

[COM4513-6513] Lab 4: Viterbi and Beam search

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The goal of this lab session is to accelerate the structured perceptron using two methods, Viterbi and Beam search. You should use the same data as in Lab 3.

- [2 marks] Implement the **Viterbi algorithm** to accelerate the **argmax** operation during both training and testing. Recall that the Viterbi is an **exact search algorithm**, i.e. it should return exactly the same result as the naive implementation, only faster. To ensure this happens, pay attention to **random seeds**, use of (ordered) dictionaries, etc. Here is an adaptation of Viterbi for the structured perceptron:

: word sequence $\mathbf{x} = [x_1, \dots, x_N]$, weights \mathbf{w}

set matrix $V^{|\mathcal{Y}| \times N} = 0$ set backpointer

$$B^{|\mathcal{Y}| \times N} = 0$$

$$V[y, n] = \max_{y' \in \mathcal{Y}} V[y', n-1] + \mathbf{w} \cdot \phi(y, y', \mathbf{x})$$

$$B[y, n] = \operatorname{argmax}_{y' \in \mathcal{Y}} V[y', n-1] + \mathbf{w} \cdot \phi(y, y', \mathbf{x})$$

- [2 marks] What speed up do you get with Viterbi compared to the standard structured perceptron?
- [2 marks] Implement **beam search** to accelerate the argmax. Recall that it is an inexact search method so the **results might differ**. But given a large enough beam size it should achieve the same results as viterbi and the structured perceptron from Lab 3.
- [2 marks] Does beam search affect your accuracy? What is the effect of beam size on speed and accuracy compared to the standard structured perceptron and Viterbi? **Tip 1:** Try **three different beam sizes** and report the results. **Tip 2:** Use a **table** to summarise the results (accuracy and training time) for the standard perceptron (from the lab 3), Viterbi and Beam search.

Submission Details

You should submit a python file (lab4.py) that can be executed as:

- `python3 lab4.py -v train.txt test.txt` for Viterbi
- `python3 lab4.py -b train.txt test.txt` for Beam search

and **returns the micro F1 for the phi_1 feature set from Lab 3**. Use python's **random** library to fix the **random seed** so that your results are reproducible! Make sure your code is **Python 3** compatible. You can use **argparse** for the command line arguments **-v** and **-b**.

You also need to accompany the code with a lab4.pdf (no more than two A4 pages) answering the questions above.

This lab will be marked out of 8. It is worth 8% of your final grade in the module.

The deadline for this assignment is the beginning of the Week 9 lab and it needs to be submitted via MOLE. Standard departmental penalties for lateness will be applied.