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➤ Motivations

- ❖ Unlike images, graph data are inherently **heterogenous** (e.g. pandemics, product co-purchase relation, molecules).
- ❖ The SOTA GraphCL [1] handles the heterogeneity with ad-hoc choices of augmentations (**priors**) for every datasets.
- ❖ Rather than the **discrete** selection on prefabricated ones, can we directly learn and generate such priors, **continuously**?
- ❖ **Question**: What us the **space**, **principle** and **framework** for learnable GraphCL priors (GraphCL-LP)?

➤ Method. Graph Generative Models as Learnable Priors

- ❖ We define the **space** of GraphCL priors as the set of stochastic mappings between graph manifolds $m: \mathcal{G} \rightarrow \mathcal{G}$.
- ❖ Naturally, we adopt the recent rising graph generative models (VGAE [2] here) for the prior space **parametrization**.
- ❖ Further, **principled** reward signals are introduced to convey messages from contrastive learning to generator training.
- ❖ Above components are assembled into the bi-level optimization **framework**.

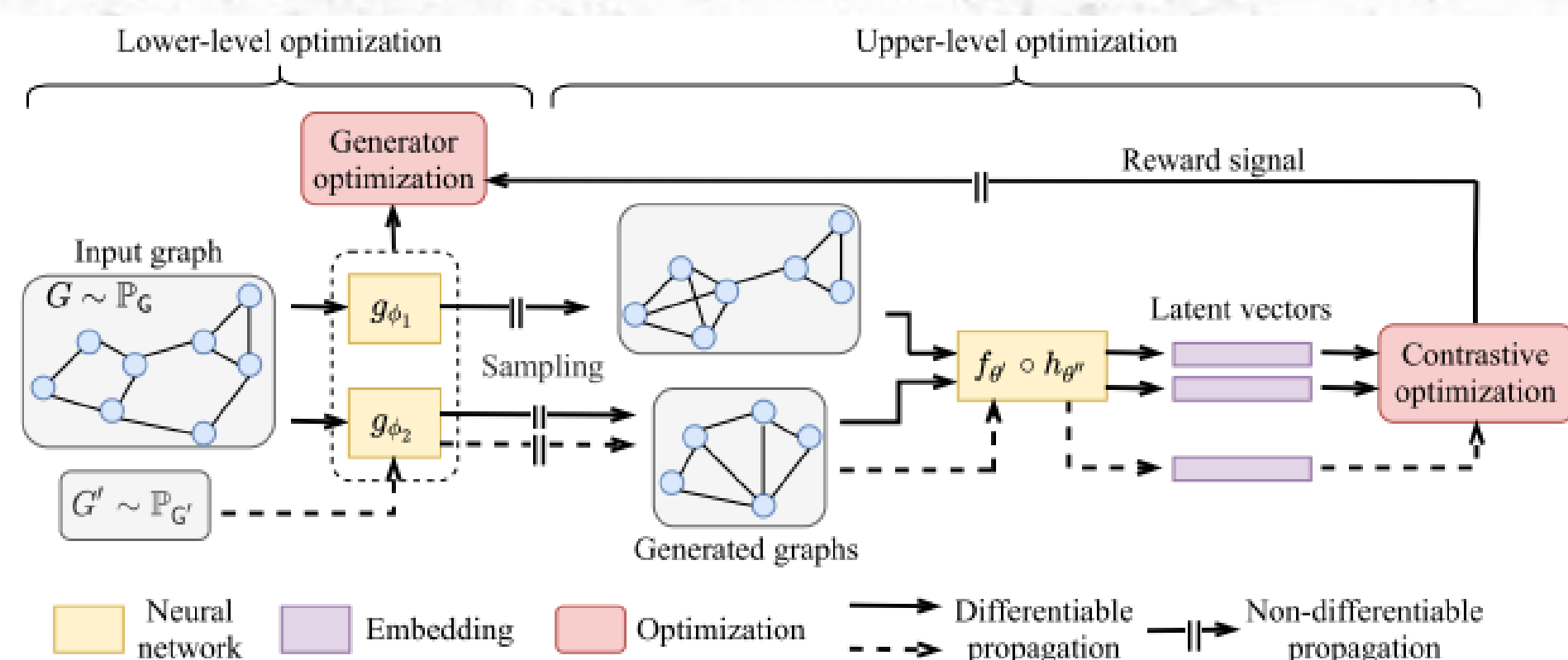


Figure 2: Pipeline of GraphCL with learned prior. Graph generative models g_{ϕ_1}, g_{ϕ_2} generate contrastive views for self-supervised contrasting, and then receive the reward for their parameter update.

[1] Yuning You et al., “Graph Contrastive Learning with Augmentations”, NeurIPS’20. [2] Thomas Kipf & Max Welling, “Variational Graph Auto-Encoders”, NeurIPS’16 Workshop. [3] Chence Shi et al., “Graphaf: a flow-based autoregressive model for molecular graph generation”, ICLR’20.

➤ References

➤ Method. Principles for Learning Priors to Contrast

- ❖ Information minimization (**InfoMin**). Encouraging contrastive views to share less mutual information.
- ❖ Information bottleneck (**InfoBN**). Diminishing the information overlap between each contrastive view and its latent representation.

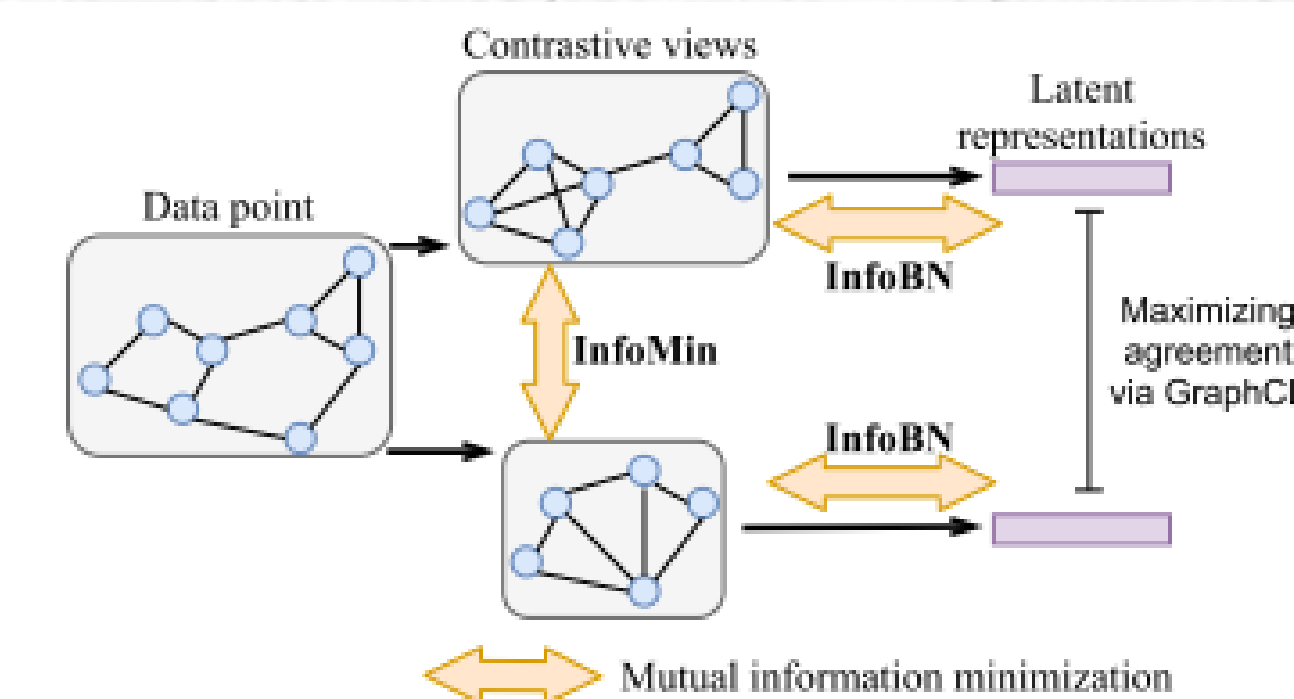


Figure 3: Schematic diagram of the InfoMin and InfoBN principles to guide the prior learning in GraphCL.

➤ Experiments

- ❖ We numerically show the competitive performance of GraphCL-LP versus SOTA methods.

Table 2: Semi-supervised learning on small-scale benchmarks from TUDataset (the first four) and large-scale ones from OGB (the last two). Shown in **red** are the best three accuracies (%) for TUDataset and the best for ogbg-ppa and F1-score (%) for ogbg-code. The SOTA results compared here are as published under the same experimental setting (- indicates that results were not available in corresponding publications).

Methods	COLLAB	RDT-B	RDT-M5K	GITHUB	ogbg-ppa	ogbg-code
No pre-train	73.71±0.27	86.63±0.27	51.33±0.44	60.87±0.17	56.01±1.05	17.85±0.60
Augmentations	74.19±0.13	87.74±0.39	52.01±0.20	60.91±0.32	-	-
GAE	75.09±0.19	87.69±0.40	53.58±0.13	63.89±0.52	-	-
Infomax	73.76±0.29	88.66±0.95	53.61±0.31	65.21±0.88	-	-
ContextPred	73.69±0.37	84.76±0.52	51.23±0.84	62.35±0.73	-	-
GraphCL	74.23±0.21	89.11±0.19	52.55±0.45	65.81±0.79	57.77±1.25	22.45±0.17
LP-InfoMin	74.66±0.14	88.03±0.46	53.00±0.26	62.71±0.54	59.10±0.88	23.50±0.22
LP-InfoBN	74.61±0.28	87.64±0.33	53.05±0.14	62.64±0.37	55.48±0.97	23.31±0.22
LP-Info(Min&BN)	74.84±0.31	87.81±0.45	53.32±0.23	63.11±0.33	57.31±0.99	23.61±0.27

- ❖ Further analysis show: graph generation quality usually aligns with downstream performance; and also molecule-specific generator (GraphAF [3] here) alone does not significantly benefit molecular datasets.

Table 5: Link prediction performance (AUROC and AUPRC, %) of VGAE generators on eight pre-training datasets. Better link prediction results are marked in **red** if accompanied with better downstream performances, as shown in Table 2, 3 and 4.

	Principles	COLLAB	RDT-B	RDT-M5K	GITHUB	ogbg-ppa	ogbg-code	Trans-Mol	Trans-PPI
AUROC (%)	InfoMin	71.28	97.32	99.08	78.68	96.53	92.39	64.54	71.20
	InfoBN	69.44	97.29	99.31	81.20	95.24	94.06	83.55	71.32
AUPRC (%)	InfoMin	80.84	96.62	98.67	78.49	95.94	90.08	64.14	69.34
	InfoBN	79.13	96.59	98.97	80.47	95.30	91.66	82.51	70.65

Table 6: Learned prior performance with different generators under the guidance of InfoMin, in the transfer learning setting on molecular datasets. **Red** numbers indicate the best performances (AUROC, %).

Methods	BBBP	Tox21	ToxCast	SIDER	ClinTox	MUV	HIV	BACE
VGAE	71.47±0.66	74.60±0.70	63.13±0.30	60.52±0.75	72.39±1.50	70.51±2.25	76.43±0.85	78.86±1.66
GraphAF	70.55±0.63	73.51±0.43	62.03±0.33	61.32±1.32	77.47±1.91	72.25±1.18	76.30±1.34	78.43±2.36