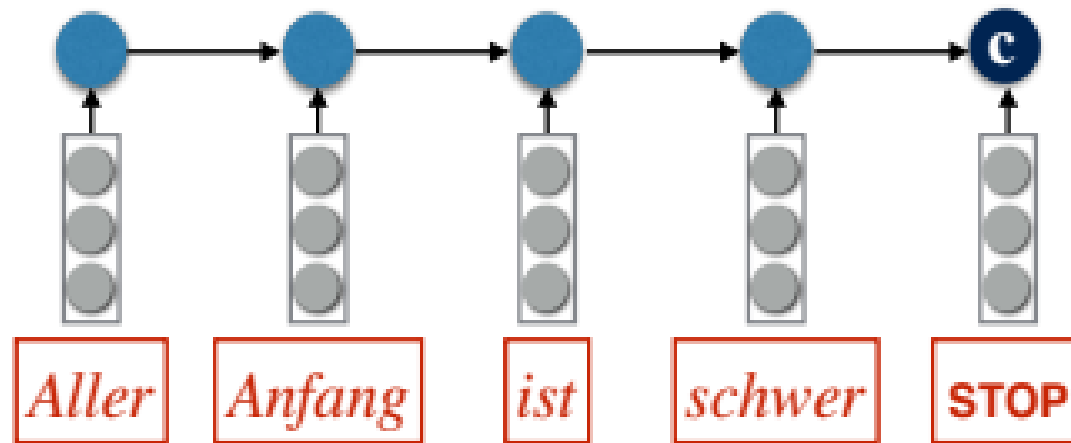


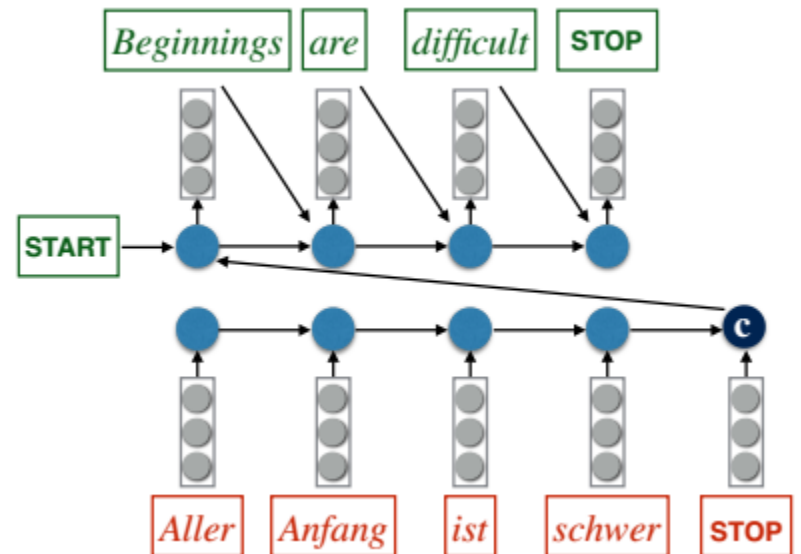
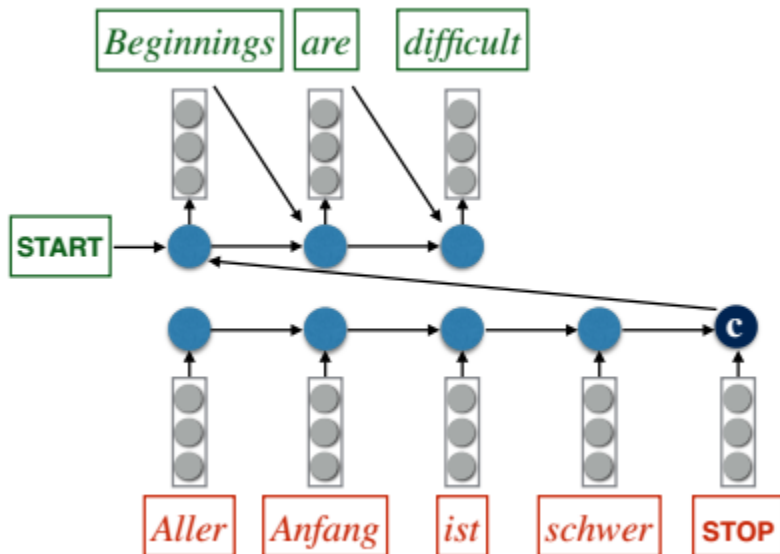
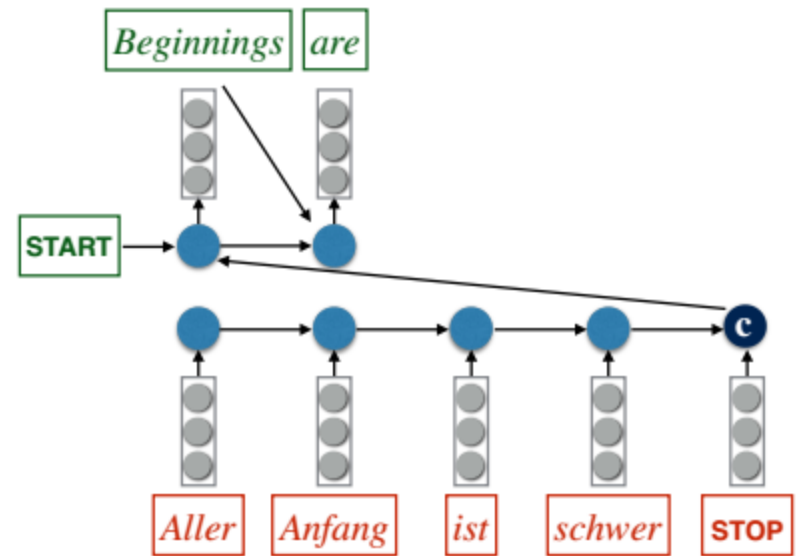
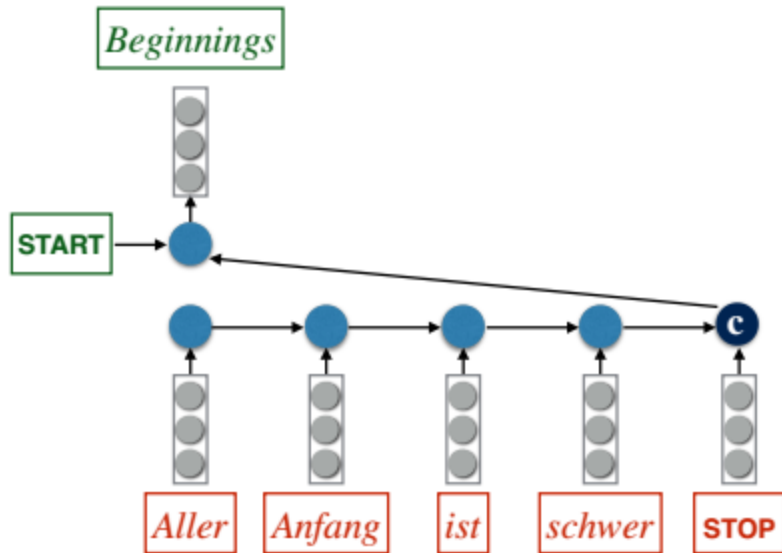
Machine translation models

Seq2Seq model1: Embed, Encode

Sentence encoder: fixed size vector



Decode from encoded vector



Pros & Cons

- **Good**

- RNNs deal naturally with sequences of various lengths
- LSTMs in principle can propagate gradients a long distance
- Very simple architecture!

- **Bad**

- The hidden state has to remember a lot of information!

Seq2Seq model2: Embed, Encode & Attention

Sentence encoding as fixed size vector

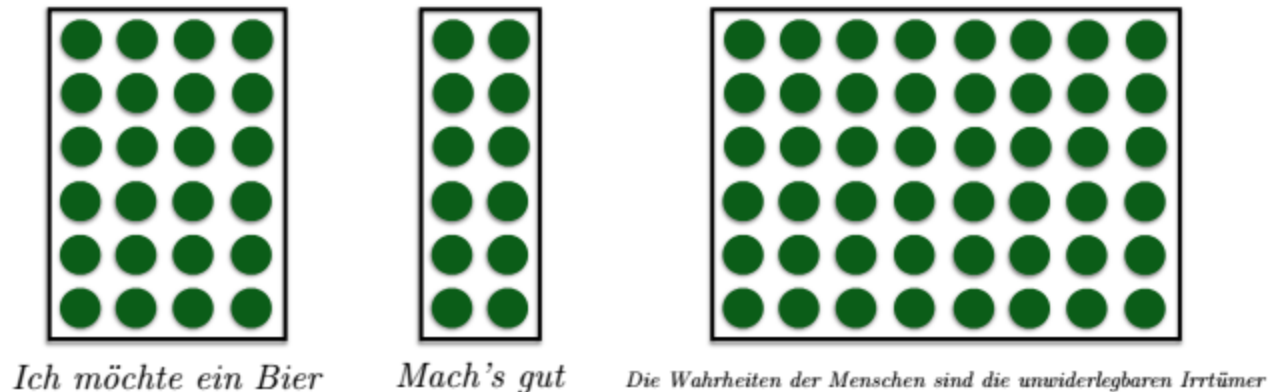
We are compressing a lot of information in a finite-sized vector.

Gradients have a long way to travel. Even LSTMs forget!

What is to be done?

Sentence encoding as matrix

- Represent a source sentence as a matrix
- Generate a target sentence from a matrix



*Each matrix has fixed number of rows, but number of columns depends on the number of words

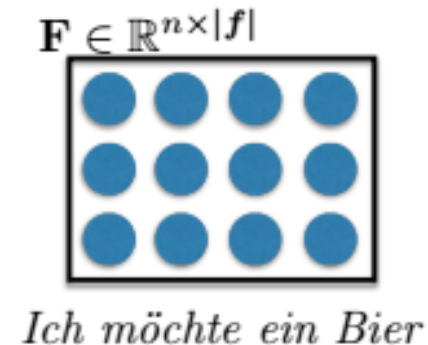
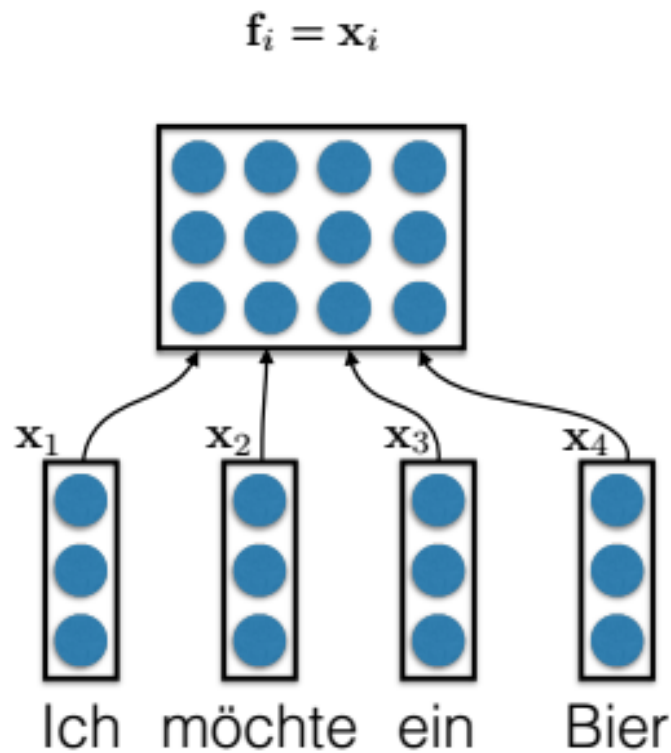
Sentence encoding as matrix

- I. Using word vectors alone
- II. Using CNN
- III. Using RNN/Bi-directional RNN

I. Sentence matrix with word vectors alone

- Each word type is represented by an n -dimensional vector
- Take all of the vectors for the sentence and concatenate them into a matrix

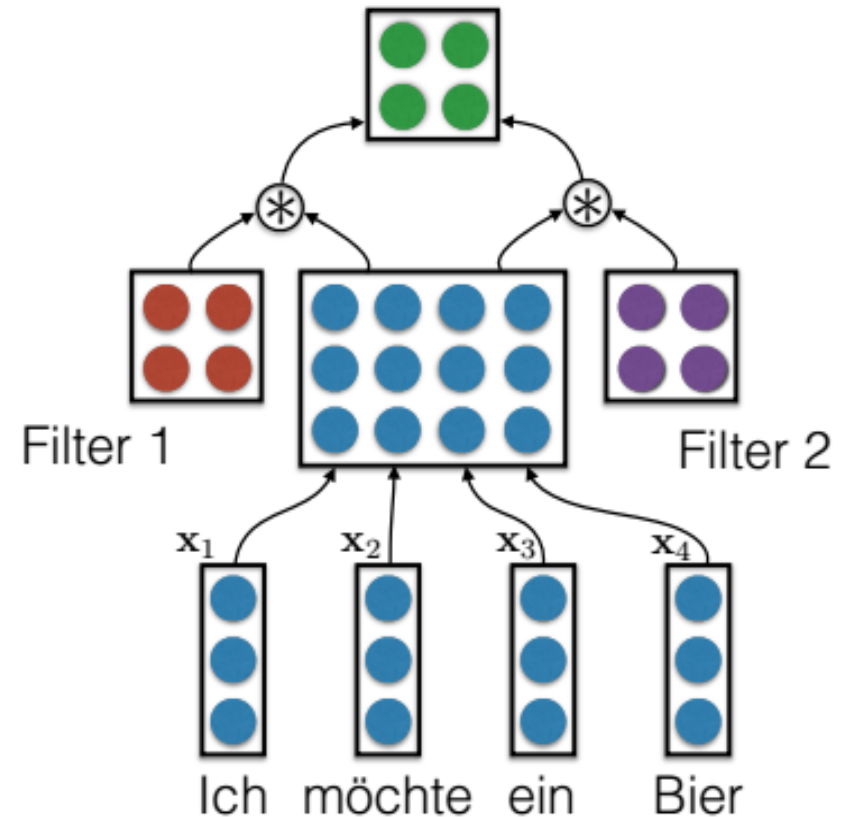
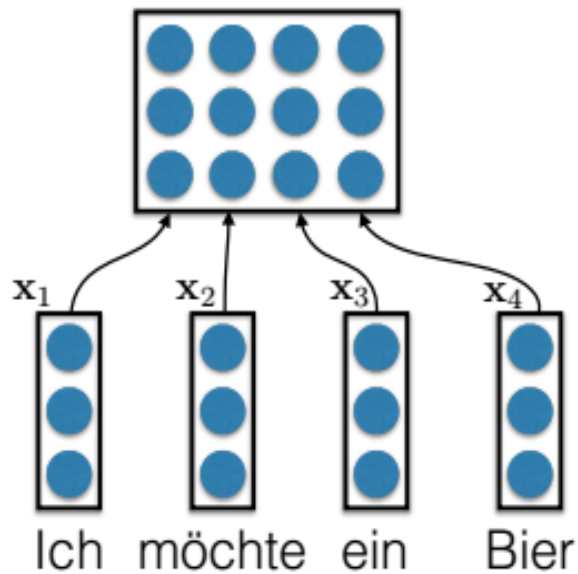
I. Sentence matrix with word vectors alone



II. Sentence matrix with CNN

- Apply convolutional networks to transform the naïve concatenated matrix to obtain a context-dependent matrix

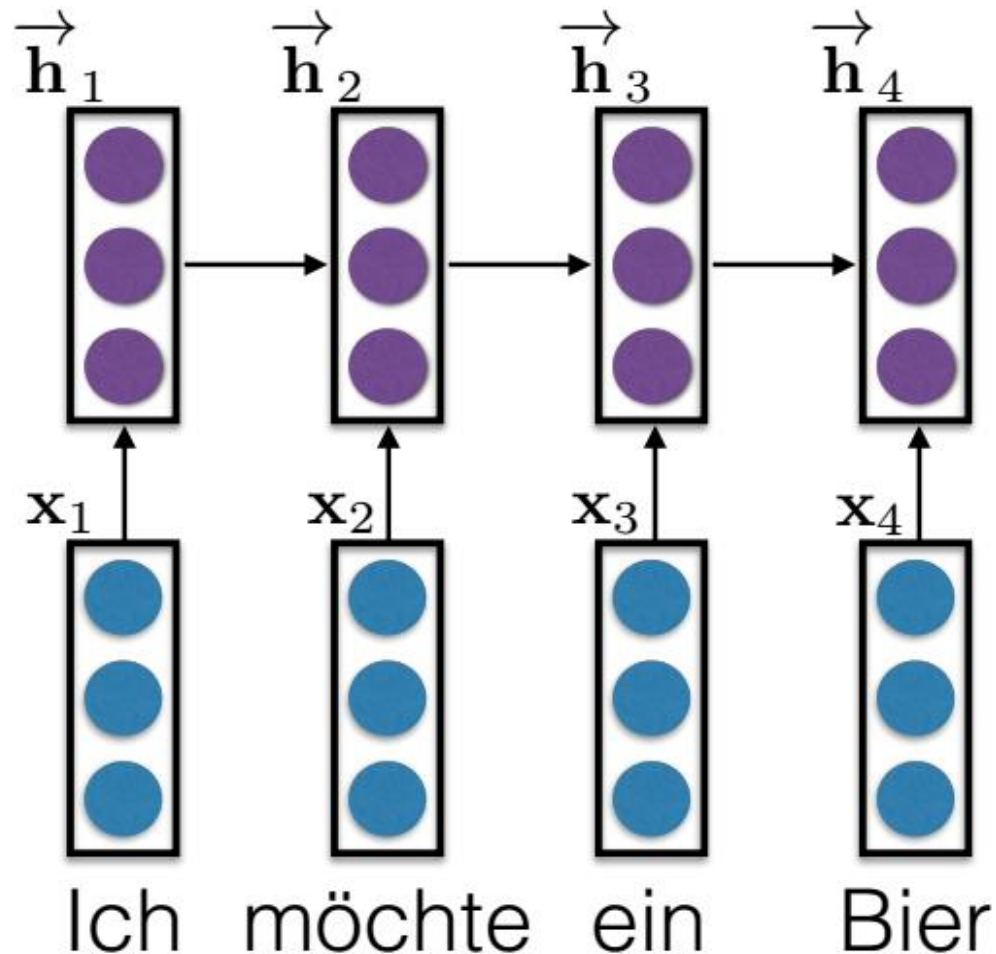
II. Sentence matrix with CNN

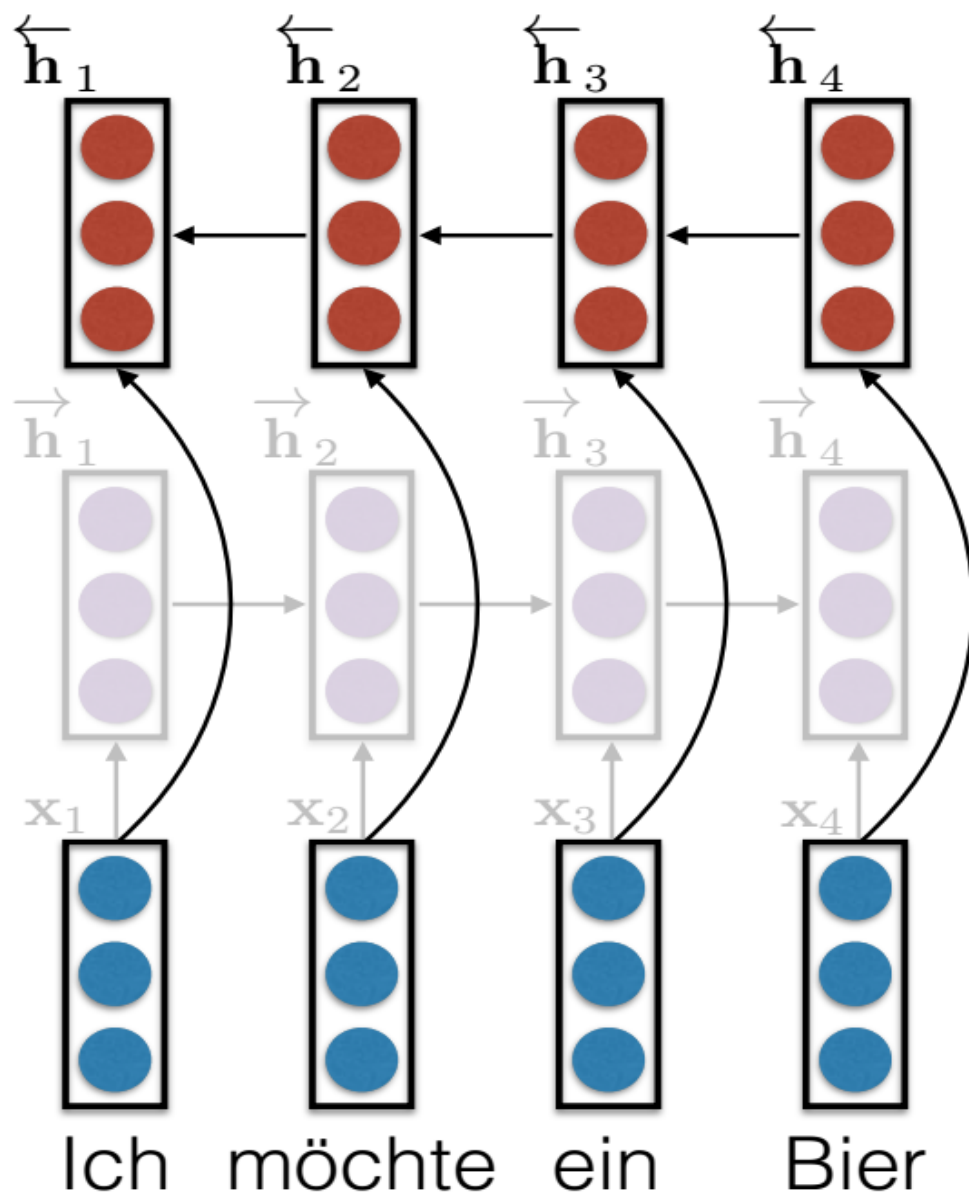


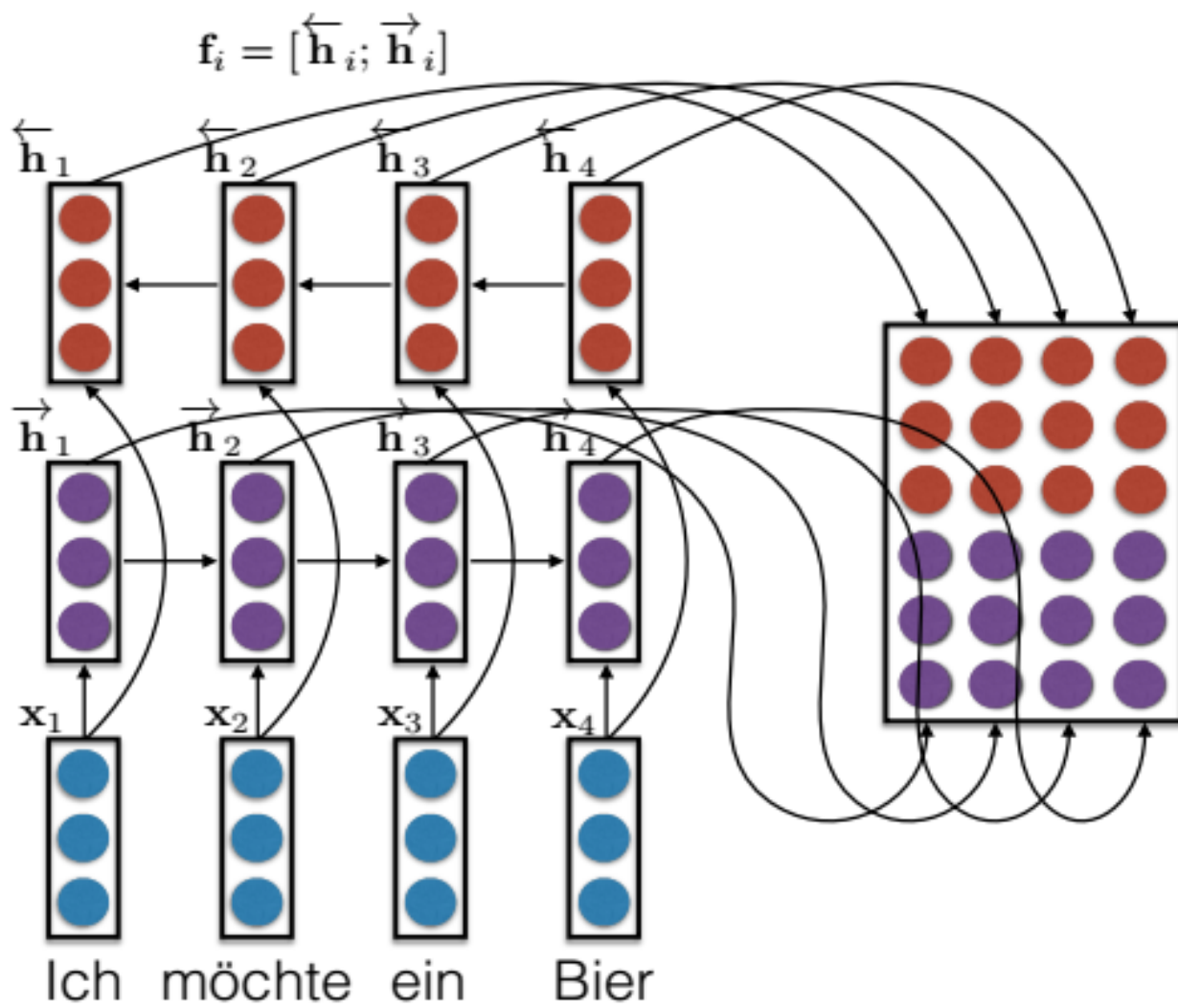
III. Sentence matrix with Bi-RNN

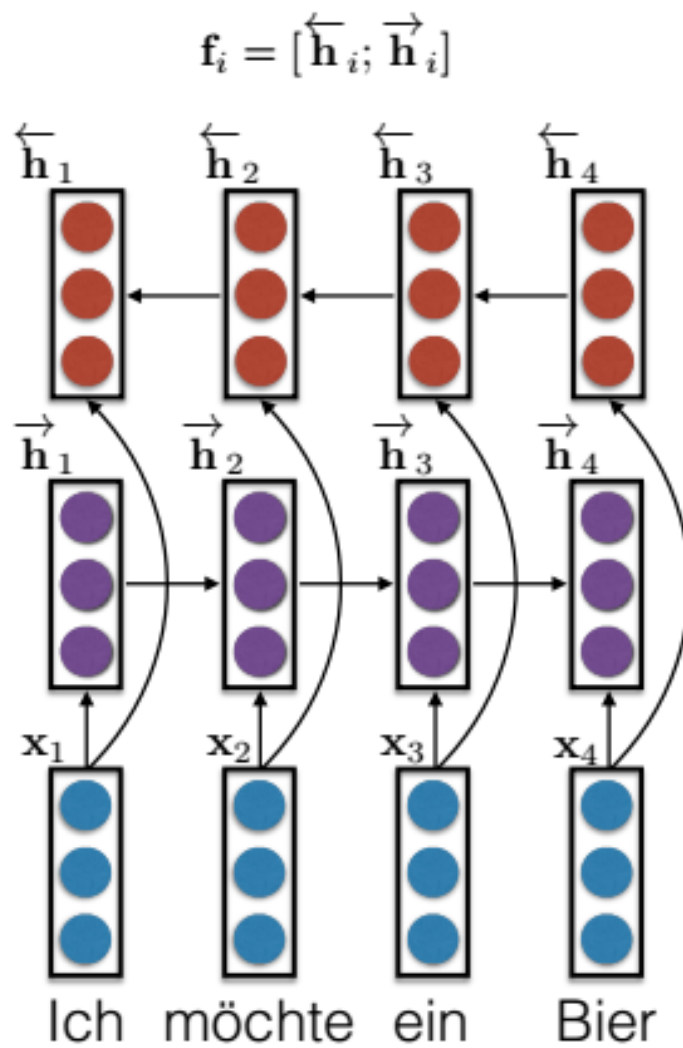
- By far the most widely used matrix representation, due to Bahdanau et al (2015)
- One column per word
- Each column (word) has two halves concatenated together:
 - a “forward representation”, i.e., a word and its left context
 - a “reverse representation”, i.e., a word and its right context
- Implementation: bidirectional RNNs (GRUs or LSTMs) to read from left to right and right to left, concatenate representations

III. Sentence matrix with Bi-RNN

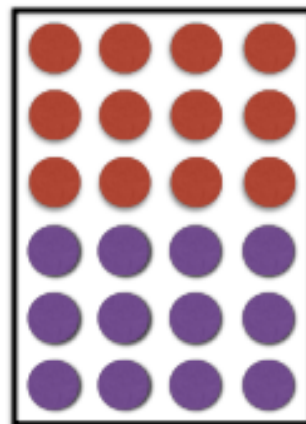






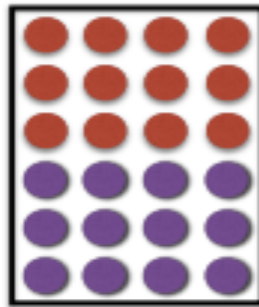
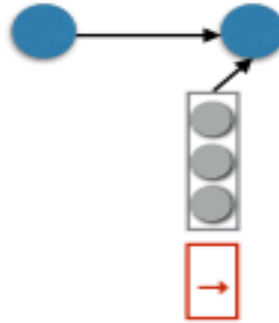


$$\mathbf{F} \in \mathbb{R}^{2n \times |f|}$$

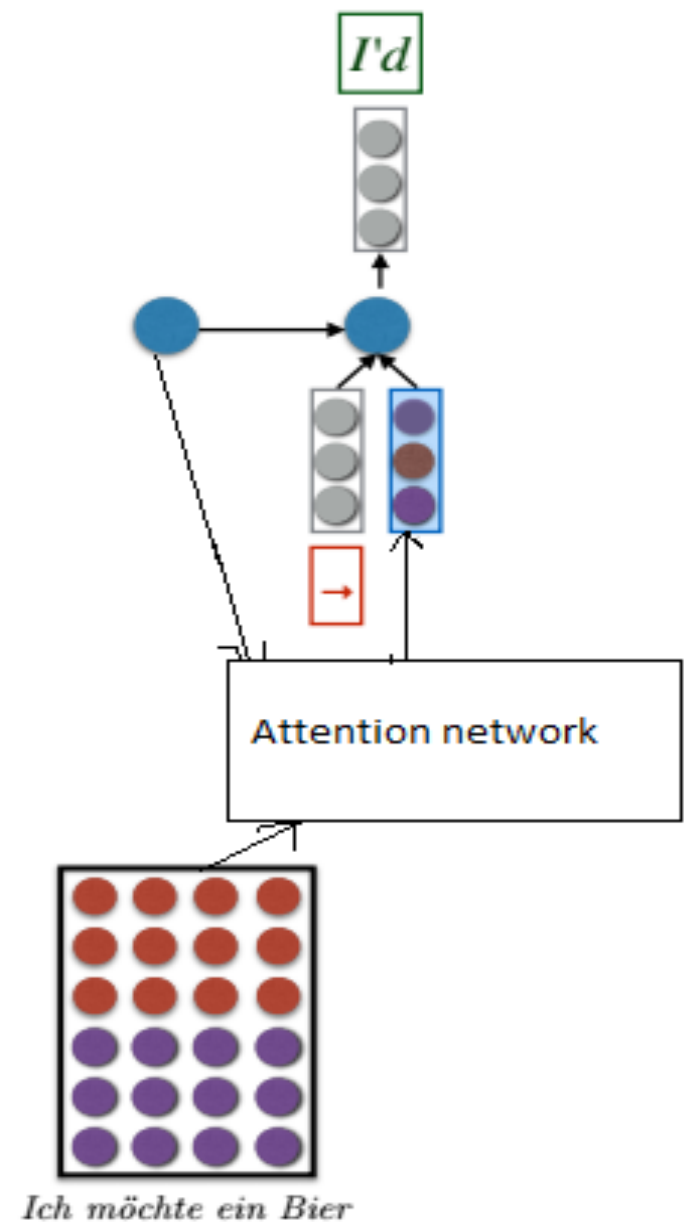
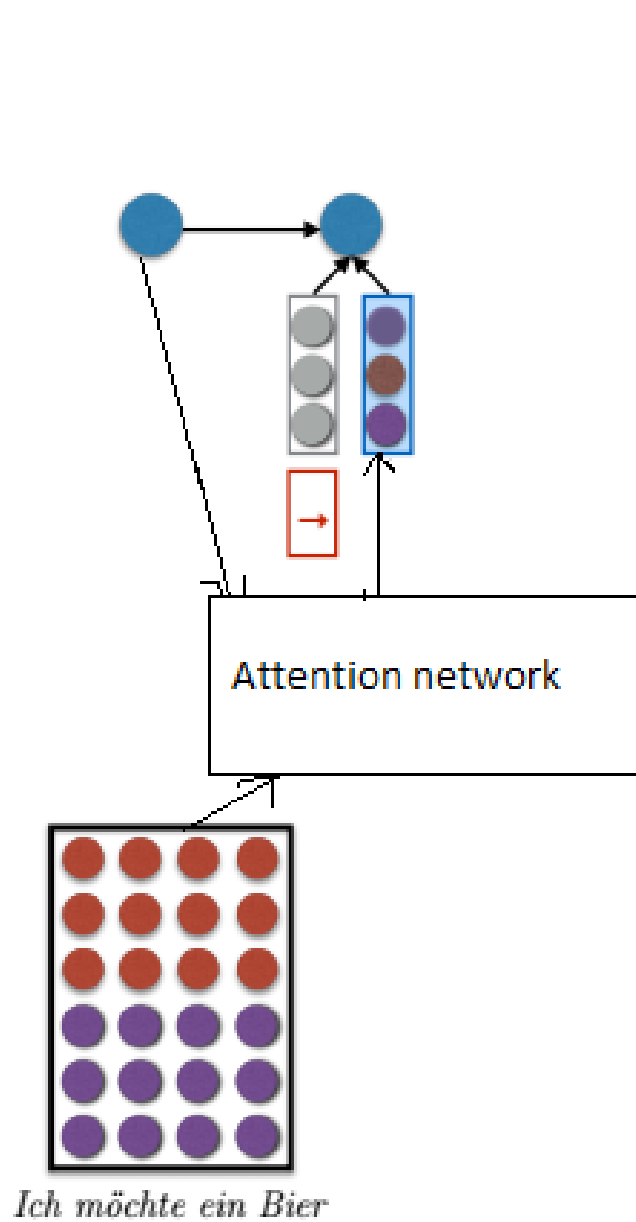


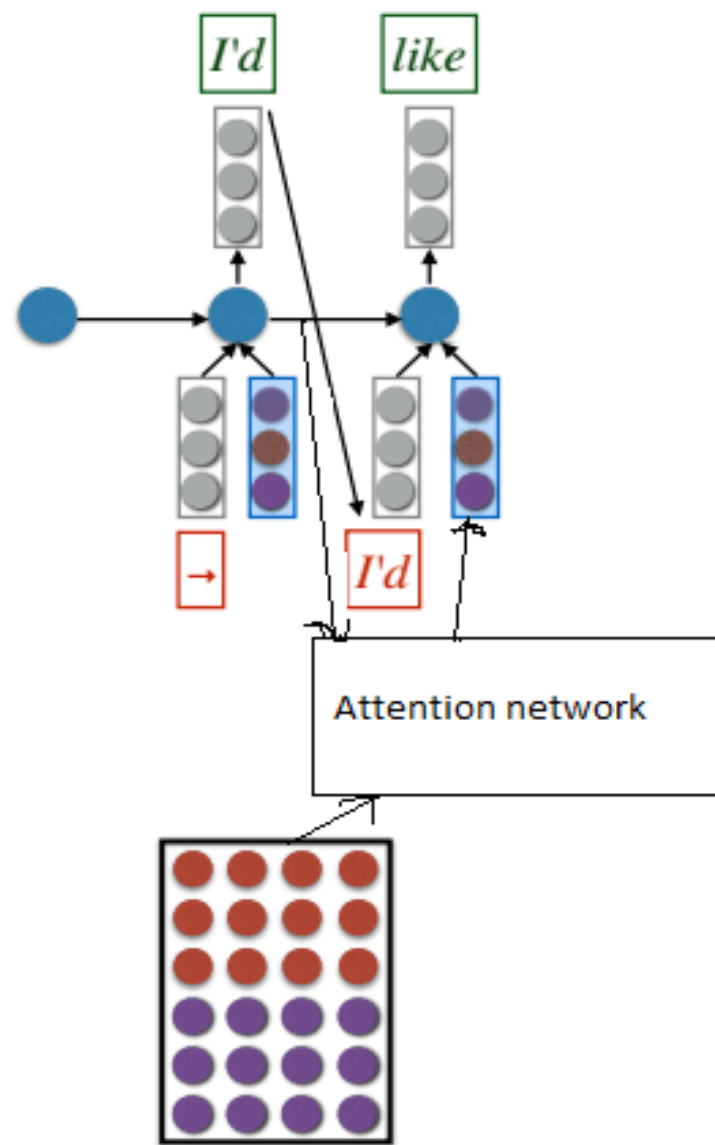
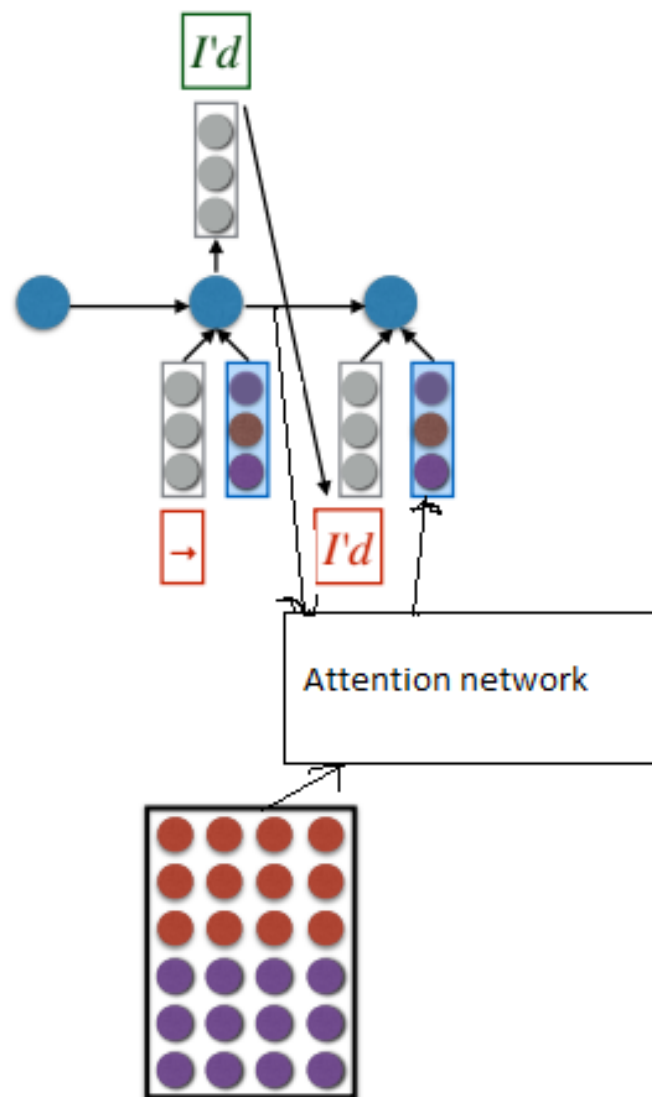
Ich möchte ein Bier

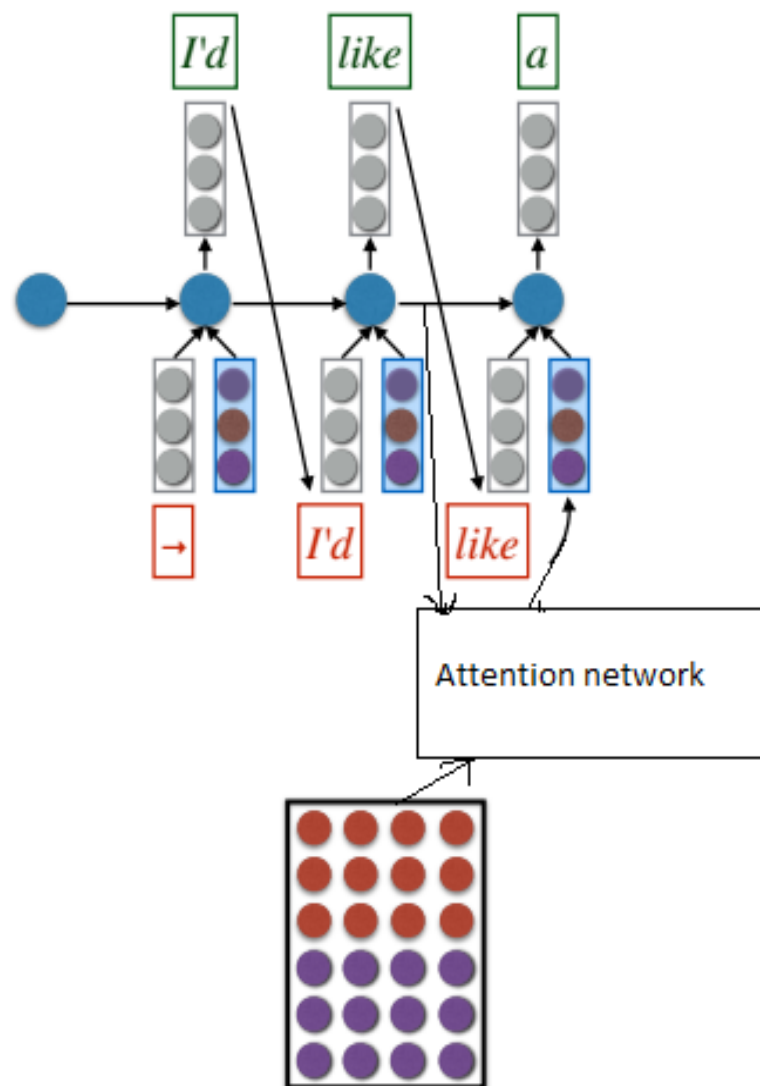
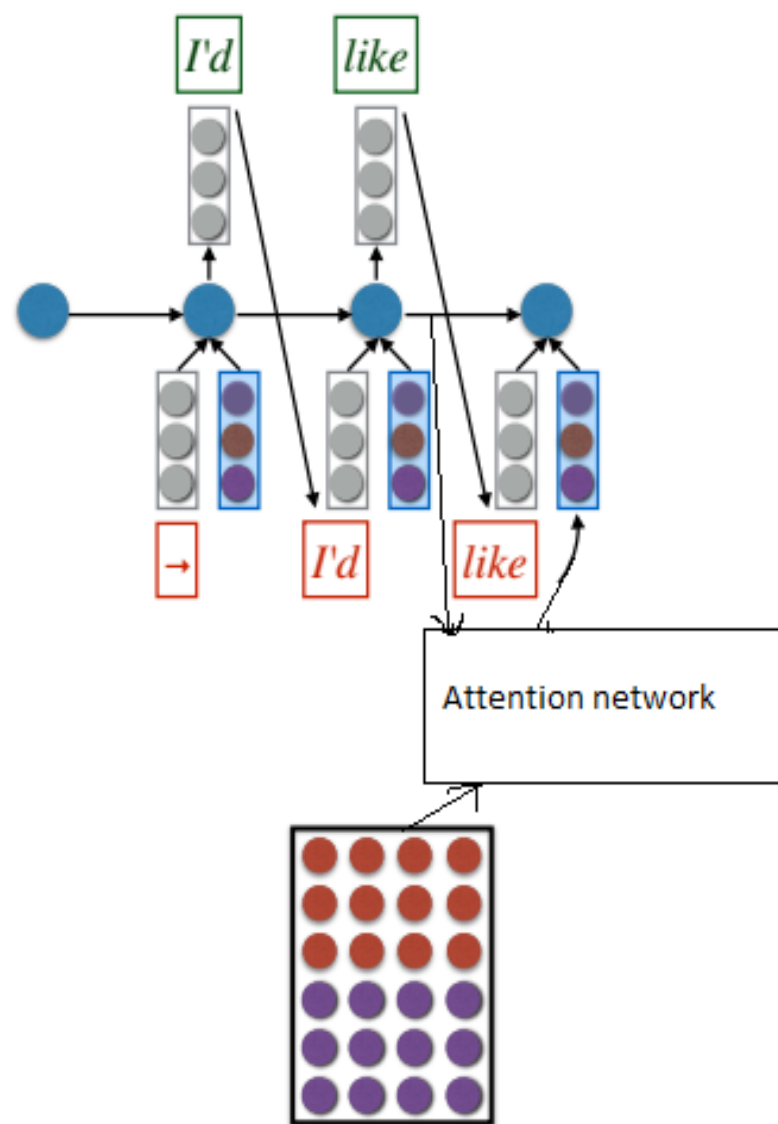
Decoder with Attention

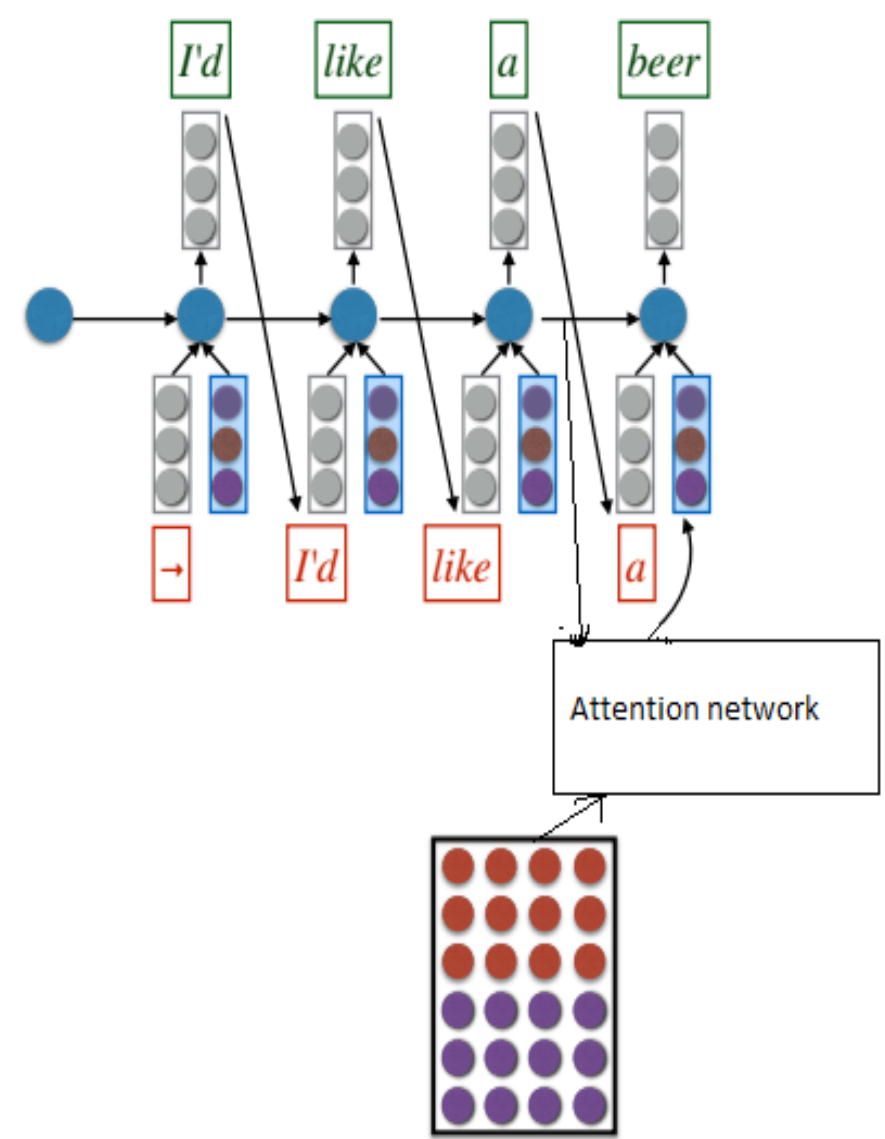
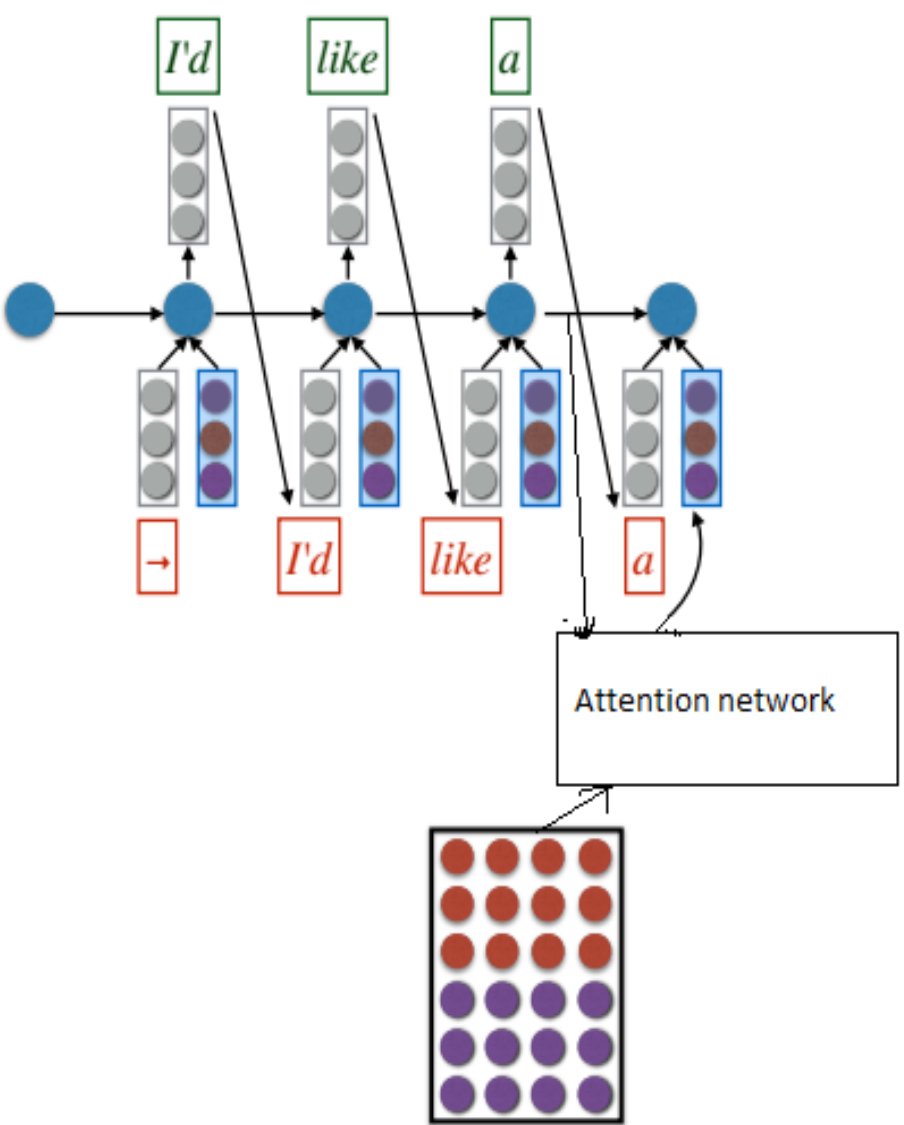


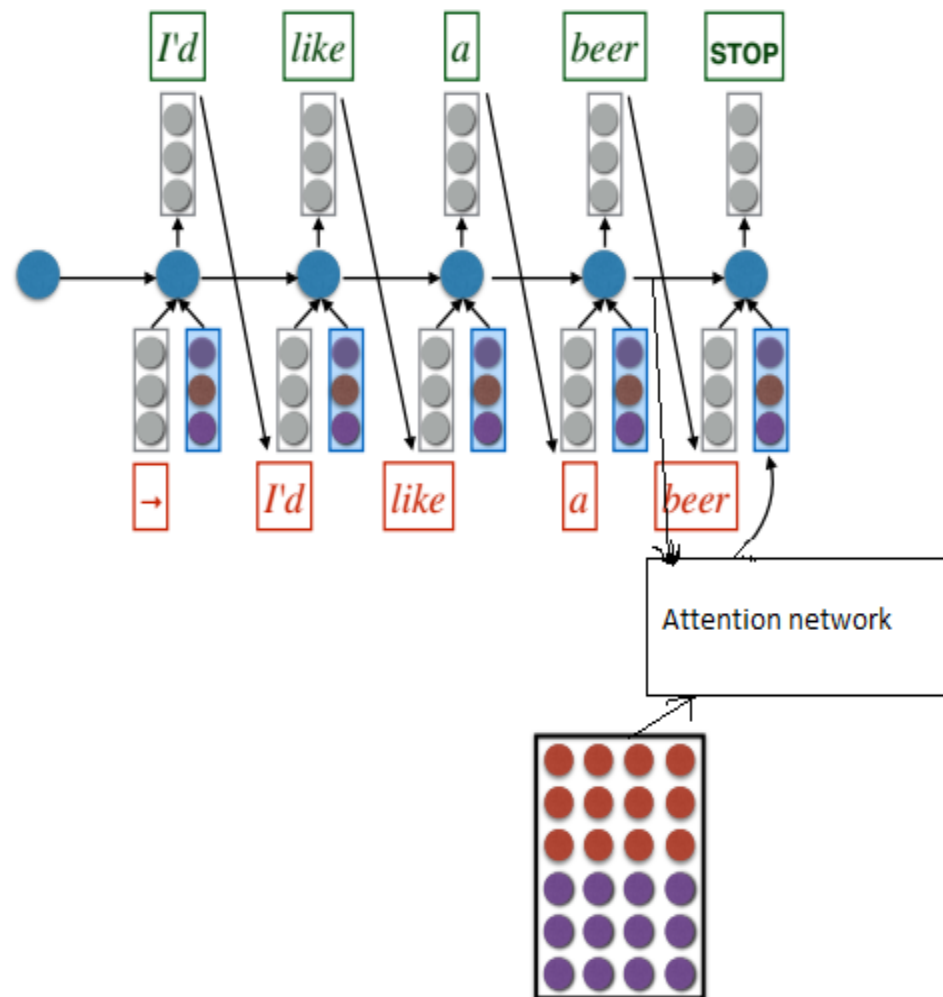
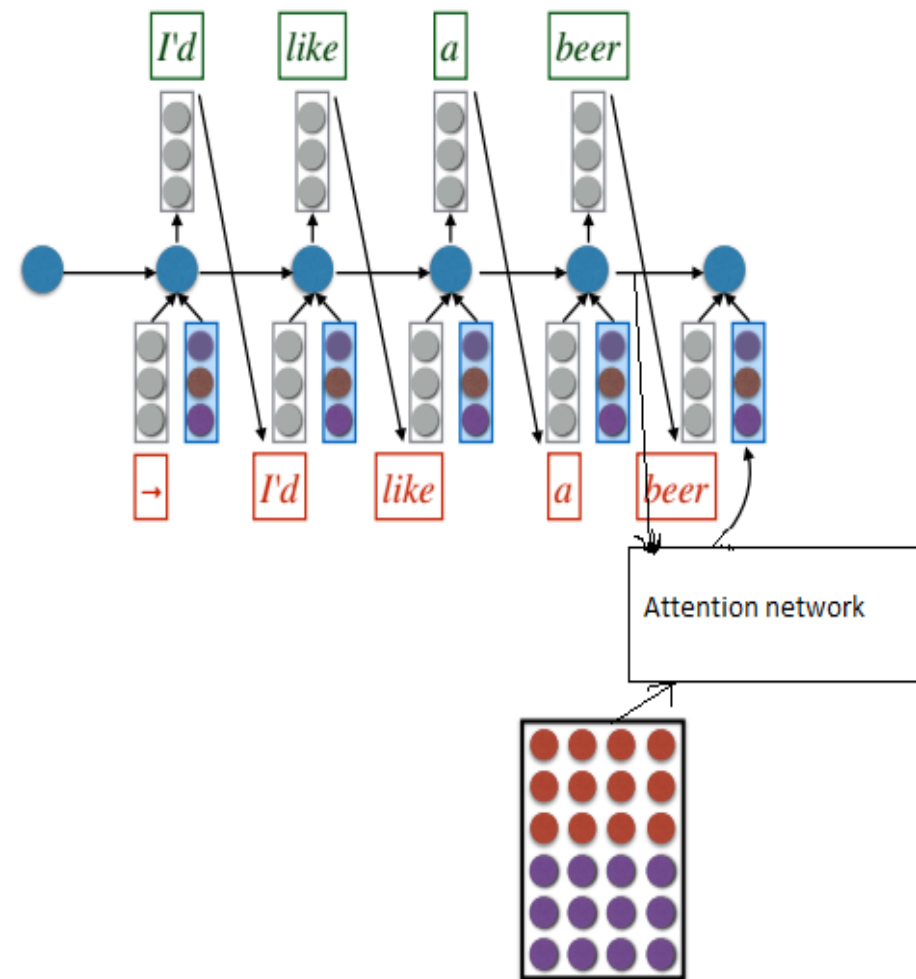
Ich möchte ein Bier











Decoding with seq2seq model

Decoding: Greedy search

Simply take the best prediction of the previous timestep as the input to the current timestep.

$$w_1^* = \arg \max_{w_1} p(w_1 \mid \mathbf{x})$$

$$w_2^* = \arg \max_{w_2} p(w_2 \mid \mathbf{x}, w_1^*)$$

$$\vdots$$

$$w_t^* = \arg \max_{w_t} p(w_t \mid \mathbf{x}, w_{<t}^*)$$

Decoding: Greedy search

- The problem with greedy search, as its name suggested, is that it may not be optimal. More concretely, the joint probability of the output sequence may not be highest possible.
- To do better than greedy search, one can do a full search. We use the recurrent networks to compute the joint probability of every possible output sequence. This can guarantee to find the best possible sequence with a large computation cost.

Decoding: Full search

In general, we want to find the most probable (MAP) output given the input, i.e.

$$\begin{aligned} \mathbf{w}^* &= \arg \max_{\mathbf{w}} p(\mathbf{w} \mid \mathbf{x}) \\ &= \arg \max_{\mathbf{w}} \sum_{t=1}^{|\mathbf{w}|} \log p(w_t \mid \mathbf{x}, \mathbf{w}_{<t}) \end{aligned}$$

This is, for general RNNs, a hard problem.

Decoding: Beam search

- Keep a list of b possible sequences sorted by the joint probability (which is computed by multiplying the output probability of each prediction in the sequence produced so far).
- During the decode procedure, any sequence that does not belong to the top- b highest joint probability will be removed from our list of candidates.
- Even though it's not guaranteed for the beam search to achieve the optimal sequence, in practice, the generated sequences in the top- b list are generally very good.

Decoding: Beam search

A slightly better approximation is to use a **beam search** with beam size b . Key idea: keep track of top b hypothesis.

E.g., for $b=2$:

$\mathbf{x} = \textit{Bier trinke ich}$
 beer drink I

$\langle s \rangle$ logprob=0

w_0

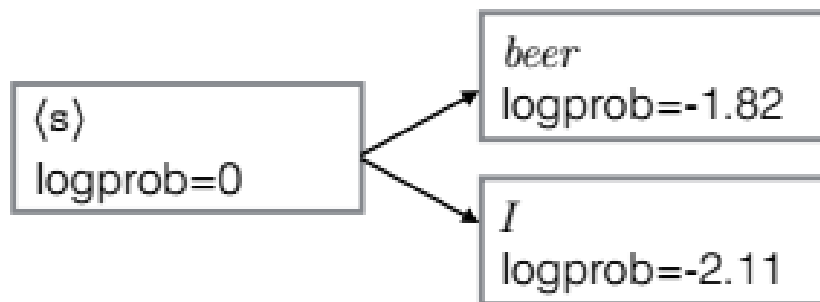
w_1

w_2

w_3

E.g., for $b=2$:

$\mathbf{x} = \textit{Bier trinke ich}$
beer drink I



w_0

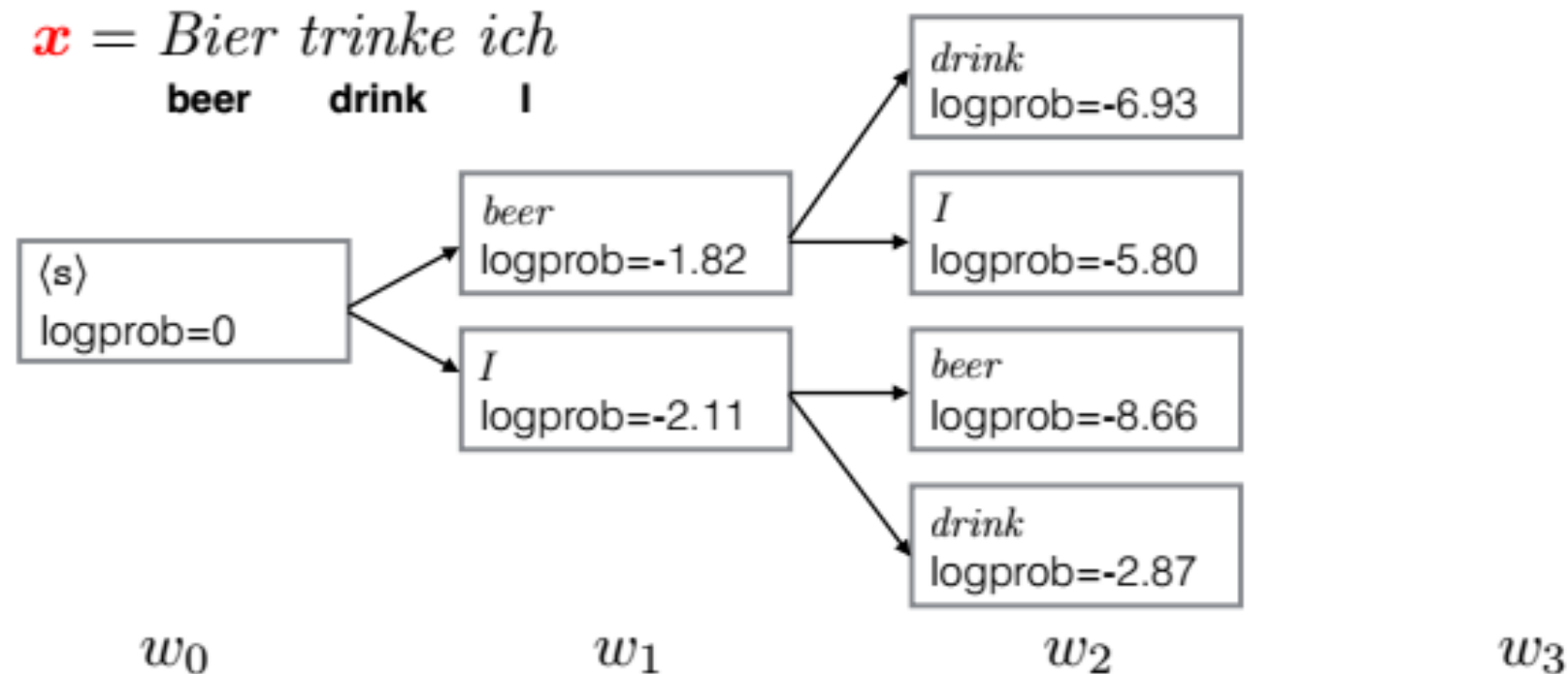
w_1

w_2

w_3

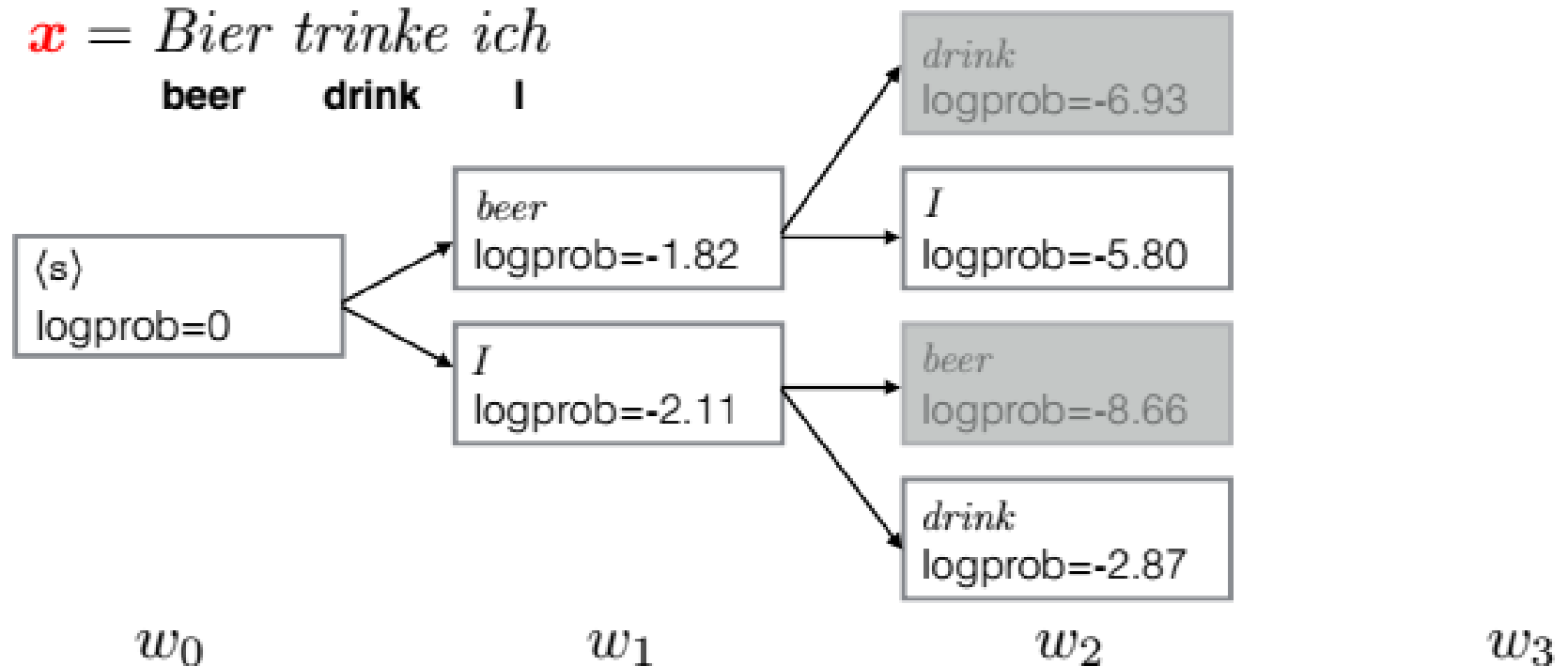
E.g., for $b=2$:

$\mathbf{x} = \textit{Bier trinke ich}$
beer drink I



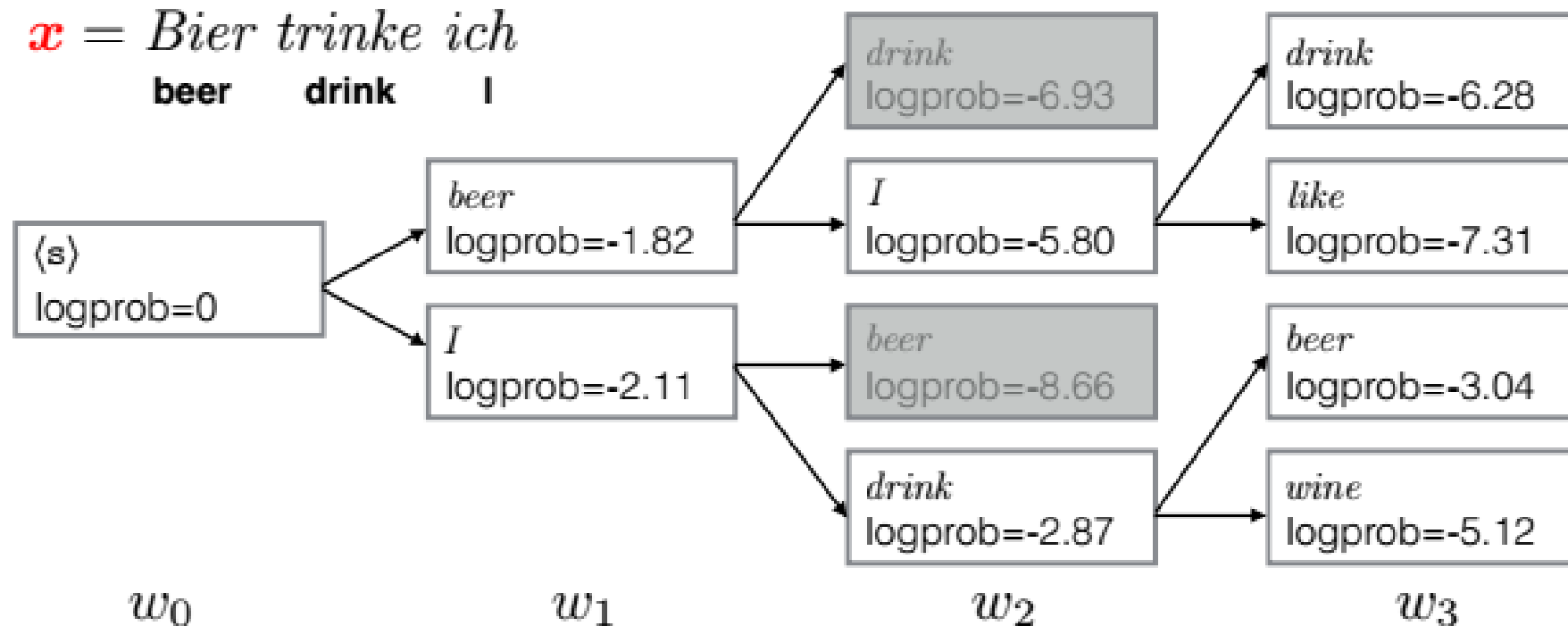
E.g., for $b=2$:

$\mathbf{x} = \textit{Bier trinke ich}$
beer **drink** **I**



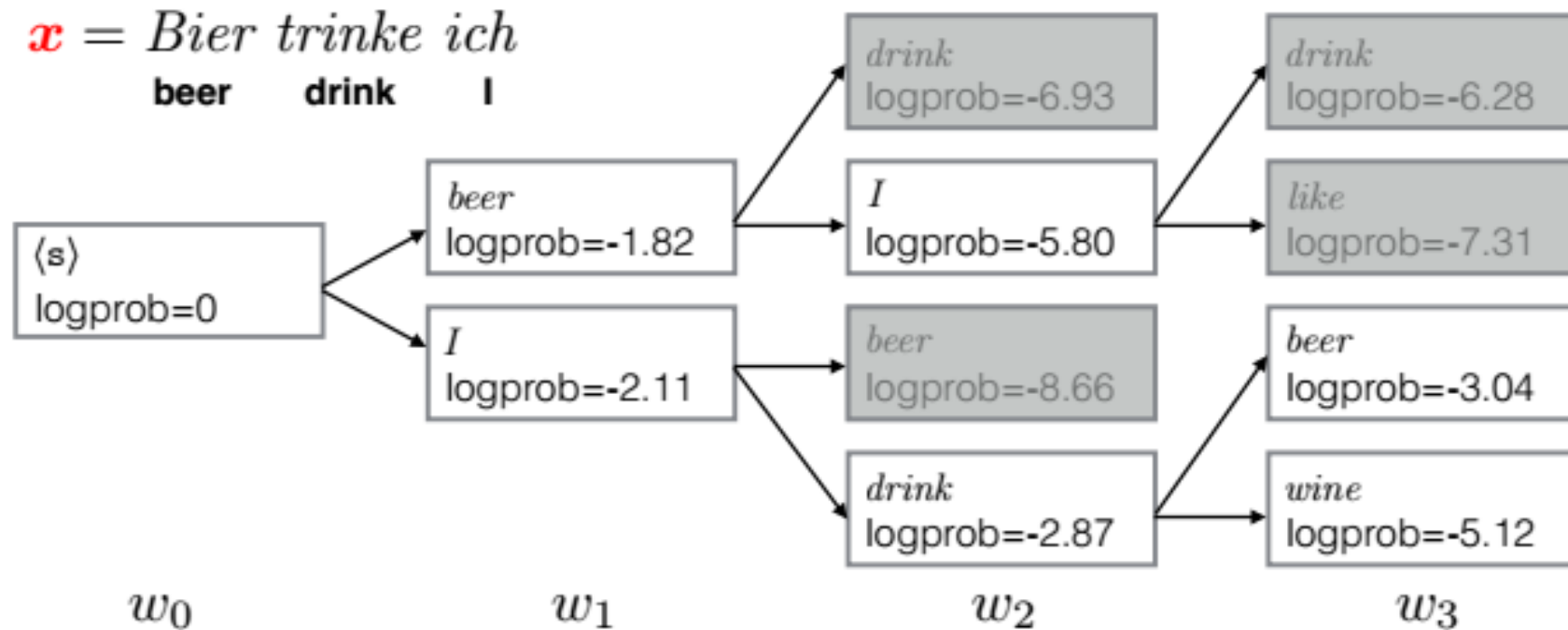
E.g., for $b=2$:

$\mathbf{x} = \text{Bier trinke ich}$
beer **drink** **I**



E.g., for $b=2$:

$\mathbf{x} = \text{Bier trinke ich}$
beer **drink** **I**



A slightly better approximation is to use a **beam search** with beam size b . Key idea: keep track of top b hypothesis.

E.g., for $b=2$:

$\mathbf{x} = \text{Bier trinke ich}$
 beer drink I

