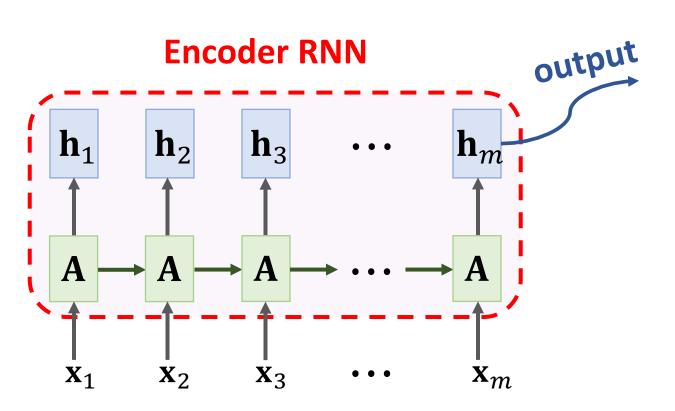
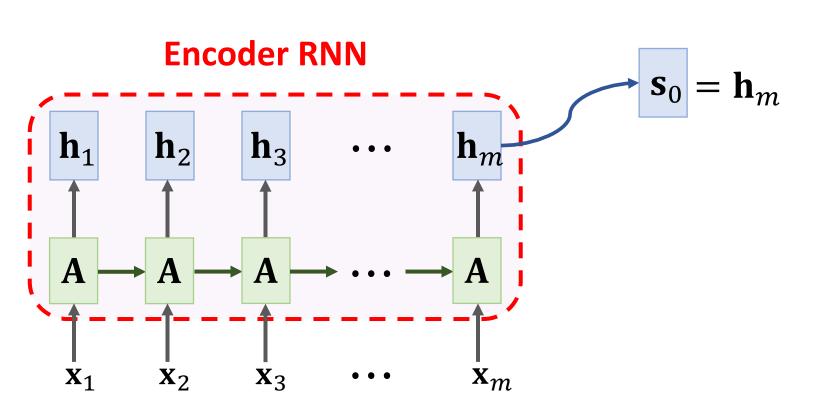
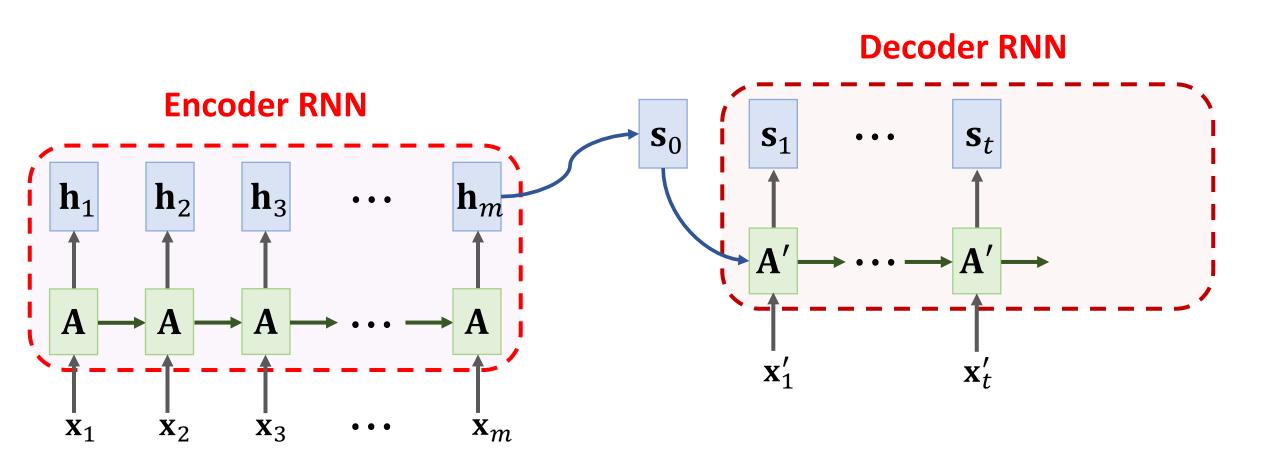
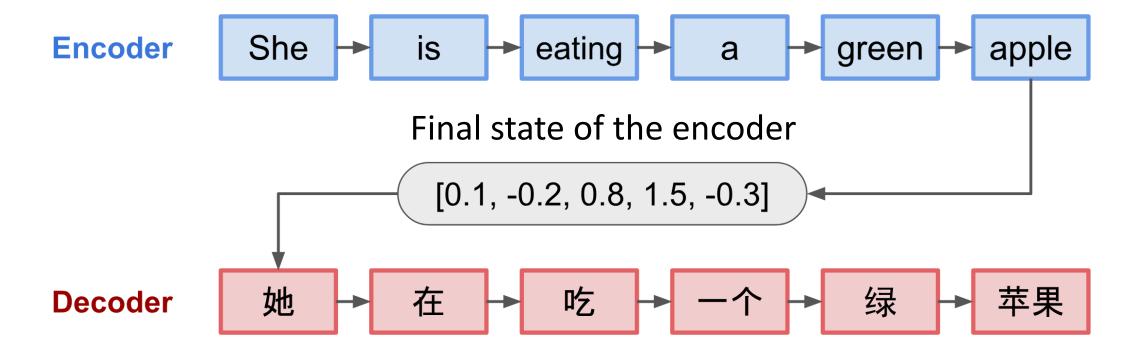
# Attention

**Shusen Wang** 



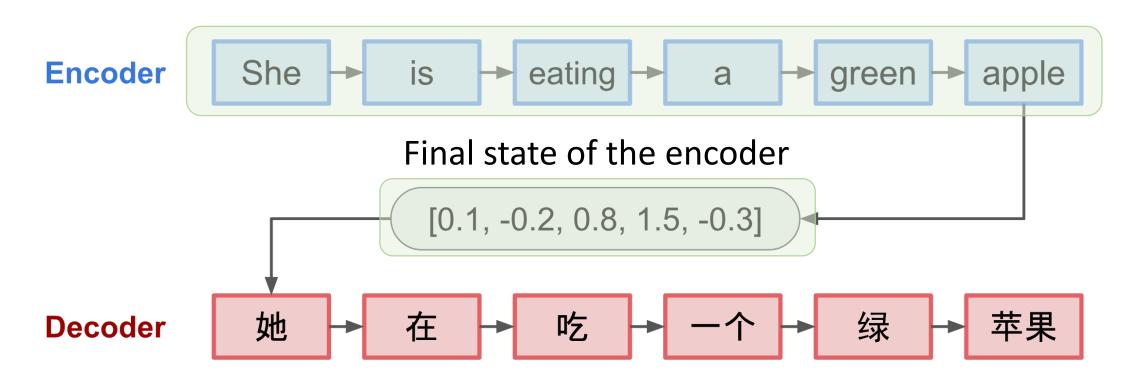






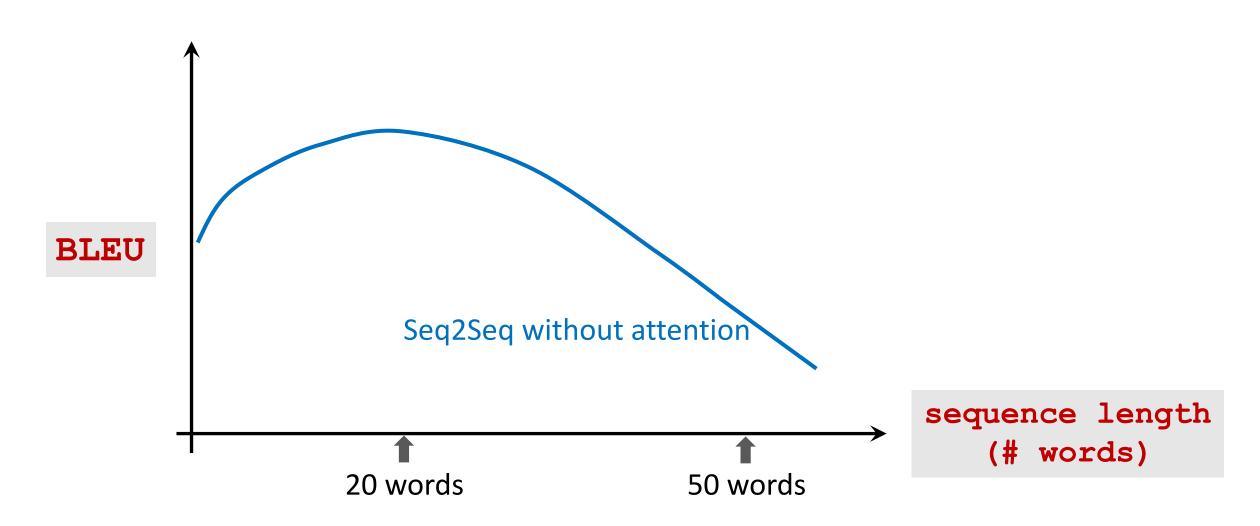
The figure is from blog lilianweng.github.io

**Shortcoming:** The final state is incapable of remembering a **long** sequence.

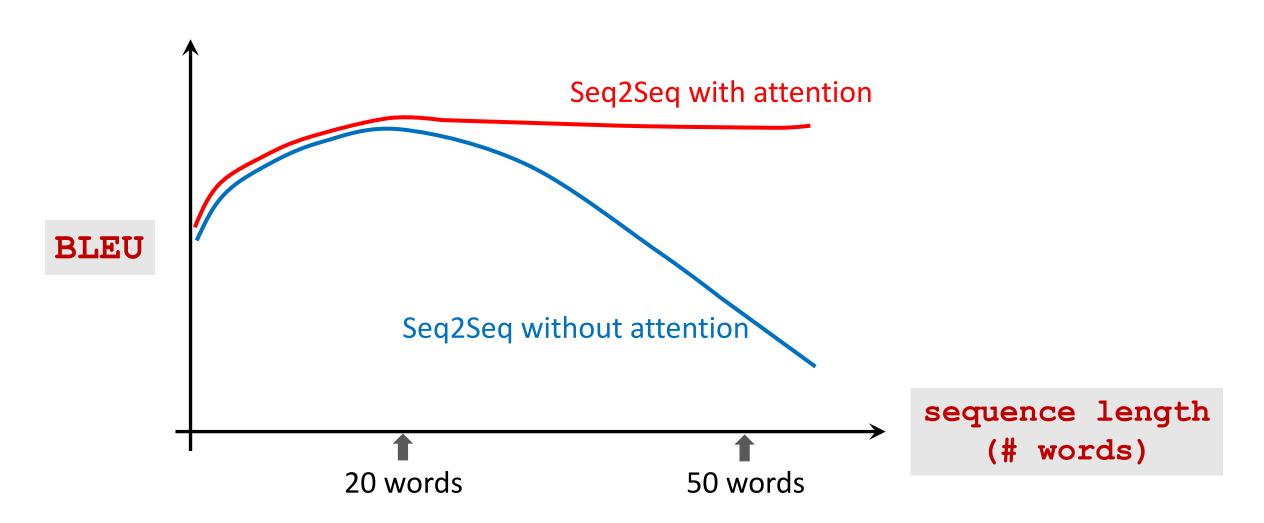


The figure is from blog lilianweng.github.io

**Shortcoming:** The final state is incapable of remembering a **long** sequence.



**Shortcoming:** The final state is incapable of remembering a **long** sequence.

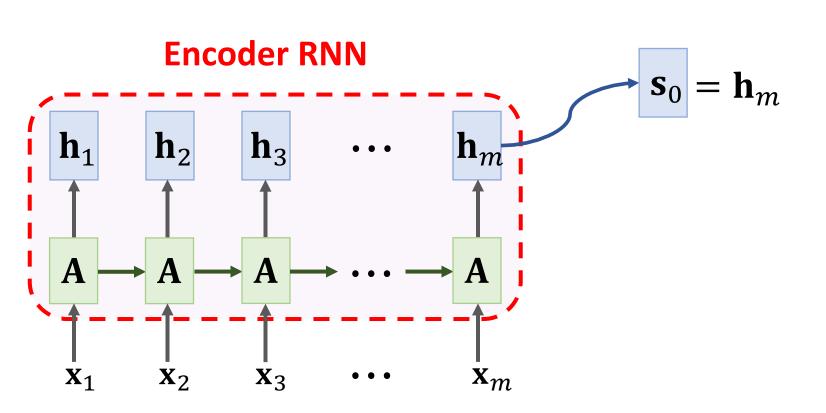


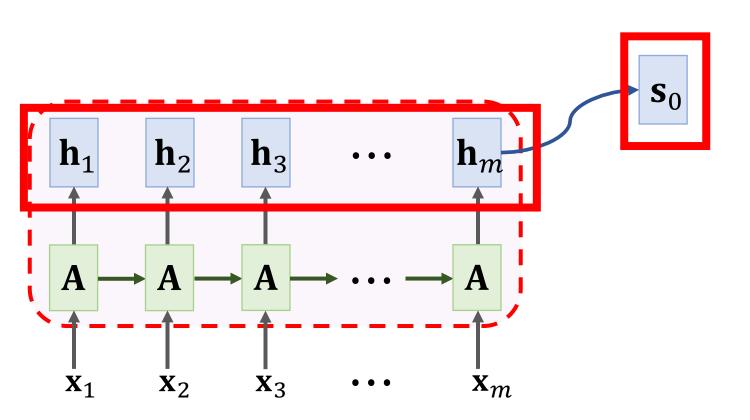
#### Seq2Seq Model with Attention

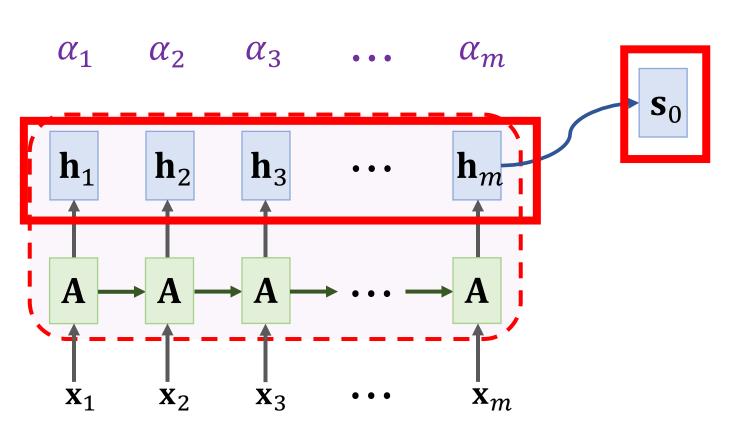
- Attention tremendously improves Seq2Seq model.
- With attention, Seq2Seq model does not forget source input.
- With attention, the decoder knows where to focus.
- Downside: much more computation.

#### **Original paper:**

Bahdanau, Cho, & Bengio. Neural machine translation by jointly learning to align and translate.
 In ICLR, 2015.

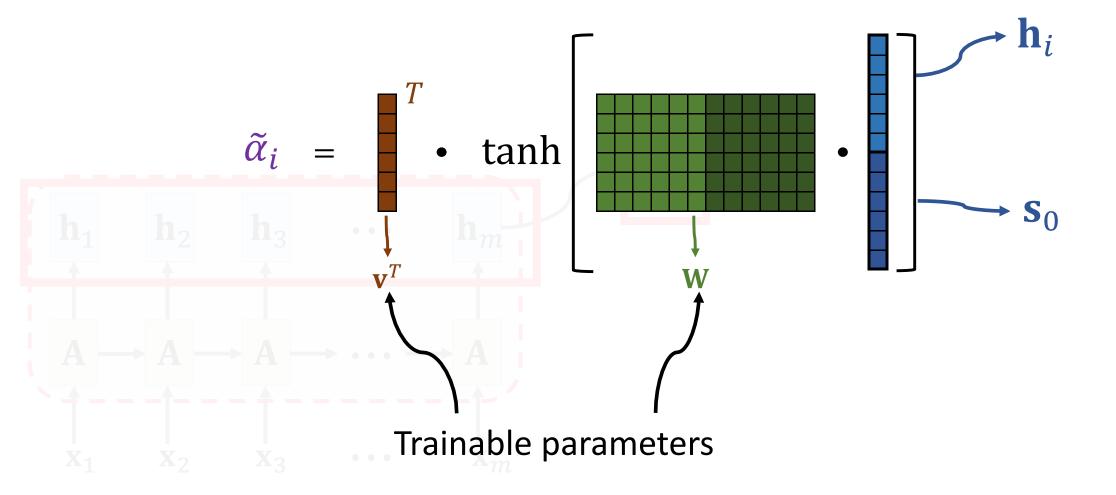






Weights:  $\alpha_i = \operatorname{align}(\mathbf{h}_i, \mathbf{s}_0)$ .

**Option 1** (used in the original paper):



Weights: 
$$\alpha_i = \operatorname{align}(\mathbf{h}_i, \mathbf{s}_0)$$
.

**Option 1** (used in the original paper):

$$\tilde{\alpha}_i = \mathbf{tanh}$$
 $\mathbf{h}_i$ 
 $\mathbf{s}_0$ 

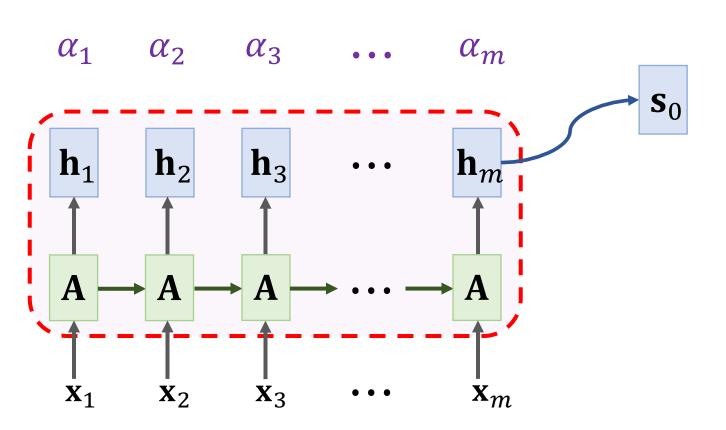
Then **normalize**  $\tilde{\alpha}_1$ ,  $\cdots$ ,  $\tilde{\alpha}_m$  (so that they sum to 1):

$$[\alpha_1, \cdots, \alpha_m] = \text{Softmax}([\tilde{\alpha}_1, \cdots, \tilde{\alpha}_m])$$

Weights: 
$$\alpha_i = \operatorname{align}(\mathbf{h}_i, \mathbf{s}_0)$$
.

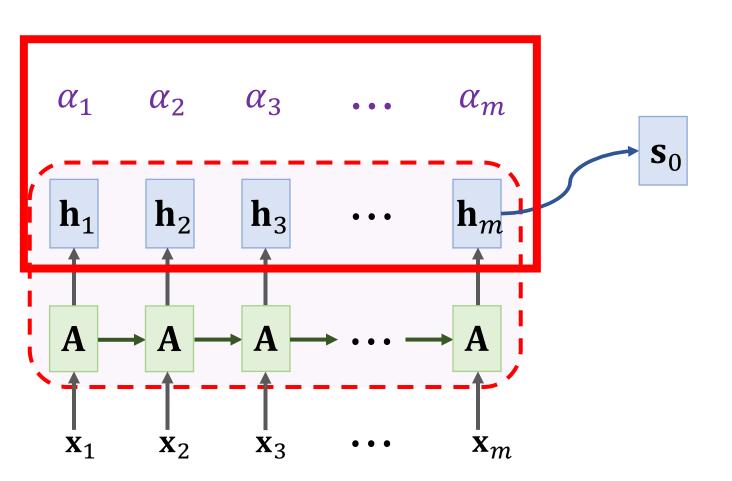
#### **Option 2** (more popular; the same to Transformer):

- 1. Linear maps:
  - $\mathbf{k}_i = \mathbf{W}_K \cdot \mathbf{h}_i$ , for i = 1 to m.
  - $\mathbf{q}_0 = \mathbf{W}_O \cdot \mathbf{s}_0$ .
- 2. Inner product:
  - $\tilde{\alpha}_i = \mathbf{k}_i^T \mathbf{q}_0$ , for i = 1 to m.
- 3. Normalization:
  - $[\alpha_1, \dots, \alpha_m] = \text{Softmax}([\tilde{\alpha}_1, \dots, \tilde{\alpha}_m])$



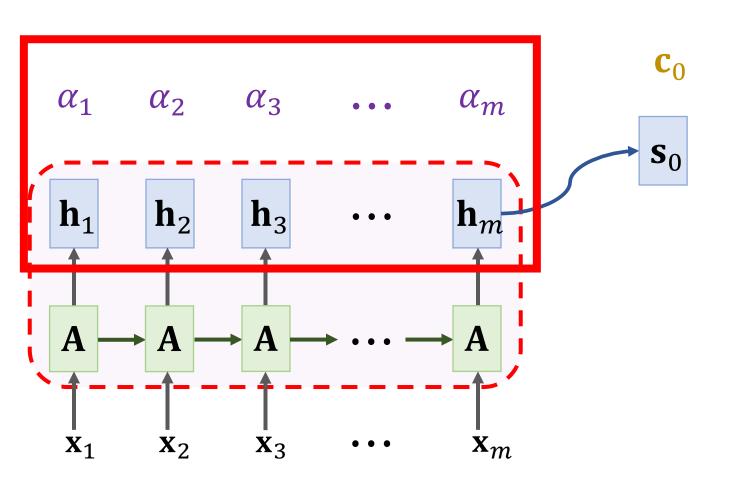
Weights:  $\alpha_i = \operatorname{align}(\mathbf{h}_i, \mathbf{s}_0)$ .

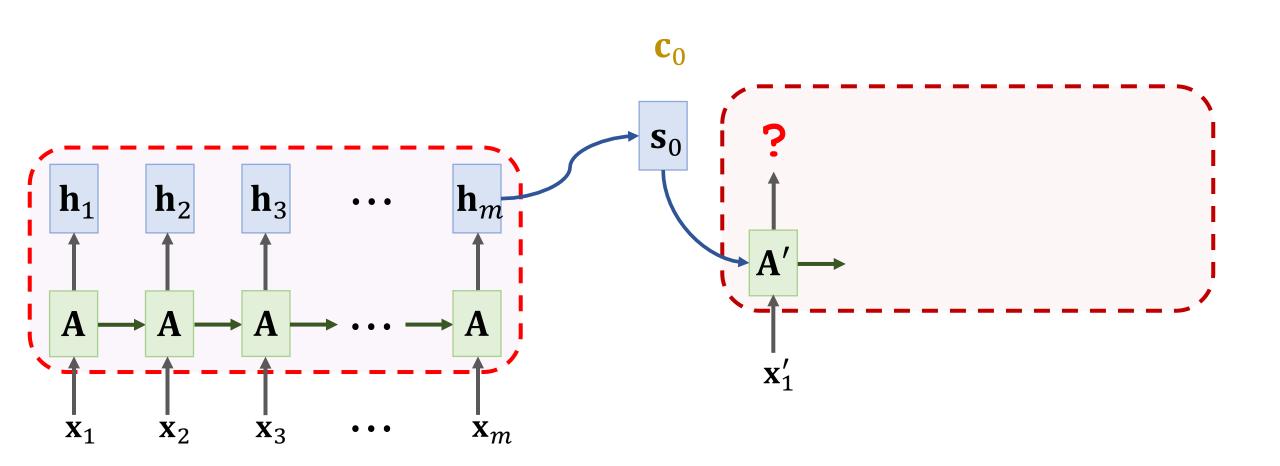
Context vector:  $\mathbf{c}_0 = \alpha_1 \mathbf{h}_1 + \cdots + \alpha_m \mathbf{h}_m$ .



Weights:  $\alpha_i = \operatorname{align}(\mathbf{h}_i, \mathbf{s}_0)$ .

Context vector:  $\mathbf{c}_0 = \alpha_1 \mathbf{h}_1 + \cdots + \alpha_m \mathbf{h}_m$ .

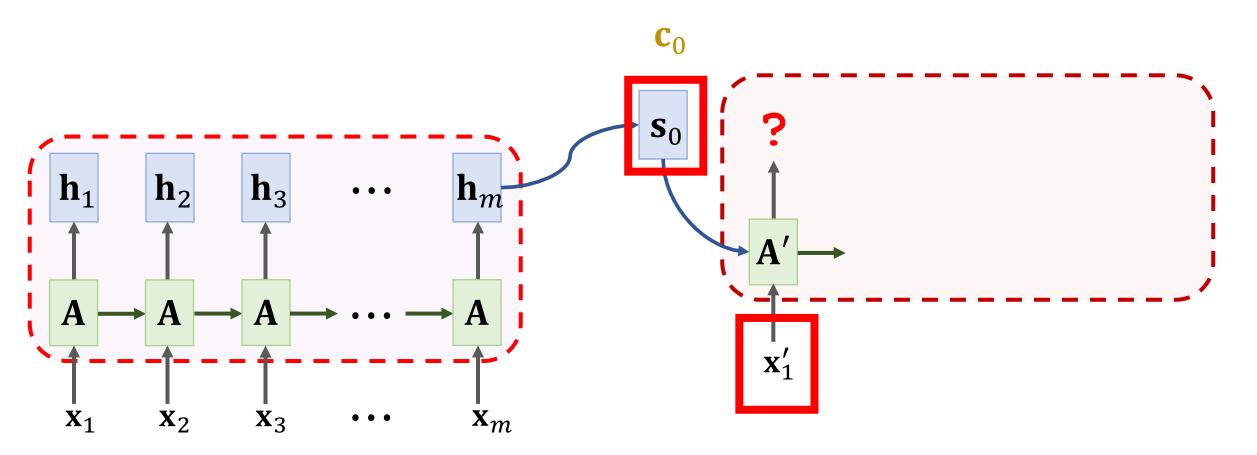


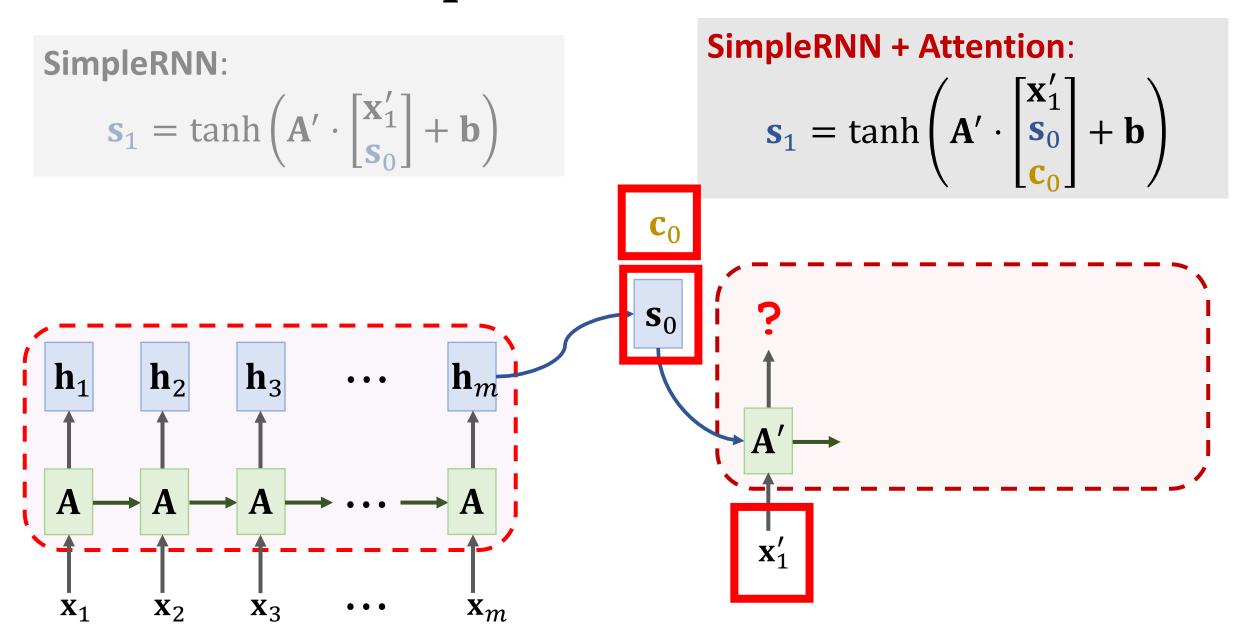


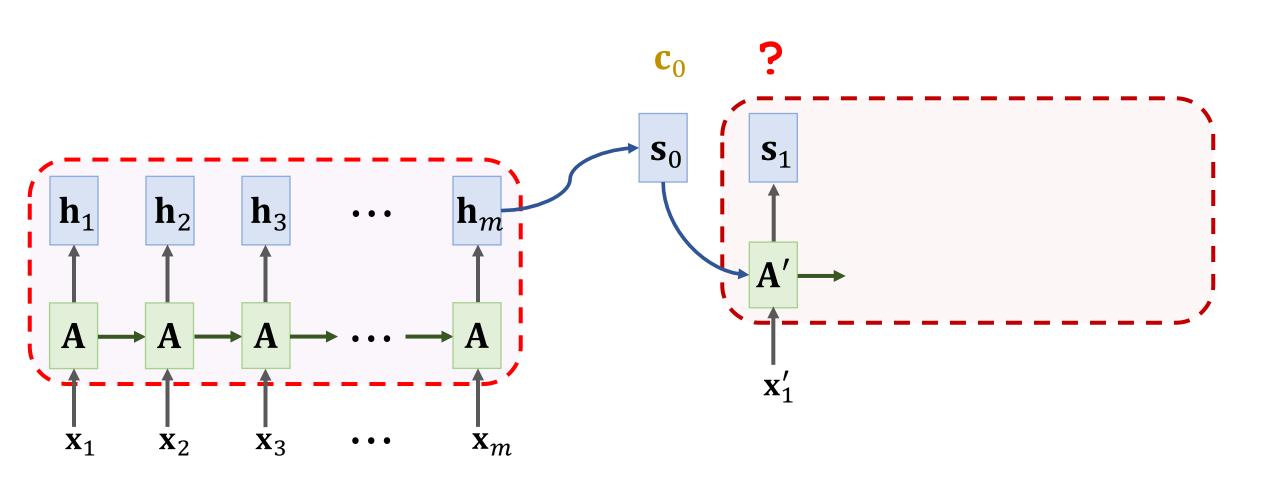
## **SimpleRNN**

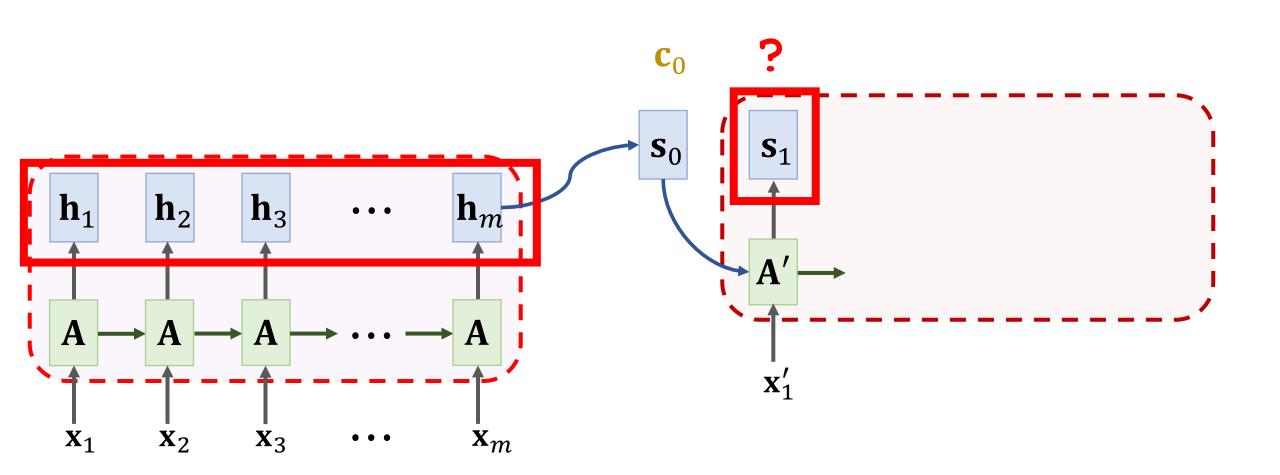
#### SimpleRNN:

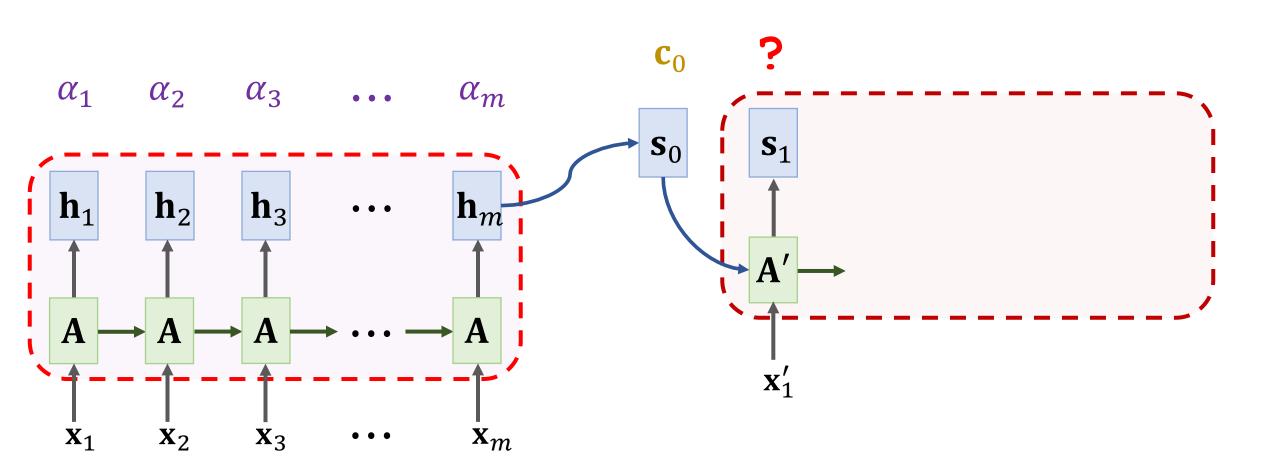
$$\mathbf{s}_1 = \tanh\left(\mathbf{A}' \cdot \begin{bmatrix} \mathbf{x}_1' \\ \mathbf{s}_0 \end{bmatrix} + \mathbf{b}\right)$$

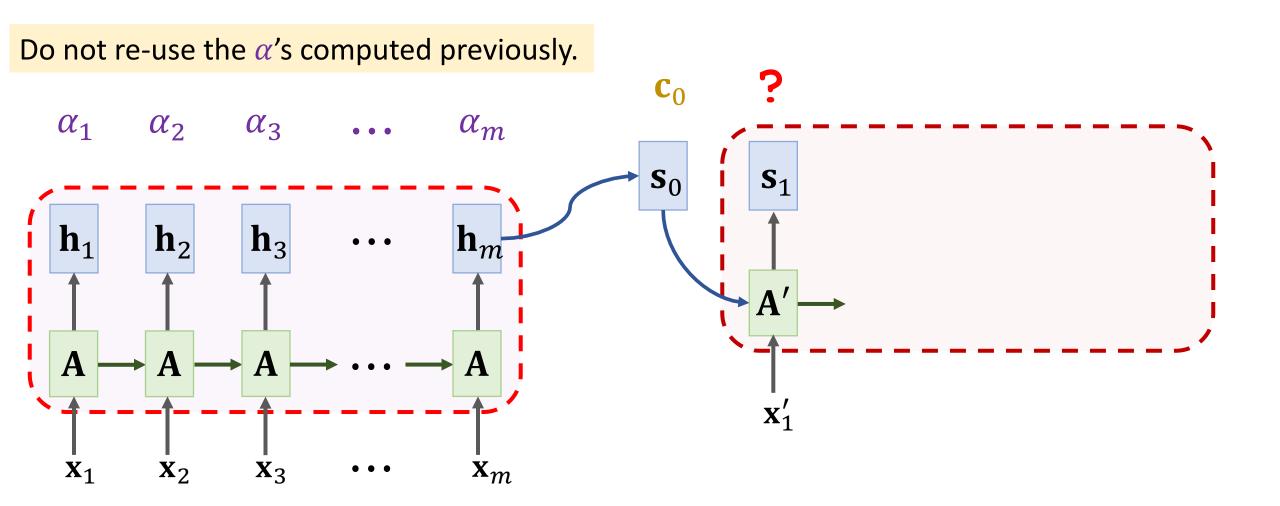


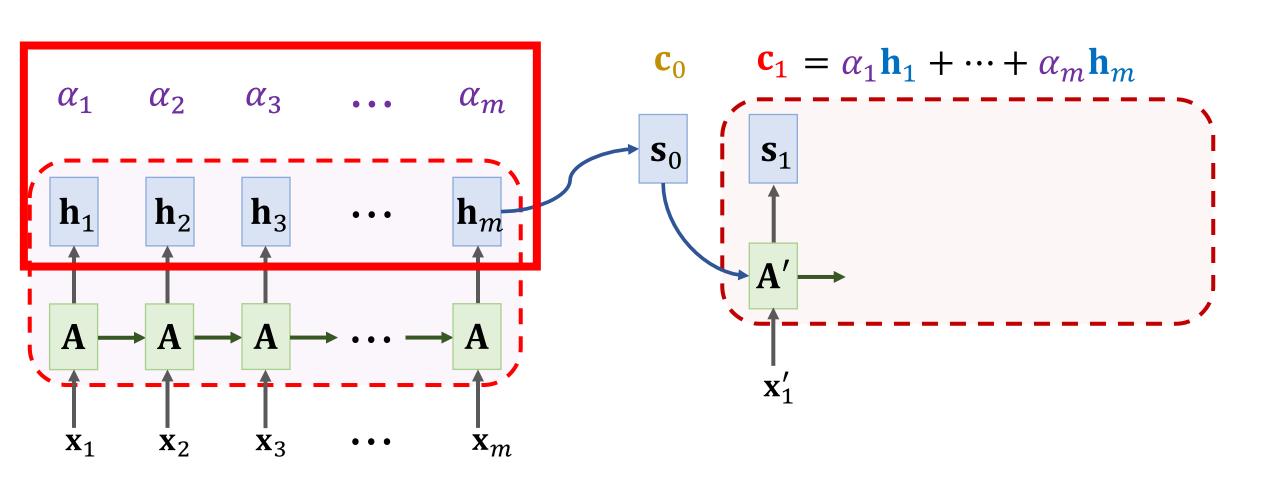




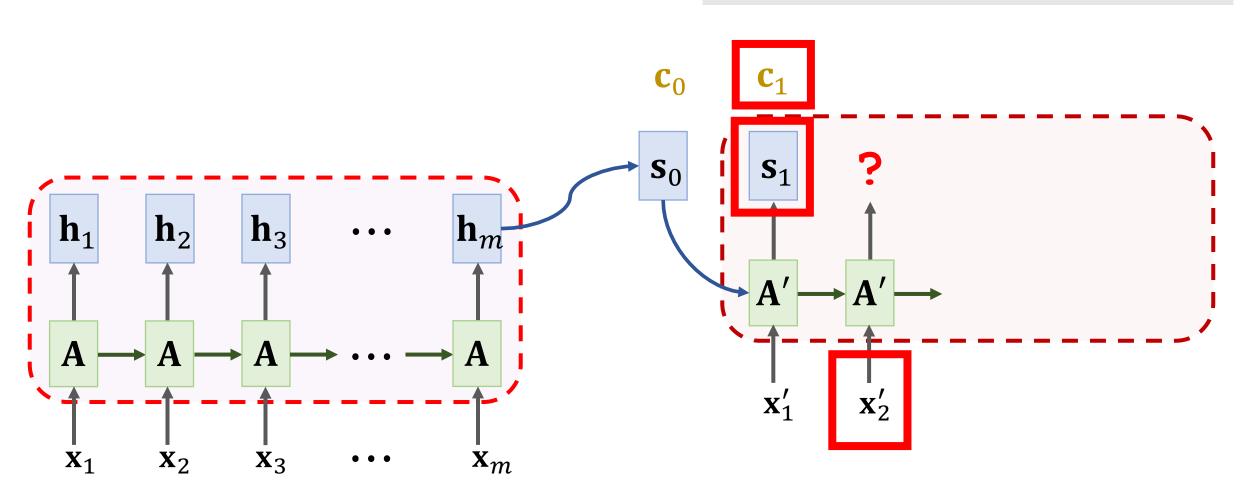


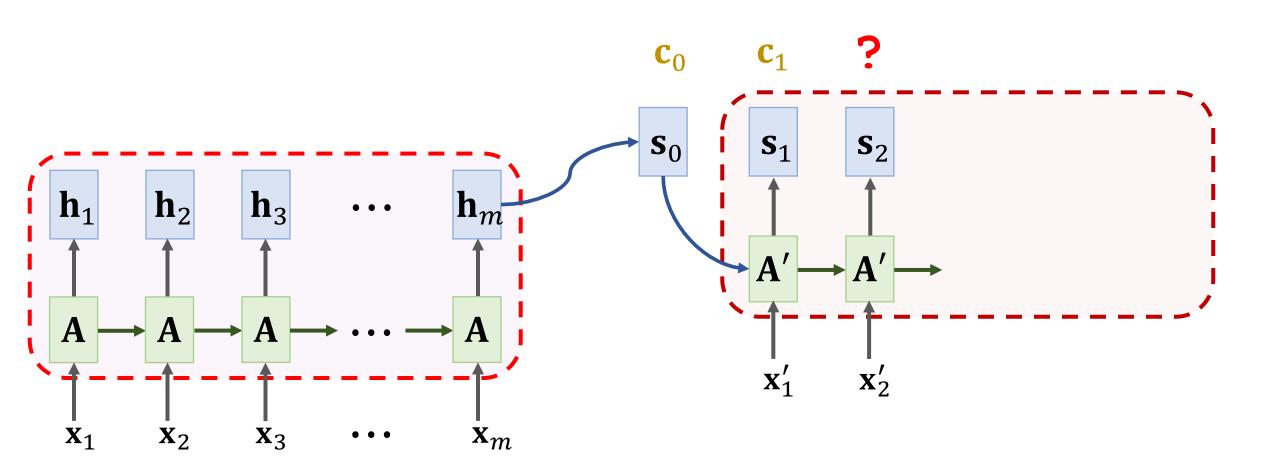


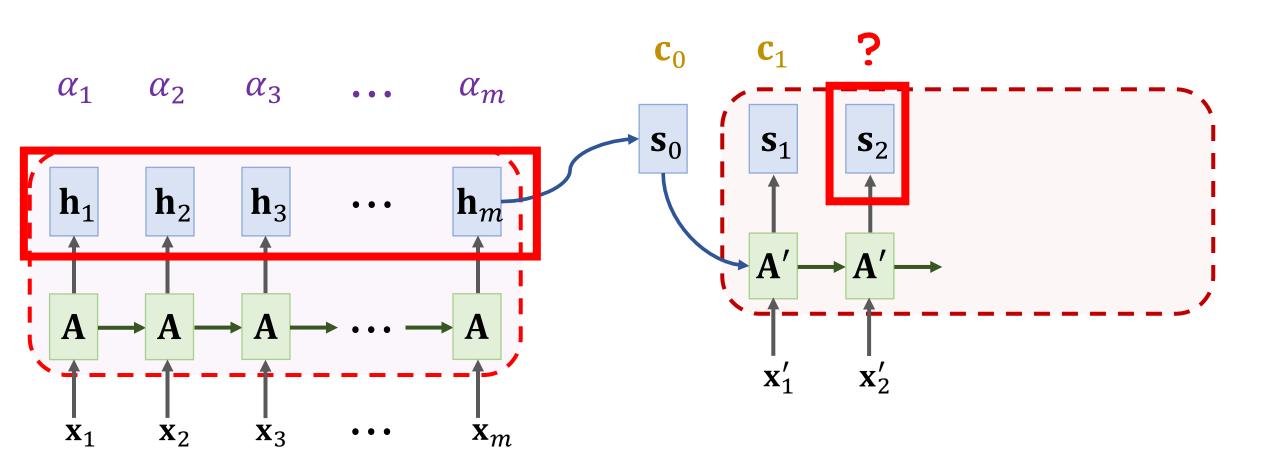


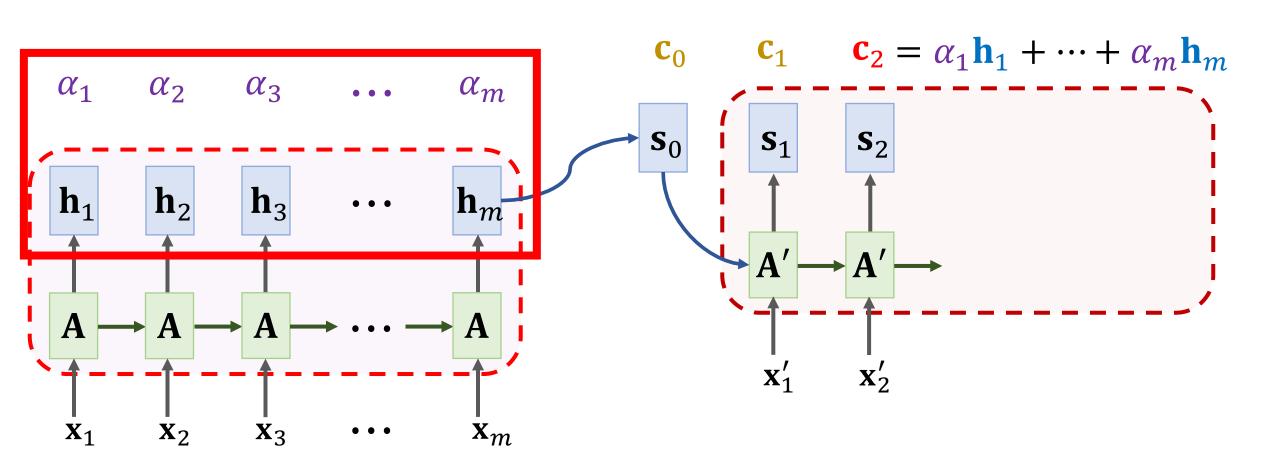


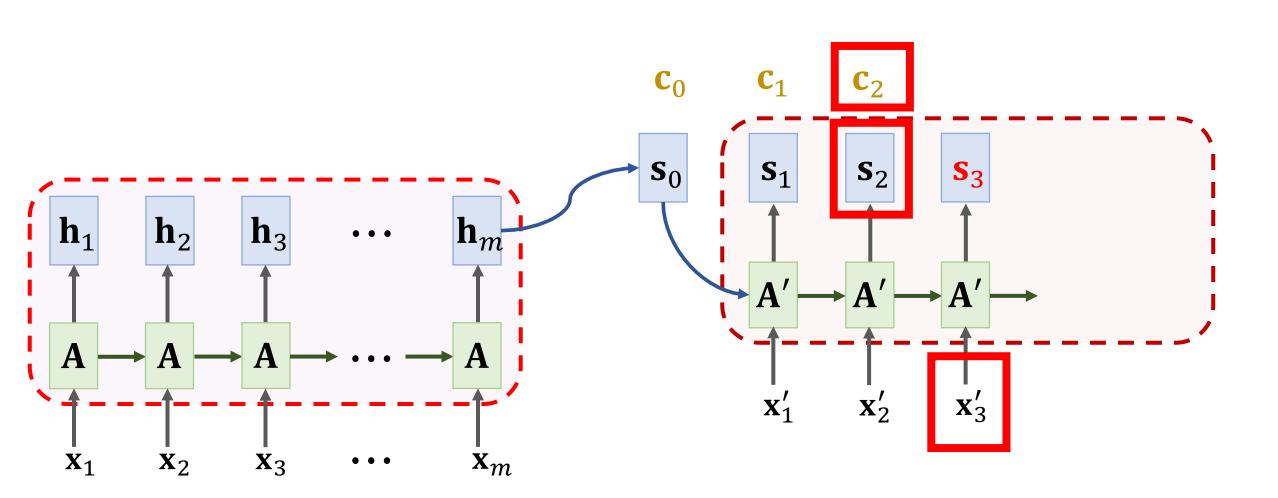
$$\mathbf{s}_2 = \tanh\left(\mathbf{A}' \cdot \begin{bmatrix} \mathbf{x}_2' \\ \mathbf{s}_1 \\ \mathbf{c}_1 \end{bmatrix} + \mathbf{b}\right)$$

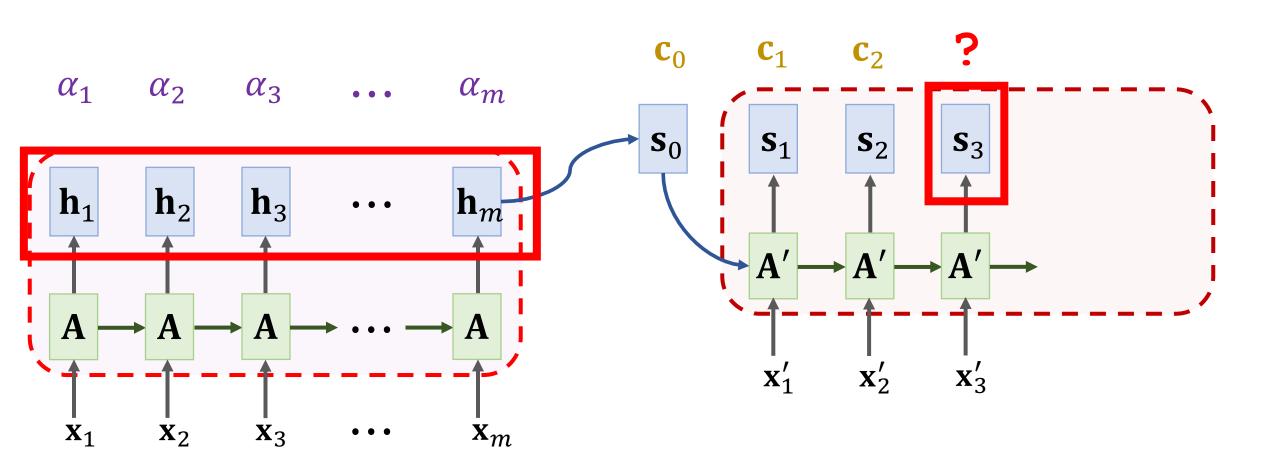


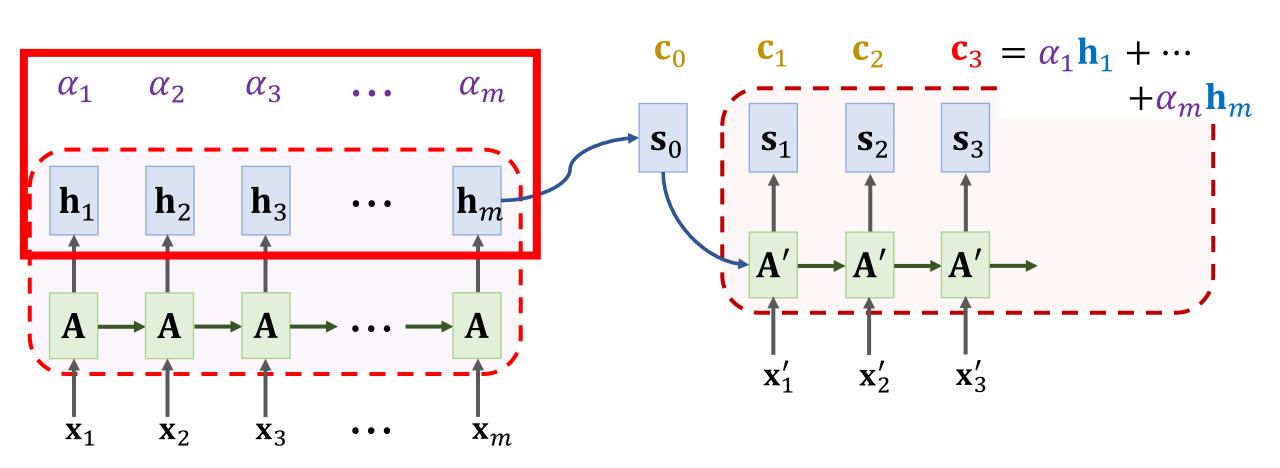


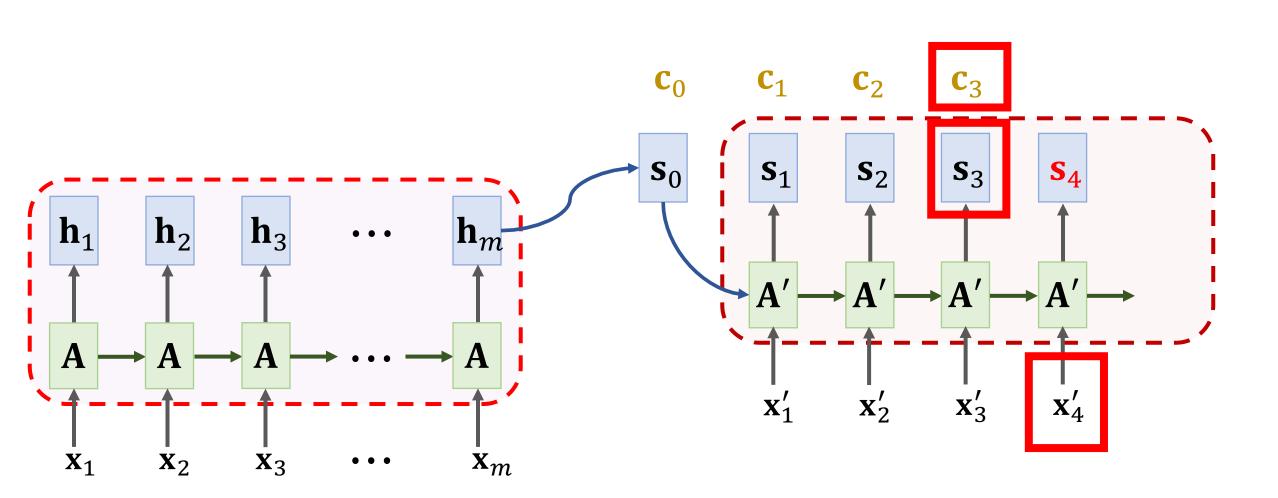


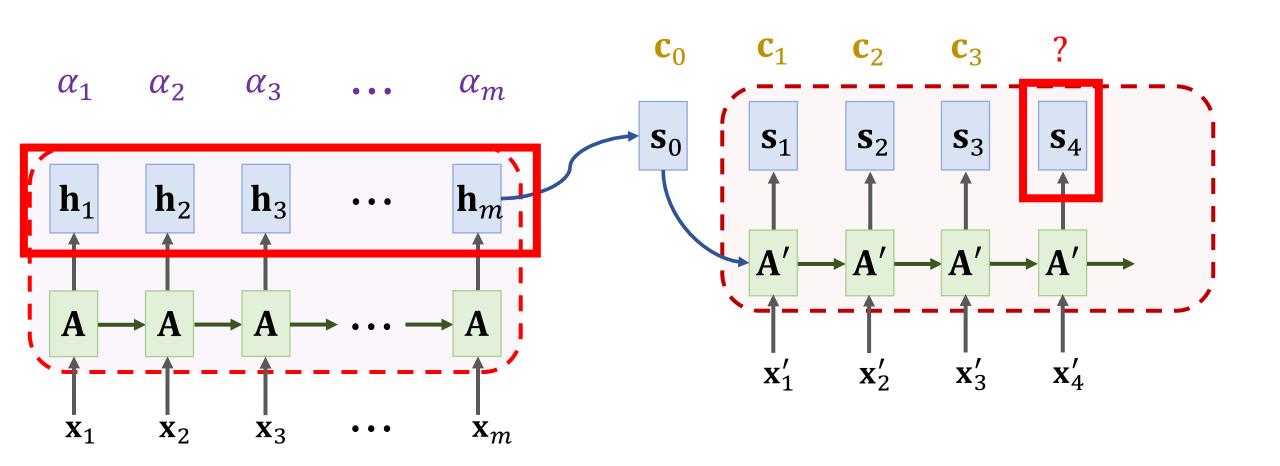


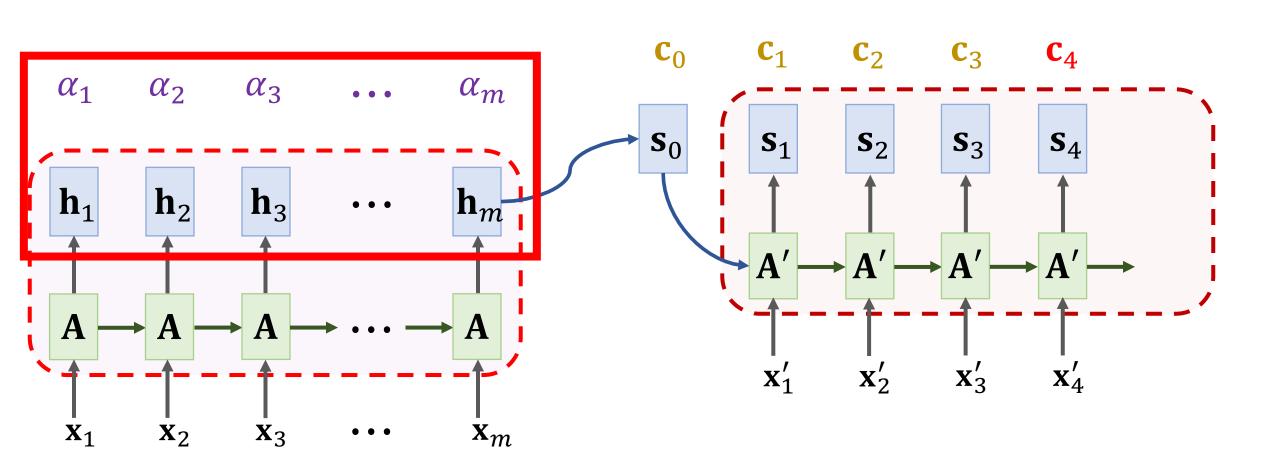


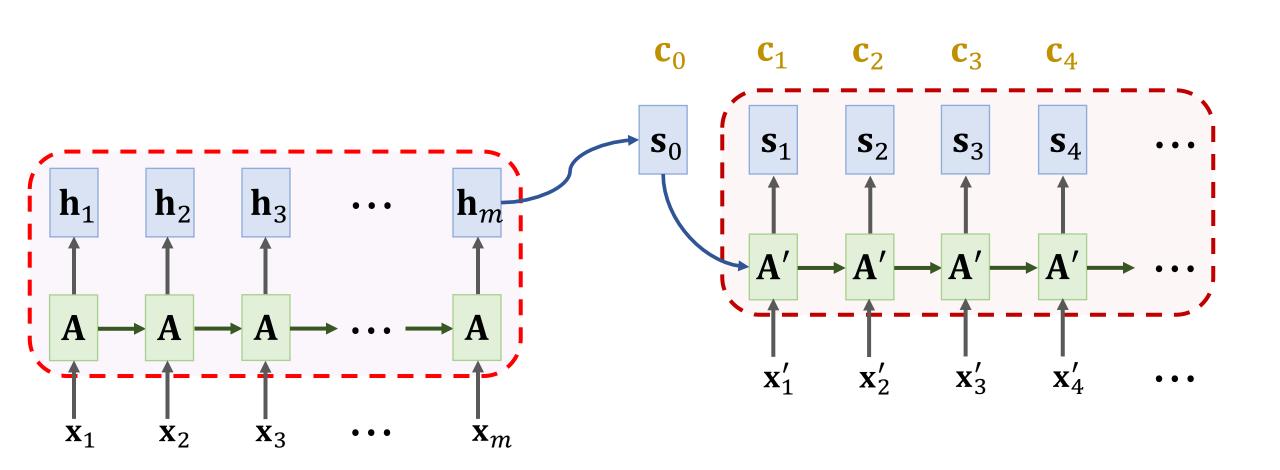




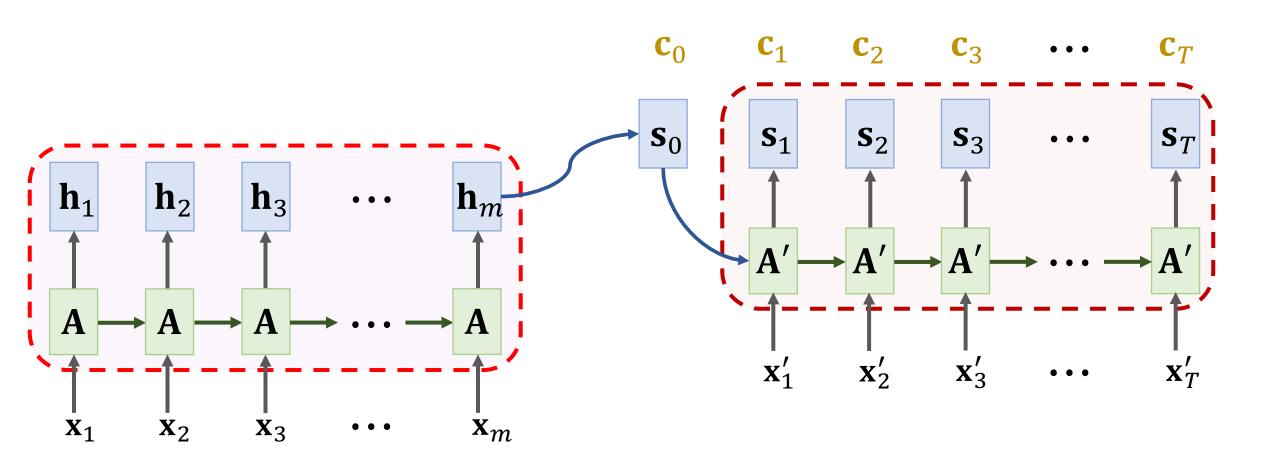






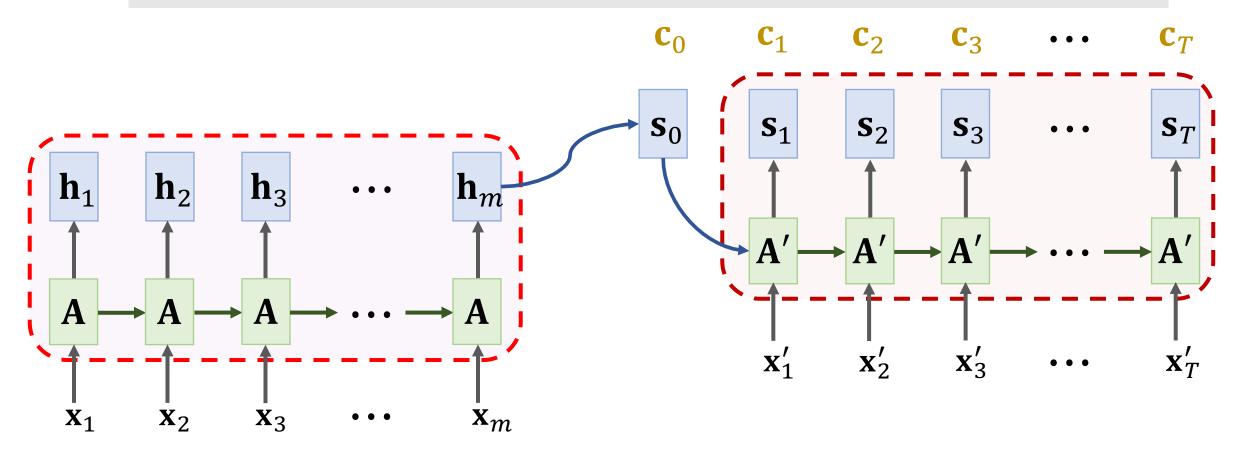


**Question:** How many weights  $\alpha_i$  have been computed?



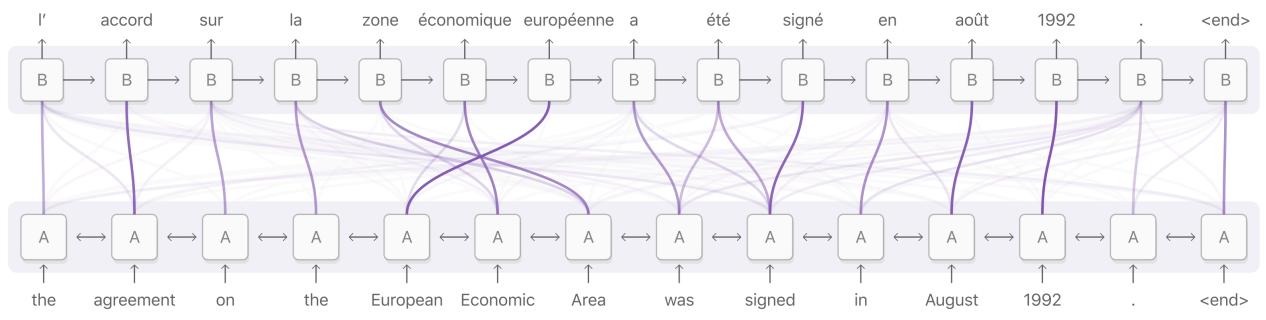
**Question:** How many weights  $\alpha_i$  have been computed?

- For every decoder state  $s_t$ , there are m weights:  $\alpha_1, \dots, \alpha_m$ .
- The decode has T states, so there are totally mT weights.



## **Attention: Weights Visualization**

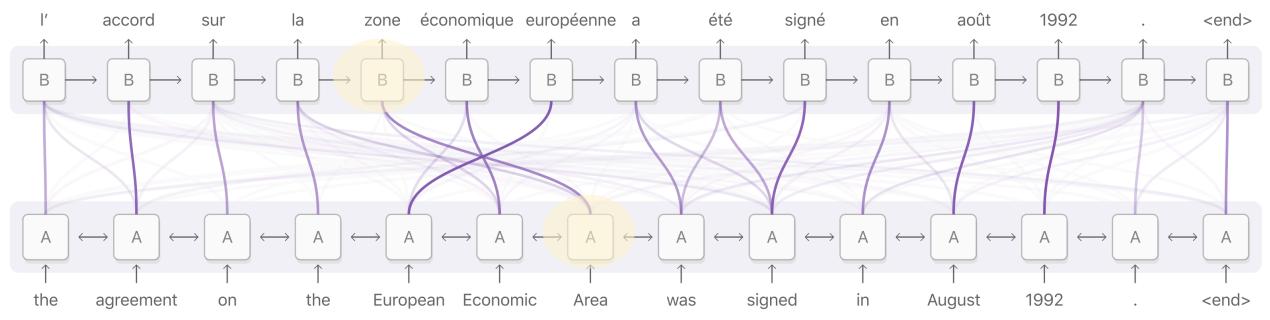
#### **Decoder RNN** (target language: French)



**Encoder RNN** (source language: English)

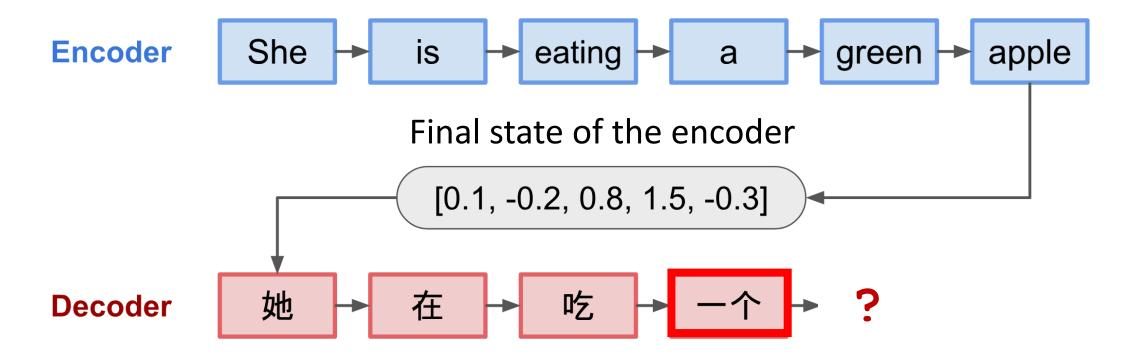
## **Attention: Weights Visualization**

#### **Decoder RNN** (target language: French)

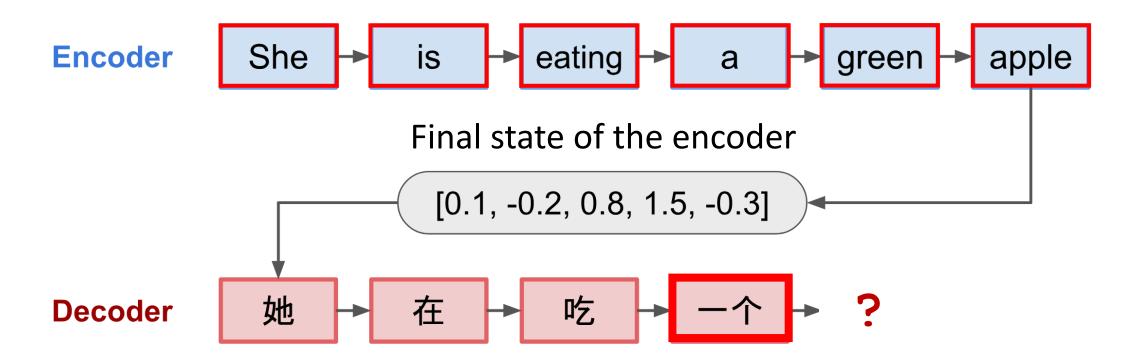


**Encoder RNN** (source language: English)

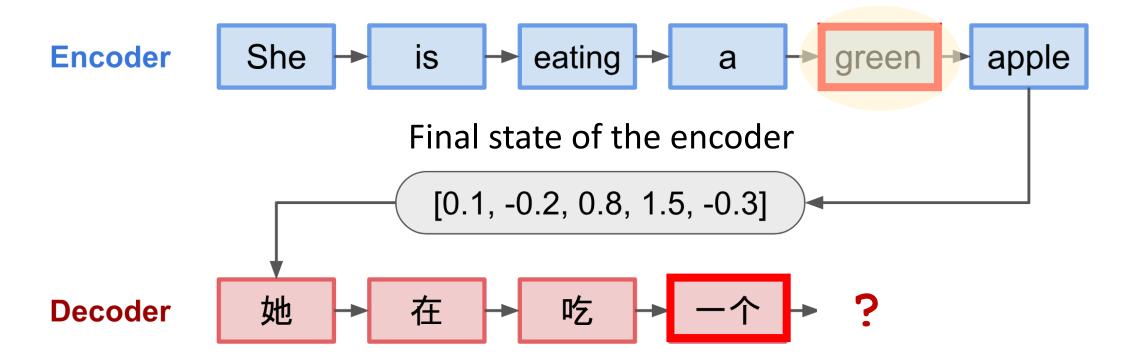
Standard Seq2Seq model: the decoder looks at only its current state.



- Standard Seq2Seq model: the decoder looks at only its current state.
- Attention: decoder additionally looks at all the states of the encoder.



- Standard Seq2Seq model: the decoder looks at only its current state.
- Attention: decoder additionally looks at all the states of the encoder.
- Attention: decoder knows where to focus on.



- Standard Seq2Seq model: the decoder looks at only its current state.
- Attention: decoder additionally looks at all the states of the encoder.
- Attention: decoder knows where to focus on.

- Downside: higher time complexity.
  - *m*: source sequence length
  - t: target sequence length
  - Standard Seq2Seq: O(m+t) time complexity
  - Seq2Seq + attention: O(mt) time complexity

Thank you!