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Energy Transformer (E.T.) is a novel architecture that is an **Energy-Based Model**, an **Associative Memory**, and a **Transformer**. Specifically, **E.T.** looks like a recurrent Transformer block that defines an attractor system (*an O.D.E. with guaranteed fixed points convergence*) that performs error correction via energy descent. **E.T.** shows excellent performance on MASKed image in-painting, graph anomaly detection, and graph classification.



E.T. is Energy-Based Associative Memory

E.T. is Interpretable by Design

Minimizing energy (inference) performs error correction on corrupted data.

Masked Input



Predictions



Visualize dynamic states (tokens and updates) at any time



E.T. is a Transformer -

 $E^{\text{ATT}} = -\frac{1}{\beta} \sum_{h} \sum_{C} \log \left(\sum_{B \neq C} \exp \left(\beta \sum_{\alpha} K_{\alpha h B} Q_{\alpha h C} \right) \right)$



Standard Transformer Block



E.T. sets SOTA on Graphs

Node anomaly detection

Graph Classification

Dataset	s Split	Top Baseline Score	ET (Ours)		
Yelp	1%	$62.1_{\pm 1.3}$	63.0 _{±0.6} ▲0.9	F	$\mathbf{g}(\mathbf{x}^{t}) \qquad \mathbf{x}^{t+1} = \mathbf{x}^{t} - \alpha \nabla_{\mathbf{g}} E$ Energy Transformer Block $\nabla_{\mathbf{g}} E^{t}$ Minimizing energy
	40%	$71.0_{\pm 0.9}$	$71.5_{\pm 0.1} \ 0.5$		
	1%	$\boxed{90.9_{\pm 0.7}}$	$89.3_{\pm 0.7}$ $\bigcirc 0.7$		
Å Amazon	40%	$92.2_{\pm 0.4}$	92.8_{±0.3} ▲0.6		
	. 1%	$84.8_{\pm 0.0}$	85 .1 $_{\pm 1.0}$ \land 0.3		
	e = 40%	$86.8_{\pm 0.0}$	88.2_{±1.0} ▲1.4		
T-Social	1%	$75.9_{\pm 0.0}$	79 .1 _{±0.7} ▲3.2		
	40%	$83.9_{\pm 0.0}$	$83.5_{\pm 0.4}$ $\bigcirc 0.4$		
Vala	1%	$75.4_{\pm 0.9}$	$73.2_{\pm 0.8}$ $\checkmark 2.2$		
reib	40%	$84.0_{\pm 0.9}$	84.9_{±0.3} ▲0.9		
	1%	$90.4_{\pm 2.0}$	91.9_{±1.0} ▲1.5	\rightarrow time	accidence miccied
	40%	$98.0_{\pm0.4}$	$97.3_{\pm 0.4}$ $\bigcirc 0.7$		assigns missing
	. 1%	$91.1_{\pm 0.0}$	92 .8 $_{\pm 1.1}$ \land 1.7		labels to nodes
' I-Fillan(e = 40%	$94.3_{\pm 0.0}$	95.0 _{±3.0} ▲0.7		
T-Social	1%	$88.0_{\pm 0.0}$	91.9 ±0.6 ▲3.9		
	40%	$95.2_{\pm 0.0}$	$93.9_{\pm 0.2}$ V 1.3		

Datasets	Top Baseline Score	ET (Ours)
PROTEINS	$84.9_{\pm 1.6}$	90.3 $_{\pm 5.4}$ $\blacktriangle 5.4$
NCI1	$87.5_{\pm 0.5}$	90 .1 $_{\pm 0.1}$ $\triangle 2.6$
NCI109	$87.4_{\pm 0.3}$	90 .5 $_{\pm 0.1}$ $\checkmark 3.1$
DD	$95.7_{\pm 1.9}$	$95.9_{\pm 0.8}$ $\land 0.2$
ENZYMES	$78.4_{\pm 0.6}$	99.8 $_{\pm 0.0}$ $\triangle 21.4$
MUTAG	$100.0_{\pm 0.0}$	$96.6_{\pm 0.2}$ V 3.4
MUTAGENICITY	$82.2_{\pm 0.6}$	98.7 $_{\pm 0.1}$ 16.5
FRANKENSTEIN	$78.9_{\pm 0.3}$	99.8 $_{\pm 0.1}$ 2 0.9





