


Lab10 Sequence to Sequence

Due Dec 2, 2018 by 11:59pm **Points** 10 **Submitting** a file upload

Download [lab10.zip \(https://canvas.ust.hk/courses/19691/files/1800542/download?wrap=1\)](https://canvas.ust.hk/courses/19691/files/1800542/download?wrap=1).

Follow the instruction file [lab10_seq2seq.pdf \(https://canvas.ust.hk/courses/19691/files/1806131/download?wrap=1\)](https://canvas.ust.hk/courses/19691/files/1806131/download?wrap=1)  [\(https://canvas.ust.hk/courses/19691/files/1806131/download?wrap=1\)](https://canvas.ust.hk/courses/19691/files/1806131/download?wrap=1)

Preview the document and complete lab10_skeleton.py.

Copy outputs (20 epochs) to a file (translation.txt, dialogue.txt) for each dataset.

Submit the python script (i.e., lab10_skeleton.py), and the two output files.