

HTK (v.3.1): Basic Tutorial

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What is HTK?

HTK is the “Hidden Markov Model Toolkit” developed by the Cambridge University Engineering Department (CUED). This toolkit aims at building and manipulating **Hidden Markov Models** (HMMs). HTK is primarily used for **speech recognition** research (but HMMs have a lot of other possible applications...)

HTK consists of a set of library modules and tools available in C source form. It is available on free download, along with a complete documentation (around 300 pages).

See: <http://htk.eng.cam.ac.uk/>.

1 Yes/No Recognition System

In this tutorial, we propose to build a 2-word recogniser with a { “Yes”, “No” } vocabulary, based on HTK tools. It’s the most basic Automatic Speech Recognition (ASR) system that can be designed...

1.1 Construction steps

The main construction steps are the following:

- Creation of a training database: each element of the vocabulary is recorded several times, and labelled with the corresponding word.
- Acoustical analysis: the training waveforms are converted into some series of coefficient vectors.
- Definition of the models: a prototype of Hidden Markov Model (HMM) is defined for each element of the task vocabulary.
- Training of the models: each HMM is initialised and trained with the training data.
- Definition of the task: the grammar of the recogniser (what can be recognised) is defined.
- Recognition of an unknown input signal.
- Evaluation: the performance of the recogniser can be evaluated on a corpus of test data.

1.2 Work space organisation

It is recommended to create a directory structure such as the following:

- `data/` : to store training and test data (speech signals, labels, etc.), with 2 sub-directories `data/train/` and `data/test/` to separate the data used to train the recogniser from the ones used for performance evaluation.
- `analysis/` : to store files that concern the acoustical analysis step.
- `training/` : to store files that concern the initialisation and training steps.
- `model/` : to store the recogniser’s models (HMMs).
- `def/` : to store files that concern the definition of the task.
- `test/` : to store files that concern the test.

1.3 Standard HTK tool options:

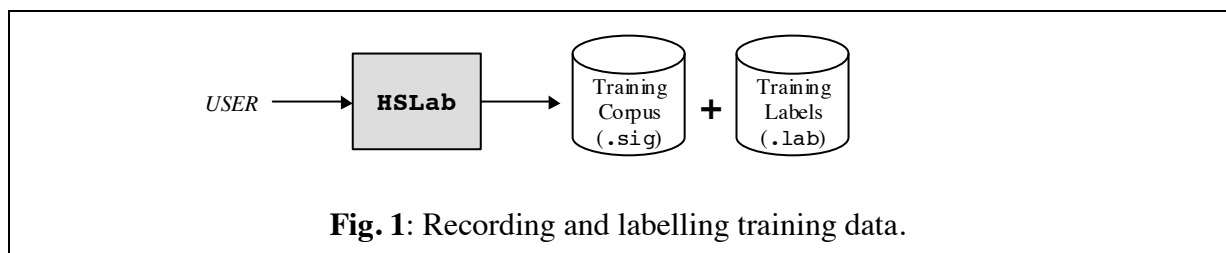
Some standard options are common to every HTK tools. In the following, we will use some of them:

- `-A` : displays the command line arguments.
- `-D` : displays configuration settings.
- `-T 1` : displays some information about the algorithm actions.

To have the complete list:

see HTK documentation, p.50 (Chap.4, The Operating Environment).

2 Creation of the Training Corpus



First, we have to record the “Yes” and “No” speech signals with which word models will be trained (the *training corpus*). Each speech signal has to be labelled, that is: associated with a text (a label) describing its content. Recording and labelling can be done with the HSLab HTK tool (but any other tool could be used).

To create and label a speech file:

HSLab any_name.sig

The tool’s graphical interface appears.

2.1 Record the Signal

Press “Rec” button to start recording the signal, then “Stop”.

A buffer file called `any_name_0.sig` is automatically created in the current directory.

(if you make a new record, it is saved in a second buffer file called `any_name_1.sig`).

Remarks:

- The signal files (`.sig`) are here saved in a specific HTK format. It is however possible to use other audio format (`.wav`, etc.):

see HTK documentation, p.68 (Chap.5, Speech Input/Output).

- The default sampling rate is 16kHz.

2.2 Label the Signal

To label the speech waveform, first press “Mark”, then select the region you want to label. When the region is marked, press “Labelas”, type the name of the label, then press *Enter*.

In this tutorial, we will only record isolated words (“Yes” or “No”) preceded and followed by a short silence. For each signal, we have to label 3 successive regions: start silence (with label `sil`), the recorded word (with label `yes` or `no`), and end silence (with label `sil`). These 3 regions cannot overlap with each other (but no matter if there is a little gap between them). When the 3 labels have been written, press “Save”: a label file called `any_name_0.lab` is created. At this point you can press “Quit”.

Remark:

The `.lab` file is a simple text file. It contains for each label a line of the type:

```
4171250 9229375 sil
9229375 15043750 yes
15043750 20430625 sil
```

where numbers indicate the start and end sample time of each label. Such a file can be manually modified (for example to adjust the start and end of a label) or even created (the use of the HSLab tool is not required).

2.3 Rename the Files

After each recording/labelling, you have to rename the `.sig` and `.lab` files to your convenience (e.g. `yes01.sig` and `yes01.lab`).

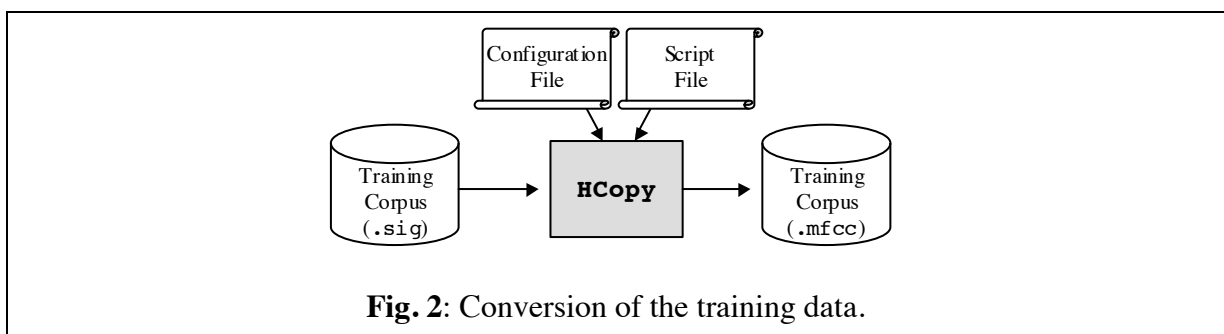
10 records for each of the 2 words should be enough for this tutorial.

The signal files should be stored in a `data/train/sig/` directory (the training corpus), the labels in a `data/train/lab/` directory (the training label set).

For more details on the HSLab graphical interface:

see *HTK documentation*, p.237 (*Reference Section*, *HSLab*).

3 Acoustical Analysis



The speech recognition tools cannot process directly on speech waveforms. These have to be represented in a more compact and efficient way. This step is called “acoustical analysis”:

- The signal is segmented in successive frames (whose length is chosen between 20ms and 40ms, typically), overlapping with each other.
- Each frame is multiplied by a windowing function (e.g. Hamming function).
- A vector of acoustical coefficients (giving a compact representation of the spectral properties of the frame) is extracted from each windowed frame.

The conversion from the original waveform to a series of acoustical vectors is done with the HCopy HTK tool:

```
HCopy -A -D -C analysis.conf -S targetlist.txt
```

analysis.conf is a configuration file setting the parameters of the acoustical coefficient extraction.

targetlist.txt specifies the name and location of each waveform to process, along with the name and location of the target coefficient files.

3.1 Configuration Parameters

The configuration file is a text file (“#” can be used to introduce a comment). In this tutorial, the following configuration file will be used:

```
#
# Example of an acoustical analysis configuration file
#
SOURCEFORMAT = HTK           # Gives the format of the speech files
TARGETKIND = MFCC_0_D_A      # Identifier of the coefficients to use

# Unit = 0.1 micro-second :
WINDOWSIZE = 250000.0        # = 25 ms = length of a time frame
TARGETRATE = 100000.0        # = 10 ms = frame periodicity

NUMCEPS = 12                 # Number of MFCC coeffs (here from c1 to c12)
USEHAMMING = T               # Use of Hamming function for windowing frames
PREEMCOEF = 0.97             # Pre-emphasis coefficient
NUMCHANS = 26                # Number of filterbank channels
CEPLIFTER = 22               # Length of cepstral liftering

# The End
```

List. 1: Analysis configuration file.

With such a configuration file, an MFCC (Mel Frequency Cepstral Coefficient) analysis is performed (prefix “MFCC” in the TARGETKIND identifier). For each signal frame, the following coefficients are extracted:

- The 12 first MFCC coefficients [c1,..., c12] (since NUMCEPS = 12)
- The “null” MFCC coefficient c0, which is proportional to the total energy in the frame (suffix “_0” in TARGETKIND)
- 13 “Delta coefficients”, estimating the first order derivative of [c0, c1,..., c12] (suffix “_D” in TARGETKIND)
- 13 “Acceleration coefficients”, estimating the second order derivative of [c0, c1,..., c12] (suffix “_A” in TARGETKIND)

Altogether, a 39 coefficient vector is extracted from each signal frame.

For more details on acoustical analysis configuration:

see HTK documentation, p.58-66 (Chap.5, Speech Input/Output).

3.2 Source / Target Specification

One or more “source file / target file” pairs (i.e. “original waveform / coefficient file”) can be directly specified in the command line of `HCopy`. When too many data are to be processed, the `-S` option is used instead. It allows to specify a script file of the form:

```
data/train/sig/yes01.sig    data/train/mfcc/yes01.mfcc
data/train/sig/yes02.sig    data/train/mfcc/yes02.mfcc
etc...
data/train/sig/no01.sig     data/train/mfcc/no01.mfcc
data/train/sig/no02.sig     data/train/mfcc/no02.mfcc
etc...
```

List. 2: Conversion script file.

Such a text file can be automatically generated (using a Perl script, for instance). The new training corpus (`.mfcc` files) is stored in the `data/train/mfcc/` directory.

For more details on the `HCopy` tool:
[see HTK documentation, p.195 \(Reference Section, HCopy\).](#)

4 HMM Definition

In this tutorial, 3 acoustical events have to be modelled with a Hidden Markov Model (HMM): “Yes”, “No” and “Silence”. For each one we will design a HMM.

The first step is to choose *a priori* a topology for each HMM:

- number of states
- form of the observation functions (associated with each state)
- disposition of transitions between states

Such a definition is not straightforward. There is actually no fixed rule for it. Here, we will simply choose the same topology for each of the 3 HMMs (Fig.3):

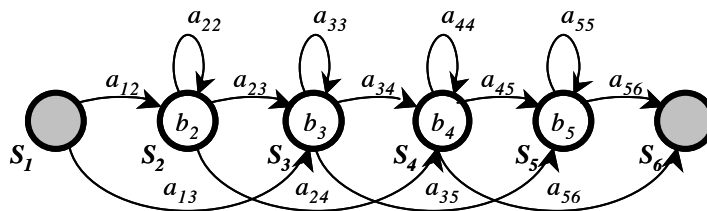


Fig. 3: Basic topology

The models consist actually of 4 “active” states $\{S_2, S_3, S_4, S_5\}$: the first and last states (here S_1 and S_6), are “non emitting” states (no observation function), only used by HTK for some implementation facilities reasons. The observation functions b_i are single gaussian distributions with diagonal matrices. The transition probabilities are quoted a_{ij} .

In HTK, a HMM is described in a text description file. The description file for the HMM depicted on Fig.3 is of the form:

```
~o <VecSize> 39 <MFCC_0_D_A>
~h "yes"
<BeginHMM>
    <NumStates> 6
    <State> 2
        <Mean> 39
            0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0
            0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0
            0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0 0.0
        <Variance> 39
            1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0
            1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0
            1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0
    <State> 3
        <Mean> 39
            0.0 0.0 (...) 0.0
        <Variance> 39
            1.0 1.0 (...) 1.0
    <State> 4
        <Mean> 39
            0.0 0.0 (...) 0.0
        <Variance> 39
            1.0 1.0 (...) 1.0
    <State> 5
        <Mean> 39
            0.0 0.0 (...) 0.0
        <Variance> 39
            1.0 1.0 (...) 1.0
    <TransP> 6
        0.0 0.5 0.5 0.0 0.0 0.0
        0.0 0.4 0.3 0.3 0.0 0.0
        0.0 0.0 0.4 0.3 0.3 0.0
        0.0 0.0 0.0 0.4 0.3 0.3
        0.0 0.0 0.0 0.0 0.5 0.5
        0.0 0.0 0.0 0.0 0.0 0.0
<EndHMM>
```

List. 3: HMM description file (prototype).

~o <VecSize> 39 <MFCC_0_D_A>
is the header of the file, giving the coefficient vector size (39 coefficients here), and the type of coefficient (MFCC_0_D_A here).

~h "yes" <BeginHMM> (...) <EndHMM>
encloses the description of a HMM called “yes”.

<NumStates> 6
gives the total number of states in the HMM, including the 2 non-emitting states 1 and 6.

<State> 2
introduces the description of the observation function of state 2. Here we have chosen to use single-gaussian observation functions, with diagonal matrices. Such a function is entirely described by a mean vector and a variance vector (the diagonal elements of the autocorrelation matrix). States 1 and 6 are not described, since they have no observation function.

<Mean> 39

0.0 0.0 (...) 0.0 (x 39)

gives the mean vector (in a 39 dimension observation space) of the current observation function. Every element is arbitrary initialised to 0: the file only gives the “prototype” of the HMM (its global topology). These coefficients will be trained later.

<Variance> 39

1.0 1.0 (...) 1.0 (x 39)

gives the variance vector of the current observation function. Every element is arbitrary initialised to 1.

<TransP> 6

gives the 6x6 transition matrix of the HMM, that is:

a_{11}	a_{12}	a_{13}	a_{14}	a_{15}	a_{16}
a_{21}	a_{22}	a_{23}	a_{24}	a_{25}	a_{26}
a_{31}	a_{32}	a_{33}	a_{34}	a_{35}	a_{36}
a_{41}	a_{42}	a_{43}	a_{44}	a_{45}	a_{46}
a_{51}	a_{52}	a_{53}	a_{54}	a_{55}	a_{56}
a_{61}	a_{62}	a_{63}	a_{64}	a_{65}	a_{66}

where a_{ij} is the probability of transition from state i to state j . Null values indicate that the corresponding transitions are not allowed. The other values are arbitrary initialised (but each line of the matrix must sum to 1): they will be later modified, during the training process.

Such a prototype has to be generated for each event to model.

In our case, we have to write a prototype for 3 HMMs that we will call “yes”, “no”, and “sil” (with headers ~h “yes”, ~h “no” and ~h “sil” in the 3 description files).

These 3 files could be named `hmm_yes`, `hmm_no`, `hmm_sil` and be stored in a directory called `model/proto/`.

For more details on HMM description files:

see HTK documentation, p.94 (Chap.7, HMM Definition Files).

5 HMM Training

The training procedure is described on Fig.4:

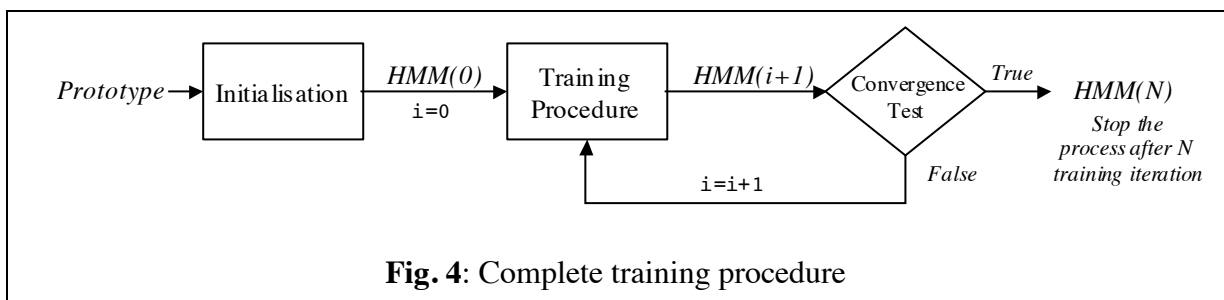
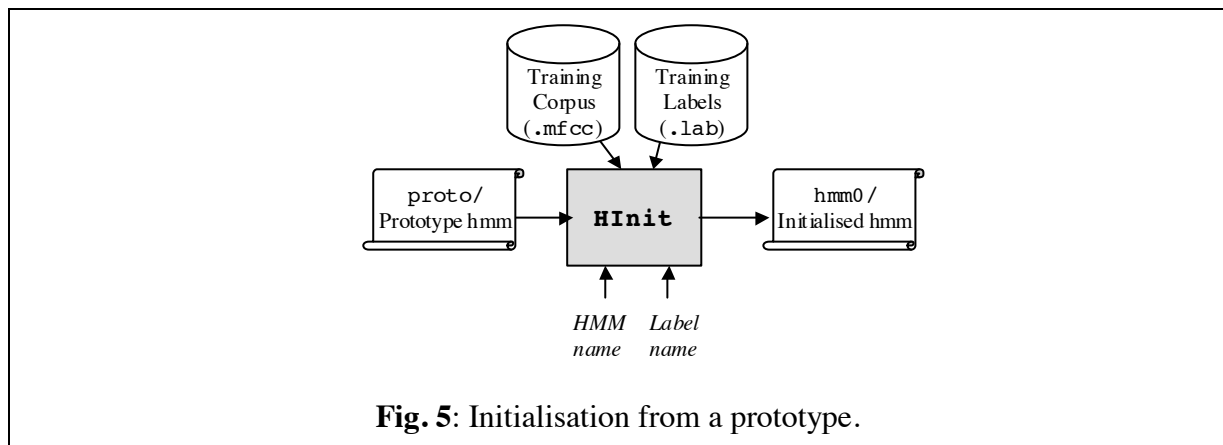


Fig. 4: Complete training procedure

5.1 Initialisation

Before starting the training process, the HMM parameters must be properly initialised with training data in order to allow a fast and precise convergence of the training algorithm.

HTK offers 2 different initialisation tools: `Hinit` and `HCompv`.



HInit

The following command line initialises the HMM by time-alignment of the training data with a Viterbi algorithm:

```
HInit -A -D -T 1 -S trainlist.txt -M model/hmm0 \  
-H model/proto/hmmfile -l label -L label_dir nameofhmm
```

nameofhmm is the name of the HMM to initialise (here: yes, no, or sil).

hmmfile is a description file containing the prototype of the HMM called nameofhmm (here: proto/hmm_yes, proto/hmm_no, or proto/hmm_sil).

trainlist.txt gives the complete list of the .mfcc files forming the training corpus (stored in directory data/train/mfcc/).

label_dir is the directory where the label files (.lab) corresponding to the training corpus (here: data/train/lab/).

label indicates which labelled segment must be used within the training corpus (here: yes, no, or sil because have used the same names for the labels and the HMMs, but this is not mandatory...)

model/hmm0 is the name of the directory (must be created before) where the resulting initialised HMM description will be output.

This procedure has to be repeated for each model (hmm_yes, hmm_no, hmm_sil).

Remark:

The HMM file output by HInit has the same name as the input prototype.

HCompv

The HCompv tool performs a “flat” initialisation of a model. Every state of the HMM is given the same mean and variance vectors: these are computed globally on the whole training corpus. The initialisation command line is in this case:

```
HCompv -A -D -T 1 -S trainlist.txt -M model/hmm0flat \  
-H model/proto/hmmfile -f 0.01 nameofhmm
```

nameofhmm , **hmmfile** , **trainlist.txt** : see HInit.

model/hmm0flat : the output directory must be different from the one used with **HInit** (to avoid overwrite).

Remark: the **label** option can also be used. In that case the estimation of the global mean and variance is based on the corresponding labelled parts of the training corpus only.

We won't use **HCompv** to initialise our models (it was already done with **HInit**).

But **HCompv** also output, along with the initialised model, an interesting file called **vFloors**, which contains the global variance vector multiplied by a factor (see List.4). This factor can be set with the **-f** option (here: **0.01**).

```
~v varFloor1
<Variance> 39
5.196781e-001 2.138549e-001 (...) 3.203219e-003
```

List. 4: Variance floors macro file, **vFloors**.

The values stored in **varFloor1** (called the “variance floor macro”) can be used later during the training process as floor values for the estimated variance vectors.

During training iterations, it can happen that the number of training frames associated with a particular HMM state is very low. The estimated variance for that state may then have a very small value (variance is even null if only one training frame is available). The floor values can be used instead in that case, preventing the variance from being too small (and possibly causing computation errors).

Here, we will use **HCompv** only once, with any of our HMM prototype, in order to compute the **varFloor1** macro described above. The corresponding **vFloors** file is output in directory **model/hmm0flat/**.

5.2 Training

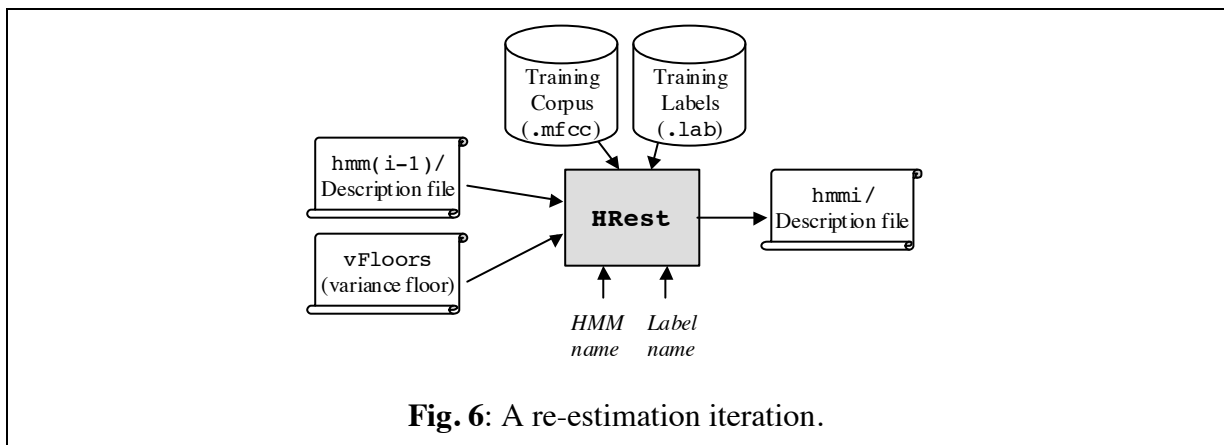


Fig. 6: A re-estimation iteration.

The following command line performs one re-estimation iteration with HTK tool **HRest**, estimating the optimal values for the HMM parameters (transition probabilities, plus mean and variance vectors of each observation function):

```
HRest -A -D -T 1 -S trainlist.txt -M model/hmmi -H vFloors \  
-H model/hmmi-1/hmmfile -l label -L label_dir nameofhmm
```

nameofhmm is the name of the HMM to train (here: `yes`, `no`, or `sil`).

hmmfile is the description file of the HMM called `nameofhmm`. It is stored in a directory whose name indicates the index of the last iteration (here `model/hmmi-1/` for example).

vFloors is the file containing the variance floor macro obtained with HCompv.

trainlist.txt gives the complete list of the `.mfcc` files forming the training corpus (stored in directory `data/train/mfcc/`).

label_dir is the directory where the label files (`.lab`) corresponding to the training corpus (here: `data/train/lab/`).

label indicates the label to use within the training data (`yes`, `no`, or `sil`)

model/hmm*i*, the output directory, indicates the index of the current iteration *i*.

This procedure has to be repeated several times for each of the HMM to train.

Each time, the `HRest` iterations (i.e. iterations within the current re-estimation iteration...) are displayed on screen, indicating the convergence through the change measure. As soon as this measure do not decrease (in absolute value) from one `HRest` iteration to another, it's time to stop the process. In our example, 2 or 3 re-estimation iterations should be enough.

The final word HMMs are then: `hmm3/hmm_yes`, `hmm3/hmm_no`, and `hmm3/hmm_sil`.

6 Task Definition

Every files concerning the task definition should be stored in a dedicated `def/` directory.

6.1 Grammar and Dictionary

Before using our word models, we have to define the basic architecture of our recogniser (the *task grammar*). We will first define the most simple one: a start silence, followed by a single word (in our case “*Yes*” or “*No*”), followed by an end silence.

In HTK, the task grammar is written in a text file, according to some syntactic rules. In our case, the grammar is quite simple:

```
/*
 * Task grammar
 */

$WORD = YES | NO;

( { START_SIL } [ $WORD ] { END_SIL } )
```

List. 5: Basic task grammar.

The `WORD` variable can be replaced by `YES` or `NO`.

The brackets `{ }` around `START_SIL` and `END_SIL` denotes zero or more repetitions (a long silence segment, or no silence at all before or after the word are then allowed).

The brackets `[]` around `$WORD` denotes zero or one occurrence (if no word is pronounced, it's possible to recognise silence only).

For more details on HTK syntactic rules:

see HTK documentation, p.163 (Chap.12, Networks, Dictionaries and Language Models).

The system must of course know to which HMM corresponds each of the grammar variables YES, NO, START_SIL and END_SIL. This information is stored in a text file called the *task dictionary*. In such a simple task, the correspondence is straightforward, and the task dictionary simply encloses the 4 entries:

YES	[yes]	yes
NO	[no]	no
START_SIL	[sil]	sil
END_SIL	[sil]	sil

List. 6: Task dictionary.

The left elements refer to the names of the task grammar variables. The right elements refer to the names of the HMMs (introduced by ~h in the HMM definition files). The bracketed elements in the middle are optional, they indicate the symbols that will be output by the recogniser: the names of the labels are used here (by default, the names of the grammar's variables would have been used.)

Remark:

Don't forget the new line at the end of the file (if not, the last entry is ignored).

6.2 Network

The task grammar (described in file **gram.txt**) have to be compiled with tool HParse, to obtain the *task network* (written in **net.slf**):

```
HParse -A -D -T 1 gram.txt net.slf
```

At this stage, our speech recognition task (Fig.7), completely defined by its **network**, its **dictionary**, and its **HMM set** (the 3 models stored in model/hmm3/), is ready for use.

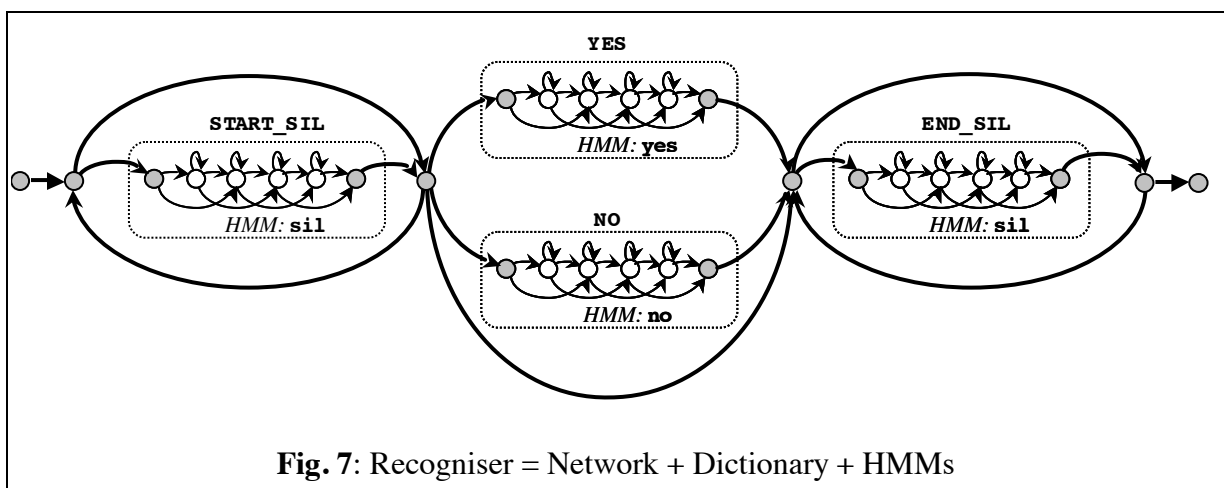


Fig. 7: Recogniser = Network + Dictionary + HMMs

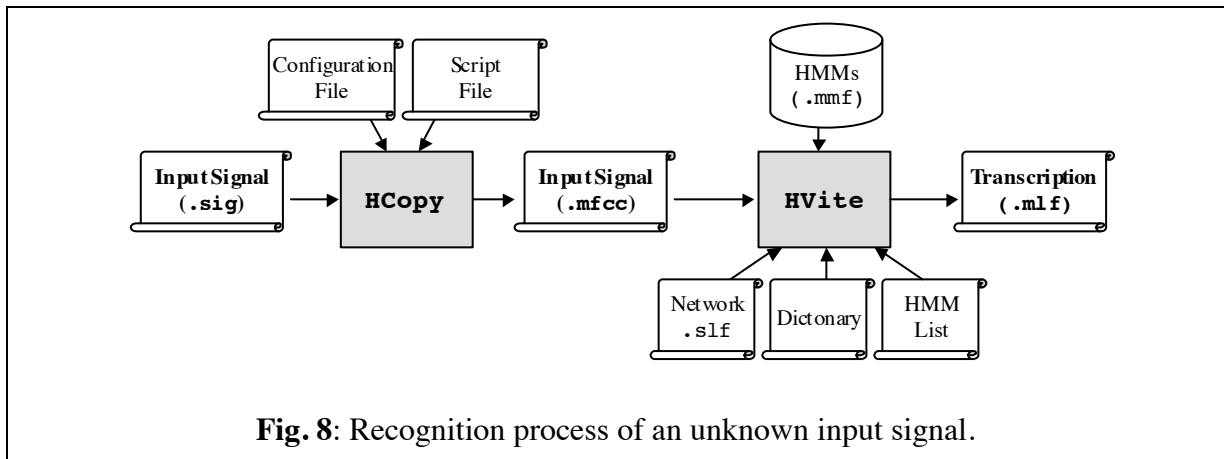
Remark:

To be sure that no mistakes were made when writing the grammar, the tool HSGen can be used to test it:

```
HSGen -A -D -n 10 -s net.slf dict.txt
```

Where **dict.txt** is the task dictionary. Option **-n** indicates that 10 grammatically conform sentences (i.e. 10 possible recognition results) will be randomly generated and displayed. Of course, it's not very useful here, but when grammars get more complicated, that can be very helpful...

7 Recognition



Let's come to the recognition procedure itself:

- An input speech signal `input.sig` is first transformed into a series of “acoustical vectors” (here MFCCs) with tool `HCopy`, in the same way as what was done with the training data (Acoustical Analysis step). The result is stored in an `input.mfcc` file (often called the *acoustical observation*).
- The input observation is then process by a Viterbi algorithm, which matches it against the recogniser's Markov models. This is done by tool `HVite`:

```
HVite -A -D -T 1 -H hmmsdef.mmf -i reco.mlf -w net.slf \  
dict.txt hmmlist.txt input.mfcc
```

input.mfcc is the input data to be recognised.

hmmlist.txt lists the names of the models to use (yes, no, and sil). Each element is separated by a new line character. Don't forget to insert a new line after the last element.

dict.txt is the task dictionary.

net.slf is the task network.

reco.mlf is the output recognition transcription file.

hmmsdef.mmf contains the definition of the HMMs. It is possible to repeat the **-H** option and list our different HMM definition files, in our case:

```
-H model/hmm3/hmm_yes -H model/hmm3/hmm_no -H model/hmm3/hmm_sil
```

but it is more convenient (especially when there are more than 3 models) to gather every definitions in a single file (called a Master Macro File, with extension `.mmf`). Such a file is simply obtained by copying each definition after the other in a single file, without repeating the header information (see List.7).

```

~o <VecSize> 39 <MFCC_0_D_A>
~h "yes"
<BeginHMM>
                (definition...)
<EndHMM>
~h "no"
<BeginHMM>
                (definition...)
<EndHMM>
~h "sil"
<BeginHMM>
                (definition...)
<EndHMM>

```

List. 7: Master Macro File (several HMM description in 1 file).

The output is stored in a file (`reco.mlf`) which contains the transcription of the input. If we use the file `data/train/mfcc/yes01.mfcc` as input data, for instance, we will get in `reco.mlf` such an output:

```

#!MLF!#
"./data/train/mfcc/yes01.rec"
0 4900000 SIL -2394.001465
4900000 12000000 YES -5159.434570
12000000 18300000 SIL -3289.197021
.

```

List. 8: Recognition output (recognised transcription).

In this example, 3 successive “word hypotheses” are recognised within the input signal. The target word “Yes” is correctly recognised. The start and end points of each hypothesis are given, along with their acoustic scores, resulting from the Viterbi decoding algorithm (right column).

A more interactive way of testing the recogniser is to use the “direct input” options of `HVite`:

```

HVite -A -D -T 1 -C directin.conf -g -H hmmsdef.mmf \
-w net.slf dict.txt hmmlist.txt

```

No input signal argument or output file are required in this case: a prompt `READY[1]>` appears on screen, indicating that the first input signal is recorded. The signal recording is stopped by a key-press. The recognition result is then displayed, and a prompt `READY[2]>` waiting for a new input immediately appears.

-g option allows to replay each input signal once.

directin.conf is a configuration file for `HVite`, allowing the use of direct audio input:

```

#
# HVite Configuration Variables for DIRECT AUDIO INPUT
#

# Parameters of the input signal
SOURCERATE = 625.0      # = 16 kHz
SOURCEKIND = HAUDIO
SOURCEFORMAT = HTK

```

```

# Conversion parameters of the input signal
TARGETKIND = MFCC_0_D_A      # Identifier of the coefficients to use
WINDOWSIZE = 250000.0        # = 25 ms = length of a time frame
TARGETRATE = 100000.0        # = 10 ms = frame periodicity
NUMCEPS = 12                  # Number of MFCC coeffs (here from c1 to c12)
USEHAMMING = T                # Use of Hamming function for windowing frames
PREEMCOEF = 0.97              # Pre-emphasis coefficient
NUMCHANS = 26                 # Number of filterbank channels
CEPLIFTER = 22                # Length of cepstral liftering

# Defines the signal to be used for remote control
AUDIOSIG = -1                 # Negative value = key-press control

# The End

```

List. 9: Configuration file for direct input recognition.

In order to allow direct extraction of the acoustical coefficients from the input signal, this file must contain the acoustical analysis configuration parameters previously used with the training data.

8 Performance Test

The recognition performance evaluation of an ASR system must be measured on a corpus of data different from the training corpus. A separate test corpus, with new “Yes” and “No” records, can be created in the `data/test/` directory as it was previously done with the training corpus. Once again, these data (stored in sub-directory `test/sig`) have to be hand-labelled (storage in `test/lab`) and converted (storage in `test/mfcc`).

(If you don’t feel excited by the perspective of a new fastidious recording and labelling session, the training corpus may be used as test corpus here, since this tutorial’s goal is just to learn how to use some HTK tools, not to get relevant performance measures...)

8.1 Master Label Files

Before measuring the performance, we need to create 2 files (called Master Label Files, with extension `.mlf`):

- The first one will contain the “correct” transcriptions of the whole test corpus, that is, the transcriptions obtained by hand-labelling. Let’s call `ref.mlf` these reference transcriptions.
- The second one will contain the recognised transcriptions of the whole test corpus, that is, the hypothesised transcriptions yielded by the recogniser. Let’s call `rec.mlf` these recognised transcriptions.

The performance measures will just result from the comparison between the reference transcription and the recognition hypothesis of each data.

A Master Label File has the following structure:


```

#!MLF!#
"path/data01.ext"
Label1
Label2
.
"path/data02.ext"
Label1
Label2
Label3
Label4
.
(ETC...)

```

List. 10: Master Label File (several transcriptions in 1 file).

Each transcription is introduced by a file name and terminated by a period “.”. A transcription consists of a sequence of labels separated by new-line characters. Optionally, each label can be preceded by start and end time indexes and /or followed by a recognition score (see List.8).

There is no HTK tool to create the reference transcription file `ref.mlf`. It must be written manually or with a script (see the `MakeRefMLF.pl` Perl script that I wrote, for instance). The content of each label file (e.g. `yes01.lab`) has to be copied sequentially in `ref.mlf`, between the line giving the file name (e.g. `*/yes01.lab`): the path can be here replaced by a `*`) and the closing period.

The recognised transcription file `rec.mlf` can be obtained directly with `HVite`. This time `HVite` does not take a single input file name as argument, but the file names (`.mfcc`) of the entire test corpus, listed into a text file:

```

HVite -A -D -T 1 -S testlist.txt -H hmmsdef.mmf -i rec.mlf \
      -w net.slf dict.txt hmmlist.txt

```

hmmlist.txt, **dict.txt**, **net.slf**, **hmmsdef.mmf**: the same as previously.

rec.mlf is the output recognition transcription file.

testlist.txt lists the names of the test files (`data/test/*.mfcc`).

After execution of the command, `rec.mlf` contains a series of transcription such as the one listed in List.8. Each transcription is introduced by the corresponding file name with a different extension (`.rec` instead of `.lab`).

8.2 Error Rates

The `ref.mlf` and `rec.mlf` transcriptions are compared with the HTK performance evaluation tool, `HResults`:

```

HResults -A -D -T 1 -e ??? sil -I ref.mlf \
         labellist.txt rec.mlf > results.txt

```

results.txt contains the output performance statistics (example: List.11).

rec.mlf contains the transcriptions of the test data, as output by the recogniser.

labellist.txt lists the labels appearing in the transcription files (here: `yes`, `no`, and `sil`).

ref.mlf contains the reference transcriptions of the test data (obtained by hand-labelling).

-e ??? sil option indicates that the **sil** labels will be ignored when computing the performance statistics (since we are interested in the recognition rate of words “*Yes*” and “*No*” only).

```
===== HTK Results Analysis =====  
Date: Tue Dec 03 19:12:58 2002  
Ref : .\ref.mlf  
Rec : .\rec.mlf  
----- Overall Results -----  
SENT: %Correct=80.00 [H=8, S=2, N=10]  
WORD: %Corr=80.00, Acc=80.00 [H=8, D=0, S=2, I=0, N=10]  
=====
```

List. 11: Results of a performance test.

List.11 shows an example of the kind of results that can be obtained. The first line (SENT) gives the sentence recognition rate (%Correct=80.00), the second one (WORD) gives the word recognition rate (%Corr=80.00).

In our case, the 2 rates are the same because our task grammar only allows “sentences” with one single word (apart from silences). It is an *isolated words* recognition task. Only the first line (SENT) should be considered here. H=8 gives the number of test data correctly recognised, S=2 the number of substitution errors (a “*Yes*” recognised as “*No*”, or a “*No*” as “*Yes*”) and N=10 the total number of test data.

The statistics given on the second line (WORD) only make sense with more sophisticated types of recognition systems (e.g. *connected words* recognition tasks). For more details: see HTK documentation, p.232 (Reference Section, HResults).