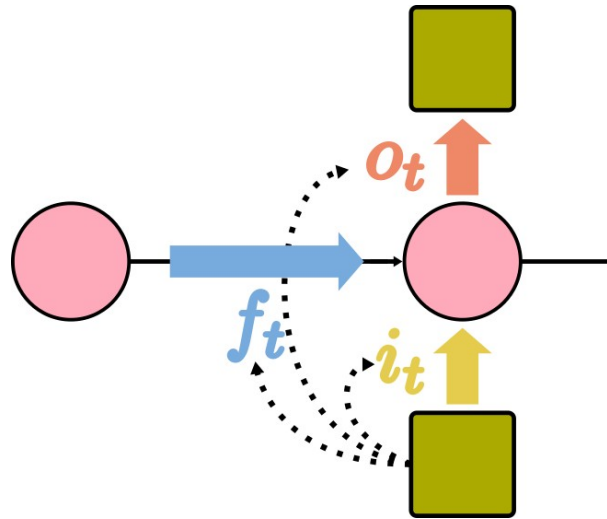


pLSTM: parallelizable Linear Source Transition Mark networks

Korbinian Pöppel, ASAP Seminar
13.08.2025

Introduction: Linear RNNs

qLSTM, DeltaNet, LRU, RWKV-4/5/6, GLA, Mamba, Griffin, mLSTM, Gated DeltaNet, TTT, DeltaProduct..



No previous (hidden) state dependence
→ Parallelization by unrolling the sum

Vision Mamba: Efficient Visual Representation Learning with Bidirectional State Space Model

Lianghui Zhu^{1*} Bencheng Liao^{2,1*} Qian Zhang³ Xinlong Wang⁴ Wenyu Liu¹ Xinggang Wang¹

VISION-RWKV: EFFICIENT AND SCALABLE VISUAL PERCEPTION WITH RWKV-LIKE ARCHITECTURES

Yuchen Duan^{1,2*}, Weiyun Wang^{3,2*}, Zhe Chen^{4,2*}, Xizhou Zhu^{5,2,6}, Lewei Lu⁶,
Tong Lu⁴, Yu Qiao², Hongsheng Li¹, Jifeng Dai^{5,2}, Wenhai Wang^{1,2✉}

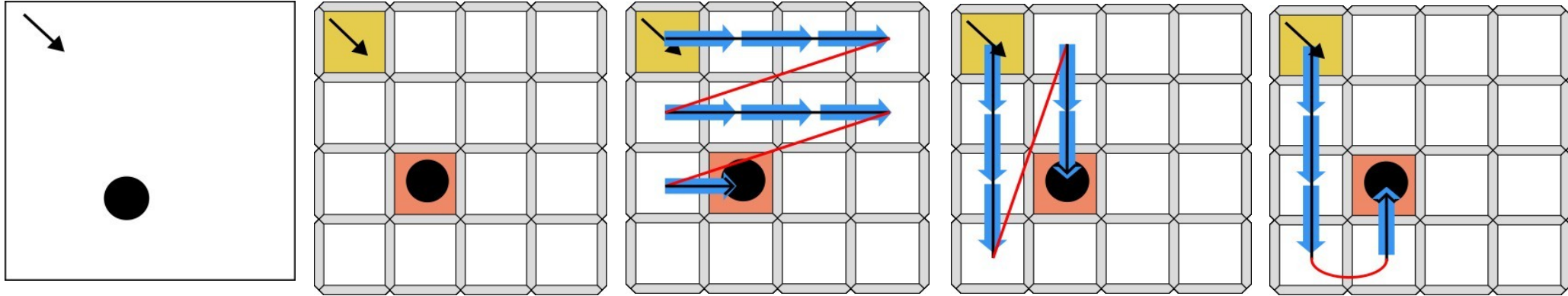
¹The Chinese University of Hong Kong, ²Shanghai AI Laboratory, ³Fudan University,

⁴Nanjing University, ⁵Tsinghua University, ⁶SenseTime Research

Vision-LSTM: xLSTM as Generic Vision Backbone

Benedikt Alkin^{1,2} Maximilian Beck^{1,3} Korbinian Pöppel^{1,3}
Sepp Hochreiter^{1,2,3} Johannes Brandstetter^{1,2}

Multi-Dimensional Data Arrow Pointing Task



Multi-Dimensional RNNs

Multi-Dimensional RNNs / LSTMs (Graves et al. 2007)

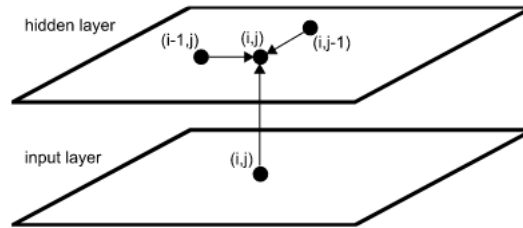


Figure 1: 2D RNN Forward pass.

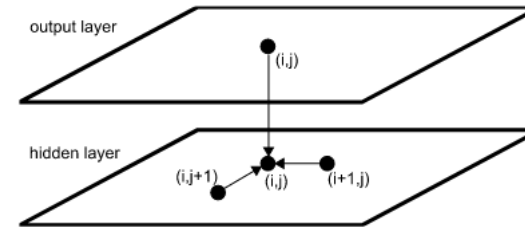
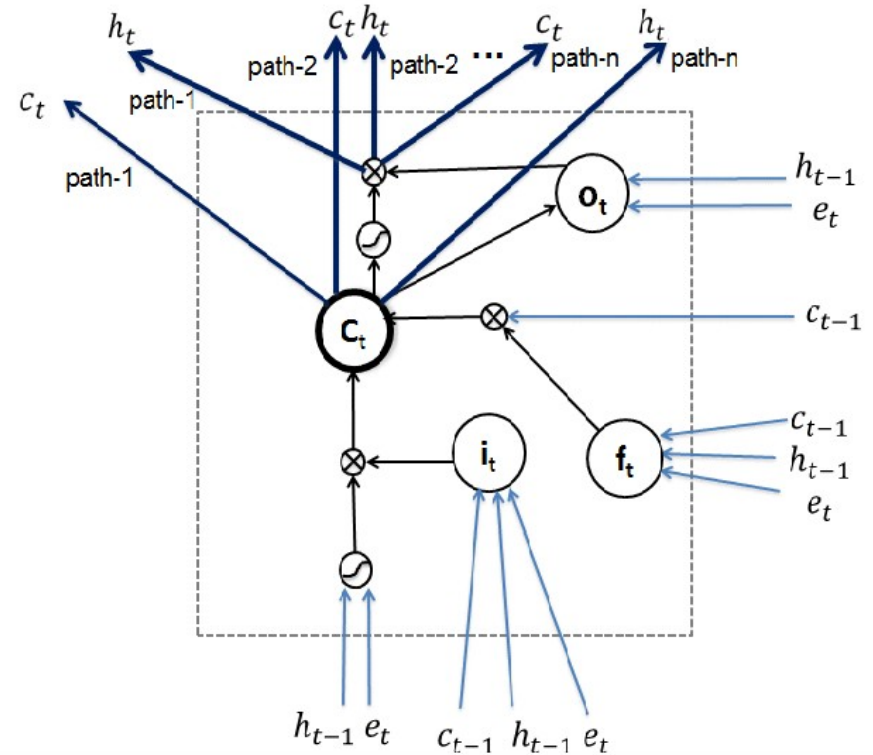


Figure 2: 2D RNN Backward pass.

DAG-LSTM

DAG-Structured Long
Short-Term Memory
for Semantic Compositionality
(Zhu et al. 2016)

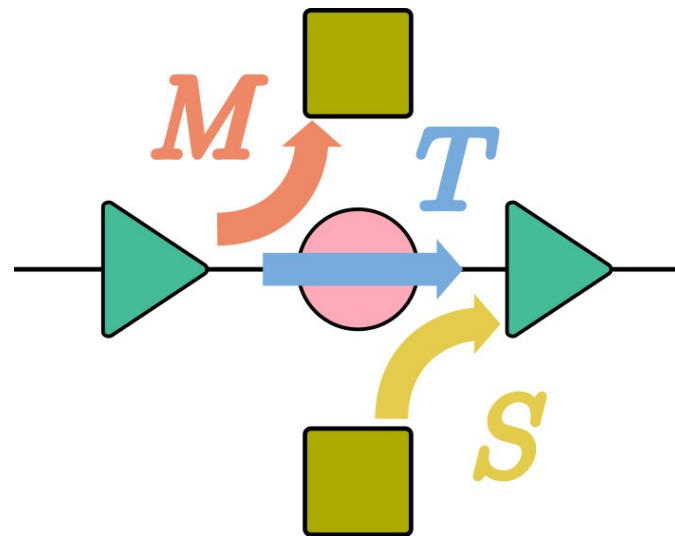
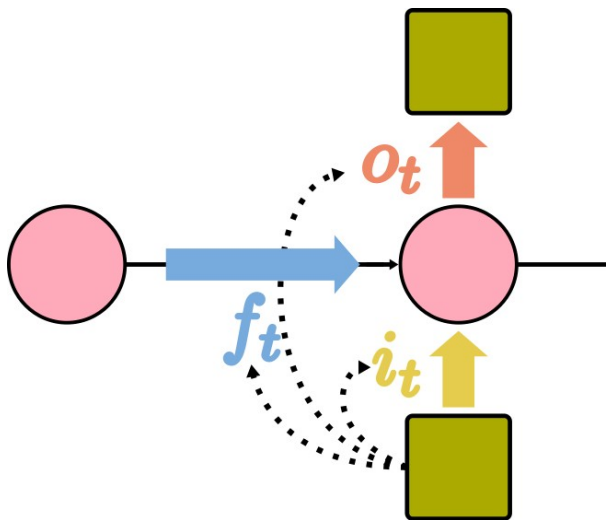


Can we combine linear RNNs with MD-RNNs?

- with parallelization ☒
- with long-range stability ☒
- with directional propagation ☒

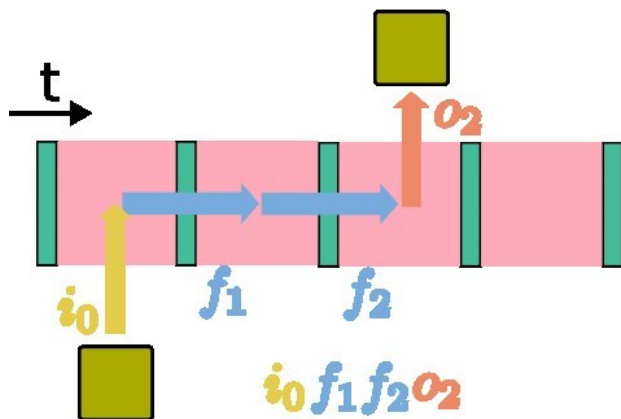
From Linear RNNs to pLSTMs

- Input, forget, output gate
- Source, Transition, Mark

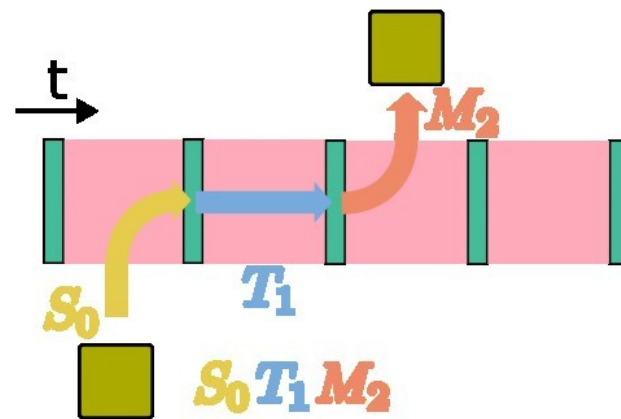


Linear RNN vs pLSTM on a sequence

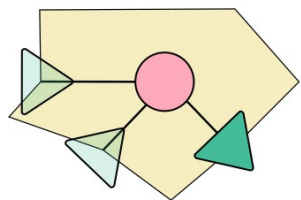
linear RNN



pLSTM

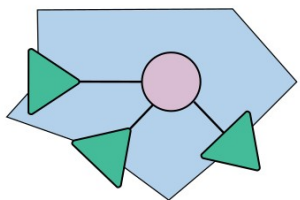


pLSTM on a DAG



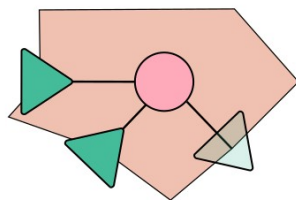
$S_{e'}^n$

Source



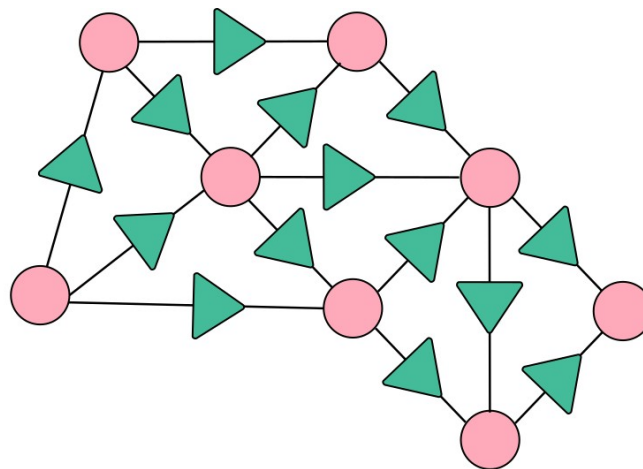
$T_{e'e}$

Transition



M_e^n

Mark



Recurrent Computation: Topological Order

- At current node:
 - Add source contributions to out-going edge states
 - Add transitions from in-coming to out-going edge states
 - Compute Mark from in-coming edge states for network output at current node
- Iterate to next node in the topological order sequence

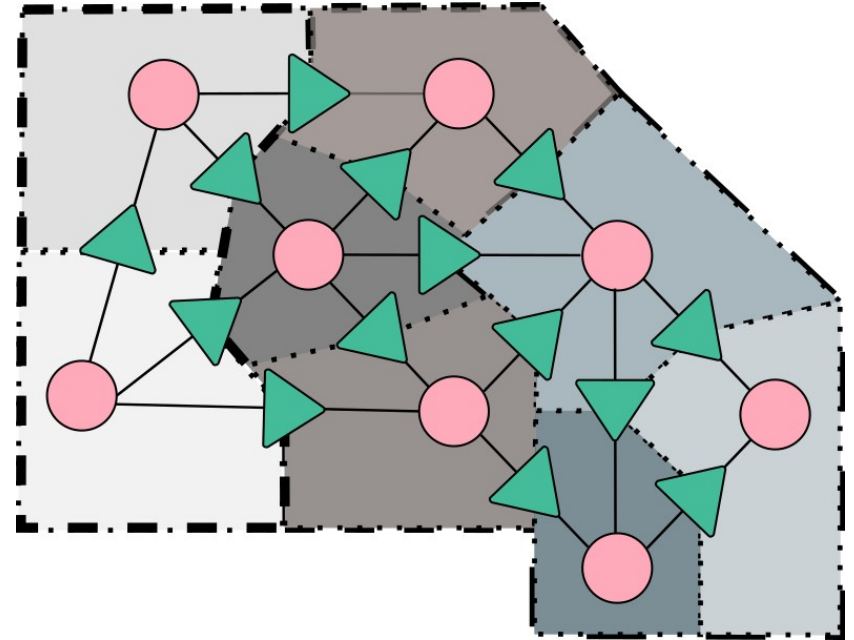
Naive Parallel Computation: Sum All Paths

- Everything is linear
- In between every pair of nodes:
 - Sum all paths in between

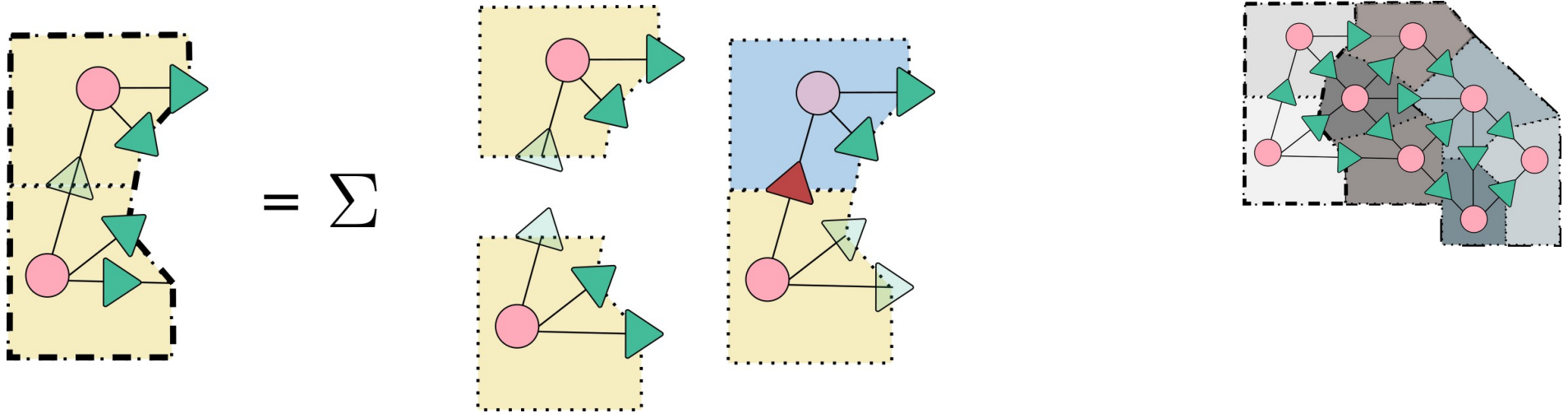
Problem: exponentially many paths (for multi-dim.)

Loosely self-similar DAG

- Low in- and out-degree
- Can be partitioned recursively
- Similar structure for all sub-partitions

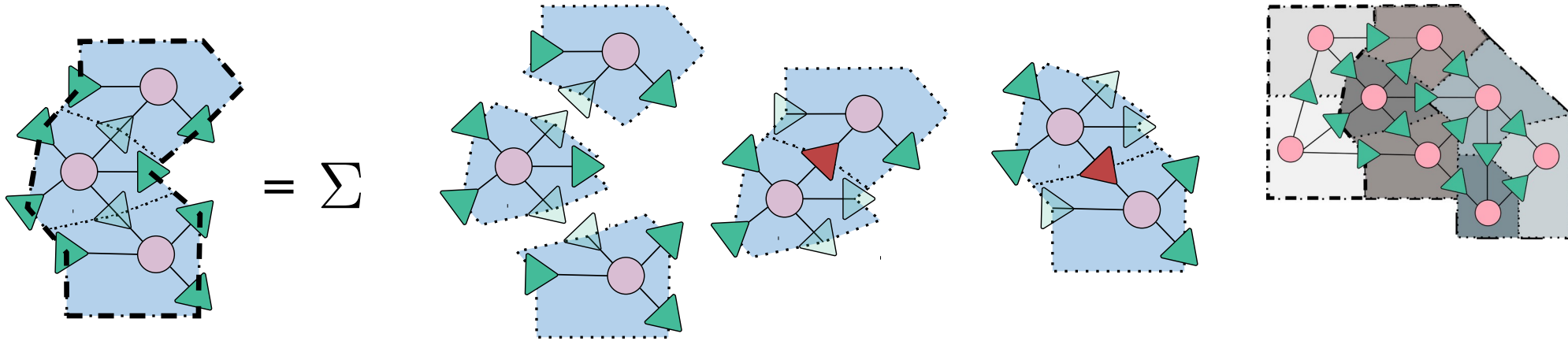


Combining Source and Transition



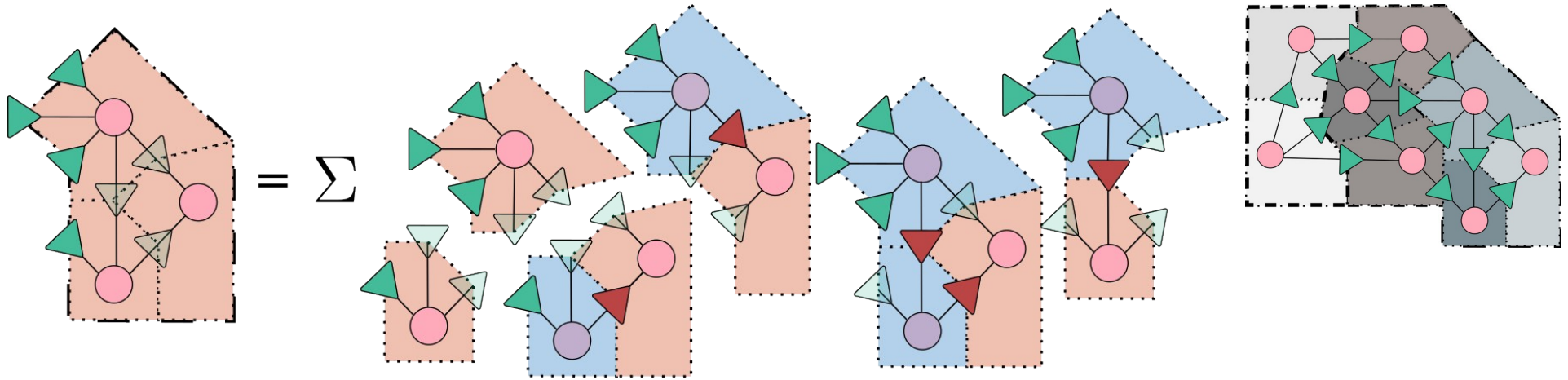
Source (Level n) = Combination of Source (Level n-1) + Transitions (Level n-1)

Combining Transitions



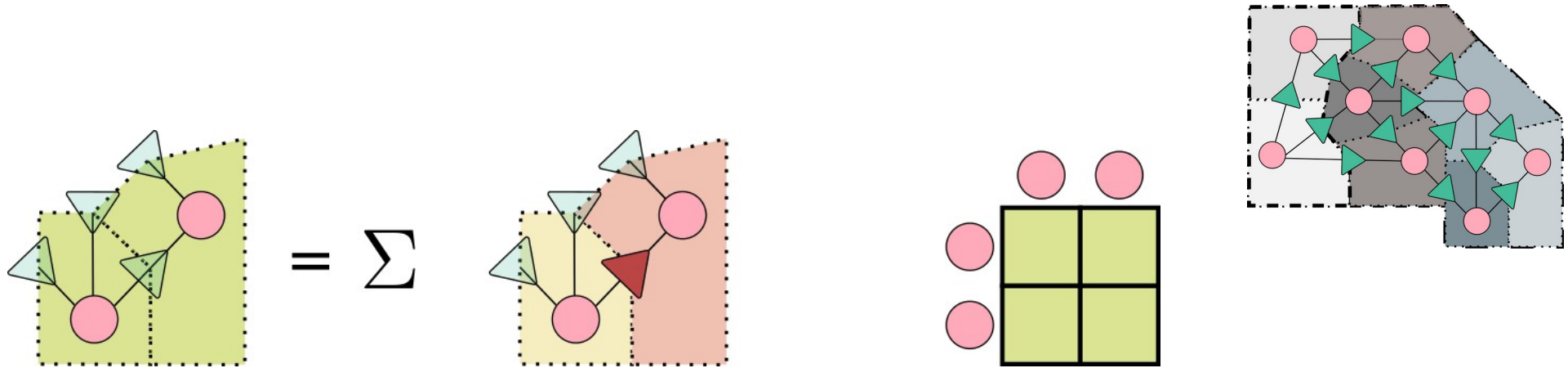
Transition (Level n) = Combination of Transitions (Level n-1)

Combining Source and Transition



Mark (Level n) = Combination of Transitions (Level $n-1$) and Mark (Level $n-1$)

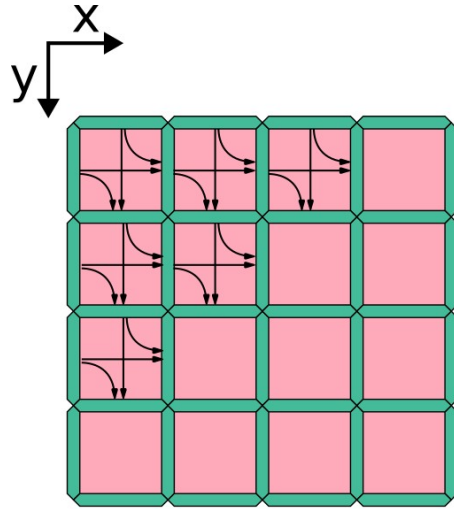
Source and Mark: Gating Matrix



Gating Matrix Entries (like an input dependent “Attention Mask”)

pLSTM on Regular Grids: Images

- Create a DAG: Go Top-Left to Bottom-Right
+ 3 other combinations

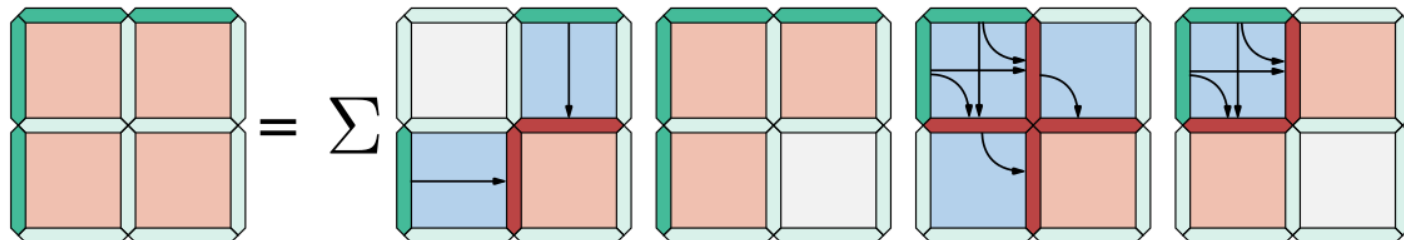
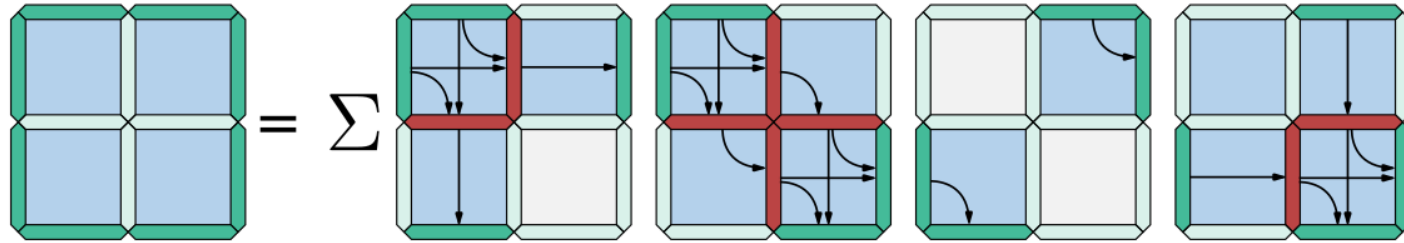
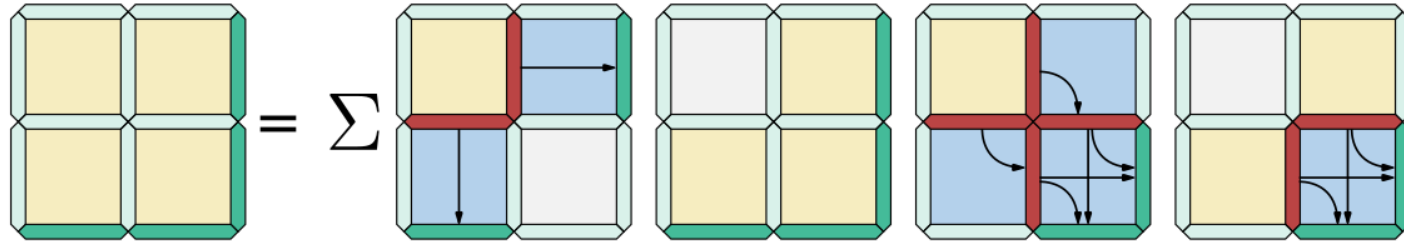


$$S = \begin{bmatrix} S_x & S_y \end{bmatrix}$$

$$T = \begin{bmatrix} T_{xx} & T_{xy} \\ T_{yx} & T_{yy} \end{bmatrix}$$

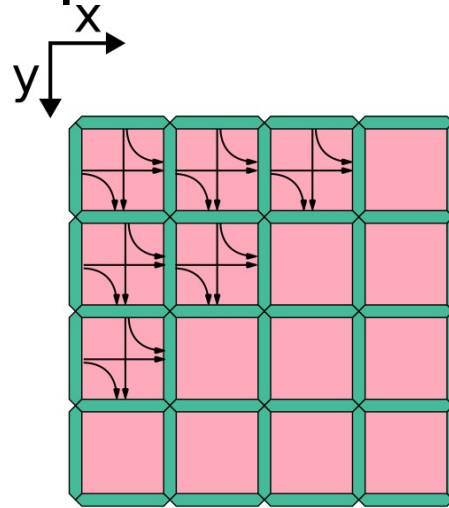
$$M = \begin{bmatrix} M_x \\ M_y \end{bmatrix}$$

Parallelization 2D



Long-Range Stability

- Exponentially many paths
 - from every edge two options
 - Activation blowup



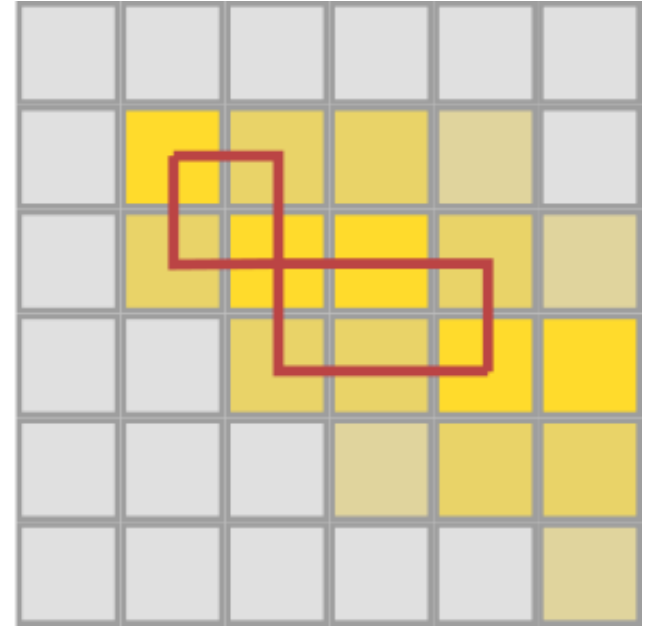
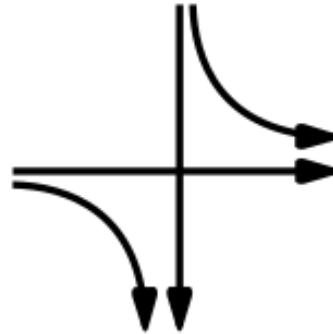
$$S = \begin{bmatrix} S_x & S_y \end{bmatrix}$$
$$T = \begin{bmatrix} T_{xx} & T_{xy} \\ T_{yx} & T_{yy} \end{bmatrix}$$
$$M = \begin{bmatrix} M_x \\ M_y \end{bmatrix}$$

P-mode: L_1 Norm on Transitions

$$|T_{xx}| + |T_{yx}| = 1$$

$$|T_{xy}| + |T_{yy}| = 1$$

$$T = \begin{bmatrix} \alpha & \alpha \\ 1 - \alpha & 1 - \alpha \end{bmatrix}$$

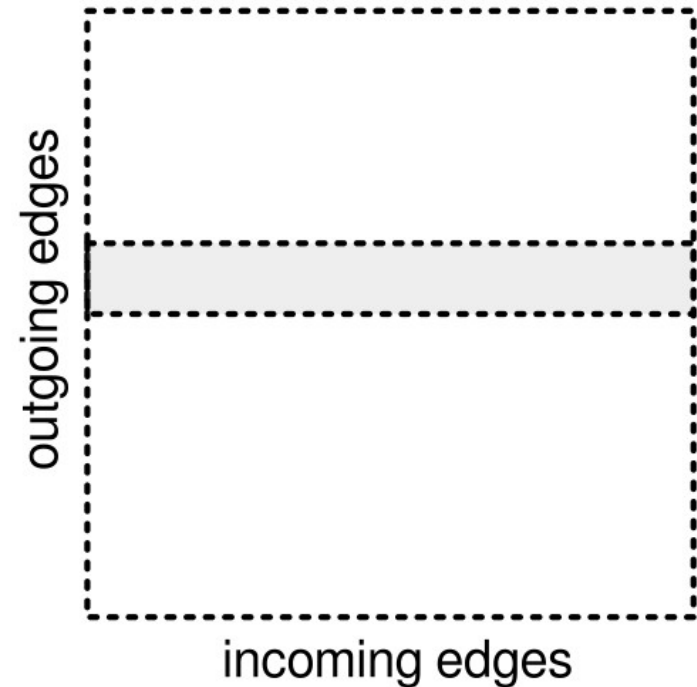


→ Directional Propagation

similar: Mamba2D
(Baty et. al. 2024)
but there fixed 45° angle 21

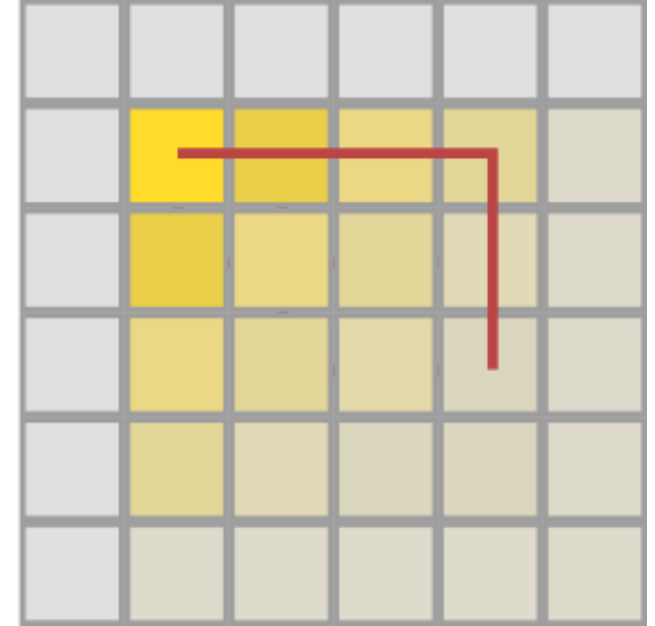
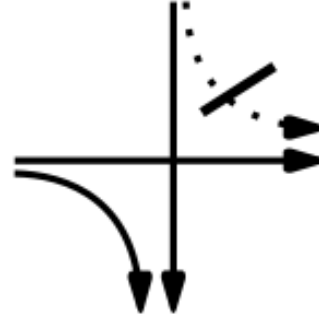
P-mode on a DAG

- L1 norm of the line-graph adjacency matrix composed of transition entries



D-mode: Single Connecting Path

$$T = \begin{bmatrix} \gamma_x & 1 \\ 0 & \gamma_y \end{bmatrix}$$

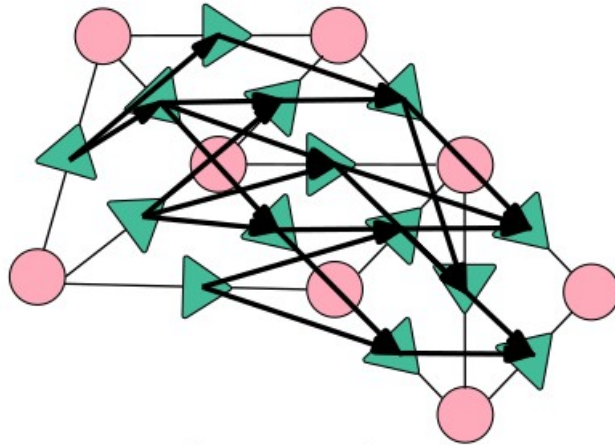


→ Diffusive Distribution

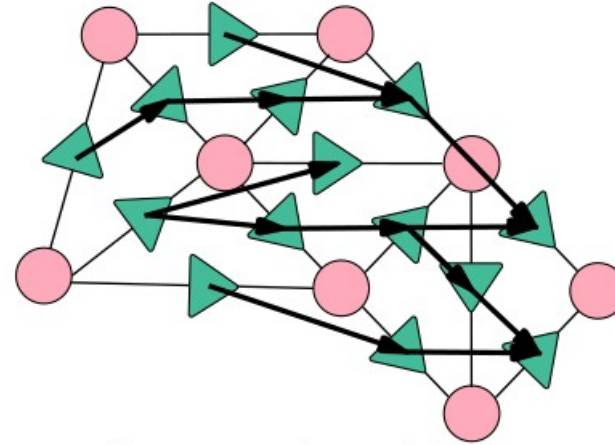
similar: 2DMamba
(Zhang et. al. 2024)

D-mode on a DAG

- Line graph is pruned to a multi-tree



line graph

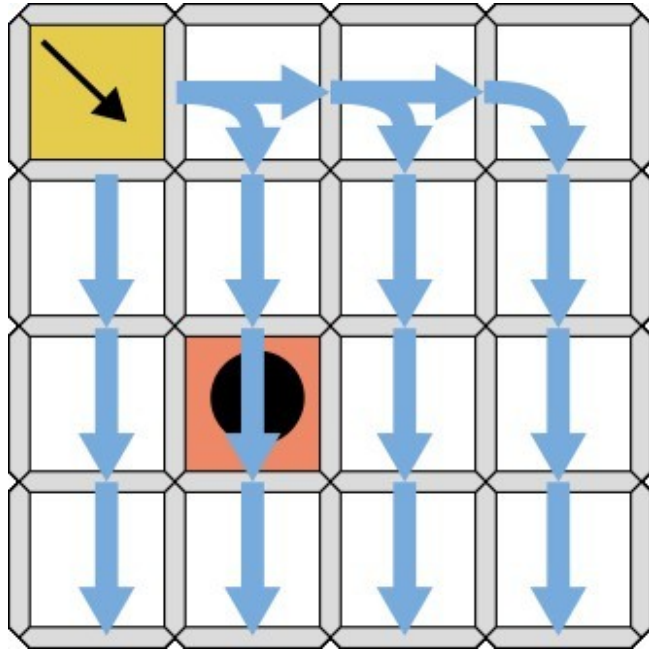


line graph multitree

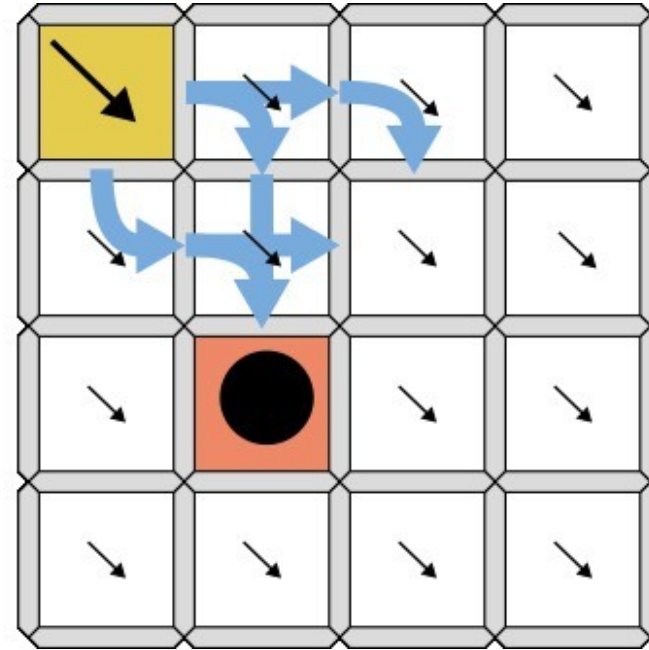
General Computational Graphs

- every finite computational graph can be unrolled to a DAG
- Back-Propagation uses gradients = linear approximations of the graph
- pLSTM analysis holds for all neural networks

pLSTM for Arrow Pointing

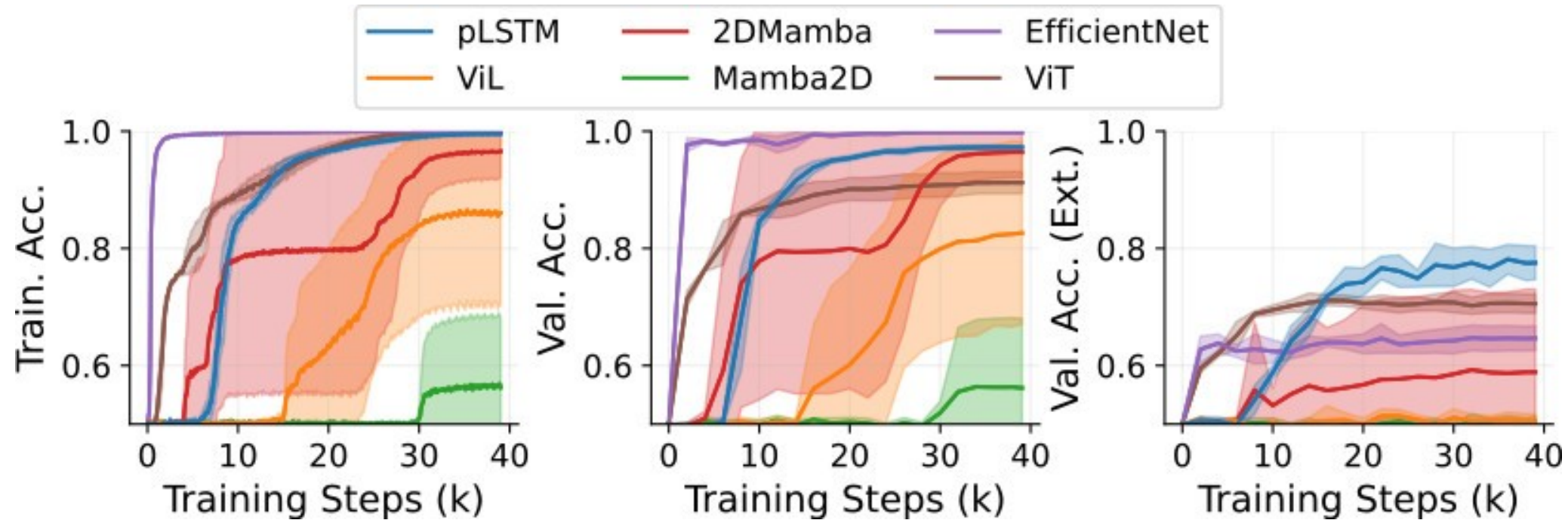


D-mode



P-mode

Arrow Pointing Extrapolation Results



ImageNet-1K

- pLSTM roughly on par with existing methods
- room for improvement by optimizing training schedule + backbone

Model	Epochs	#Params	FLOPS	IN-1K
EfficientNet-B0 [Tan and Le, 2019]	?	5M	0.39G	77.1
DeiT-T [Touvron et al., 2021]	300	6M	1.3G	72.2
DeiT-III-T (reimpl.) [Touvron et al., 2022]	800+20	6M	1.1G	75.4
VRWKV-T [Duan et al., 2024]	300	6M	1.2G	75.1
Vim-T [Zhu et al., 2024]	300	7M	1.5G	76.1
ViL-T [Alkin et al., 2025]	800+20	6M	1.3G	78.3
pLSTM-Vis-T	800+20	6M	1.4G	75.2
EfficientNet-B4 [Tan and Le, 2019]	?	19M	1.8G	82.9
DeiT-S [Touvron et al., 2021]	300	22M	4.6G	79.8
DeiT-III-S (reimpl.) [Touvron et al., 2022]	400+20	22M	4.6G	80.3
ConvNeXt-S (<i>iso.</i>) [Liu et al., 2022]	300	22M	4.3G	79.7
VRWKV-S [Duan et al., 2024]	300	24M	4.6G	80.1
Vim-S [Zhu et al., 2024]	300	26M	5.3G	80.5
Mamba2D-T [Baty et al., 2024]	300	27M	-	82.4
2DMamba-T [Zhang et al., 2024a]	?	30M	4.9G	82.8
ViL-S [Alkin et al., 2025]	400+20	23M	4.7G	81.5
pLSTM-Vis-S	400+20	23M	4.9G	80.7
EfficientNet-B6 [Tan and Le, 2019]	?	43M	19G	84.0
DeiT-B [Touvron et al., 2021]	300	86M	17.6G	81.8
DeiT-III-B (reimpl.) [Touvron et al., 2022]	400+20	87M	16.8G	83.5
ConvNeXt-B (<i>iso.</i>) [Liu et al., 2022]	300	87M	16.9G	82.0
VRWKV-B [Duan et al., 2024]	300	94M	18.2G	82.0
2DMamba-S [Zhang et al., 2024a]	?	50M	8.8G	83.8
ViL-B [Alkin et al., 2025]	400+5	89M	17.9G	82.4
pLSTM-Vis-B	400+20	89M	18.2 G	82.5

pLSTM on molecule graphs

model	MUTAG	NCI1	PROTEINS	PTC_FM	AVG
GAT [Veličković et al., 2018]	0.7822 $\pm$ 0.09	0.7968 $\pm$ 0.03	0.7215 $\pm$ 0.03	0.6105 $\pm$ 0.05	0.7277 $\pm$ 0.06
GCN [Kipf and Welling, 2017]	0.7234 $\pm$ 0.08	0.7852 $\pm$ 0.02	0.7395 $\pm$ 0.03	0.6162 $\pm$ 0.04	0.7161 $\pm$ 0.05
GIN [Xu et al., 2018]	0.8251 $\pm$ 0.10	0.8175 $\pm$ 0.02	0.7350 $\pm$ 0.04	0.6097 $\pm$ 0.10	0.7468 $\pm$ 0.08
LSTM GNN [Liang et al., 2016]	0.7450 $\pm$ 0.11	0.7951 $\pm$ 0.02	0.7503 $\pm$ 0.04	0.6076 $\pm$ 0.04	0.7245 $\pm$ 0.06
MPNN [Gilmer et al., 2017]	0.7450 $\pm$ 0.09	0.8012 $\pm$ 0.02	0.7350 $\pm$ 0.04	0.5786 $\pm$ 0.07	0.7149 $\pm$ 0.06
PLSTM	0.8512 $\pm$ 0.06	0.7324 $\pm$ 0.03	0.7502 $\pm$ 0.05	0.6133 $\pm$ 0.08	0.7368 $\pm$ 0.06

- pLSTM roughly on par with existing methods
- room for improvement by including more inductive biases

Conclusion

- Contributions:
 - translating linear RNNs to multiple dimensions + general DAGs, including a parallelization scheme
 - Long-range stabilization scheme for linear networks on DAGs with P- and D-mode
 - Arrow-Pointing Extrapolation (APE) task as synthetic benchmark for long-range directional capabilities
- Limitations / Future work:
 - Mechanistically investigate solution modes of pLSTM (and others) on APE
 - Improve results on common benchmark, by integrating further inductive biases
 - Harder real-world benchmarks that need long-range multi-dimensional relationships (e.g. maps?)

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