1) We have a lot of predictive power, as humans, when it comes to language. Suppose you're asked to guess the next letter in the sequence:

WA_

Likely you don't have a uniform probability over letters. Letters like S or N are quite likely (in English), but Q or E are pretty unlikely (this, despite the fact that E is generally a common letter).

2) More context can sharpen the probability:

GEORGE WA _ BEES WA _ WHAT _ DO _ YOU WA _

3) A language model over an alphabet A is a probability distribution over letter and A given the previous letters in a sequence a, az, ..., an

Pr(an+1 | a, az, ,, an)

4 Language models are frequently used in natural (anguage processing to determine how "Fluent" a sequence sounds (relative to a particular language). For instance, a good tanguage model (LM) for English should have

Pr("N" | "W", "A") > Pr("E" | "W", "A")

5) A sequence's probability can be scored as:

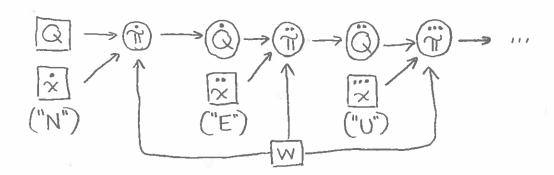
Pr(a,, ..., a_k) = TT Pr(a_k | a_1, ..., a_{k-1}) A good LM for English should have Pr ("THERE IS A") >> Pr ("IHRA S EET")

6) RNNs provide an effective way to create LMs.

The main difference between RNN LMs and the RNNs We've seen so far is that RNN LMs don't just take an input character at each time step, they also emit an output character.

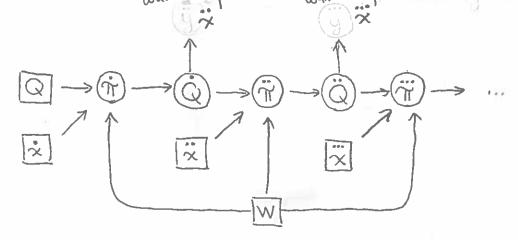
RNN LANGUAGE MODELS

3) Consider our standard RNN:



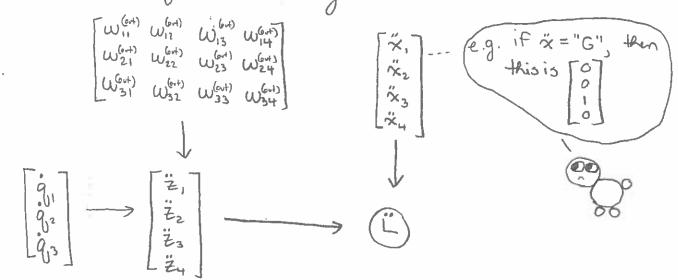
As it reads in letters ("NEU..."), it updates the state vector Q.

18) To make this into a language model, we will try to predict the next letter from each state vector:



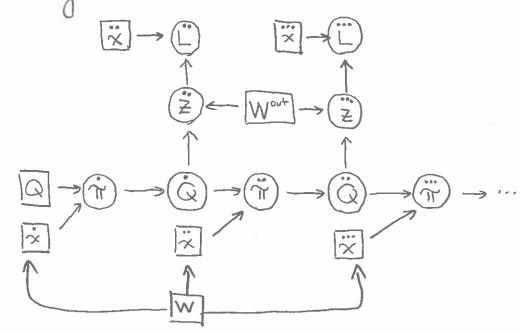
To show some simple concrete examples, let's assume we're learning a language model over DNA sequences (so our alphabet is $\{3A,C,G,T\}$).

To predict the next letter from the current state vector Q, we can use a simple danse layer:



where! L(z, x)=-log(xT·softmax(z)) z=WTg

10) Putting this into our RNN:

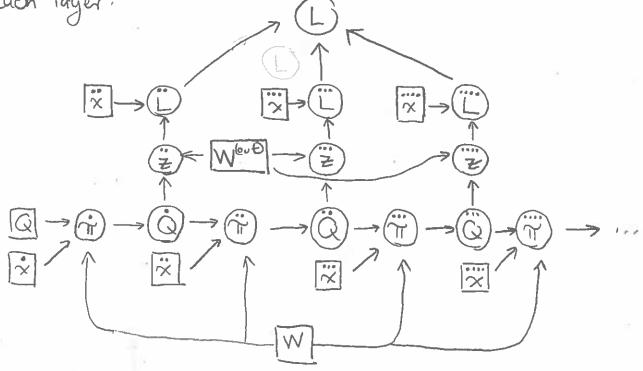


1) It's mildly awkward that our loss is distributed across each layer, but it's easy to reconcile using probability and logs. Essentially, we want to maximize the predicted probability of the actual sequence:

This is equivalent to maximizing the log probability

Or alternatively, minimizing the negative lag probability:

12) So the overall loss is just the sum of the losses at each layer:



13) From this depiction, it is straightforward (though arducus by hand) to compute 2L and 2L DW (CM)

for each sequence x, x, x, in our training data

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