \ {\rm the} \ \ {\rm dataset} \ \ {\rm structure}$ 

DataStem.txtbag default bag or list format for linkBags(1) command of DCABags. Word indices offset by 0.

DataStem.words tokens/words in the dictionary, one per line.

## 5 Output Files

The following files are output when the system checkpoints or at the end of the run. These are built by adding a suffix to the report stem, RepStem. The first set of files are:

RepStem.log Log file created if -e option not used.

RepStem.mu The mu matrix of dimension E by T. The file is in standard sparse format (each line is "e k val" for e=epoch index, k=topic index and the value). Generated with appropriate use of the -1mu option. Note the probabilities are expected to be normalised along the topic dimension.

RepStem.par Parameter and dimensions file in simple "var = value" format. These are detailed in the next section.

RepStem.phiNNN The phi matrix written for NNN is a 3-digit representation of the topic index (so, "000", "001", ...). The file is in standard sparse format (each line is "w k val" for w=word index, k=topic index and the value). Generated with appropriate use of the -lphi option. Note the probabilities are expected to be normalised along the word dimension.

RepStem.top A simple text report giving the top word indices for each topic. If a hierarchical model in use, then the "-1" topic is for the base distribution of words. Word indices are offset from 0.

RepStem.tpk Topic-word information per epoch.

Note the RepStem.phiNNN and RepStem.mu files are also used as input with the -rphi and -rmu options respectively.

The second set of files gives the actual runtime statistics. Output matrices are in a simple readable sparse vector format the same as the DataStem.docword format.

RepStem.cdt Document by topic table counts.

RepStem.cpet Epoch by topic table counts (the actual data counts are totaled from RepStem.cdt).

RepStem.istats Debugging output optionally switched on giving details for each topic of how much the epoch affects others.

RepStem.mevt Epoch by word by topic counts, statistics for phi.

RepStem.sevt Epoch by word by topic table counts corresponding to RepStem.mevt.

RepStem.zt With no burstiness, gives topic index (offset by 0), one per line. With burstiness, gives one "z,r" per line where "z" is the topic index (offset by 0) and "r" is the burst table indicator, which is 1 if the word contributes to standard LDA statistics, and 0 if burstiness modelling says the word is a burst so does not contribute to LDA statistics.

These files along with RepStem.par are input on a restart using -r.

### 6 The Parameter File

The parameter file has the following dimensions:

N – number of words in the full collection, summed over all documents.

NT – number of words in the training set, summed over all training documents.

W – number of words in the dictionary.

D – number of documents in total.

TRAIN – number of documents to train on, is always the the first ones in the file.

TEST – number of documents to test on, is always the the last ones in the file.

T – maximum number of topics.

ITER – number of major cycles made last.

In addition, the float parameters allowed to be specified with the **-F** and **-G** options are also given. This includes all the hyper-parameters for distribution  $\mu$ ,  $\phi$  and  $\theta$  (am, bm0, bm1, b0m, at, bt, ab, bb, ap0, ap1, b0p, bp1).

## 7 Examples

These examples work as is on late model Linux, Macs and Windows. However, you need to replace the executable, tca, by the system dependent one, from the install directory where the data/ directory is. For instance, on Windows that might be tca/tca.exe.

Run the topic model from ? with fixed hyper-parameters on the full dataset and no testing, sending logging to stderr.

```
tca -v -e -V -K20 -C100 data/ch c1
```

The parameters as they are sampled will be printed on a line beginning wth "Par:". At the end, the top 20 words will be printed and the final training perplexity printed. This is based on the posterior probability, not word probability estimates. Restart and build a topic probability vector for each document, as well as sparsity mappings for the words in data/ch.smap file.

```
tca -v -rtca -e -lsp,2,1 -ltheta,2,1 -C20 data/ch c1
```

Now view the sparsity report at c1.smap and the topic probabilities at c1.theta, and the values saved in the parameter file c1.par. Rerun the model fitting on the full dataset and testing on the final 100 documents. Logging now to c1.log. Checkpoint every 20 cycles (note, we usually only do this for cycles taking over 10 minutes each).

```
tca -v -V -K20 -C100 -c20 -T100 data/ch c1
```

View the end of the log file to get the test perplexity, which is printed after "log\_2(test perpML)" Now rerun but use document completion (every 4th word), not the default likelihood calculation.

```
tca -v -V -rtca -hdoc,4 data/ch c1
```

View the end of the log file to get the test perplexity, which is printed after "log\_2(test perpHold)". Note it is also recorded in the parameter file. Restart, no sampling this time using  $-\mathbf{C}\theta$ , just record the PMI and the classification details on test data.

```
tca -v -V -rtca -C0 -Llike,0,0 -Lclass,10,1 -p data/ch c1
```

Note the  $-\mathbf{L}like, 0, 0$  option is used to prevent it doing test likelihood calculations, which are potentially slow on larger data sets. The PMI data has a value printed for each topic as well as a final average. It bases its calcuations on the matrix  $\mathtt{data/ch/pmi.gz}$  created explicitly for this test set. For other datasets, you will need to down load prepared PMI matrices from the project homepage.

To relax the assumptions on the fixed hyperparameters, one can run with the optimisation options. For instance, if we allow the word-topic distribution (phi) to evolve at different speed for different topics, we can sample the concentration parameter bp0 and bp1 as the following example,

```
tca -v -v -K20 -C200 -Gbp0,2,2 -Gbp1,2,2 data/ch c1
```

The hyperparameter sampling slows it down quite a bit but seems to make a significant difference. Unused topics sometimes get a very low concentration.

### 8 Errors

There is some error reporting on failure.

If the software quits during a run on larger data with an error message like:

```
S_V(N,M,A) tagged 'XXX' hit bounds (BN,BM)
```

for integers N,M and label XXX then you need to increase the bounds. If the tag XXX is "SX, docXtopic PYP" then increase the bound BM using the option  $-\mathbf{N}BM$  with an increased BM. If the tag XXX is "SX, docXtopic PYP" then increase both bounds BN,BM using the option  $-\mathbf{N}BM,BN$  (note the order of the bounds).

For other errors, please report to the maintainer.

### 9 See Also

The command linkBags(1) is available from http://mloss.org in the software DCABags. The extended library libstb, parts of which are included, is available individually from http://mloss.org. Other supporting software and data sets are available at the project home page at https://github.com/wbuntine/topic-models

### 10 Version

This programme is version 0.4 of September 16, 2014. This incorporates parts of the library *libstb* version 1.8 also of September 16, 2014.

# 11 License and Copyright

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### 12 Author

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Some parts also done by Dr. Jinjing Li and Mr. Swapnil Mishra.

### References

Buntine, W. and Li, J. (2014). Dynamic Topic Models using Hierarchical Two-parameter Poisson-Dirichlet Process