

Large Language Models

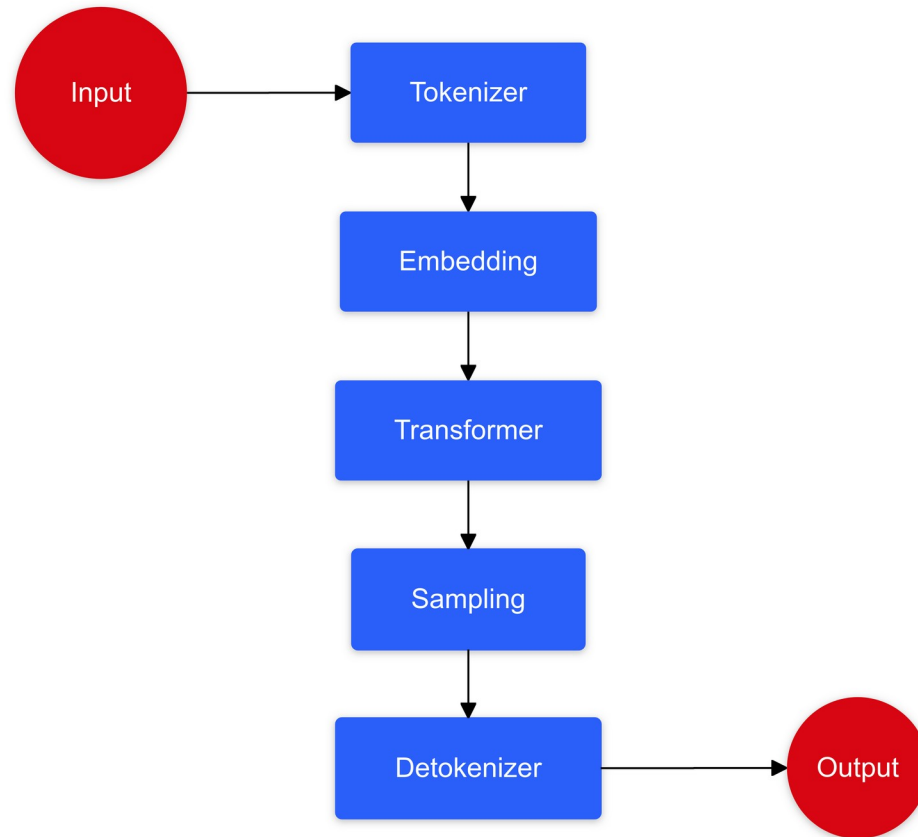
Fundamentos e aplicações

Github

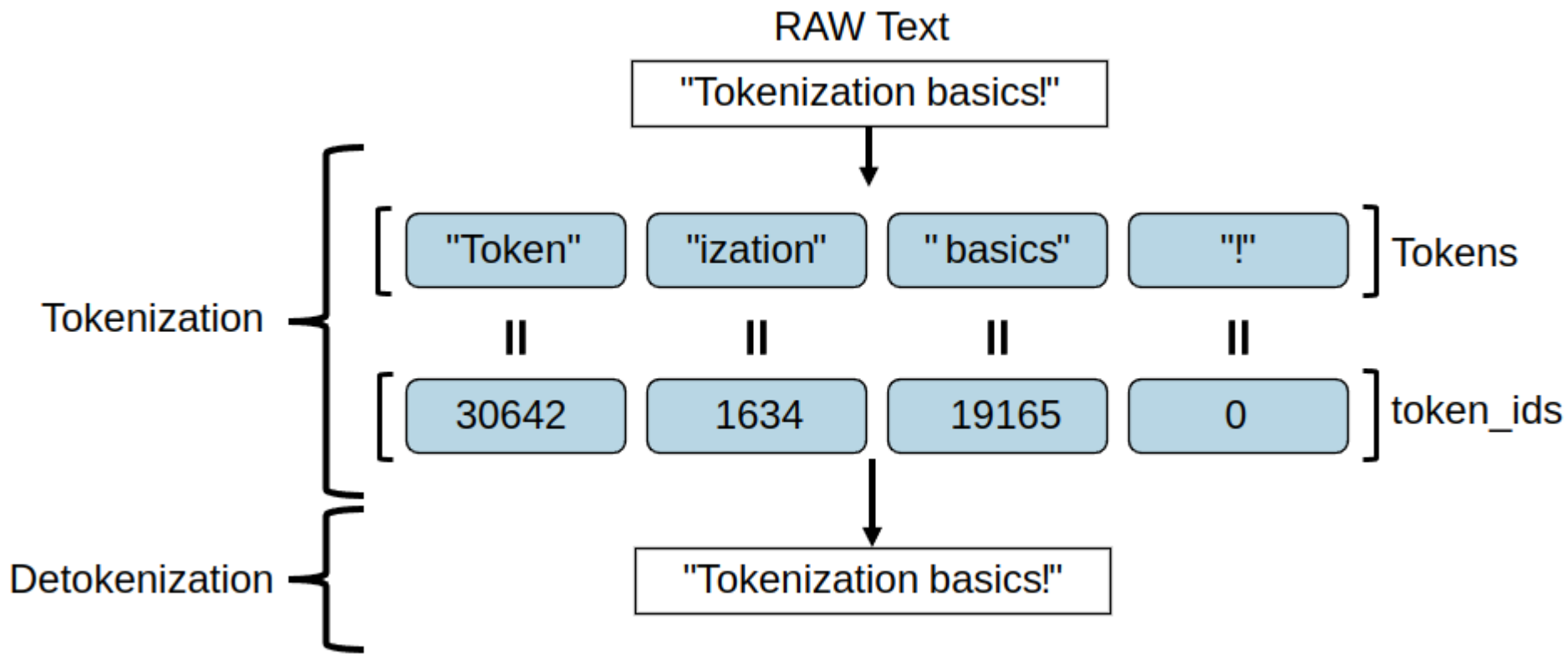
<https://github.com/mucciaccia/eri-es-llm>

Os códigos em Python utilizados durante a oficina estão disponíveis no link do Github acima.

Partes de uma LLM



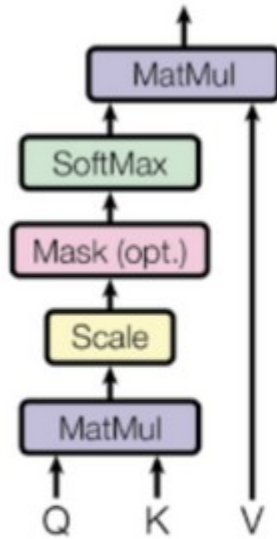
Tokenizer



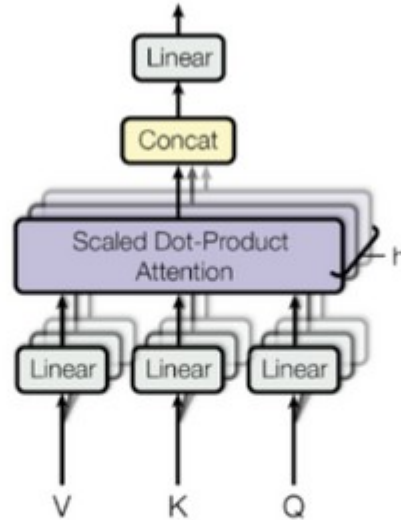
Embedding

Token String	Token ID	Embedded Token Vector
'<s>' ->	0 ->	[0.1150, -0.1438, 0.0555, ...]
'<pad>' ->	1 ->	[0.1149, -0.1438, 0.0547, ...]
'</s>' ->	2 ->	[0.0010, -0.0922, 0.1025, ...]
'<unk>' ->	3 ->	[0.1149, -0.1439, 0.0548, ...]
'.'	4 ->	[-0.0651, -0.0622, -0.0002, ...]
' the'	5 ->	[-0.0340, 0.0068, -0.0844, ...]
','	6 ->	[0.0483, -0.0214, -0.0927, ...]
' to'	7 ->	[-0.0439, 0.0201, 0.0189, ...]
' and'	8 ->	[0.0523, -0.0208, -0.0254, ...]
' of'	9 ->	[-0.0732, 0.0070, -0.0286, ...]
' a'	10 ->	[-0.0194, 0.0302, -0.0838, ...]
...		

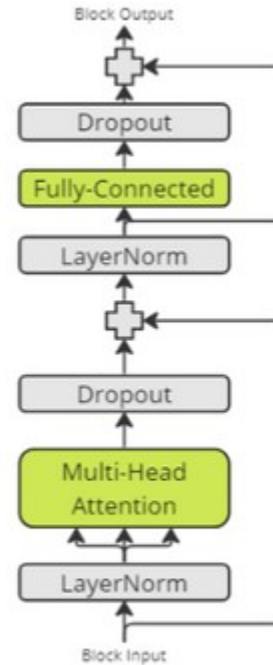
Transformer



Scale Causal Attention



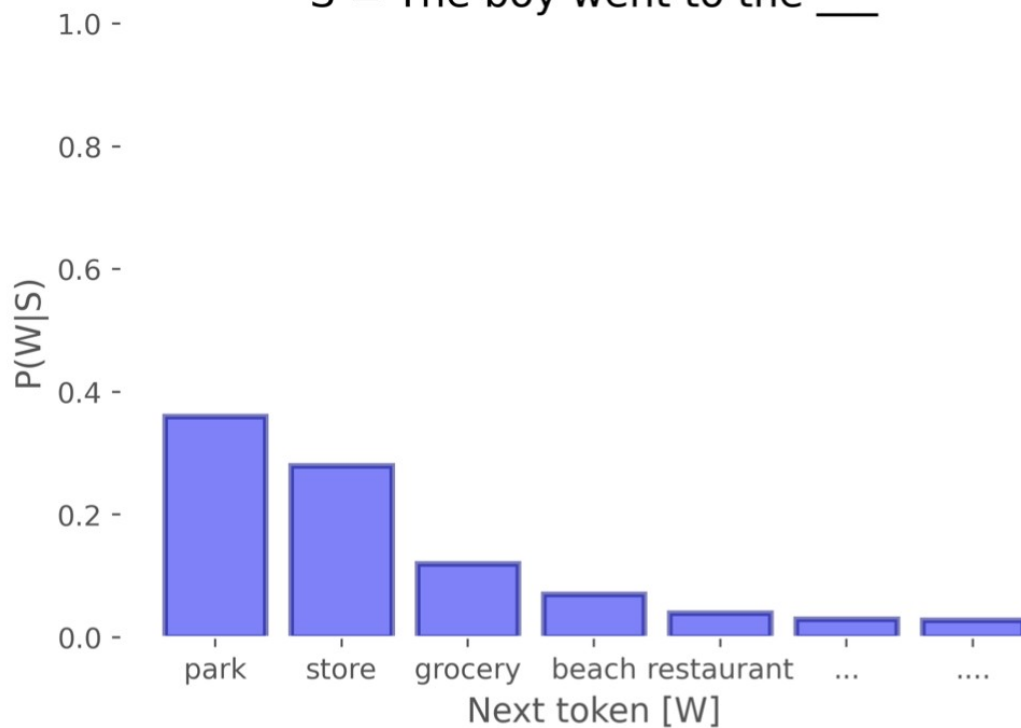
Multi-Head Attention



Transformer Block

Sampling

S = The boy went to the ____



Tipos das saídas

- Input: String
- Tokenizer: Vetor de inteiros ($N \times 1$)
- Embedding: Matriz de pontos flutuantes ($N \times M$)
- Transformer: Vetor de probabilidades ($T \times 1$)
- Sampling: Inteiro (1×1)
- Detokenizer: String

NLP Tasks

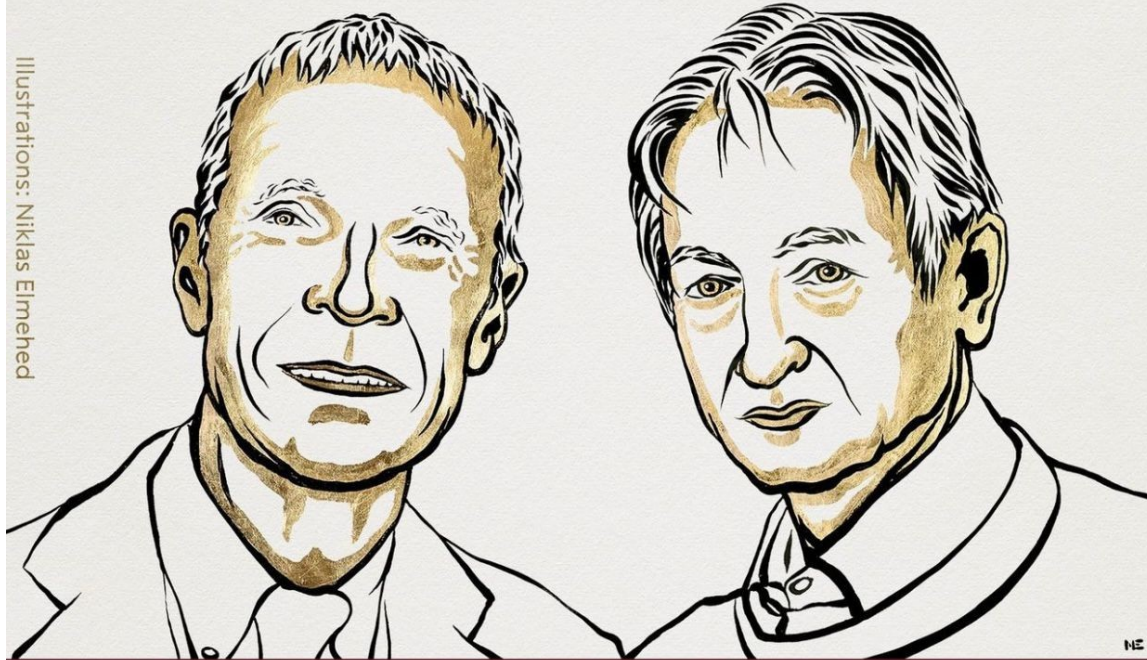
- Sentiment Analysis
- Sumarization
- Question answering
- Translation
- Grammar check
- Text completion

Curiosidade

- Prêmios Nobel de 2024 de física e química foram dados a pesquisadores de inteligência artificial

THE NOBEL PRIZE IN PHYSICS 2024

Illustrations: Niklas Elmehed



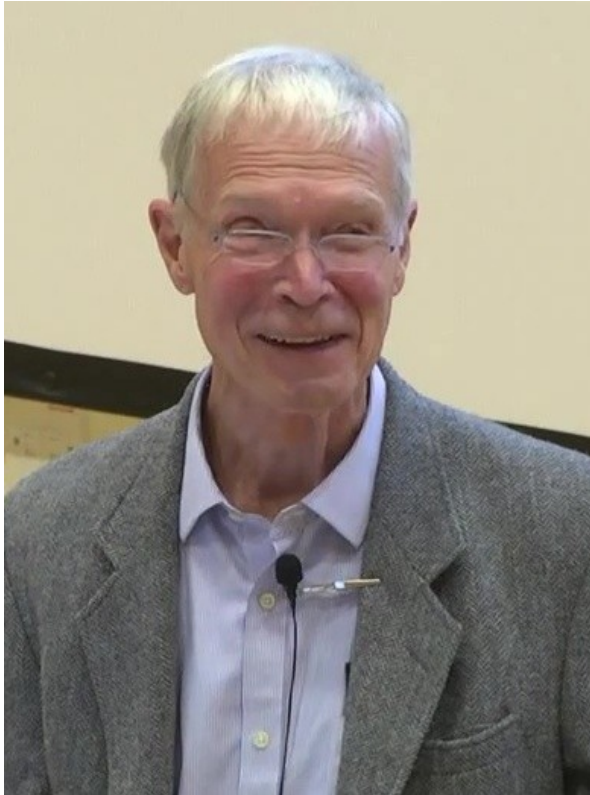
John J. Hopfield

Geoffrey E. Hinton

"for foundational discoveries and inventions
that enable machine learning
with artificial neural networks"

THE ROYAL SWEDISH ACADEMY OF SCIENCES

John Hopfield



- Ph.D. em Física
- Hopfield Network (1982)
- Energy Minimization Approach

Geoffrey Hinton



- Ph.D. em Inteligência Artificial
- Backpropagation (1986)
- Deep Learning
- Dropout (2012)

THE NOBEL PRIZE IN CHEMISTRY 2024



Illustrations: Niklas Elmehed

**David
Baker**

"for computational
protein design"

**Demis
Hassabis**

"for protein structure prediction"

**John M.
Jumper**

THE ROYAL SWEDISH ACADEMY OF SCIENCES

David Baker



- Ph.D. em Bioquímica
- Diretor do Institute for Protein Design
- Rosetta Software (1990)

Demis Hassabis



- PhD em neurociência cognitiva
- DeepMind (2010)
- AlphaGo (2016)
- AlphaZero (2017)
- AlphaFold (2020)

John Jumper



- Ph.D. em Física e Biologia Computacional
- AlphaFold 1 (2018)
- AlphaFold 2 (2020)

Fim