



Graph Convolutional Policy Network for Goal-Directed Molecular Graph Generation

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Preliminaries & Related Work

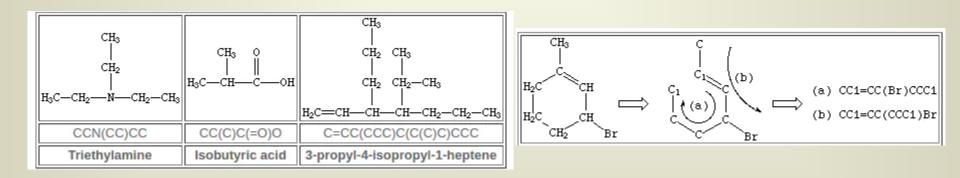
A Brief Introduction of Molecule Graph Generation



Representation of Molecules

- Sequential representation: SMILES
- Graph representation: structural formula

(recap high school chemistry)



Graph ConvNets (GCNs)



GCN Convolves over node neighborhoods.

• As Matrix Multiplication:

 $H^{(t+1)} = \sigma \left(A H^{(t)} W^{(t)} \right)$

• As Neighborhood Aggregation:

$$H_u^{(t+1)} = \sigma\left(\sum_{v \in N(u)} H_v^{(t)} W^{(t)}\right)$$

(two flips of one coin)

MolGraph Generation



Related work in MolGraph Generation (GCN based).

MolGraph + VAE = GraphVAE (2018) MolGraph + VAE + structural priors = JT-VAE (2018) MolGraph + GAN = MolGAN (2018) MolGraph + Flow = GraphNVP (2019); GraphAF (2020) MolGraph + RL + GAN = GCPN (2018)

(arXiv year)

Markovian Graph Generation (MDP)

Motivations & Basic Schemes Behind GCPN

MolGraph Gen. as MDP



Markovian Decision Process

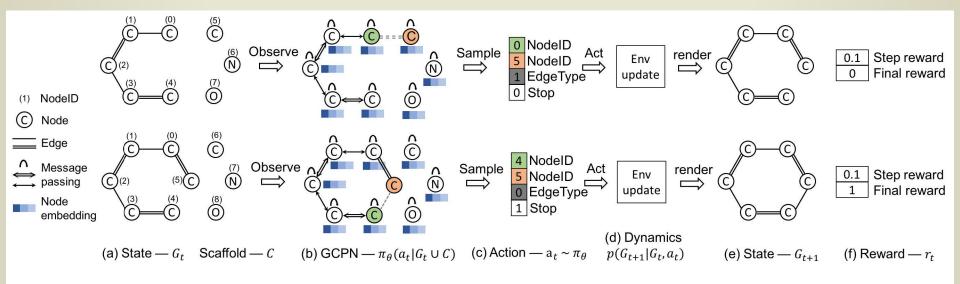
Markovian Condition:

$$p(s_{t+1}|s_t, s_{t-1}, \cdots) = p(s_{t+1}|s_t)$$

 In GCPN, this condition means in the generation process, each step is determined fully by the <u>intermediately</u> <u>generated molecule</u>.

GCPN Overview



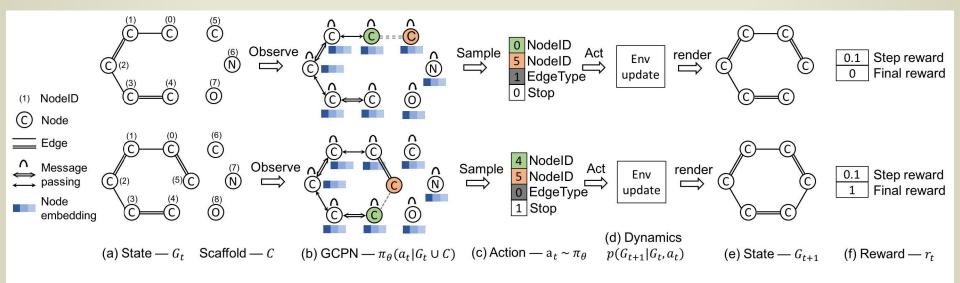


Overall techniques: RL + Adversarial Loss

- Sequentially constructing a MolGraph Using MDP
- The agent gain rewards according to chem. rules and GAN validity

GCPN Overview





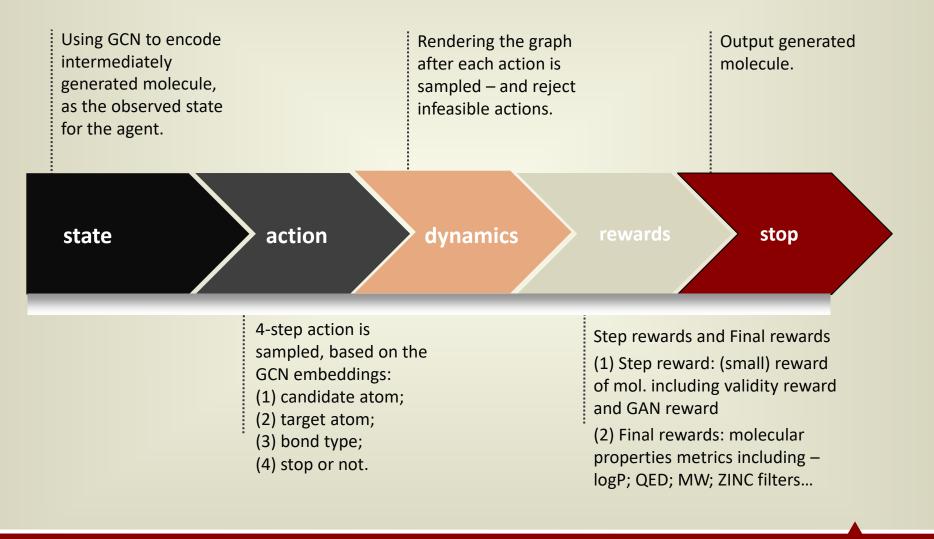
In each step, an RL agent

- either connect a new atom to an existing one
- or connect two existing atoms.
- (as well as) choose whether to stop.

Model Details: RL setup in GCPN

Details *(brief, actually)* in the RL Implementation of GCPN

Generation Scheme



State



(Heterogenous) GCN state encoder.

•
$$H^{(l+1)} = \operatorname{AGG}(\operatorname{ReLU}(\{\tilde{D}_i^{-\frac{1}{2}}\tilde{E}_i\tilde{D}_i^{-\frac{1}{2}}H^{(l)}W_i^{(l)}\}, \forall i \in (1, ..., b)))$$

Note: the AGG(·) is conducted for different bond types.
(distinguish with graph pooling / neighborhood aggregation)

Action



Four-step action generated from states. Generated & sampled sequentially.

$$\begin{aligned} a_{t+1} &= \operatorname{CONCAT}(a_{\operatorname{first}}, a_{\operatorname{second}}, a_{\operatorname{edge}}, a_{\operatorname{stop}}) \\ f_{\operatorname{first}}(s_t) &= \operatorname{SOFTMAX}(m_f(X)), \\ f_{\operatorname{second}}(s_t) &= \operatorname{SOFTMAX}(m_s(X_{a_{\operatorname{first}}}, X)), \\ f_{\operatorname{edge}}(s_t) &= \operatorname{SOFTMAX}(m_e(X_{a_{\operatorname{first}}}, X_{a_{\operatorname{second}}})), \\ f_{\operatorname{stop}}(s_t) &= \operatorname{SOFTMAX}(m_t(\operatorname{AGG}(X))), \end{aligned}$$

 $a_{\text{first}} \sim f_{\text{first}}(s_t) \in \{0, 1\}^n$ $a_{\text{second}} \sim f_{\text{second}}(s_t) \in \{0, 1\}^{n+c}$ $a_{\text{edge}} \sim f_{\text{edge}}(s_t) \in \{0, 1\}^b$ $a_{\text{stop}} \sim f_{\text{stop}}(s_t) \in \{0, 1\}$





Produce ("render") the graph according to the sampled action. Reject infeasible actions. (Deterministic state trans. dynamics.)

Reward



Step (small) & Final (large) Reward

- Step reward 1: validity. Penalize infeasible actions.
- Step reward 2: -1 * GAN Loss. Penalize "weird" samples.
- Final reward: molecular properties
 - logP: octanol-water partition coefficient (脂水分配系数)
 - QED: druglikeness (类药性)
 - MW: molecular weights (分子量)
 - Penalty over unrealistic molecules (经验标准)
 - Zinc fn. group filters: (生医标准,来自于商用药物分子库 ZINC)

Training



Policy Gradient + Pretraining.

• **Proximal Policy Optimization (PPO)**

We will not unpack it here.See ref. J. Schulman, F. Wolski, P. Dhariwal, A. Radford, and O. Klimov.Proximal policy optimization algorithms. CoRR, abs/1707.06347, 2017.

Pretraining using generation trajectory of true molecules.

(could be very important !)

Experiments & Observations

Some Brief Results of GCPN

Experiments



Property optimization

Gen. mols with some property optimized.

Property Targeting

Gen. mols with some property closed to a given target.

Constraint Property Optimization

Conduct property optimization with the molecule

contains a given substructure.

Comp. v.s. SOTA (JT-VAE / ORGAN)



Table 1: Comparison of the top 3 property scores of generated molecules found by each model.

Method	Penalized logP				QED			
	1st	2nd	3rd	Validity	1st	2nd	3rd	Validity
ZINC	4.52	4.30	4.23	100.0%	0.948	0.948	0.948	100.0%
ORGAN	3.63	3.49	3.44	0.4%	0.896	0.824	0.820	2.2%
JT-VAE	5.30	4.93	4.49	100.0%	0.925	0.911	0.910	100.0%
GCPN	7.98	7.85	7.80	100.0%	0.948	0.947	0.946	100.0%

Table 2: Comparison of the effectiveness of property targeting task.

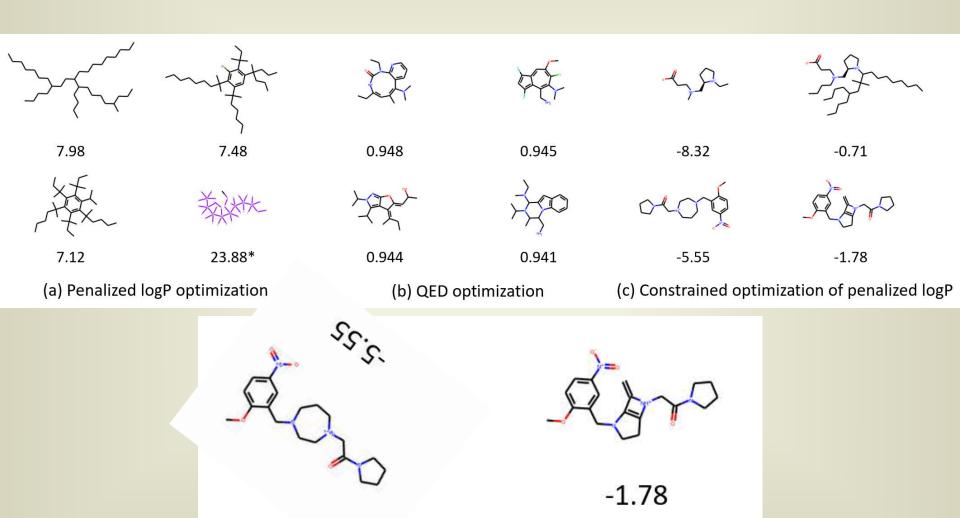
Method -	$-2.5 \le \log P \le -2$		$5 \le \log P \le 5.5$		$150 \leq \mathrm{MW} \leq 200$		$500 \le \mathrm{MW} \le 550$	
	Success	Diversity	Success	Diversity	Success	Diversity	Success	Diversity
ZINC	0.3%	0.919	1.3%	0.909	1.7%	0.938	0	_
JT-VAE	11.3%	0.846	7.6%	0.907	0.7%	0.824	16.0%	0.898
ORGAN	0	_	0.2%	0.909	15.1%	0.759	0.1%	0.907
GCPN	85.5%	0.392	54.7%	0.855	76.1%	0.921	74.1 %	0.920

Table 3: Comparison of the performance in the constrained optimization task.

δ		JT-VAE		GCPN			
	Improvement	Similarity	Success	Improvement	Similarity	Success	
0.0	1.91 ± 2.04	0.28 ± 0.15	97.5%	4.20 ± 1.28	0.32 ± 0.12	100.0%	
0.2	1.68 ± 1.85	0.33 ± 0.13	97.1%	4.12 ± 1.19	0.34 ± 0.11	100.0 %	
0.4	0.84 ± 1.45	0.51 ± 0.10	83.6%	2.49 ± 1.30	0.47 ± 0.08	100.0 %	
0.6	0.21 ± 0.71	0.69 ± 0.06	46.4%	$\boldsymbol{0.79 \pm 0.63}$	0.68 ± 0.08	$\mathbf{100.0\%}$	

Some demo.





Summary of Observation



- Indeed, GCPN produced very "valid" results.
- This is probably the results of its elaborated process of generation, including the trajectory pretraining & GAN loss, while the diversity is somehow harmed.
- No chem. or bio. motivation is adequately combined in the generation process. Therefore, some actions of the agent seems bizarre.

Thanks

<u>References</u>



(All available in arXiv so only arXiv ID provided.)

- 1707.06347 PPO
- 1802.03480 GraphVAE
- 1802.04364 JT-VAE
- 1805.11973 MolGAN
- 1806.02473 GCPN
- 1905.11600 GraphNVP
- 2001.09382 GraphAF