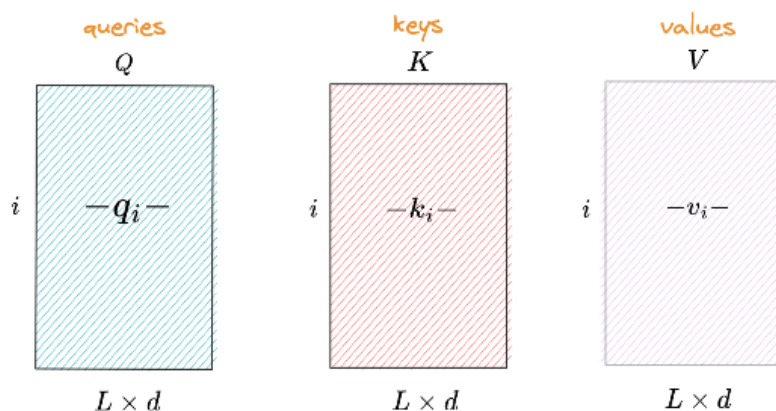


Performer

Attention: Recap

Attention

L Sequence Length Regime of interest: $L \gg d$
 d Embed Dim.



Attention
Matrix

$$A_{i,j} = \underbrace{\exp(q_i^\top k_j)}_{=:\beta_{i,j}} = \text{relevance of } x_j \text{ to } x_i \quad i, j \in \{1, \dots, T\}$$

$$\bar{A}_{i,j} = \underbrace{\exp(q_i^\top k_j) / \sum_{\ell} \exp(q_i^\top k_{\ell})}_{=:\alpha_{i,j}} = \text{normalized relevance of } x_j \text{ to } x_i \quad i, j \in \{1, \dots, T\}$$

$$A = \exp \left(\begin{array}{c|c} Q & K^\top \\ \hline i & k_j \\ -q_i- & -k_j- \\ j & \end{array} \right) \quad \bar{A} = \text{Softmax} \left(\begin{array}{c|c} Q & K^\top \\ \hline i & k_j \\ -q_i- & -k_j- \\ j & \end{array} \right)$$

Attention (cont.)

$$A = \left(\begin{array}{c|c} Q & K^\top \\ \hline i & j \end{array} \right) \begin{pmatrix} \beta_{1,1} & \cdots & \beta_{1,L} \\ \vdots & & \vdots \\ \beta_{L,L} & \cdots & \beta_{L,1} \end{pmatrix} \quad \bar{A} = \text{Softmax} \left(\begin{array}{c|c} Q & K^\top \\ \hline i & j \end{array} \right) = \begin{pmatrix} \alpha_{1,L} & \cdots & \alpha_{1,L} \\ \vdots & & \vdots \\ \alpha_{L,1} & \cdots & \alpha_{L,L} \end{pmatrix}$$

$$\beta_{i,j} = \frac{\alpha_{i,j}}{\sum_{\ell} \alpha_{i,\ell}} \longrightarrow \bar{A} = D^{-1}A \quad D = \begin{pmatrix} \sum_{\ell} \beta_{1,\ell} & & \\ & \ddots & \\ & & \sum_{\ell} \beta_{L,\ell} \end{pmatrix}$$

$$D = A \mathbf{1}_L = Q K^\top \mathbf{1}_L$$

Attention (cont.)

$$Z = \text{Softmax} \left(\begin{array}{c|c} Q & K^\top \\ \hline i & j \end{array} \right) \cdot V$$

$$= \bar{A}V = D^{-1}AV$$

Computation complexity: $O(L^2d)$

(prohibitive for long sequences)

space complexity: $O(L^2)$

Efficient transformers

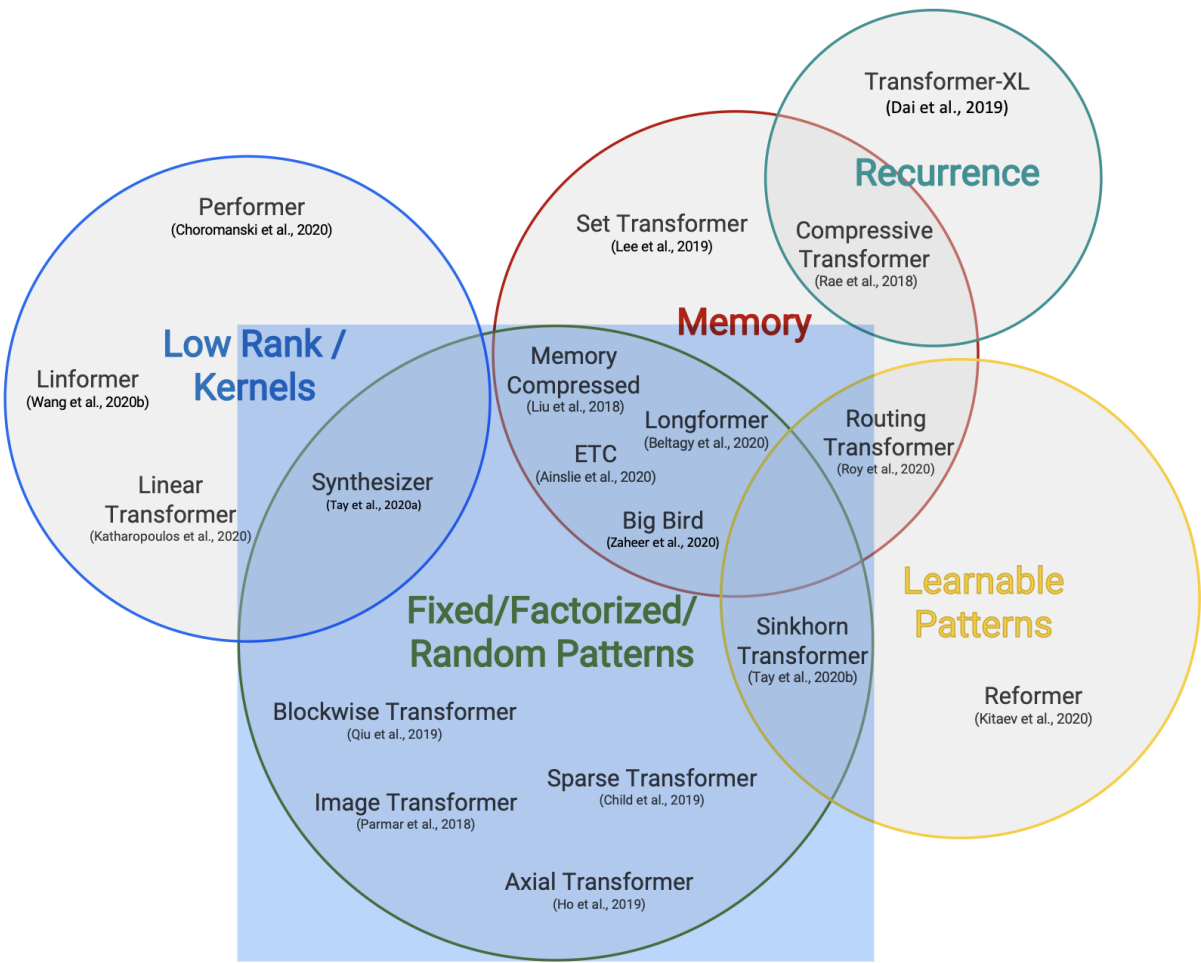
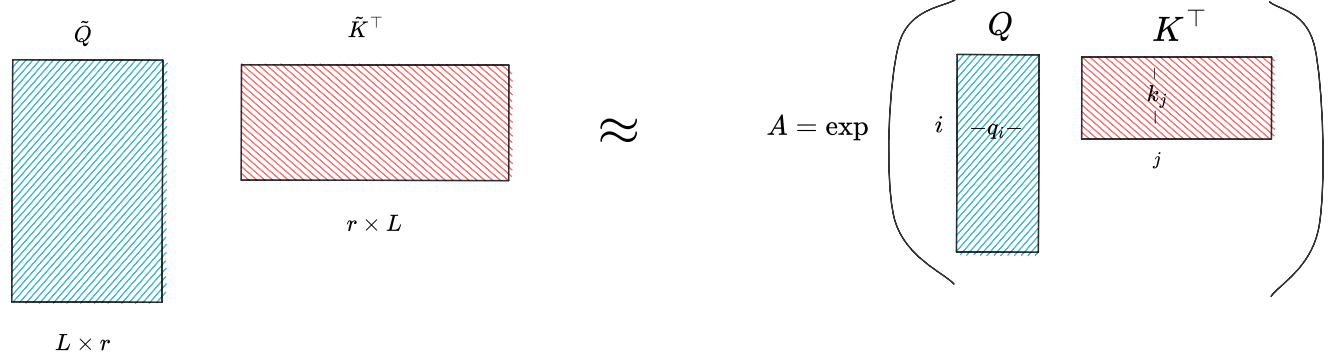
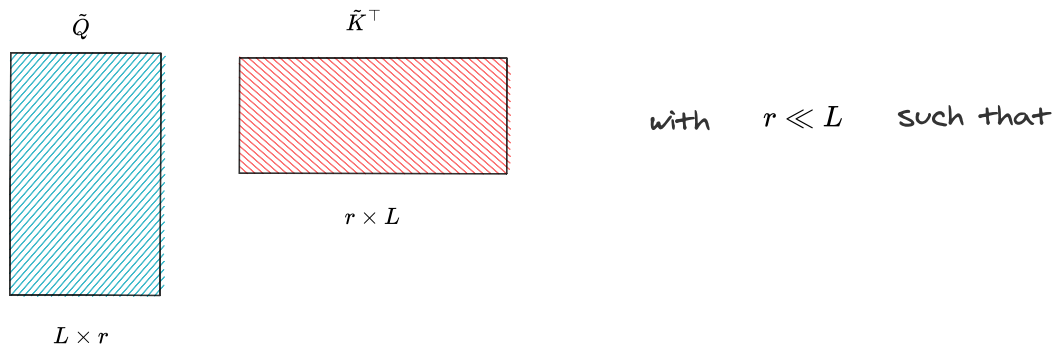


Figure 2: Taxonomy of Efficient Transformer Architectures.

Performer

main idea: use random features to construct



Performer (cont.)

computational and space savings

$$\tilde{\mathcal{Z}} = \tilde{D}^{-1} \tilde{Q} \left(\tilde{K}^\top V \right)$$

\tilde{D}^{-1} $L \times L$ (diag)

\tilde{Q} $L \times r$

\tilde{K}^\top $r \times L$

V $L \times d$

$\tilde{D} = \tilde{Q} \tilde{K}^\top 1_L$

Computation complexity: $O(Lr^2)$

space complexity: $O(Lr)$

Random Features

$$k(x, y) = \exp(x^\top y) \quad \text{kernel/similarity function}$$

$$A_{i,j} = \exp(q_i^\top k_j) = k(q_i, k_j)$$

Construction for $r=1$

draw $w_1, \dots, w_r \sim \mathcal{N}(0, I)$

$$x \xrightarrow{\phi_w} \exp(w^\top x - \|x\|^2/2)$$

Lemma

$$\mathbb{E}[\phi_w(x)^\top \phi_w(y)] = \exp(x^\top y)$$

Amplification ($r>1$)

draw $w_1, \dots, w_r \sim \mathcal{N}(0, I) \quad r = \tilde{\Theta}(d/\epsilon^2)$

$$x \xrightarrow{\phi} \frac{1}{\sqrt{r}}(\phi_{w_1}(x), \dots, \phi_{w_r}(x))$$

Theorem

w.h.p. for all x, y (simultaneously)

$$|\phi(x)^\top \phi(y) - k(x, y)| \leq \epsilon$$

$$\tilde{Q} = \begin{bmatrix} \text{---} & \phi(q_1) & \text{---} \\ & \circ & \\ & \circ & \\ & \circ & \\ \text{---} & \phi(q_L) & \text{---} \end{bmatrix} \quad \tilde{K} = \begin{bmatrix} \text{---} & \phi(k_1) & \text{---} \\ & \circ & \\ & \circ & \\ & \circ & \\ \text{---} & \phi(k_L) & \text{---} \end{bmatrix} \longrightarrow \|\tilde{Q}\tilde{K}^\top - A\|_\infty \leq \epsilon$$

Random Features: Proof

$$k(x, y) = \exp(x^\top y) \quad \text{kernel/similarity function}$$

$$\text{draw } w_1, \dots, w_r \sim \mathcal{N}(0, I)$$

$$x \xrightarrow{\phi_w} \exp(w^\top x - \|x\|^2/2)$$

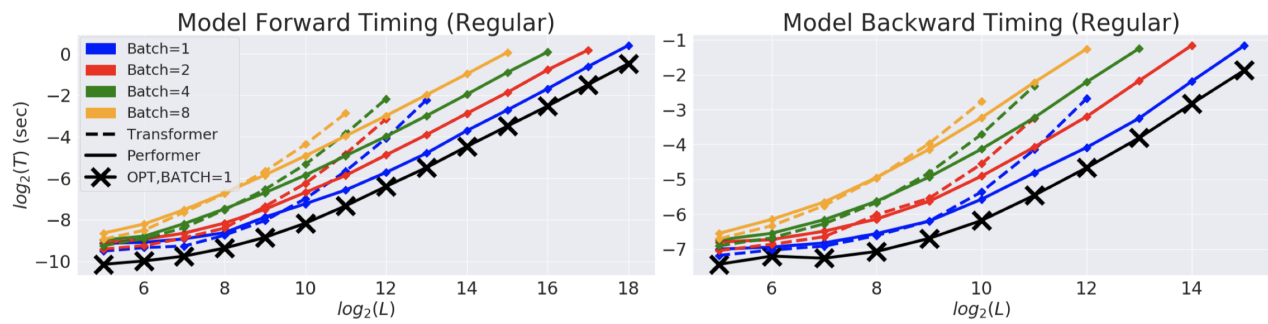
Lemma

$$\mathbb{E}[\phi_w(x)^\top \phi_w(y)] = \exp(x^\top y)$$

Proof sketch

$$\begin{aligned} \mathbb{E}_w[\phi_w(x)^\top \phi_w(y)] &\approx \int_w \phi_w(x) \phi_w(y) \exp(-\|w\|^2/2) dw \\ &= \int_w \exp(-\|x\|^2/2) \exp(-\|y\|^2/2) \exp(w^\top (x + y)) \exp(-\|w\|^2/2) dw \\ &= \int_w \exp(\|w - (x + y)\|^2) \exp(x^\top y) dw \\ &= \exp(x^\top y) \underbrace{\int_w \exp(\|w - (x + y)\|^2) dw}_{\approx 1} \end{aligned}$$

Experiments



TrEMBL: predicting interactions among groups of proteins by concatenating protein sequences

