weekly report: amino acid seq

Jitian Zhao

University of Wisconsin Madison

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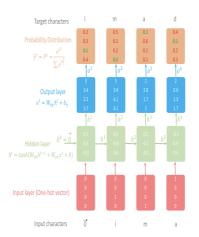
1/9

Outline

- low performance of Char-RNN
- code modification of Transformer

char-RNN:recap

- training: $\hat{X}_n = sample(P(X_n|X_1,...,X_{n-1}))$
- generating: $\hat{X}_n = sample(P(X_n|\hat{X}_1,...,\hat{X}_{n-1}))$



Source: https://towardsdatais.unce.training.processlanguage-model-1439f5dd87fe

low performance of Char-RNN

- low reconstruction accuracy for test dataset: 33%
- low accuracy was not due to error propagation
- model insensitive to the given information

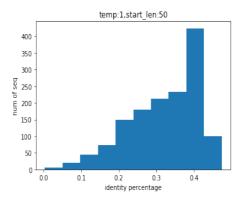


Figure: reconstruction accuracy

possible reasons of why it doesn't work

- simple structure: 1 gru layer + 1 fc layer
- does not learn from the position
- sequence diversity or data clustering
- ...

Transformer

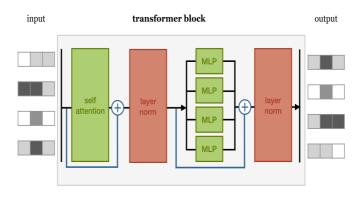


Figure: structure of transformer

self-attention mechanism

- input: $X_1, ..., X_t$
- output: $Y_i = \sum_i w_{ij} X_j$
- weight: w_{ij} = softmax $(w_{ij}^{'}), w_{ij}^{'} = X_i^T X_j$
- one input vector X_i serves 3 needs: contribute to its own weight (query), other inputs weight (key), final weighted sum (value)

7/9

catch the position information: multi-head attention

- example: 'mary', 'gives', 'jane', 'a', 'flower'
- neighbor matters!
- different attention heads have different sets of matrices, which gives it power of discriminating neighbors.

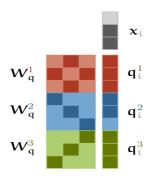


Figure: multi-head attention

Source: http://peterbloem.nl/blog/transformers

tailored modification for a.a. seq

- sequence padding
- ullet random trunk \longrightarrow randomly choose one sequence
- word encoding: different vocabulary

9/9