



Towards New Transformers' Revolution

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"Rethinking Attention with Performers" Google Al Blog Post code

to appear @ ICLR 2021, oral presentation, top **40** accepted papers

Why we need better Memorization & Attention in ML?



Fig. 1 DeepMind policy navigating simply by "sight".



• *Lifelong Learning* requires going beyond purely-reactive Robotics tasks:

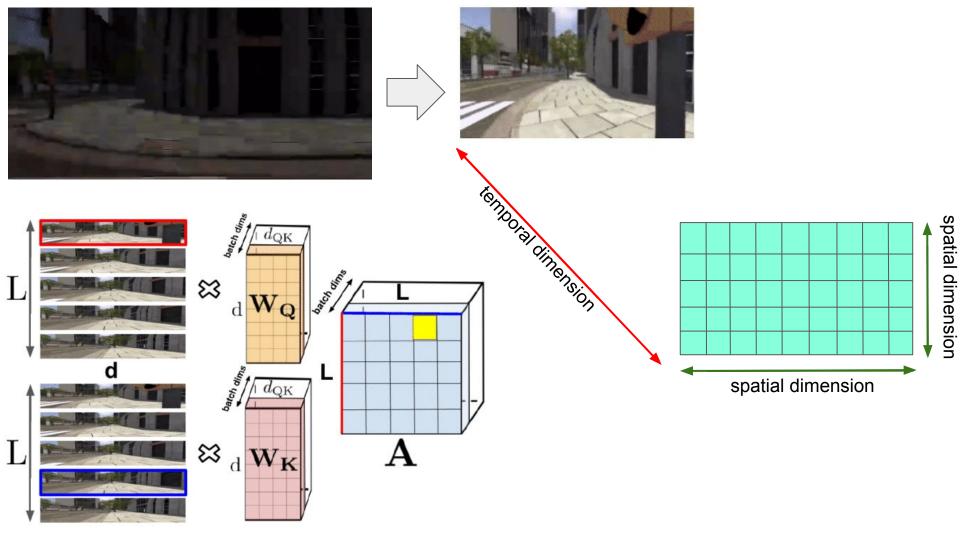
"Developmental Robotics: A Complex Dynamical System with Several Spatiotemporal scales."

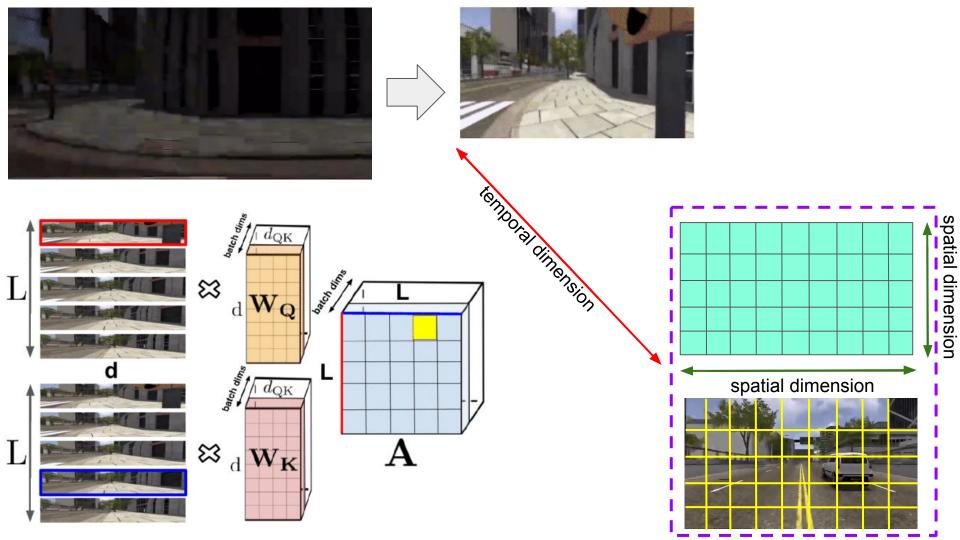
- Memory is a key to AI and currently existing sequential recurrent architectures fail to memorize well.
- Lets learn how to attend to the world is <u>Attention</u> all you need ? Different attention dimensions: **spatial** & **temporal**.
- Standard attention mechanisms are effectively parallelizable and avoid catastrophic forgetting, but are not scalable.

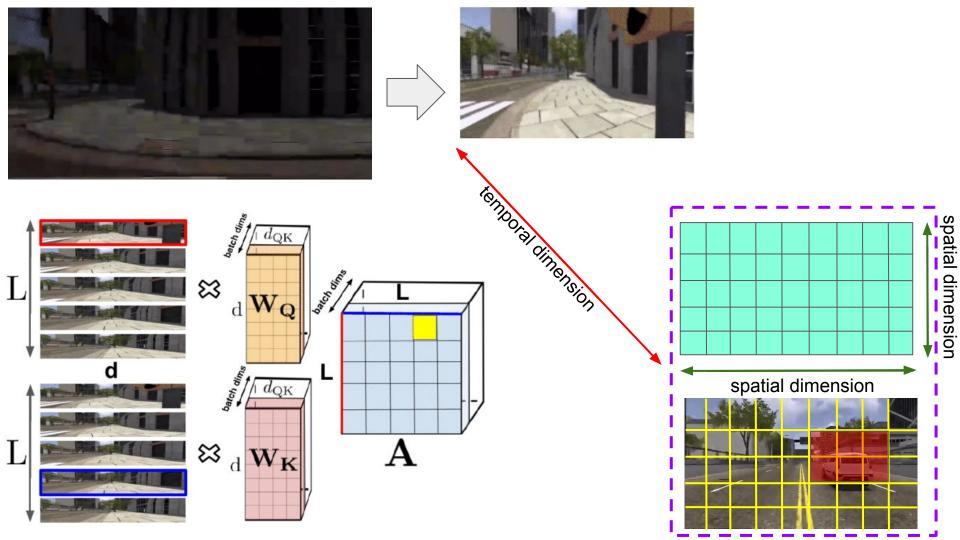
"It uses more memory and more computation per real interaction..." [DeepMind nav by sight]

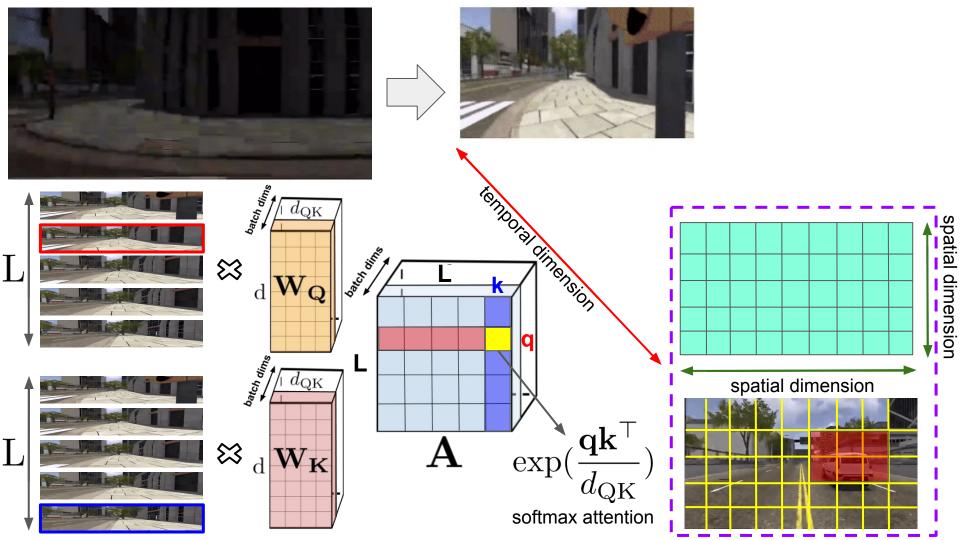
• Lifelong Learning Robotics requires long range contexts with **no attention priors**.

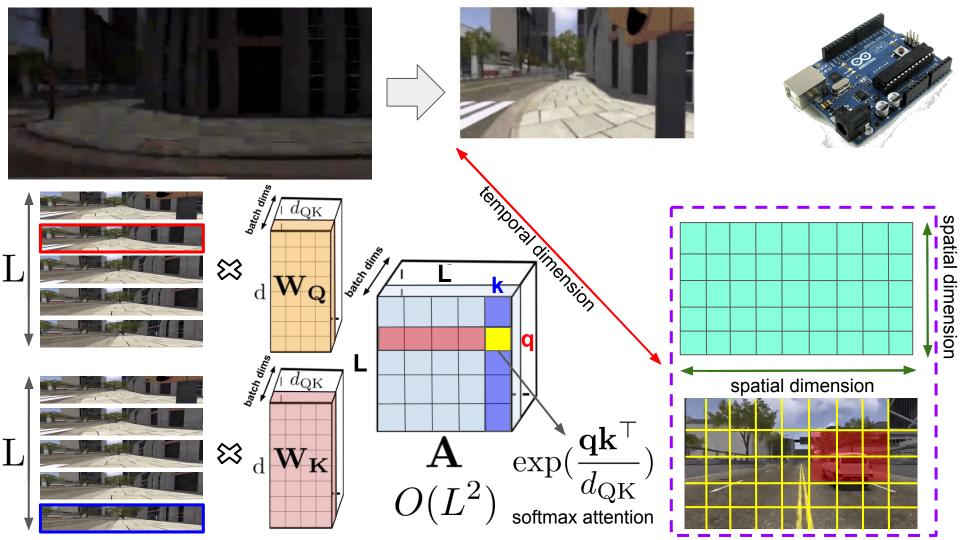
Fig. 2 Robotic arm "solving" Hanoi towers.

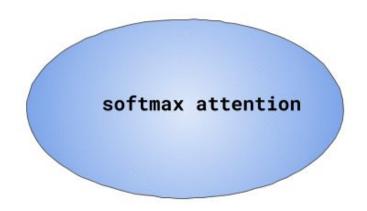












Performers = Linear Attention Modules Capable of Modeling Different Attention Kernels (Also **SOFTMAX** !)

Performers performing on the Long Range Arena Long Range Arena Paper



Model	ListOps	Text	Retrieval	Image	Pathfinder	Path-X	Avg
Transformer	36.37	64.27	57.46	42.44	71.40	FAIL	54.39
Local Attention	15.82	52.98	53.39	41.46	66.63	FAIL	46.06
Sparse Trans.	17.07	63.58	59.59	44.24	71.71	FAIL	51.24
Longformer	35.63	62.85	56.89	42.22	69.71	FAIL	53.46
Linformer	35.70	53.94	52.27	38.56	76.34	FAIL	51.36
Reformer	37.27	56.10	53.40	38.07	68.50	FAIL	50.67
Sinkhorn Trans.	33.67	61.20	53.83	41.23	67.45	FAIL	51.39
Synthesizer	<u>36.99</u>	61.68	54.67	41.61	69.45	FAIL	52.88
BigBird	36.05	64.02	<u>59.29</u>	40.83	74.87	FAIL	55.01
Linear Trans.	16.13	65.90	53.09	42.34	75.30	FAIL	50.55
Performer	18.01	<u>65.40</u>	53.82	<u>42.77</u>	77.05	FAIL	51.41
Task Avg (Std)	29 (9.7)	61 (4.6)	55 (2.6)	41 (1.8)	72 (3.7)	FAIL	52 (2.4)

Table 1: Experimental results on Long-Range Arena benchmark. Best model is in boldface and second best is underlined. All models do not learn anything on Path-X task, contrary to the Pathfinder task and this is denoted by FAIL. This shows that increasing the sequence length can cause seriously difficulties for model training. We leave Path-X on this benchmark for future challengers but do not include it on the Average score as it has no impact on relative performance.

	Steps per second					Peak Memory Usage (GB)				
Model	1 K	2K	3K	4K	1K	2K	3K	4K		
Transformer	8.1	4.9	2.3	1.4	0.85	2.65	5.51	9.48		
Local Attention	9.2 (1.1x)	8.4 (1.7x)	7.4 (3.2x)	7.4 (5.3x)	0.42	0.76	1.06	1.37		
Linformer	<u>9.3</u> (1.2x)	9.1 (1.9x)	8.5 (3.7x)	7.7 (5.5x)	0.37	0.55	0.99	0.99		
Reformer	4.4(0.5x)	2.2 (0.4x)	1.5(0.7x)	1.1(0.8x)	0.48	0.99	1.53	2.28		
Sinkhorn Trans	9.1 (1.1x)	7.9 (1.6x)	6.6 (2.9x)	5.3 (3.8x)	0.47	0.83	1.13	1.48		
Synthesizer	8.7 (1.1x)	5.7 (1.2x)	6.6 (2.9x)	1.9(1.4x)	0.65	1.98	4.09	6.99		
BigBird	7.4 (0.9x)	3.9 (0.8x)	2.7 (1.2x)	1.5(1.1x)	0.77	1.49	2.18	2.88		
Linear Trans.	9.1 (1.1x)	<u>9.3</u> (1.9x)	8.6(3.7x)	<u>7.8</u> (5.6x)	0.37	<u>0.57</u>	0.80	1.03		
Performer	9.5 (1.2x)	9.4 (1.9x)	8.7 (3.8x)	8.0 (5.7x)	0.37	0.59	<u>0.82</u>	1.06		

Table 2: Benchmark results of all Xformer models with a consistent batch size of 32 across all models. We report relative speed increase/decrease in comparison with the vanilla Transformer in brackets besides the steps per second. Memory usage refers to per device memory usage across each TPU device. Benchmarks are run on 4x4 TPU V3 Chips.

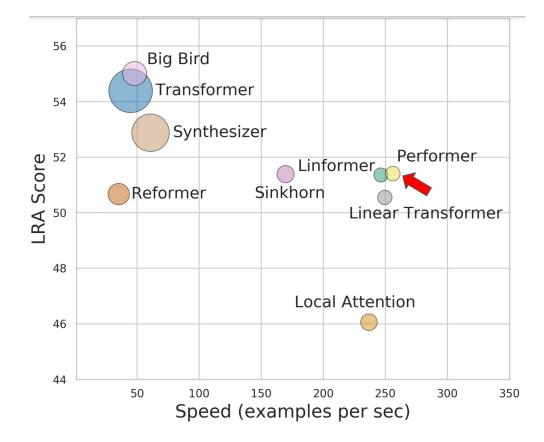


Figure 3: Performance (y axis), speed (x axis), and memory footprint (size of the circles) of different models.

Performers in Vision

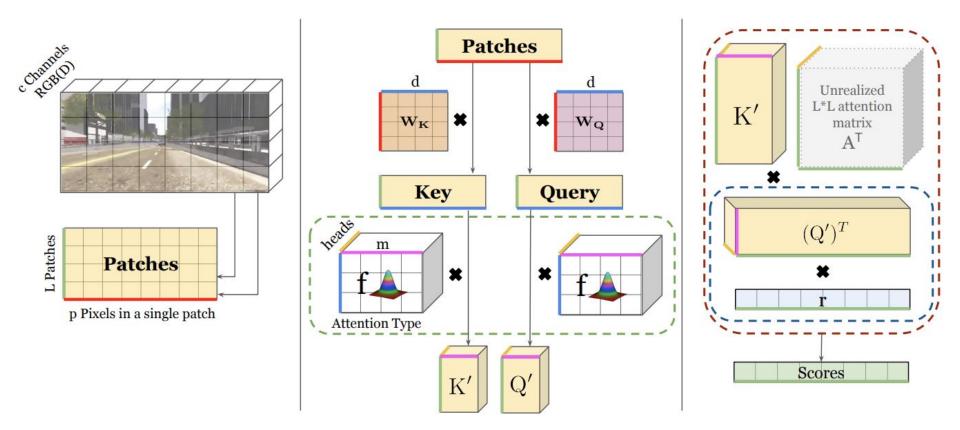
<u>Tokens-to-Token ViT: Training Vision Transformers from Scratch on ImageNet</u> Yuan et al.

Models	Top1-Acc (%)	Params (M)	ns MAC (G)		
ResNet50 [15]	76.2	25.5	4.3		
ResNet50*	79.1	25.5	4.3		
T2T-ViT-14	80.6	21.4	4.8		
T2T-ViT _t -14	80.7	21.5	5.2		
ResNet101 [15]	77.4	44.6	7.9		
ResNet101*	79.9	44.6	7.9		
T2T-ViT-19	81.2	39.0	8.0		
T2T-ViT _t -19	81.4	39.0	8.4		
ResNet152 [15]	78.3	60.2	11.6		
ResNet152*	80.8	60.2	11.6		
T2T-ViT-24	81.8	63.9	12.6		
T2T-ViT _t -24	82.2	64.1	13.2		

Models	Tokens-to-Token module				T2T-ViT backbone			Model size	
	T2T Transformer	Depth	Hidden dim	MLP size	Depth	Hidden dim	MLP size	Params (M)	MACs (G)
ViT-S/16 [12]	-	-	-	-	8	786	2358	48.6	10.1
ViT-B/16 [12]	-		-	-	12	786	3072	86.8	17.6
ViT-L/16 [12]	-	1.77	-	-	24	1024	4096	304.3	63.6
T2T-ViT _t -14	Transformer	2	64	64	14	384	1152	21.5	5.2
T2T-ViT _t -19	Transformer	2	64	64	19	448	1344	39.0	8.4
T2T-ViT _t -24	Transformer	2	64	64	24	512	1536	64.1	13.2
T2T-ViT-14	Performer	2	64	64	14	384	1152	21.4	4.8
T2T-ViT-19	Performer	2	64	64	19	448	1344	39.0	8.0
T2T-ViT-24	Performer	2	64	64	24	512	1536	63.9	12.6
T2T-ViT-7	Performer	2	64	64	8	256	512	4.2	0.9
T2T-ViT-10	Performer	2	64	64	10	256	512	5.6	1.2
T2T-ViT-12	Performer	2	64	64	12	256	512	6.8	1.4

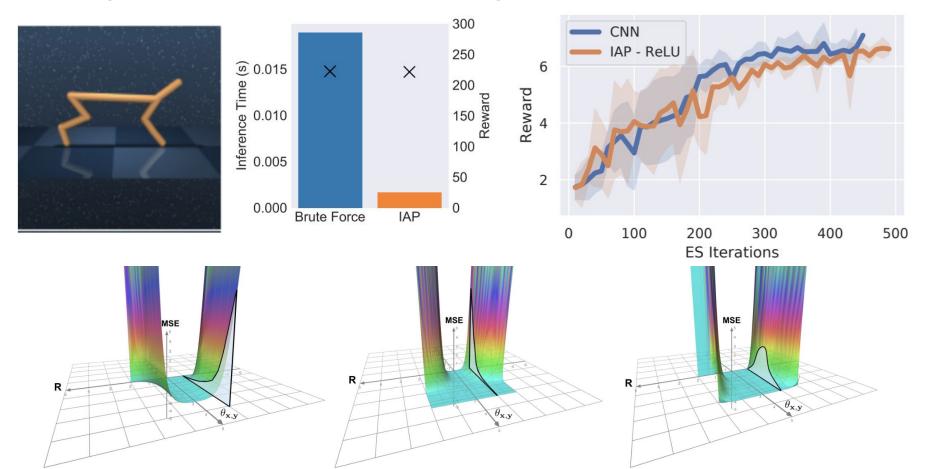
Performers for Vision in Reinforcement Learning (IAP)

Unlocking Pixels for Reinforcement Learning via Implicit Attention Choromanski et al.



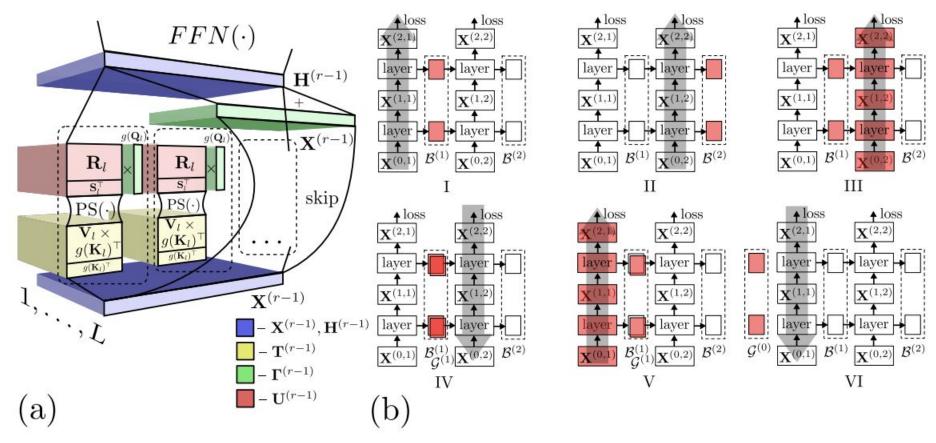
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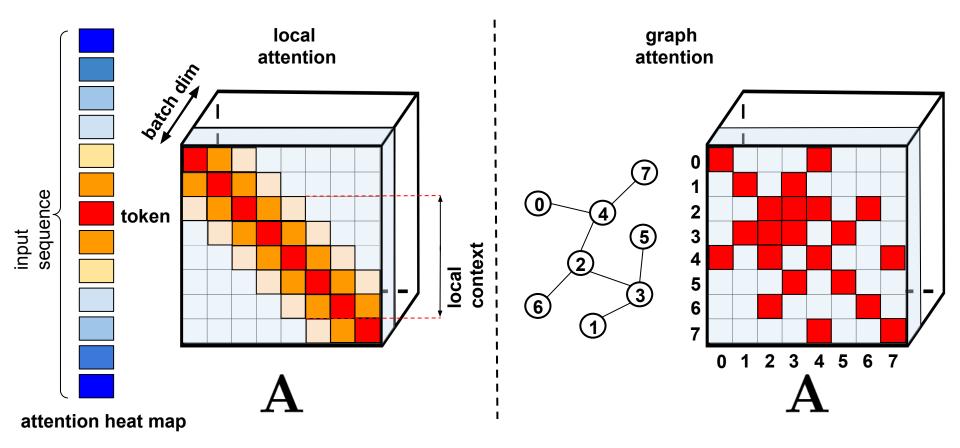


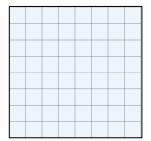
Sub-Linear Memory: How To Make Performers Slim

Likhosherstov et al.

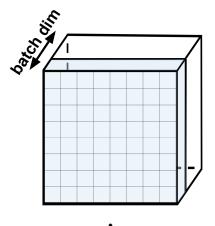


What we Do NOT DO: Sparsifying Attention

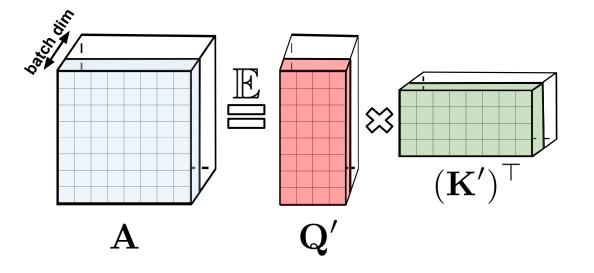


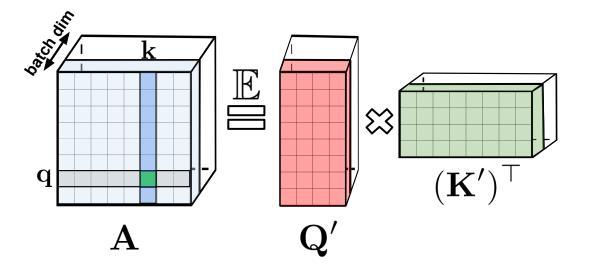


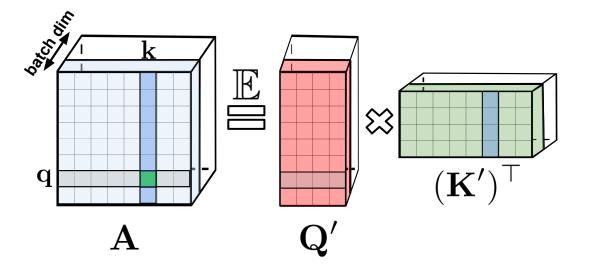
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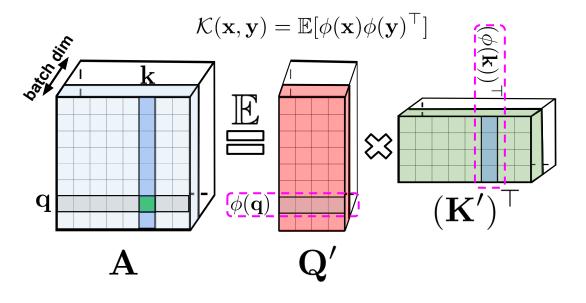


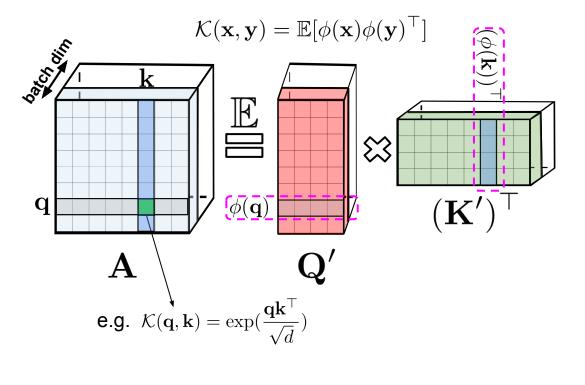


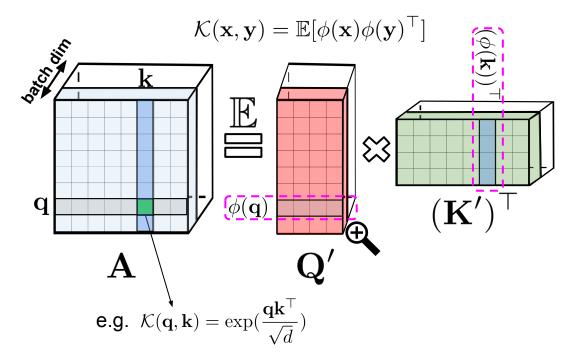


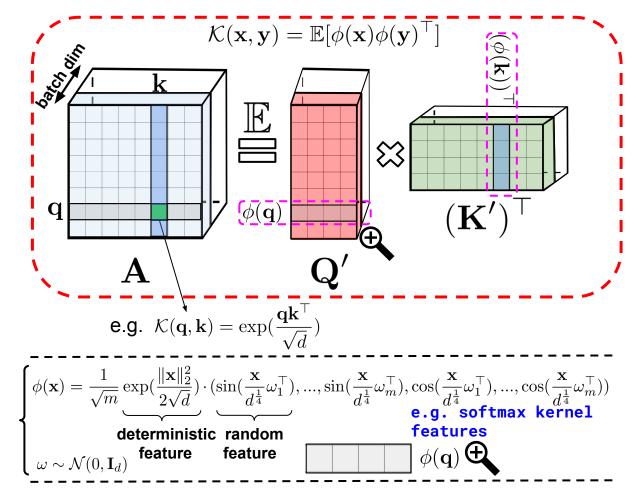




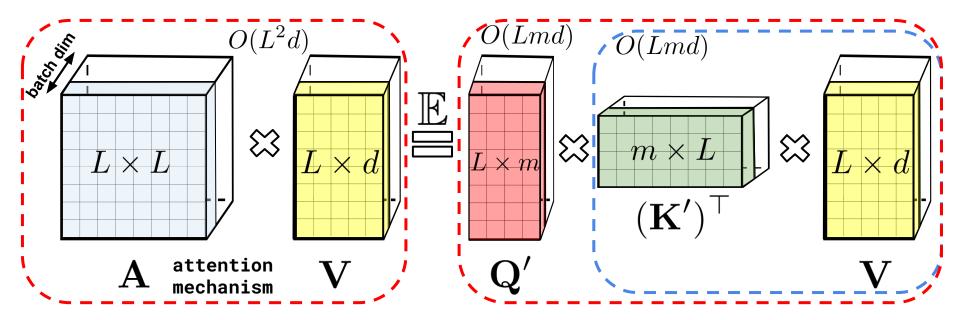




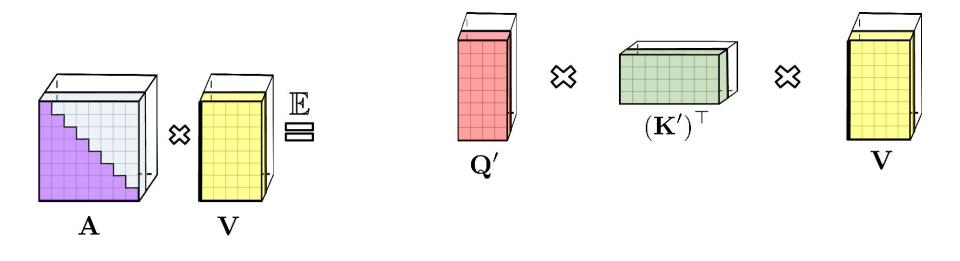




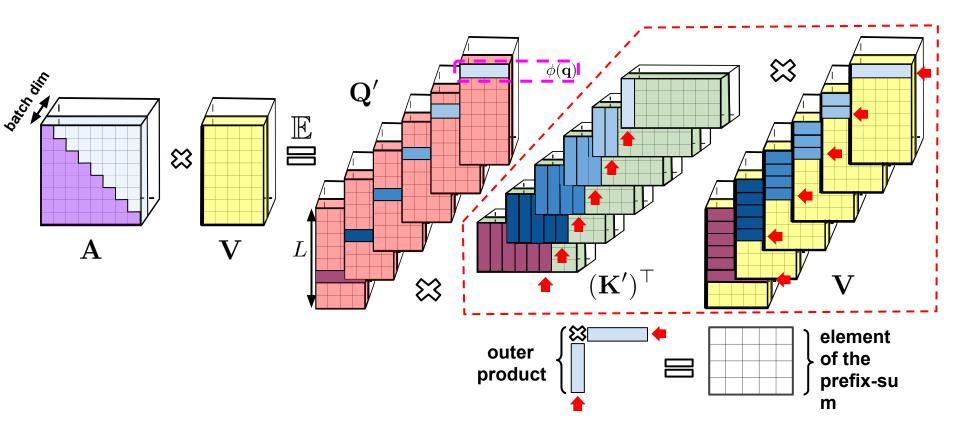
Associativity for Speedups and Space Compression



Causal Transformers as Prefix-Sums Calculators

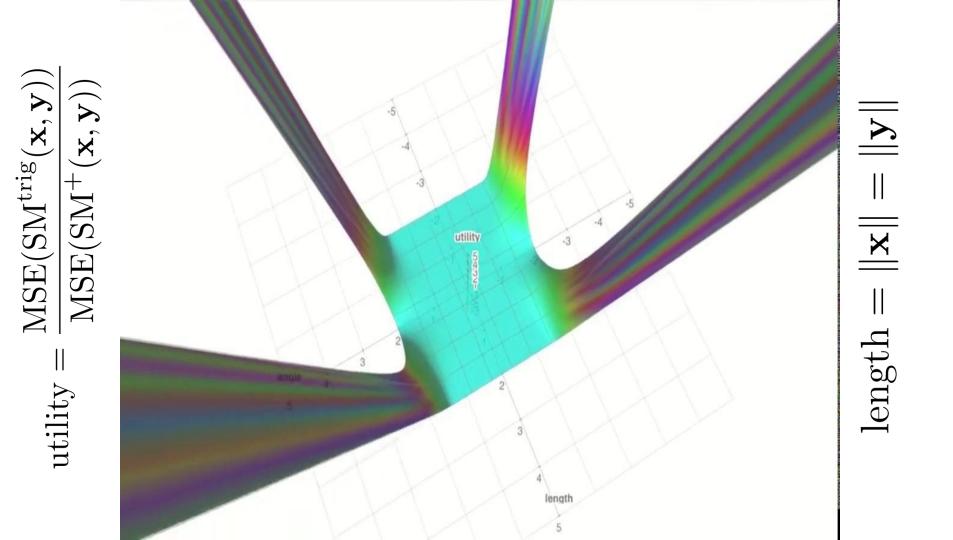


Causal Transformers as Prefix-Sums Calculators

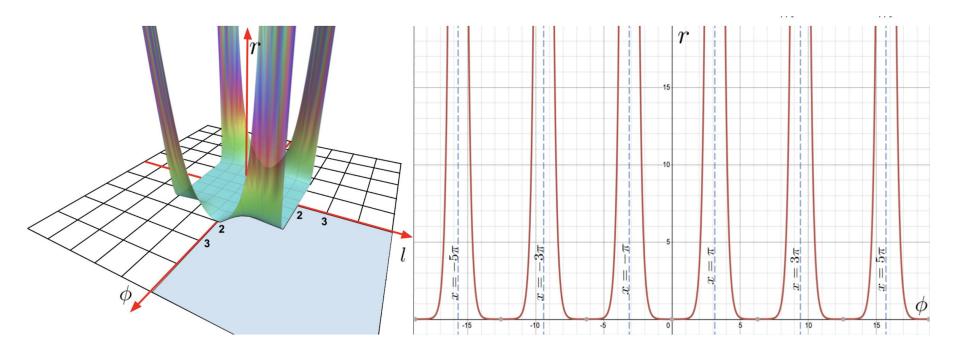


Why this is not the end of the story ?

FAVOR+

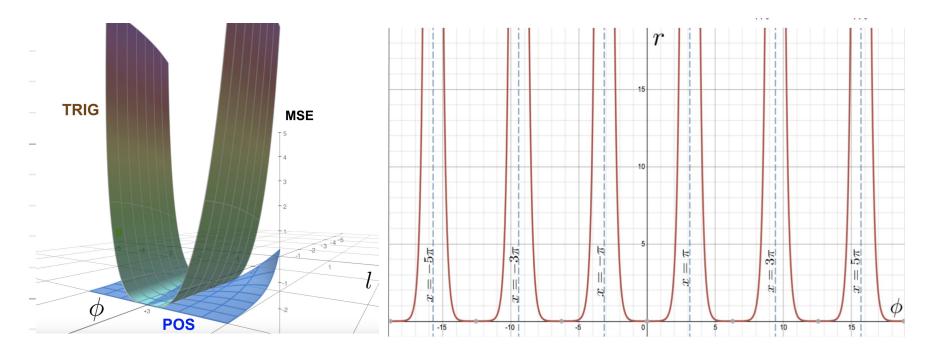


Do we need to go beyond trigonometric features ?

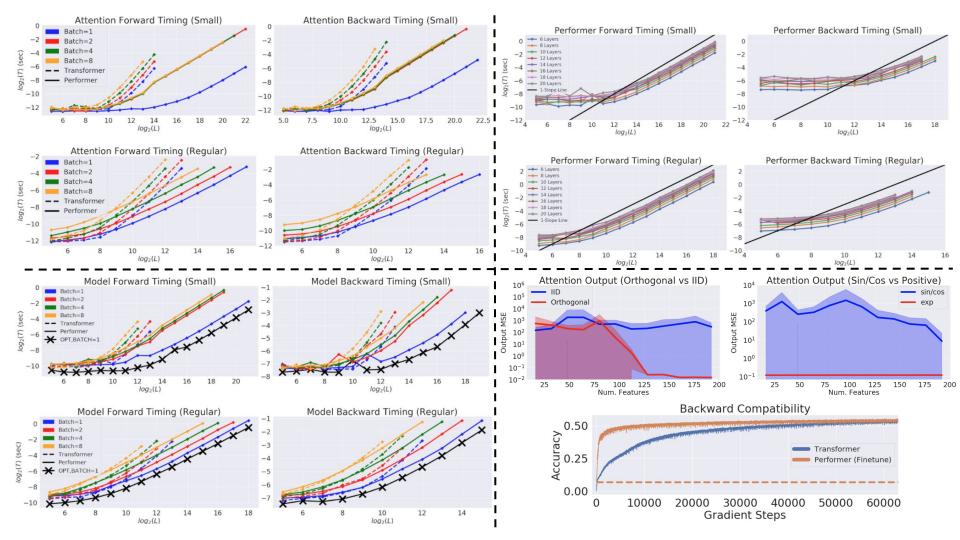


Left: Symmetrized (around origin) utility function r (defined as a ratio of the mean squared errors of estimators built on: trigonometric and positive random features) as a function of the angle φ (in radians) between input feature vectors and their lengths I. Larger values indicate regions of (φ , I)-space with better performance of positive random features. We see that for critical regions with φ large enough (small enough softmax-kernel values) our method is arbitrarily more accurate than trigonometric random features. Plot presented for domain [$-\pi$, π] × [-2, 2]. **Right:** The slice of function r for fixed I = 1 and varying angle φ .

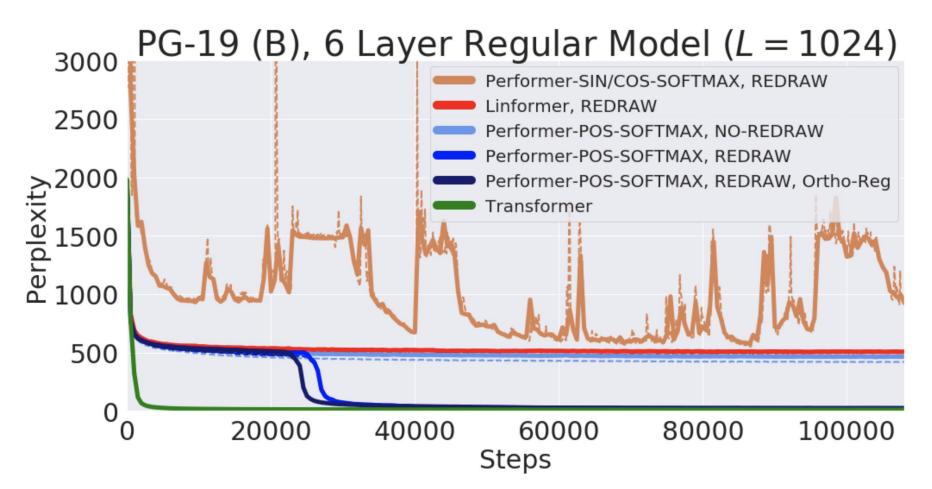
Do we need to go beyond trigonometric features ?



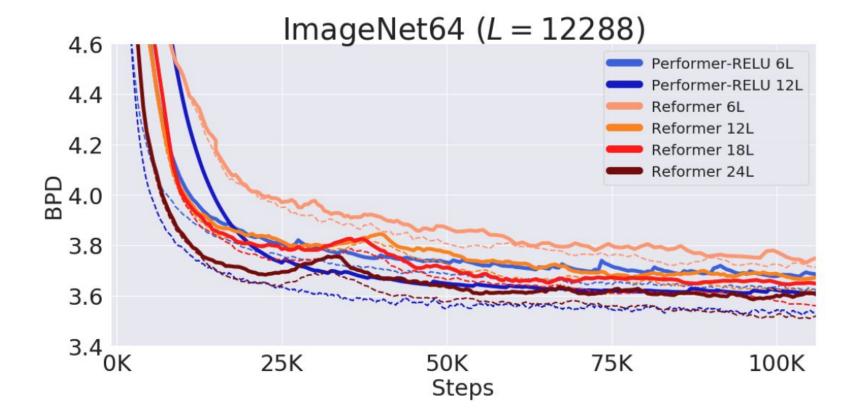
Left: Comparison of the mean squared errors (MSEs) of the estimators applying trigonometric random features (TRIG) and the one leveraging the mechanism of positive random features (POS) in the region of small softmax-kernel values. **Right:** The slice of function r for fixed I = 1 and varying angle φ .



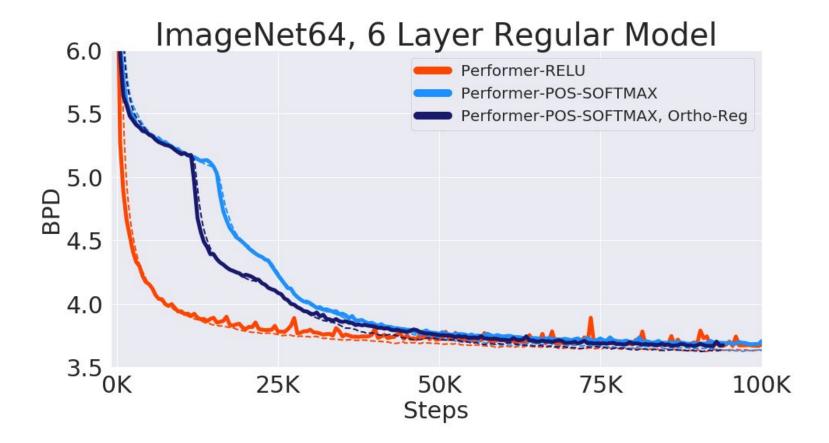
Positive vs trigonometric random features in practice



Performers on ImageNet64 - pixel predictions models



Performers on ImageNet64 - Approx. Softmax vs ReLU

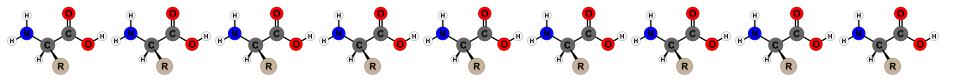


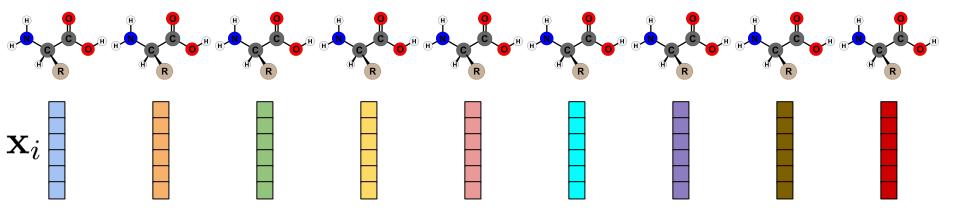


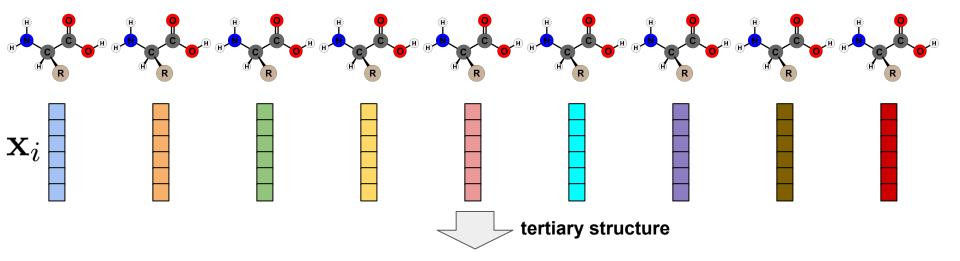
Performers for Protein Design

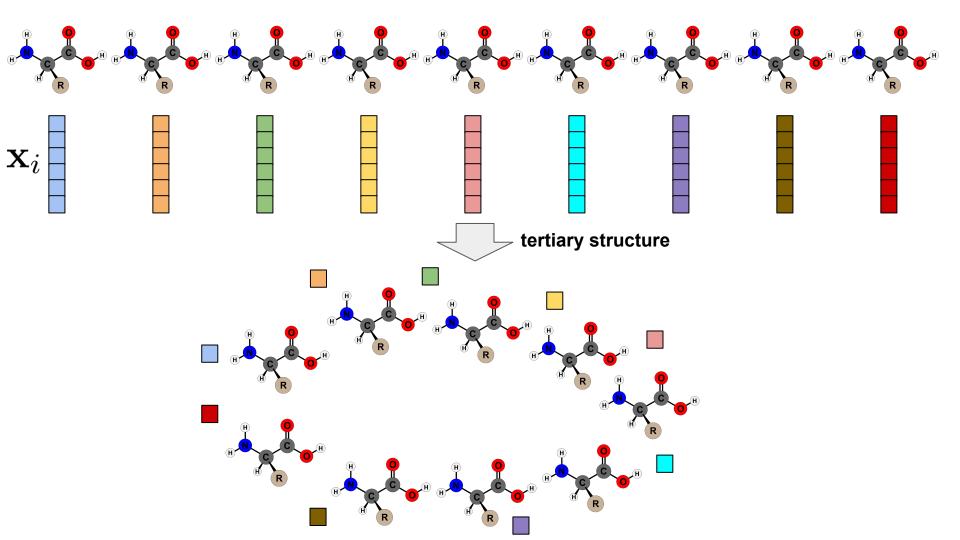


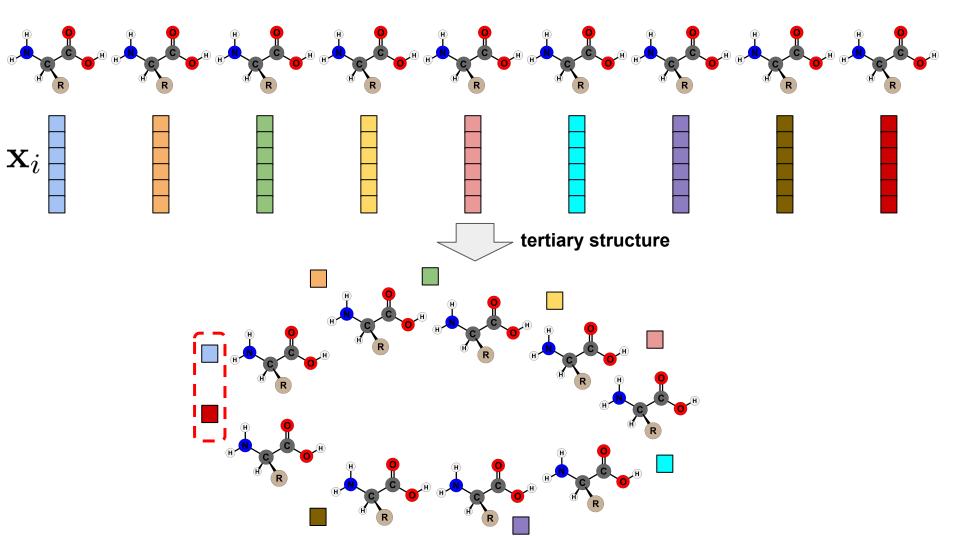
Performers for Protein Design







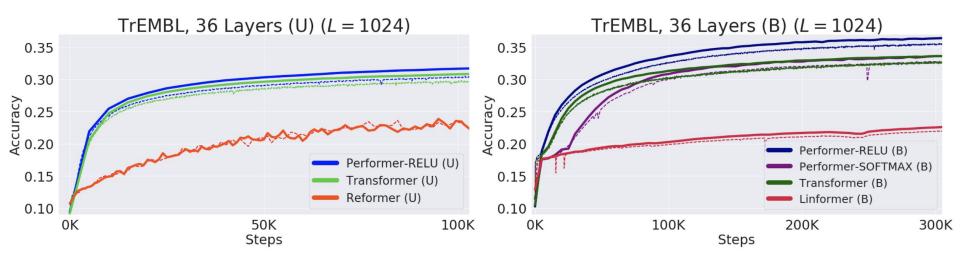




What we have already done...

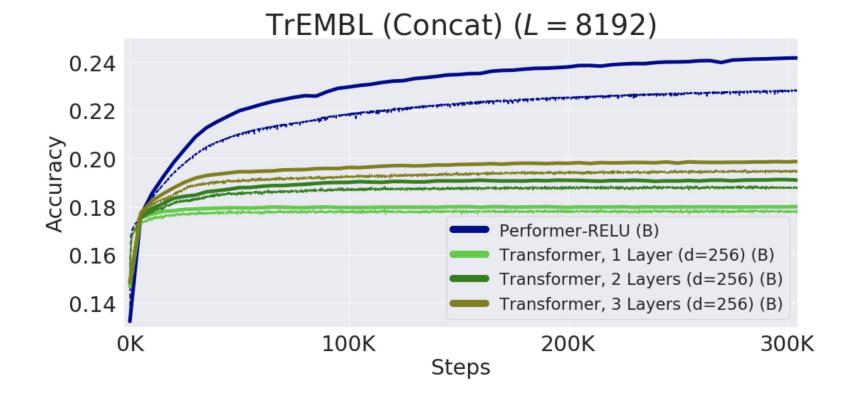


Performers on moderate-size biological sequences

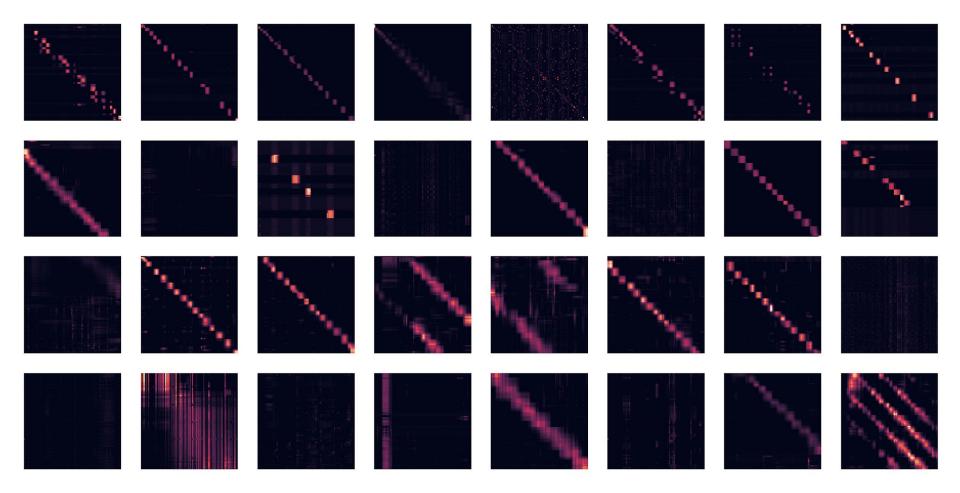


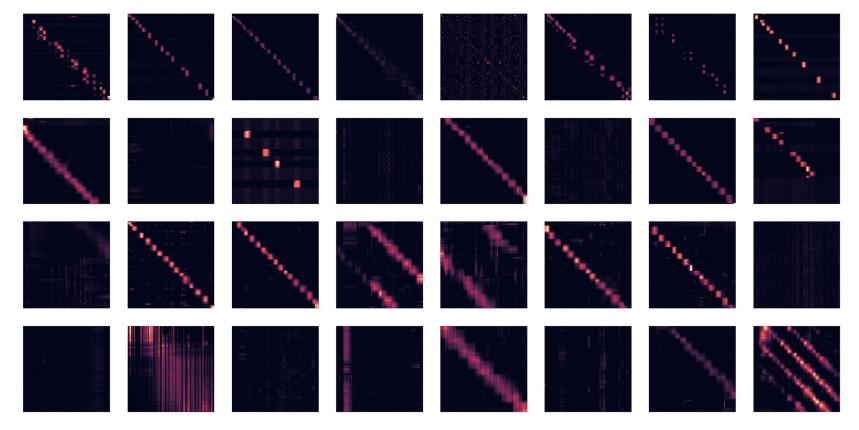
Train = Dashed, Validation = Solid, Unidirectional = (U), Bidirectional = (B). For TrEMBL, we used the exact same model parameters (nheads, nlayers, dff, d) = (8, 36, 1024, 512) from (Madani et al., 2020) for all runs. For fairness, all TrEMBL experiments used 16x16 TPU-v2's. Batch sizes were maximized for each separate run given the compute constraints. Hyperparameters & extended results including dataset statistics, out of distribution evaluations, and visualizations will appear soon in the extended version of the paper.

Performers on long biological sequences: towards modeling complexes of proteins - proof of concept

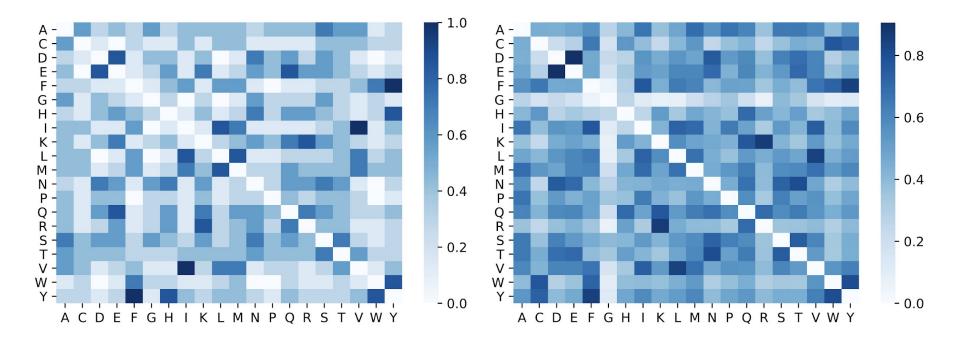


What do Performers attend to ?





We show the attention matrices for the first 4 layers and all 8 heads (each row is a layer, each column is head index, each cell contains the attention matrix across the entire BPT1_BOVIN protein sequence). Note that many heads show a diagonal pattern, where each node attends to its neighbors, and some heads show a vertical pattern, where each head attends to the same fixed positions.



Amino acid similarity matrix estimated from attention matrices aggregated across a small subset of sequences, as described in Vig et al. (Vig et al., 2020). The sub-figures correspond respectively to: (1) the normalized BLOSUM matrix, (2) the amino acid similarity estimated via a trained Performer model. Note that the Performer recognizes highly similar amino acid pairs such as (D, E) and (F, Y).

Thank you for the Attention !

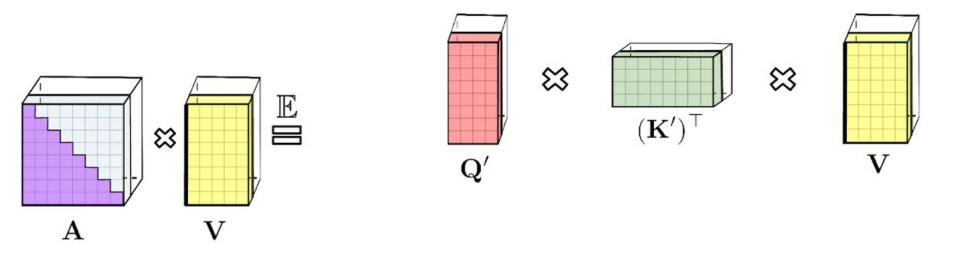


Fig. Linearized softmax causal attention as a prefix-sum computation engine.