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How to use sequences in Deep Learning

## Sequences, could you be even more vague?



MEACCMELVKC

TACCTTGGC...

- Typically:
- Proteins
- DNA
- RNA
- Generally:
- Order important
- Text representation



Trimming
Appending


- Select sequence length depended of model layers
- E.q. LSTM < 400
- Append with content
- Truncate now, join later
006
017
006

000 $\quad$| 0.8 |
| :--- |
| 0.7 |
| 0.6 |
| 0.0 |

## MEACCMELVKC

- Problems:
- When using one hot encoding, network needs to learn properties of each letter
- Each letter has no context information


```
Arg
Arg -
Asn -2
Asp -2 -2 1
Cys
Gln
Glu
His
His 
Ile 
Leu
Lys
Mhe 
```



```
Pro 
Ser
Thr 
Tyr
Ala Arg Asn Asp Cys Gln Glu Gly His lle Leu Lys Met Phe Pro Ser Thr Trp Tyr Val
```

- Potential solutions:
- Encode properties (measurements, Blosum62 ...)
- Use different embedding per letter dependent on context

 Input
[CLS] the man [MASK] to the store [SEP] penguin [MASK] are flightless birds [SEP]
- Idea we treat our sequence as language
- Words < Sentence < Document


## http://jalammar.github.io/illustrated-bert/



## MEACCMELVKC



- Idea we treat our sequence as language
- Words < Sentence < Document
- Aminoacid < Proteindomain < Protein
- Triplet < multiple Triplets < Sequence

- Trained on UniRef50
- Word length 1, stride 1, sentence 1024 words
- Trained on DNA of same proteins
- Word length 3 , stride 3 , sentence about 80 words

DNA/RNA/Proteins != human language
What comes next?

Needs to rethink training and architecture

Thank you for your attention


