

2023 Winter Lab Seminar Big Bird: Transformers for Longer Sequences (2020, NeurIPS)

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- Transformers [91], such as BERT [22, 63],
 - wildly successful for a wide variety of Natural Language
 Processing (NLP) tasks and consequently are mainstay of modern
 NLP research.
 - versatility and robustness -> wide-scale adoption of Transformers
 - diverse range of sequence based tasks as a seq2seq model for translation [91], summarization [66], generation [15], etc. or as a standalone encoders for sentiment analysis [83], POS tagging [65], machine reading comprehension [93], etc. and it is known to vastly outperform previous sequence models like LSTM [37].

Transformers [91]



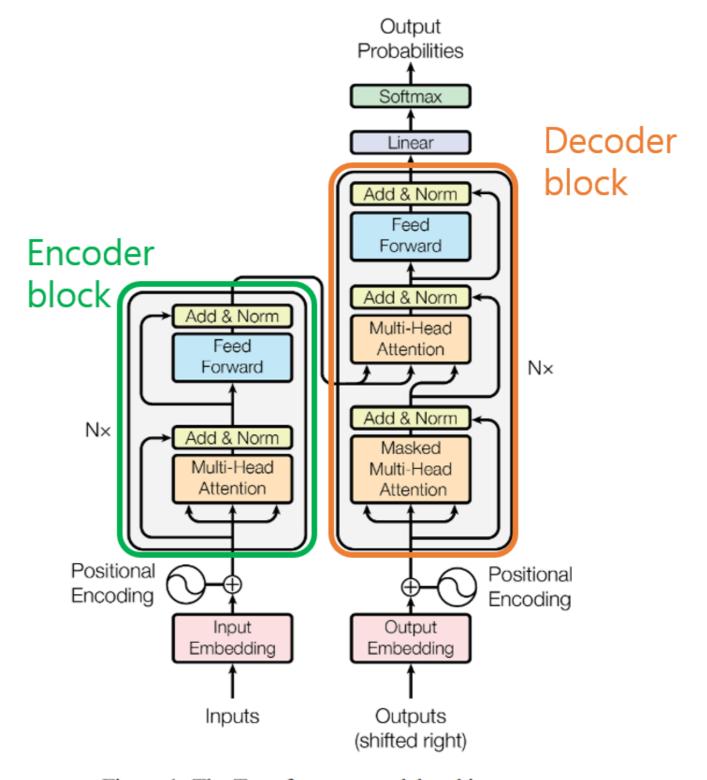


Figure 1: The Transformer - model architecture.

- key innovation in Transformers: introduction of a self-attention mechanism
 - can be evaluated in parallel for each token of the input sequence,
 - eliminating the sequential dependency in recurrent neural networks, like LSTM
 - Parallelism -> 1. leverage the full power of modern SIMD hardware accelerators like GPUs/TPUs, 2. facilitating training of NLP models on datasets of unprecedented size

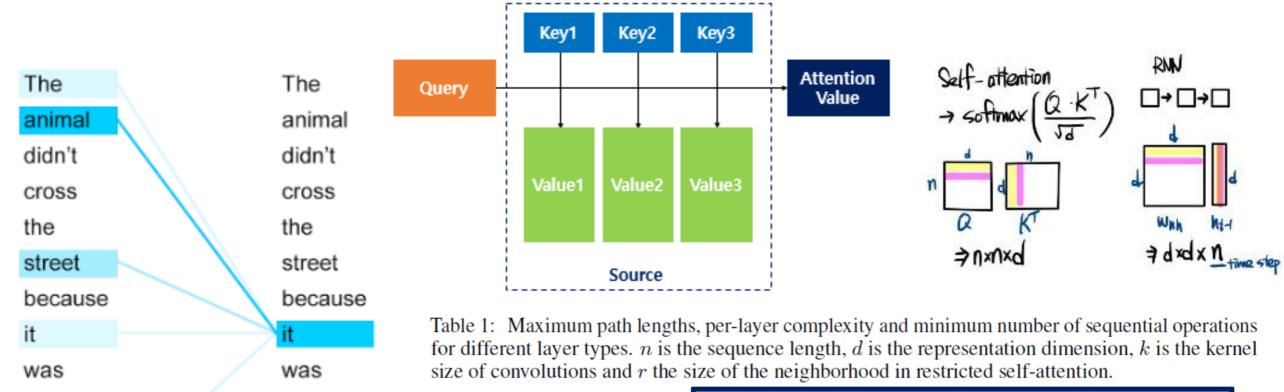
- Transformers [91], such as BERT [22, 63],
 - Self-attention

too

tired

too

tired



Layer Type	Complexity per Layer	Sequential	Maximum Path Length
7 71	1 71 7	Operations	
Self-Attention	$O(n^2 \cdot d)$	O(1)	O(1)
Recurrent	$O(n \cdot d^2)$	O(n)	O(n)
Convolutional	$O(\hat{k} \cdot n \cdot \hat{d}^2)$	O(1)	$O(log_k(n))$
Self-Attention (restricted)	$O(r \cdot n \cdot d)$	O(1)	O(n/r)
			_

- self-attention mechanism
 - Operation: allowing each token in the input sequence to attend independently to every other token in the sequence
 - Problem: full self-attention
 - quadratic in the sequence length => computational and memory requirement
 - Limitation: input sequences of length 512 tokens
 - QA [60], document classification, etc => reduces its direct applicability

- theoretical understanding about self-attention and Transformers
 - 1. What aspects of the self-attention model are <u>necessary</u> for its performance?
 - 2. What can we say about the <u>expressivity</u> of Transformers and similar models?

- Pérez et al. [72]
 - full transformer is Turing Complete
 - (i.e. can simulate a full Turing machine)
 - 1. Can we achieve the empirical benefits of a fully quadratic selfattention scheme using fewer inner-products?
 - 2. Do these sparse attention mechanisms preserve the expressivity and flexibility of the original network?

- interesting attempts, alleviating the quadratic dependency of Transformers
- 1. embraces the length limitation and develops method around it
 - Simplest way: employ sliding window [93]
 - general most work: call transformer block multiple time with different contexts each time
 - SpanBERT [42], ORQA [54], REALM [34], RAG [57]
 - Limitation: require significant engineering efforts (like back prop through large scale nearest neighbor search) and are hard to train

- SpanBERT [42]
 - Improving Pre-training by Representing and Predicting Spans

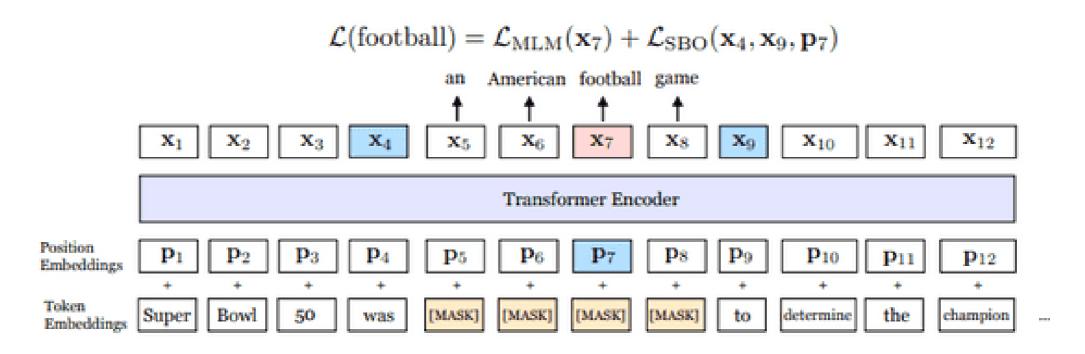
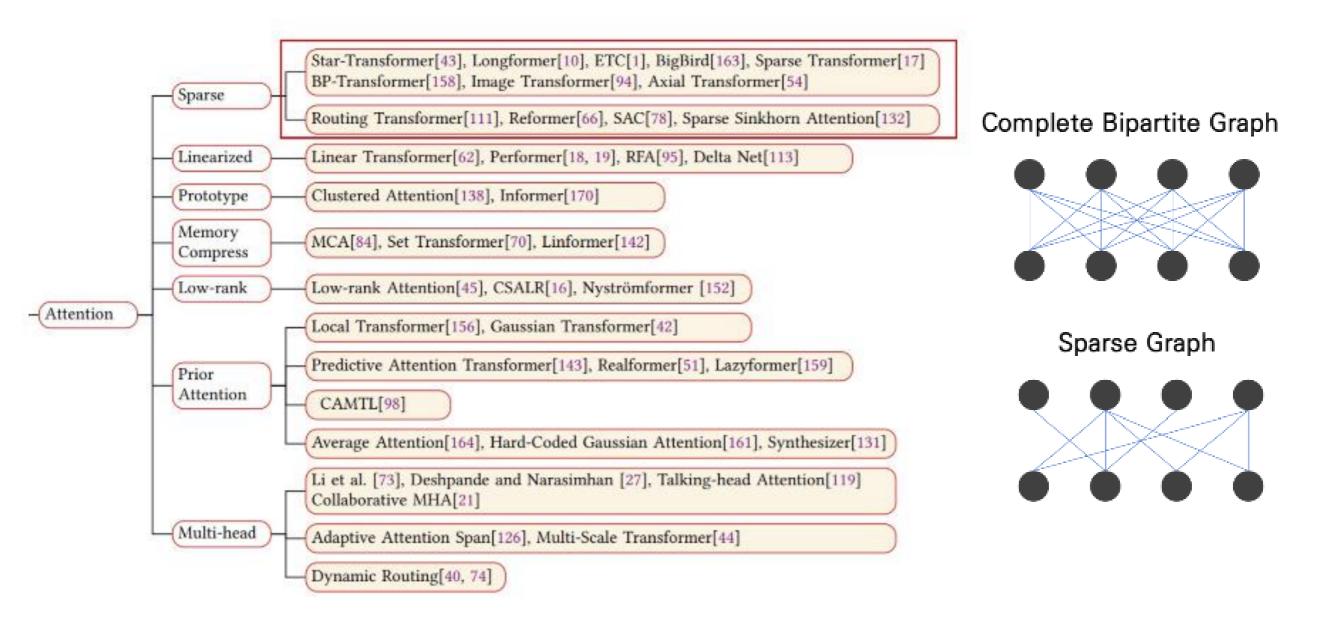


Figure 1: An illustration of SpanBERT training. In this example, the span an American football game is masked. The span boundary objective then uses the boundary tokens was and to to predict each token in the masked span.

2. approaches that do not require full attention



2. approaches that do not require full attention

- Dai et al. [21], Sukhbaatar et al. [82], Rae et al. [74]
 - auto-regresive models
 - work well for left-to-right language modeling but suffer in tasks which require bidirectional context.
- Child et al. [16]
 - sparse model that reduces the complexity to $O(n \sqrt{n})$
- Kitaev et al. [49]
 - reduced the complexity to O(n log(n)) by using LSH to compute nearest neighbors. (Reformer)
- Longformer [8]: introduced a localized sliding window based mask with few global mask to reduce computation and extended BERT to longer sequence based tasks. 13

- 2. approaches that do not require full attention
- Extended Transformers Construction [4]
 - designed to encode structure in text for transformers
 - global tokens was used extensively by them to achieve their goals

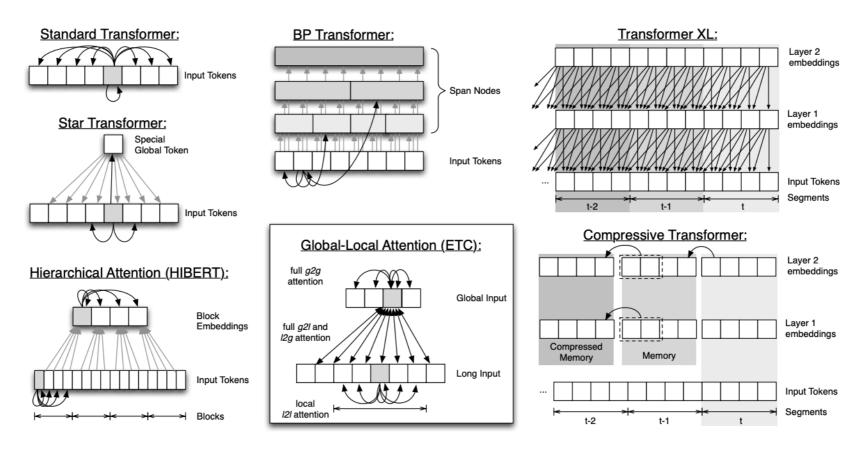
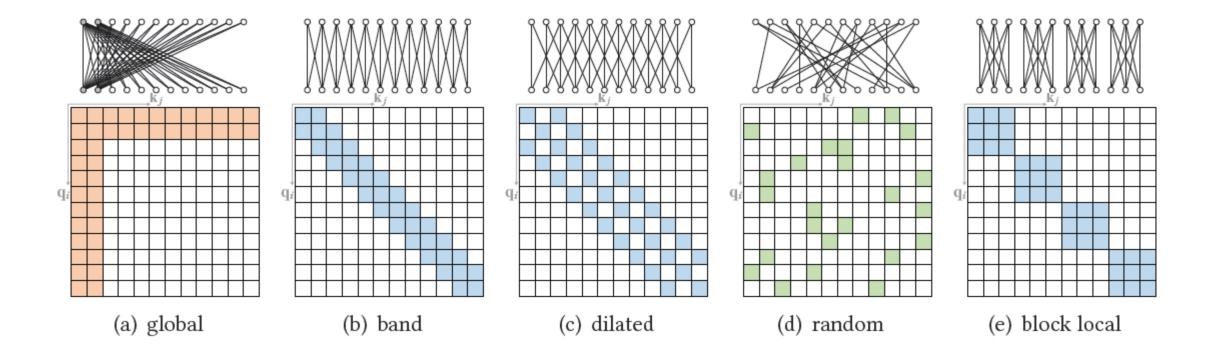


Figure 1: An illustration of mechanisms to scale attention to long inputs, including our proposed model, ETC.

- 2. approaches that do not require full attention
 - A Survey of Transformers(Lin, Tianyang, et al. Al Open (2022))
 - Sparse attention
 - Computational cost: $O(n^2) \rightarrow O(n)$
 - Long range Dependency: Global attention
 - Method
 - Position-based
 - Content-based

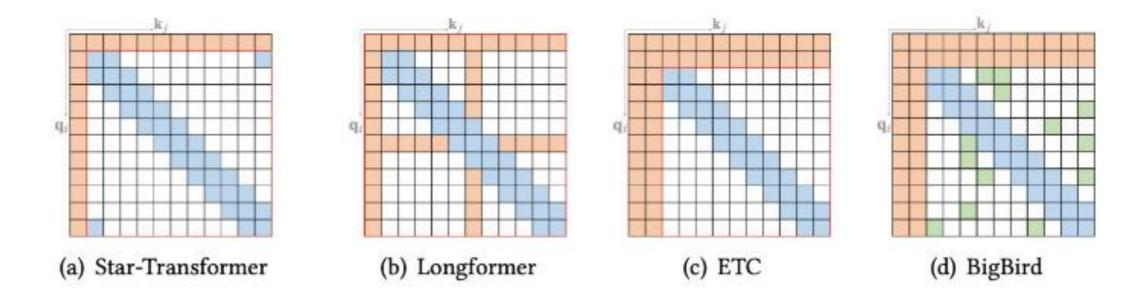
2. approaches that do not require full attention

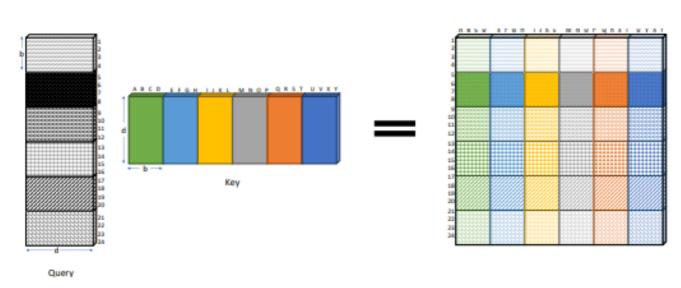
- A Survey of Transformers(Lin, Tianyang, et al. Al Open (2022))
- Method: Position-based



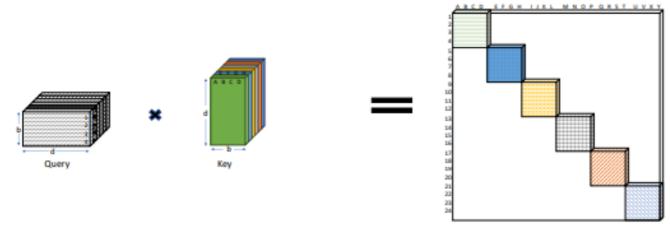
2. approaches that do not require full attention

- A Survey of Transformers(Lin, Tianyang, et al. Al Open (2022))
 - Star-Transformer: Band Attention + Global Attention
 - Longformer: Band Attention + Dilated Attention + Global Attention
 - ETC: Band Attention + Global Attention
 - BIGBIRD: Band Attention + Global Attention + Random Attention

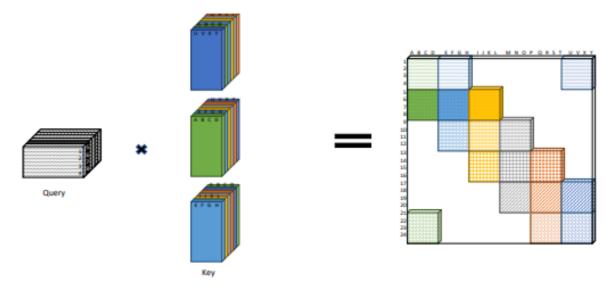




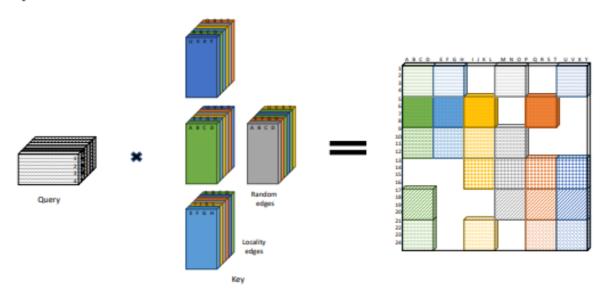
(a) Full all pair attention can be obtained by direct matrix multiplication between the query and key matrix. Groupings just shown for guidance.



(b) Block diagonal attention can be computed by "blockifying" the query and key matrix



(c) Window local attention obtained by "blockifying" the query/key matrix, copying key matrix, and rolling the resulting key tensor (Obtaining rolled key-block tensor is illustrated in detail in Fig. 5). This ensures that every query attends to at least one block and at most two blocks of keys of size b on each side.

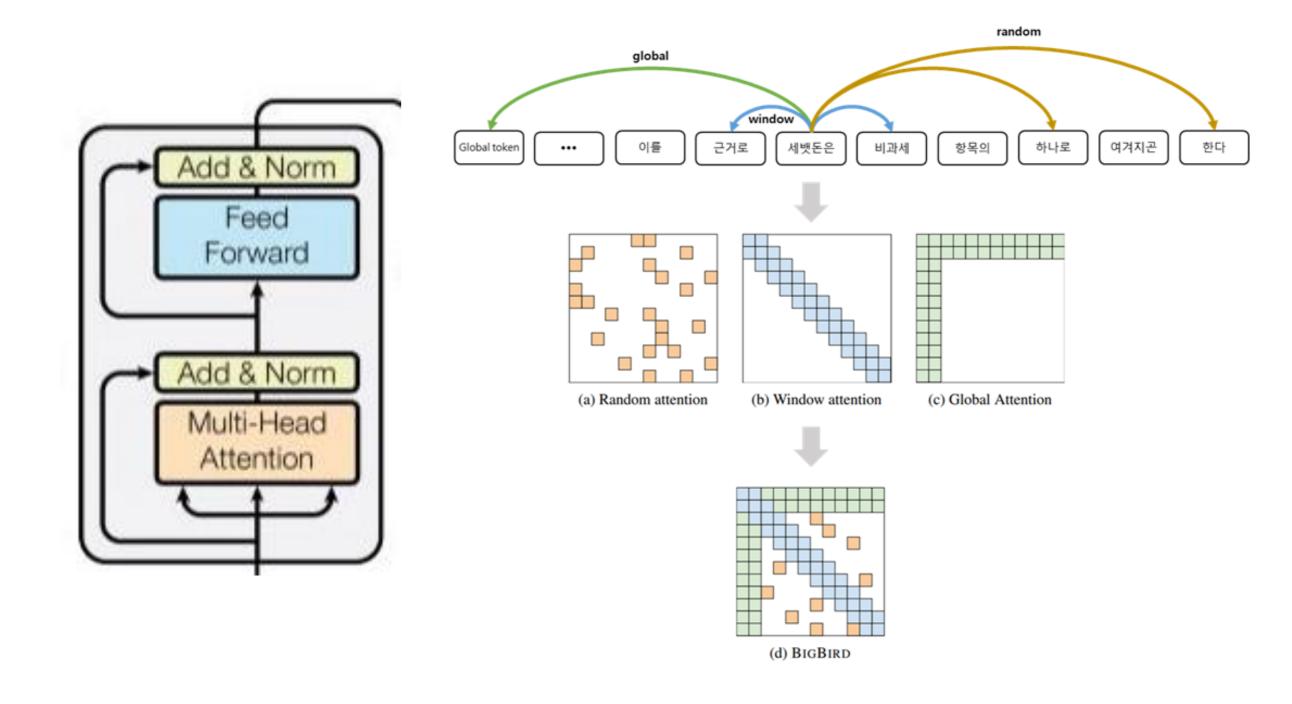


(d) Window + Random attention obtained by following the procedure above along with gathering some random key blocks.

Figure 4: Idea behind fast sparse attention computation in BIGBIRD.

- Star Transformer (NACCL, 2019)
- Longformer(2020)
- ETC(EMNLP, 2020)
- BIGBIRD(NeurIPS, 2020)
- Reformer(ICLR, 2020)

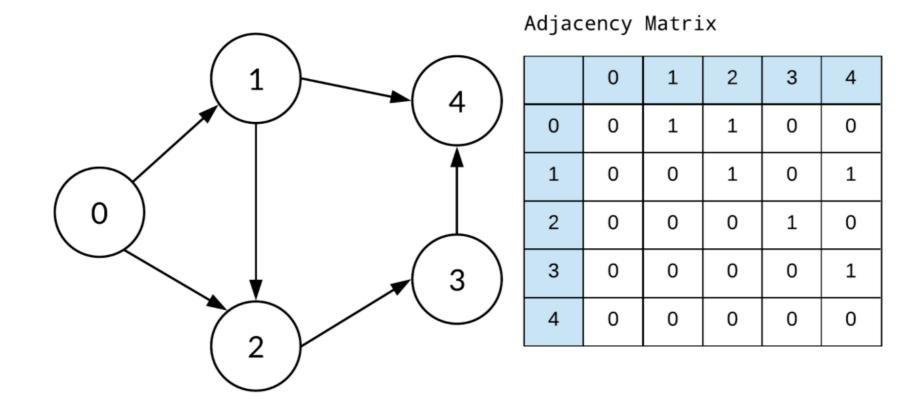




- Generalized attention mechanism
 - input sequence $oldsymbol{X} = (oldsymbol{x}_1,...,oldsymbol{x}_n) \in \mathbb{R}^{n imes d}$
 - directed graph D, vertex set is [n] = {1, . . . , n}.
 - N(i): the out-neighbors set of node i in D
 - i th output vector of the generalized attention mechanism

$$ATTN_D(\boldsymbol{X})_i = \boldsymbol{x}_i + \sum_{h=1}^{H} \sigma\left(Q_h(\boldsymbol{x}_i) K_h(\boldsymbol{X}_{N(i)})^T\right) \cdot V_h(\boldsymbol{X}_{N(i)})$$

- generalised attention mechanism
 - operate on the adjacency matrix A of the graph D

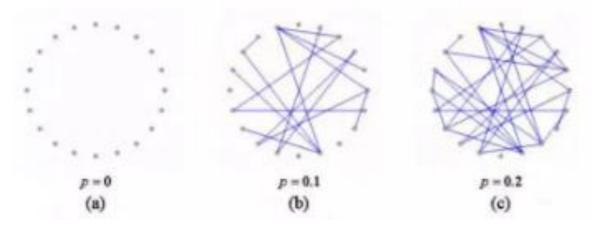


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$$Attention(Q, K, V) = \operatorname{softmax}\left(\frac{QK^{T}}{\sqrt{d_{k}}}\right)V$$

- generalised attention mechanism
 - quadratic complexity
 - Since all tokens attend on every other token
 - The problem of reducing the quadratic complexity is graph sparsification problem
 - Erdos-Rényi model,



each edge is independently chosen with a fixed probability.

sparse random graph for attention mechanism should have two desiderata

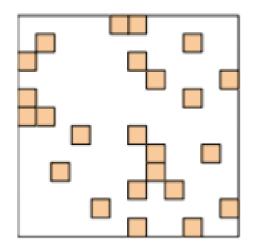
1. small average path length between nodes

- Erdos-Rényi model,
 - each edge is independently chosen with a fixed probability.
 - random graph approximates the complete graph spectrally
 - its second eigenvalue (of the adjacency matrix) is quite far from the first eigenvalue
 - -> rapid mixing time for random walks in the grpah

sparse random graph for attention mechanism should have two desiderata

1. small average path length between nodes

- sparse attention
 - where each query attends over r random number of keys
 - i.e. $A(i, \cdot) = 1$ for r randomly chosen keys



(a) Random attention

sparse random graph for attention mechanism should have two desiderata

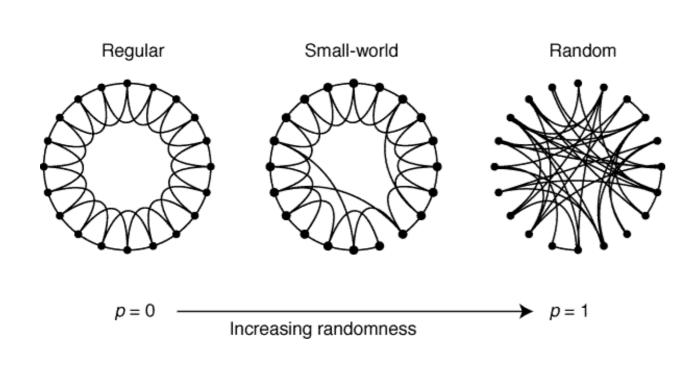
2. locality of reference

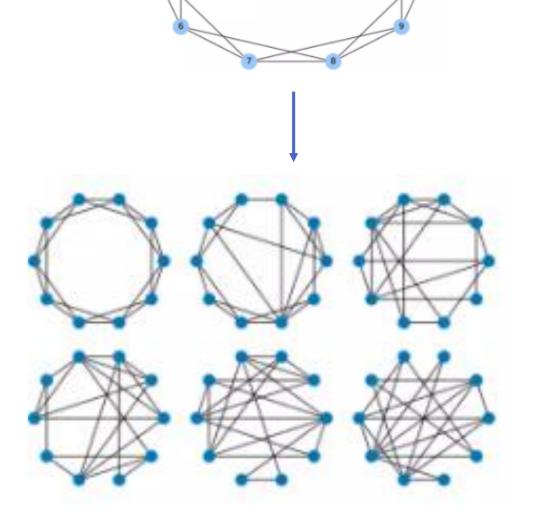
- Locality: proximity of tokens in linguistic structure
- graph theory =>
 - <u>clustering coefficient</u> is a measure of locality of connectivity,
 - high when the graph contains many cliques or near-cliques
- Erdos-Rényi random graphs
 - do not have a high clustering coefficient [84], but a class of random graphs, known as <u>small world graphs</u>, exhibit high clustering coefficient [94]

sparse random graph for attention mechanism should have two desiderata

2. locality of reference

Generative process (Watts and Strogatz [94])





sparse random graph for attention mechanism should have two desiderata

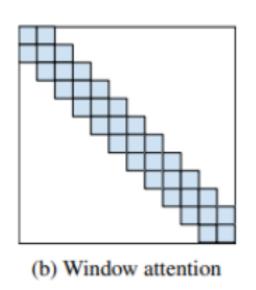
2. locality of reference

- "sliding window on the nodes"
 - random subset (k%), other (100 k)%
- Operation
 - width w, query at location i
 - attends from i w/2 to i + w/2 keys
 - A(i, i-w/2 : i+w/2) = 1

sparse random graph for attention mechanism should have two desiderata

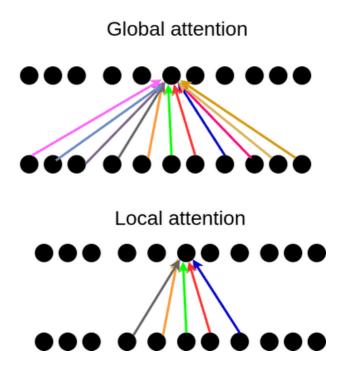
2. locality of reference

- basic experiments
 - random blocks and local window were insufficient

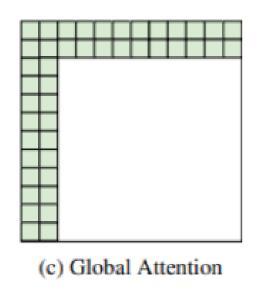


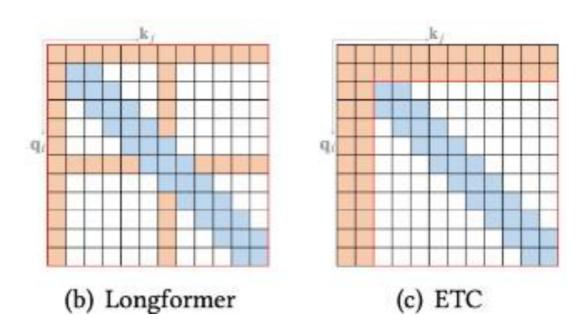
Model	MLM	SQuAD	MNLI
BERT-base	64.2	88.5	83.4
Random (R)	60.1	83.0	80.2
Window (W)	58.3	76.4	73.1
R + W	62.7	85.1	80.5

Table 1: Building block comparison @512



- Global Tokens
 - 1. BIGBIRD-ITC: In internal transformer construction (ITC)
 - make some existing tokens "global", which attend over the entire sequence
 - 2. BIGBIRD-ETC: In extended transformer construction (ETC)
 - include additional "global" tokens such as CLS





- final attention mechanism for BIGBIRD
 - queries attend to r random keys,
 - each query attends to w/2 tokens to the left of its location and w/2 to the right of its location
 - they contain g global tokens

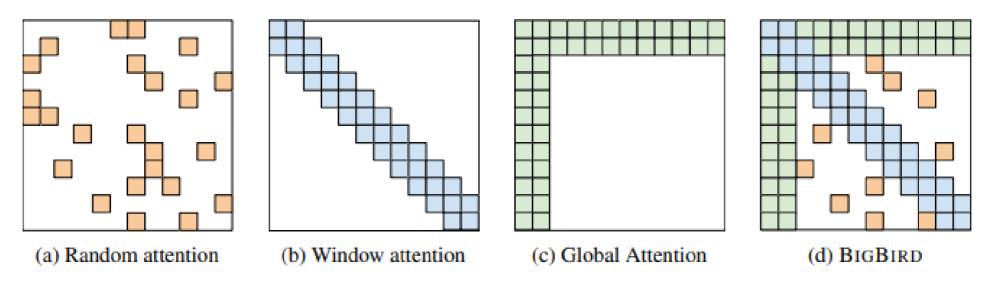
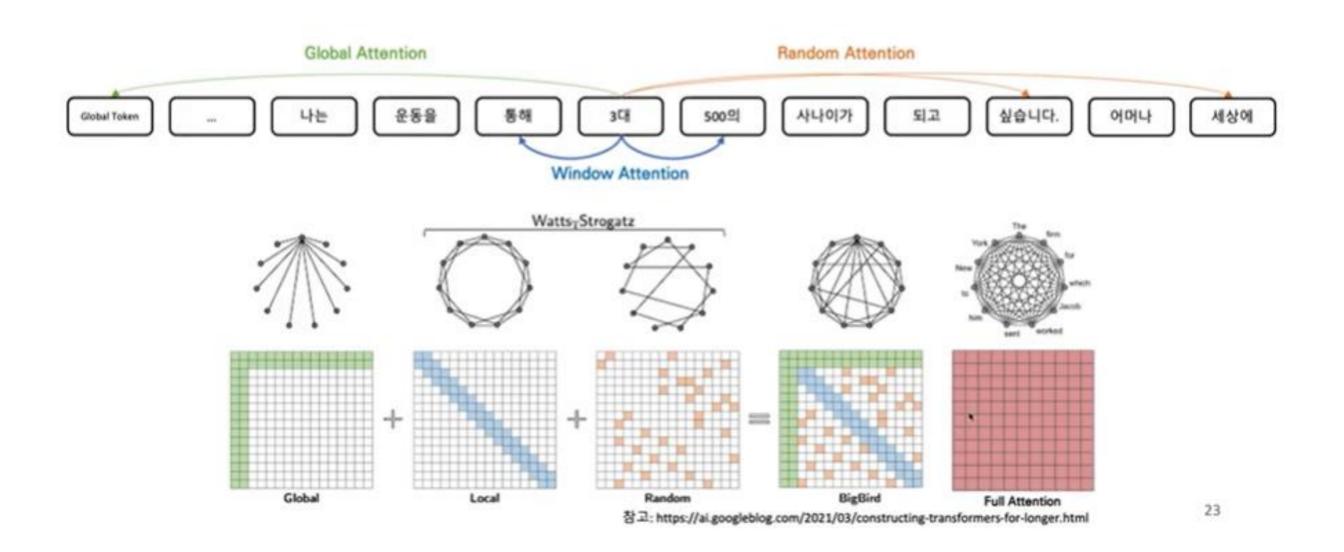


Figure 1: Building blocks of the attention mechanism used in BIGBIRD. White color indicates absence of attention. (a) random attention with r=2, (b) sliding window attention with w=3 (c) global attention with g=2. (d) the combined BIGBIRD model.

final attention mechanism for BIGBIRD



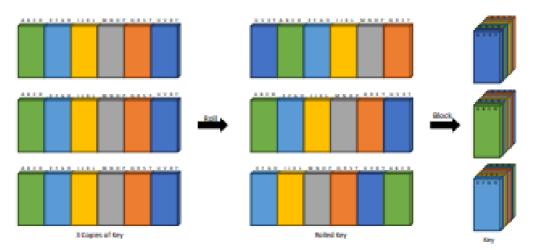


Figure 5: Construction of rolled key-block tensor. Make w copies of the key matrix. Index the copies as $-(w-1)/2 \le j \le (w-1)/2$. Roll jth copy by j blocks. Positive roll means circular shift entries left and likewise for negative roll corresponds to right shift. Finally, reshape by grouping the blocks along a new axis to obtain the key-blocked tensor. For illustration purpose w=3 is chosen.

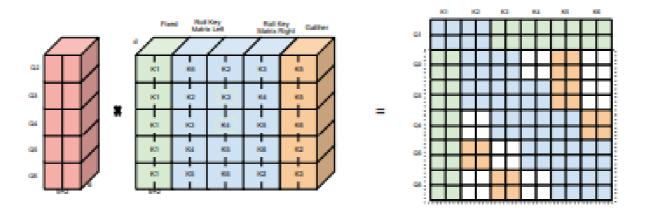


Figure 6: Overview of BIGBIRD attention computation. Structured block sparsity helps in compactly packing our operations of sparse attention, thereby making our method efficient on GPU/TPU. On the left, we depict the transformed dense query and key tensors. The query tensor is obtained by simply blocking and reshaping while the final key tensor by concatenating three transformations: The first green columns, corresponding to global attention, is fixed. The middle blue columns correspond to window local attention and can be obtained by appropriately rolling as illustrated in Fig. 5. For the final orange columns, corresponding to random attentions, we need to use computationally inefficient gather operation. Dense multiplication between the query and key tensors efficiently calculates the sparse attention pattern (except the first row-block, which is computed by direct multiplication), using the ideas illustrated in Fig. 4. The resultant matrix on the right is same as that shown in Fig. 3d.

Theoretical Results about Sparse Attention Mechanism

- https://velog.io/@gyfla1512/Big-Bird-Transformers-for-Longer-Sequences
- sparse attention mechanisms -> Universal Approximators of sequence to sequence functions in the style of Yun et al. [104]
- sparse encoder-decoder transformers are Turing Complete
- moving to a sparse-attention mechanism incurs a cost

Experiments: Natural Language Processing

- 1. masked language modeling (MLM; Devlin et al. 22)
 - to check if better contextual representations can be learnt by utilizing longer contiguous sequences.
- 2. Question Answering(QA, with supporting evidence)
 - to handle longer sequence would allow us to retrieve more evidence using crude systems like TF-IDF/BM25
- 3. long document classification
 - where discriminating information may not be located in first 512 tokens
- using sequence length 4096

1. Pretraining and MLM

Parameter	BIGBIRD-ITC	BIGBIRD-ETC
Block length, b	64	84
# of global token, g	$2 \times b$	256
Window length, w	3 imes b	$3 \times b$
# of random token, r	3 imes b	0
Max. sequence length	4096	4096
# of heads	12	12
# of hidden layers	12	12
Hidden layer size	768	768
Batch size	256	256
Loss	MLM	MLM
Activation layer	gelu	gelu
Dropout prob	0.1	0.1
Attention dropout prob	0.1	0.1
Optimizer	Adam	Adam
Learning rate	10^{-4}	10^{-4}
Compute resources	$8 \times 8 \text{ TPUv3}$	$8 \times 8 \text{ TPUv3}$

Table 8: Hyperparameters for the two BIGBIRD base models for MLM.

- 1. Pretraining and MLM
 - Task: predicting a random subset of tokens which have been masked out.
 - four standard data-sets for pretraining(Tab 9)
 - compare performance in predicting the masked out tokens in terms of bits per character(Tab 10)

Dataset	# tokens	Avg. doc len.
Books [110]	1.0B	37K
CC-News [34]	7.4B	561
Stories [89]	7.7B	8.2K
Wikipedia	3.1B	592

Table 9: Dataset used for pre training.

Model	Base	Large
RoBERTa (sqln: 512)	1.846	1.496
Longformer (sqln: 4096)	1.705	1.358
BIGBIRD-ITC (sqln: 4096)	1.678	1.456
BIGBIRD-ETC (sqln: 4096)	1.611	1.274

Table 10: MLM performance on held-out set.

2. Question Answering(QA)

	Instan	ces	Instance l	Instance Length		
Dataset	Training	Dev	Median	Max		
HotpotQA-distractor [100]	90447	7405	1227	3560		
Natural Questions [52]	307373	7830	3258	77962		
TriviaQA [41]	61888	7993	4900	32755		
WikiHop [95]	43738	5129	1541	20337		

Table 11: Question Answering Datasets

Parameter	Hotp	otQA	Natu	ıralQ	Trivi	aQA	Wiki	Нор
Global token location	ITC	ETC	ITC	ETC	ITC	ETC	ITC	ETC
# of global token, g	128	256	128	230	128	320	128	430
Window length, w	192	252	192	252	192	252	192	252
# of random token, r	192	0	192	0	192	0	192	0
Max. sequence length	4096	4096	4096	4096	4096	4096	4096	4096
# of heads	12	12	12	12	12	12	12	12
# of hidden layers	12	12	12	12	12	12	12	12
Hidden layer size	768	768	768	768	768	768	768	768
Batch size	32	32	128	128	32	32	64	64
Loss		cross-entropy golden spans		cross-entropy golden spans		entropy ans [18]	cross-entropy ans choices	
Compute resources	4×2	4×2 TPUv3		$4 \times 8 \text{ TPUv3}$		TPUv3	$4\times 4~\text{TPUv3}$	

Table 12: Hyperparameters of base BIGBIRD model used for Question Answering i.e. the numbers reported in Tab. 2

- 2. Question Answering(QA)
 - Natural Questions [52], HotpotQA-distractor [100], TriviaQA-wiki
 [41], WikiHop [95]
 - Experiment result (Tab 2)

Model	ŀ	IotpotQ	A	Natu	ıralQ	TriviaQA	WikiHop
	Ans	Sup	Joint	LA	SA	Full	MCQ
RoBERTa	73.5	83.4	63.5	-	-	74.3	72.4
Longformer	74.3	84.4	64.4	-	-	75.2	75.0
BIGBIRD-ITC	75.7	86.8	67.7	70.8	53.3	79.5	75.9
BIGBIRD-ETC	75.5	87.1	67.8	73.9	54.9	78.7	75.9

Table 2: QA Dev results using Base size models. We report accuracy for WikiHop and F1 for HotpotQA, Natural Questions, and TriviaQA.

- 2. Question Answering(QA)
 - Experiment result (Tab 3)
 - compare to top-3 entries from the leaderboard

Model	I	IotpotQ	A	Natu	NaturalQ		TriviaQA	
	Ans	Sup	Joint	LA	SA	Full	Verified	MCQ
HGN [26]	82.2	88.5	74.2	-	-	-	-	-
GSAN	81.6	88.7	73.9	-	-	-	-	-
ReflectionNet [32]	-	-	-	77.1	64.1	-	-	-
RikiNet-v2 [61]	-	-	-	76.1	61.3	-	-	-
Fusion-in-Decoder [39]	-	-	-	-	-	84.4	90.3	-
SpanBERT [42]	-	-	-	-	-	79.1	86.6	-
MRC-GCN [87]	-	-	-	-	-	-	-	78.3
MultiHop [14]	-	-	-	-	-	-	-	76.5
Longformer [8]	81.2	88.3	73.2	-	-	77.3	85.3	81.9
BIGBIRD-ETC	81.2	89.1	73.6	77.8	57.9	84.5	92.4	82.3

Table 3: Fine-tuning results on **Test** set for QA tasks. The Test results (F1 for HotpotQA, Natural Questions, TriviaQA, and Accuracy for WikiHop) have been picked from their respective leaderboard. For each task the top-3 leaders were picked not including BIGBIRD-etc. **For Natural Questions**Long Answer (LA), TriviaQA, and WikiHop, BIGBIRD-ETC is the new state-of-the-art. On HotpotQA we are third in the leaderboard by F1 and second by Exact Match (EM).

- 3. Classification
 - Experiments: datasets of different lengths and contents,
 specifically various document classification and GLUE tasks.

Parameter	IMDb	Arxiv	Patents	Hyperpartisan	Yelp-5
Batch size	64	64	64	32	32
Learning rate	1×10^{-5}	3×10^{-5}	5×10^{-5}	5×10^{-6}	2×10^{-5}
Num epochs	40	10	3	15	2
TPUv3 slice	4×4	4×4	4×4	4×2	4×8
# of heads			12		16
# of hidden layers			12		24
Hidden layer size			768		1024
Block length, b			64		
Global token location			ITC		
# of global token, g			2 imes b		
Window length, w			$3 \times b$		
# of random token, r			$3 \times b$		
Max. sequence length			4096		
Vocab size			50358		
Activation layer			gelu		
Dropout prob			0.1		
Attention dropout prob			0.1		
Loss			cross-entrop	у	
Optimizer			Adam		

Table 14: Hyperparameters for document classification.

- 3. Classification
 - Experiments: datasets of different lengths and contents,
 specifically various document classification and GLUE tasks.

Model	IMDb [64]	Yelp-5 [108]	Arxiv [35]	Patents [53]	Hyperpartisan [47]
# Examples	25000	650000	30043	1890093	645
# Classes	2	5	11	663	2
Excess fraction	0.14	0.04	1.00	0.90	0.53
SoTA	[88] 97.4	[3] 73.28	[69] 87.96	[69] 69.01	[40] 90.6
RoBERTa	95.0 ± 0.2	71.75	87.42	67.07	87.8 ± 0.8
BIGBIRD	95.2 ± 0.2	72.16	92.31	69.30	92.2 ± 1.7

Table 15: Classification results. We report the F1 micro-averaged score for all datasets. Experiments on smaller IMDb and Hyperpartisan datasets are repeated 5 times and the average performance is presented along with standard deviation.

System	MNLI-(m/mm)	QQP	QNLI	SST-2	CoLA	STS-B	MRPC	RTE
	392k	363k	108k	67k	8.5k	5.7k	3.5k	2.5k
BERT	84.6/83.4	71.2	90.5	93.5	52.1	85.8	88.9	66.4
XLNet	86.8/-	91.4	91.7	94.7	60.2	89.5	88.2	74.0
RoBERTa	87.6/-	91.9	92.8	94.8	63.6	91.2	90.2	78.7
BIGBIRD	87.5/87.3	88.6	92.2	94.6	58.5	87.8	91.5	75.0

Table 16: GLUE Dev results on base sized models. Number of training examples is reported below each task. MCC score is reported for CoLA, F1 score is reported for MRPC, Spearman correlation is reported for STS-B, and accuracy scores are reported for the other tasks.

- 4. Encoder-Decoder Tasks
 - Suffering: quadratic complexity due to the full self attention
 - the sparse attention mechanism of BIGBIRD only at the encoder side
 - Experiment
 - realistic scenarios: that the median output sequence length is
 200 whereas the input sequence's median length is > 3000.
 - sparse attention -> encoder, full self-attention -> decoder.

	Instances			Input I	Length	Output Length		
Dataset	Training	Dev	Test	Median	90%-ile	Median	90%-ile	
Arxiv [20] PubMed [20] BigPatent [78]	203037 119924 1207222	6436 6633 67068	6440 6658 67072	6151 2715 3082	14405 6101 7693	171 212 123	352 318 197	

Table 18: Statistics of datasets used for summarization.

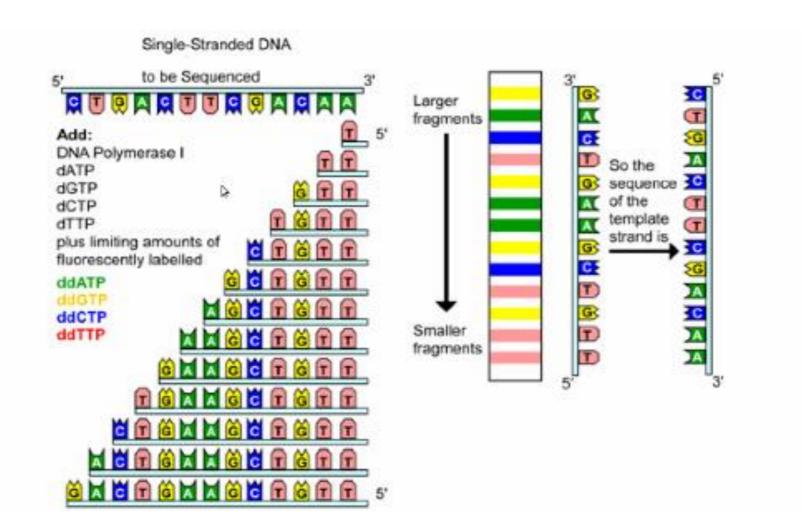
- 5. Summarization
 - using, three long document datasets
 - focus on abstractive summarization of long documents where using a longer contextual encoder should improve performance.
 - 1. the salient content can be evenly distributed in the long document, not just in first 512 tokens(BigPatents dataset [78]).
 - 2. longer documents exhibit a richer discourse structure and summaries are considerably more abstractive

- 5. Summarization
 - ROUGE(Recall-Oriented Understudy for Gisting Evaluation) Score

		Arxiv			PubMed]	BigPaten	t
Model	R-1	R-2	R-L	R-1	R-2	R-L	R-1	R-2	R-L
SumBasic [68]	29.47	6.95	26.30	37.15	11.36	33.43	27.44	7.08	23.66
LexRank [25]	33.85	10.73	28.99	39.19	13.89	34.59	35.57	10.47	29.03
LSA [97]	29.91	7.42	25.67	33.89	9.93	29.70	-	-	
	29.30	6.00	25.56	31.55	8.52	27.38	28.74	7.87	24.66
Attn-Seq2Seq [85] Pntr-Gen-Seq2Seq [77]	32.06	9.04	25.16	35.86	10.22	29.69	33.14	11.63	28.55
Long-Doc-Seq2Seq [20]	35.80	11.05	31.80	38.93	15.37	35.21	-	-	
Sent-CLF [81]	34.01	8.71	30.41	45.01	19.91	41.16	36.20	10.99	31.83
Sent-PTR [81]	42.32	15.63	38.06	43.30	17.92	39.47	34.21	10.78	30.07
Extr-Abst-TLM [81]	41.62	14.69	38.03	42.13	16.27	39.21	38.65	12.31	34.09
Dancer [31]	42.70	16.54	38.44	44.09	17.69	40.27	-	-	
Transformer	28.52	6.70	25.58	31.71	8.32	29.42	39.66	20.94	31.20
2 + RoBERTa [76]	31.98	8.13	29.53	35.77	13.85	33.32	41.11	22.10	32.58
ទ្ធ + RoBERTa [76] + Pegasus [107]	34.81	10.16	30.14	39.98	15.15	35.89	43.55	20.43	31.80
BIGBIRD-RoBERTa	41.22	<u>16.43</u>	36.96	<u>43.70</u>	19.32	39.99	<u>55.69</u>	<u>37.27</u>	45.56
υ Pegasus (Reported) [107]	44.21	16.95	38.83	45.97	20.15	41.34	52.29	33.08	41.75
Pegasus (Reported) [10/] Pegasus (Re-eval) → BIGBIRD-Pegasus	43.85	16.83	39.17	44.53	19.30	40.70	52.25	33.04	41.80
☐ BIGBIRD-Pegasus	46.63	19.02	41.77	46.32	20.65	42.33	60.64	42.46	50.01

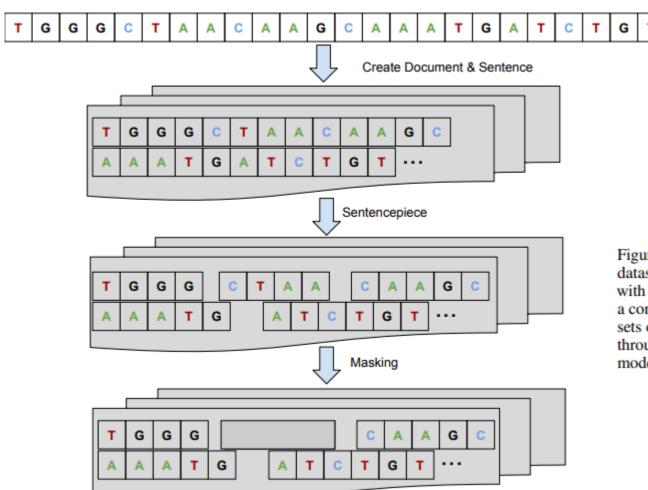
Table 4: Summarization ROUGE score for long documents.

- 5. Genomics
 - recent upsurge in using deep learning for genomics data [86, 106, 13]
 - Inputs: DNA sequence fragments



- 5. Genomics
 - recent upsurge in using deep learning for genomics data [86, 106, 13]
 - Inputs: DNA sequence fragments
 - longer input sequence handling capability of BIGBIRD would be beneficial as many functional effects in DNA are highly nonlocal [12]
 - MLM pretraining: taking inspiration from NLP, we learn powerful contextual representations for DNA fragments utilizing abundant unlabeled data
 - long input BIGBIRD along with the proposed pretraining significantly improves performances in two downstream tasks

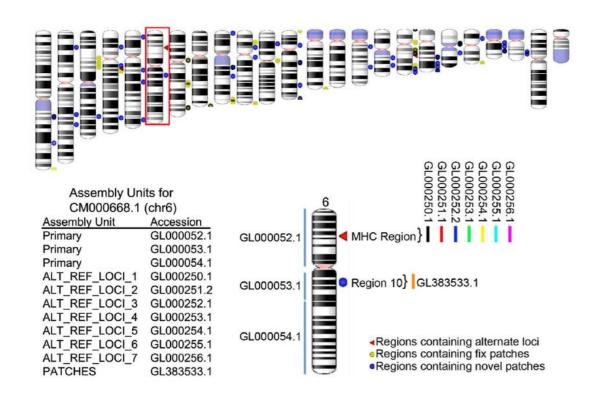
- 5. Genomics
 - 1. Pre-training and MLM
 - Liang [58], first segment DNA into tokens so as to further increase the context length



* raw DNA sequence: GRCh37

Figure 7: Visual description of how the masked language modeling data was generated from raw DNA dataset. The raw DNA sequences of GRCh37, where split at random positions to create documents with 50-100 sentences where each sentence was 500-1000 base pairs (bps). Thus each document had a continuous strand of 25000-100,000 bps of DNA. This process was repeated 10 times to create 10 sets of document for each chromosome of GRCH37. The resulting set of documents was then passed through Sentencepiece that created tokens of average 8bp. For pretraining we used masked language model and masked 10% of the tokens and trained on predicting the masked tokens.

- 5. Genomics
 - 1. Pre-training and MLM
 - Liang [58], first segment DNA into tokens so as to further increase the context length
 - raw DNA sequence: GRCh37



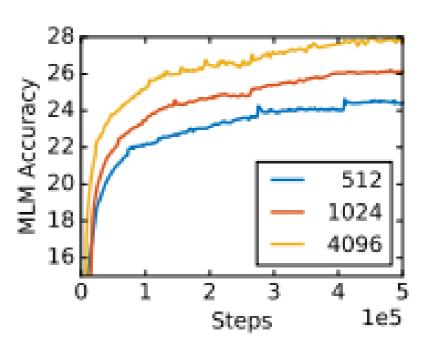


Figure 8: BIGBIRD accuracy with context length.

- 5. Genomics
 - 1. Pre-training and MLM
 - Experiments Result
 - bits per character (BPC)

Model	BPC
SRILM [58] BERT (sqln. 512)	1.57 1.23
BIGBIRD (sqln. 4096)	1.12

Table 5: MLM BPC



- 5. Genomics
 - 2. Promoter Region Prediction
 - Promoter: DNA region typically located upstream of the gene
 - Multiple methods have been proposed to identify the promoter regions in a given DNA sequence.
 - The corresponding machine learning task is to classify a given DNA fragment as promoter or non-promoter sequence.
 - Dataset: Oubounyt et al. [71]

Model	F1
CNNProm [90] DeePromoter [71]	69.7 95.6
BIGBIRD	99.9

Table 6: Comparison.

- 5. Genomics
 - 3. Chromatin-Profile Prediction

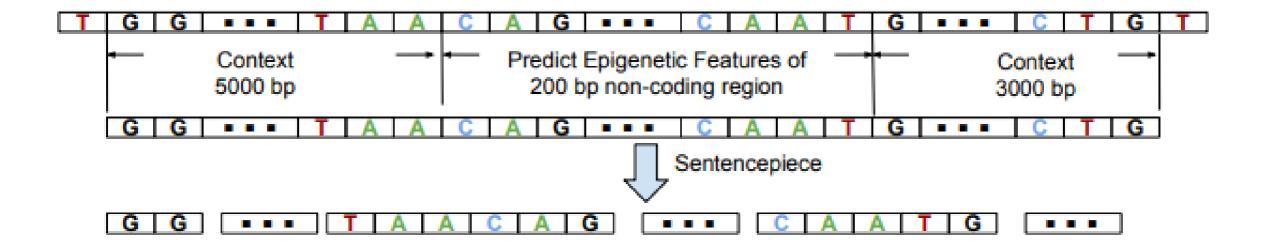


Figure 9: Visual description of the DNA segment from which we predict the chromatin profile for a given non-coding region of the raw DNA sequences of GRCh37. We take 8000 bps of DNA before and after the given non-coding region as context. The complete fragment of DNA including the context on both side, is then tokenized to form our input sequence of tokens. The task is to predict 919 chromatin profile including 690 transcription factors (TF) binding profiles for 160 different TFs, 125 DNase I sensitivity (DHS) profiles and 104 histone-mark (HM) profiles

- 5. Genomics
 - 3. Chromatin-Profile Prediction
 - understanding the functional effects of non-coding regions of DNA is a very important task
 - Method: We jointly learn 919 binary classifiers to predict these functional effects from sequence of DNA fragments.

Model	TF	HM	DHS
gkm-SVM [30]	89.6	-	-
DeepSea [109]	95.8	85.6	92.3
BIGBIRD	96.1	88.7	92.1

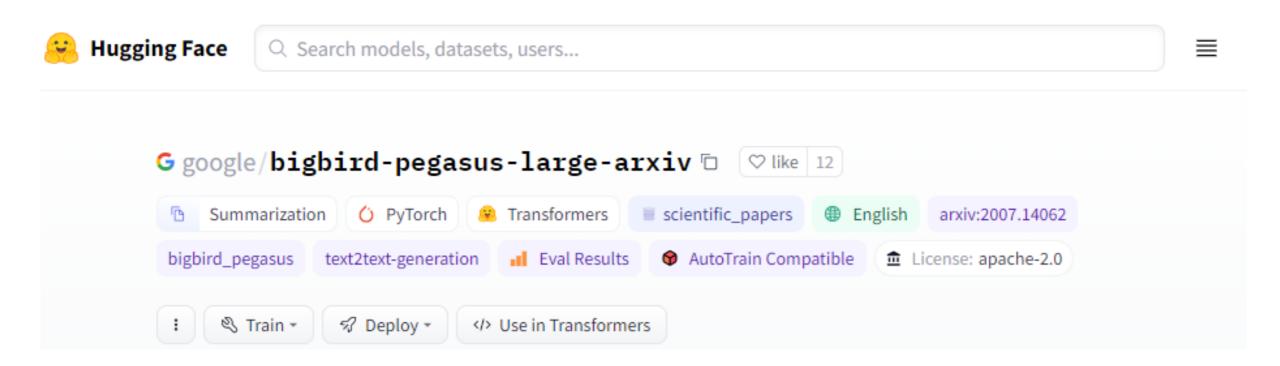
Table 7: Chromatin-Profile Prediction

Conclusion

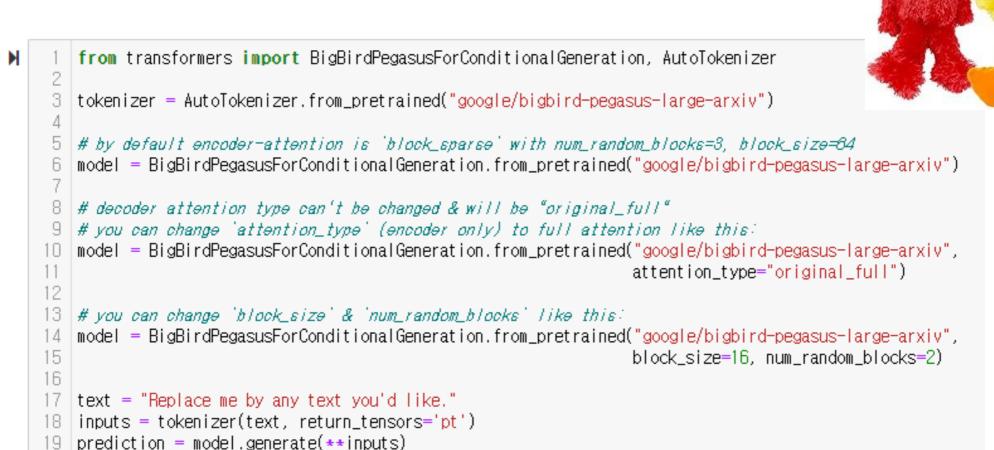
- BIGBIRD: a sparse attention mechanism that is linear in the number of tokens
 - universal approximator of sequence to sequence functions
 - is also Turing complete.
- Method
 - use the power of extra global tokens preserve the expressive powers of the model.
- BIGBIRD gives state-of-the-art performance on a number of NLP tasks such as question answering and long document classification
- introduce attention based contextual language model for DNA and fine-tune it for down stream tasks

BIGBIRDHugging Face

- https://huggingface.co/transformers/v4.7.0/model_doc/bigbird.html
 #overview
- https://huggingface.co/google/bigbird-pegasus-largearxiv/blob/main/README.md



BIGBIRD_ Hugging Face



In [13]: ▶ 1 prediction

20 | prediction = tokenizer.batch decode(prediction)

Out[13]: ['<s> we present a new method for the generation of non - abelian gauge fields from a non - abelian gauge field.<n> we show that the non - abelian gauge field can be efficiently generated from a non - abelian gauge field.<n> the method is based on the use of a non - linear coupling between the non - abelian and the abelian gauge field.<n> the non - abelian gauge field can be efficiently generated from a non - abelian gauge field. <n> @xmathO non - abelian gauge field.<n> @xmath1 non - abelian gauge field.<n> @xmath2 non - abelian gauge field.<n> @xmath5 non - abelian gauge field.<n> @xmath6 non - abelian gauge field.<n> @xmath7 non - abelian gauge field.<n> @xmath8 non - abelian gauge field.<n> @xmath9 non - abelian gauge field.<n> @xmath1 non - abelian gauge field.<n> @xmath2 non - abelian gauge field.<n> @xmath3 non - abelian gauge field.<n> @xm

Summary and Take-home message

- 좋은 점: 42쪽의 분량으로 인해서 어펜딕스가 풍부하다!!! 그 덕에 궁금한 부분이 있다면 어펜딕스에서 그 어려움을 해결할 수 있다(다만, 수학적인 내용과 기본 지식을 알고 있다는 전제 하에 논문이 작성된 것 같아서 이해가 많이 어려울 수도 있다)
- 아쉬운 점: 내용이 어렵다ㅠㅠㅠ 예전에 Attention is all you need 논문을
 3주 가량 공부했는데 그만큼 시간이 주어지지 못해 이해하지 못한 부분이
 많다는 점에서 아쉬움이 남는다
- 흥미로운 점: Transformer가 NLP에 끼친 영향이 어마어마 하다는 것은 알고 있었지만, 정말 그 이상이구나 싶었고, 그로 인한 다양한 연구들이 진행되어왔다는 현행 상황들까지 살펴볼 수 있어서 좋았다!!!

Longformer: The Long-Document Transformer

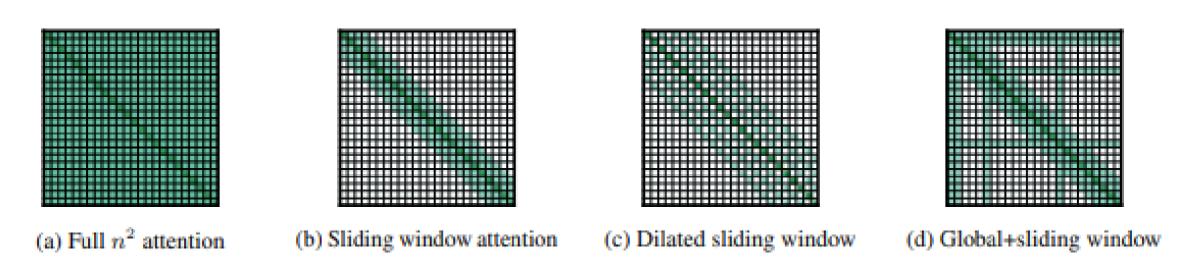
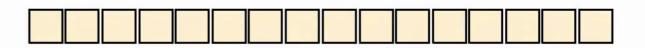


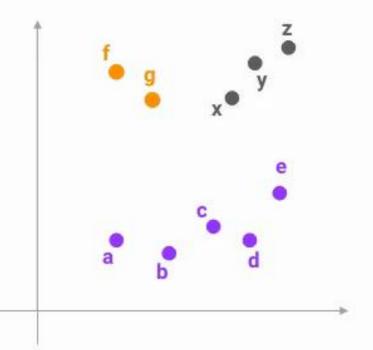
Figure 2: Comparing the full self-attention pattern and the configuration of attention patterns in our Longformer.

- Reformer: The Efficient Transformer
 - O(L log L)
 - LSH

(Locality-Sensitive-Hashing)



Locality Sensitive Hashing (LHS)



Point	Hash
a	
b	
C	
d	
е	
f	
g	
X	
у	
Z	

Reformer: The Efficient Transformer

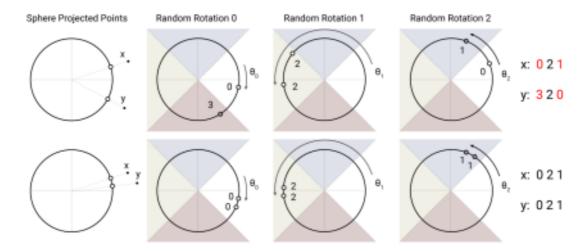


Figure 1: An angular locality sensitive hash uses random rotations of spherically projected points to establish buckets by an argmax over signed axes projections. In this highly simplified 2D depiction, two points x and y are unlikely to share the same hash buckets (above) for the three different angular hashes unless their spherical projections are close to one another (below).

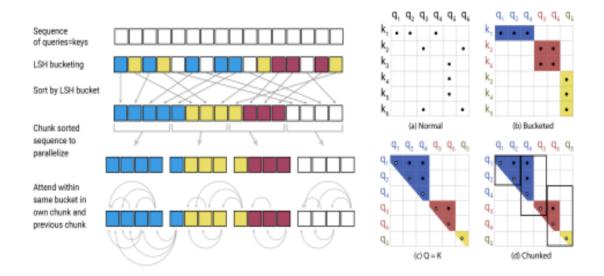


Figure 2: Simplified depiction of LSH Attention showing the hash-bucketing, sorting, and chunking steps and the resulting causal attentions. (a-d) Attention matrices for these varieties of attention.

- Extended Transformers Construction
- ETC: Encoding Long and Structured Inputs in Transformers

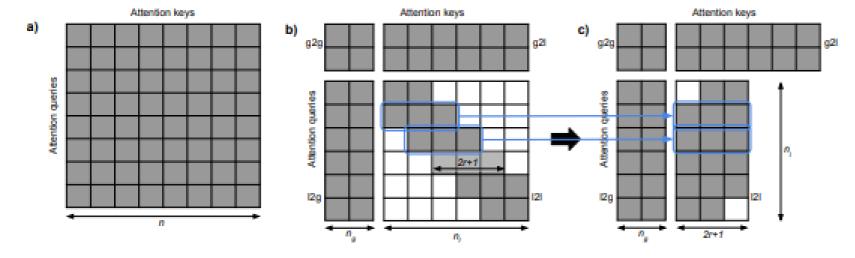


Figure 2: Sparsity diagram showing which attention queries (rows) can attend to which attention keys (columns) a) for standard Transformer attention with input size n; b) for global-local attention with input sizes n_g , n_l , and radius r; c) how the 121 attention piece is reshaped into a much smaller attention matrix, limited by local radius.

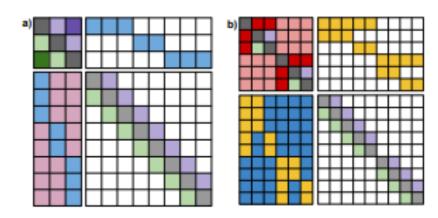


Figure 3: Example attention patterns for handling (a) long inputs and (b) structured inputs. White background means attention is masked via M, and the other colors indicate different relative position labels.

LittleBird: Efficient Faster & Longer Transformer for Question

Answering(2022, EMNLP)

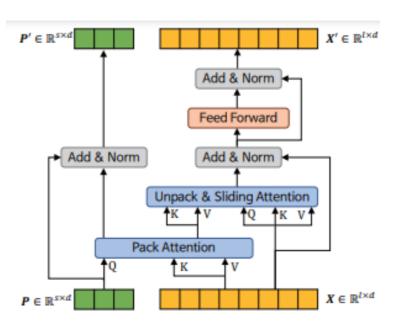


Figure 1: LittleBird Layer

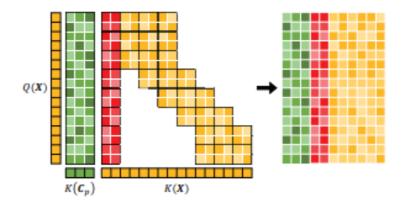


Figure 2: Unpack & Sliding Window Attention of LittleBird

