Retrosynthesis Planning via Worst-path Policy **Optimisation in Tree-structured MDPs**

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A Brief Introduction

estimated advantage.

Successful Synthetic Tree

We develop a weighted self-imitation

worst-path return when first expanding s with reaction a and subsequently

$$Q^{\pi}(s,a) = \mathbb{E}_{\tau \sim \pi} \left[\min_{p \sim P(\tau)} \sum_{t=0}^{T} \gamma^{t} r(s_{t}) \mid s_{0} = s, a_{0} = a \right]$$

$$V^{\pi}(s) = \mathbb{E}_{\tau \sim \pi} \left[\min_{p \sim P(\tau)} \sum_{t=0}^{T} \gamma^{t} r(s_{t}) \mid s_{0} = s \right]$$

quantifies the relative benefit of applying reaction a to molecule s compared to following the policy:

Our objective is to find a policy that maximises the worst-path objective. We achieve this through an iterative procedure: in each iteration i, we aim to find an improved policy π^{i+1} by imitating advantageous state-action pairs (s, a) experienced under policy π^i . The advantage $A^{\pi^i}(s,a)$ quantifies this, and the learning objective for π^{i+1} is formulated as:

$$J(\pi^{i+1}) = \mathbb{E}_{s \sim d_{-i}(\cdot), a \sim \pi^{i}(\cdot \mid s)} \left[\exp \left(\beta A^{\pi^{i}}(s, a) \right) \log \pi^{i+1}(a \mid s) \right]$$

where $\beta > 0$ is the advantage coefficient controlling the strength of advantage toward better-than-average reactions.

algorithm to optimise a worst-path objective.

Given a molecule s as the root node, a Q-function estimates the expected following policy π :

Here, s_0 and a_0 denote the root molecule and the initial reaction, respectively. Similarly, the value function $V^{\pi}(s)$ represents the expected worst-path return when all subsequent reactions follow policy π :

$$V^{\pi}(s) = \mathbb{E}_{\tau \sim \pi} \left[\min_{p \sim P(\tau)} \sum_{t=0}^{T} \gamma^{t} r(s_{t}) \mid s_{0} = s \right]$$

With these definitions in place, we can express the advantage function, which

$$A^{\pi}(s, a) = Q^{\pi}(s, a) - V^{\pi}(s)$$

A positive advantage indicates that reaction a leads to better outcomes than the policy's average behaviour.

$$J(\pi^{i+1}) = \mathbb{E}_{s \sim d_{\pi^{i}}(\cdot), a \sim \pi^{i}(\cdot \mid s)} \left[\exp\left(\beta A^{\pi^{i}}(s, a)\right) \log \pi^{i+1}(a \mid s) \right]$$

weighting, and d_{π^i} is the state distribution induced by policy π^i . In this case, reactions with higher advantages receive higher weights, guiding the policy

We propose InterRetro, a search-free approach to multi-step retrosynthesis planning.

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Algorithm 1 Interactive retrosynthesis planning (InterRetro).
Input: pre-trained one-step policy \pi_{\theta}, value def INTERRETRO(\pi_{\theta}, m):
     function V_{\phi}, training set \mathcal{D}, replay buffer \mathcal{B}.
                                                                        1: for i = 1, ..., I do
def EXPLORE(\pi_{\theta}, m):
                                                                                 while \mathcal{D} is not empty do
                                                                                    m \leftarrow \mathcal{D}.\mathsf{pop}()
 1: tree \leftarrow Tree(root = m)
 2: q \leftarrow \{m\}
                                                                                    \mathsf{tree} \leftarrow \mathsf{EXPLORE}(\pi_{\theta}, m)
                                                                                    \mathtt{brs} \leftarrow \{\}
 3: step \leftarrow 0
     while q \neq \emptyset and step < max_steps do
                                                                                    for each subtree \tau \in \mathsf{tree} \ \mathsf{do}
        s \leftarrow q.pop()
                                                                                        if \tau is successful then
        a, \mathcal{S}_r \leftarrow \pi_{\theta}.\mathtt{get\_reactants}(s)
                                                                                           brs \leftarrow brs \cup ALLBRANCHES(\tau)
         \mathtt{tree.add\_branch}(s, a, \mathcal{S}_r)
                                                                                        end if
                                                                                    end for
        # Add non-building blocks
                                                                                    \mathcal{B}.append(brs)
                                                                                   \texttt{branches} \leftarrow \mathcal{B}.\texttt{sample()}
        q \leftarrow q \cup \{ s' \in \mathcal{S}_r \mid s' \notin \mathcal{S}_{bb} \}
                                                                                    V_{\phi}.update(branches)
                                                                                                                              ⊳ Eq. 15
                                                                                    \pi_{\theta}.update(V_{\phi}, \text{ branches}) \triangleright \text{Eq. } 16
        \mathtt{step} \leftarrow \mathtt{step} + 1
13: end while
                                                                                end while
                                                                       16: end for
14: return tree
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We demonstrate that InterRetro is SOTA.

- success rate (achieving up to 100% on benchmark datasets),
- route length (reducing steps by 4.9%), and
- sample efficiency (reaching 92% of full performance with only 10% of training data).

Table 1: Performance evaluation on three benchmarks. The evaluation metrics include the success rate under different test molecules with different budgets of model calls, which are direct generation (DG), 100, 200 and 500 model calls. The DG columns are single-step model's results without search.

	Retro -190				CUEMIRT-1000				GDB17-1000				
Single-step	Search	DG	100	200	500	DG	100	200	500	DG	100	200	500
Template	MCTS	20.00	43.68	47.37	62.63	32.00	45.60	68.80	71.90	3.00	3.20	3.70	4.50
Template	Retro*	20.00	38.42	58.42	75.26	32.00	69.10	72.00	74.70	3.00	5.40	6.60	7.50
LocalRetro	MCTS	22.10	44.21	57.36	62.10	47.30	62.70	69.10	75.00	4.60	14.00	16.70	20.30
LocalRetro	Retro*	22.10	58.94	64.73	73.68	47.30	74.80	80.40	82.40	4.60	18.90	22.20	28.80
MEGAN	Retro*	8.42	60.52	62.10	73.15	38.00	71.70	75.40	79.00	6.20	37.60	45.70	57.20
Graph2Edits	Retro*	16.84	41.05	50.00	56.31	47.10	68.70	78.80	80.70	5.90	18.20	24.00	32.20
Self-improve	Retro*	_	67.37	83.16	94.74	_	_	_	81.10	_	_	_	15.00
PDVN	Retro*	_	93.68	97.37	98.95	_	_	_	83.50	_	_	_	26.90
RetroCaptioner	Retro*	5.26	68.94	72.63	85.26	3.90	72.60	76.50	78.70	3.20	56.20	68.20	75.20
DreamRetroer		32.10	78.94	88.42	90.52	31.10	78.10	81.70	83.10	4.20	27.36	28.97	33.20
InterRetro	MCTS	95.78	89.47	98.94	100.00	93.10	78.40	89.30	97.50	89.00	80.80	96.10	99.50
InterRetro	Retro*	95.78	96.31	100.00	100.00	93.10	91.40	96.20	98.20	89.00	83.80	96.50	97.20

$P(\tau) = \{ABD, ABEF, ABEGH, AC\}.$

As shown in the figure, each non-leaf node represents a molecule that is

decomposed into one or more reactants. Left tree: A successful synthetic

Examples illustrating the tree MDP formulation.

Failed Synthetic Tree

Retrosynthesis planning aims to decompose target molecules into available

building blocks, forming a synthetic tree where each internal node represents

an intermediate compound and each leaf ideally corresponds to a purchasable

reactant. However, this tree becomes invalid if any leaf node is not a valid

building block, making the planning process vulnerable to the "weakest link" in

the synthetic route. Existing methods often optimise for average performance

In this paper, we reframe retrosynthesis as a worst-path optimisation problem

within tree-structured Markov Decision Processes (MDPs). We prove that this

formulation admits a unique optimal solution and provides monotonic

improvement guarantees. Building on this insight, we introduce Interactive

Retrosynthesis Planning (InterRetro), a method that interacts with the tree

MDP, learns a value function for worst-path outcomes, and improves its policy

through self-imitation, preferentially reinforcing past decisions with high

We formulate the retrosynthesis planning

problem into tree-structured MDPs.

Building Blocks

route for target molecule A. It contains 4 root-to-leaf paths:

across branches, failing to account for this worst-case sensitivity.

Since all leaf nodes are building blocks, each path receives a value of γ^T , where T is the path length. The tree's overall value is

$$\min_{p \in P(\tau)} \{ \gamma^2, \gamma^3, \gamma^4, \gamma \} = \gamma^4,$$

determined by the longest path. Right tree: A failed synthesis attempt for molecule A. One of its paths, ABEG, terminates at G, which is not a building block. This gives path *ABEG* a value of 0, making the tree's overall value

$$\min_{p \in P(\tau)} \{ \gamma^2, \gamma^3, 0, \gamma \} = 0,$$

illustrating why a single failing path invalidates the entire route.

References

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Due to a visa delay, the first author is unable to attend the conference in person. If you have any questions about the project, please feel free to contact him at mianchu.wang@outlook.com or via WeChat by scanning the QR code.

