

LING682 FINAL PROJECT

# Controlled Molecular Generation

Using Sequential Monte Carlo Sampling with Small Language Models

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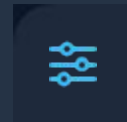
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# The Challenge



## De Novo Discovery

Modern drug discovery aims to reveal novel, valid compounds satisfying specific physicochemical properties. Traditional methods are limited by predefined rules and fragment libraries that restrict accessible chemical space.



## The Control Problem

A persistent challenge in AI-based generation is achieving reliable control over properties like **Molecular Weight (MW)**, **Lipophilicity (logP)**..etc without incurring massive computational costs.

# Sequential Monte Carlo (SMC)



## 1. Importance Sampling

Estimates expectations under an intractable target distribution by drawing samples from a proposal distribution and reweighting them effectively.



## 2. Sequential Update

Weights are updated dynamically as tokens are appended. Potential functions score how well a partial sequence matches the desired constraints.



## 3. Adaptive Resampling

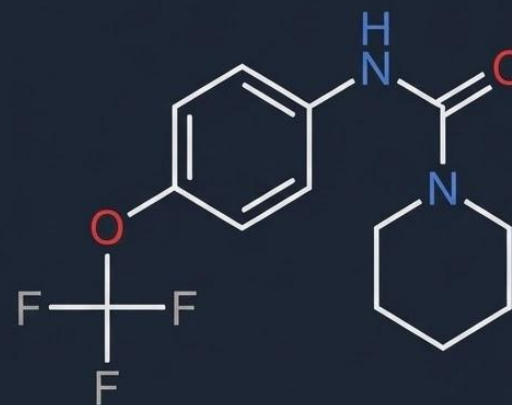
Prevents particle collapse by replicating high-weight particles and discarding low-weight ones.

# Experimental Setup

We compare a small model against a massive fine-tuned model on constrained molecules generation.

Feature	GPT2-ZINC	Smiley Llama
Size	87 Million	8 Billion
Training	ZINC (~480M)	ChEMBL (~2M)
Input	SMILES Prefix	Natural Language
Method	SMC-Guided	Fine-tuning

## Chemical Structure Input



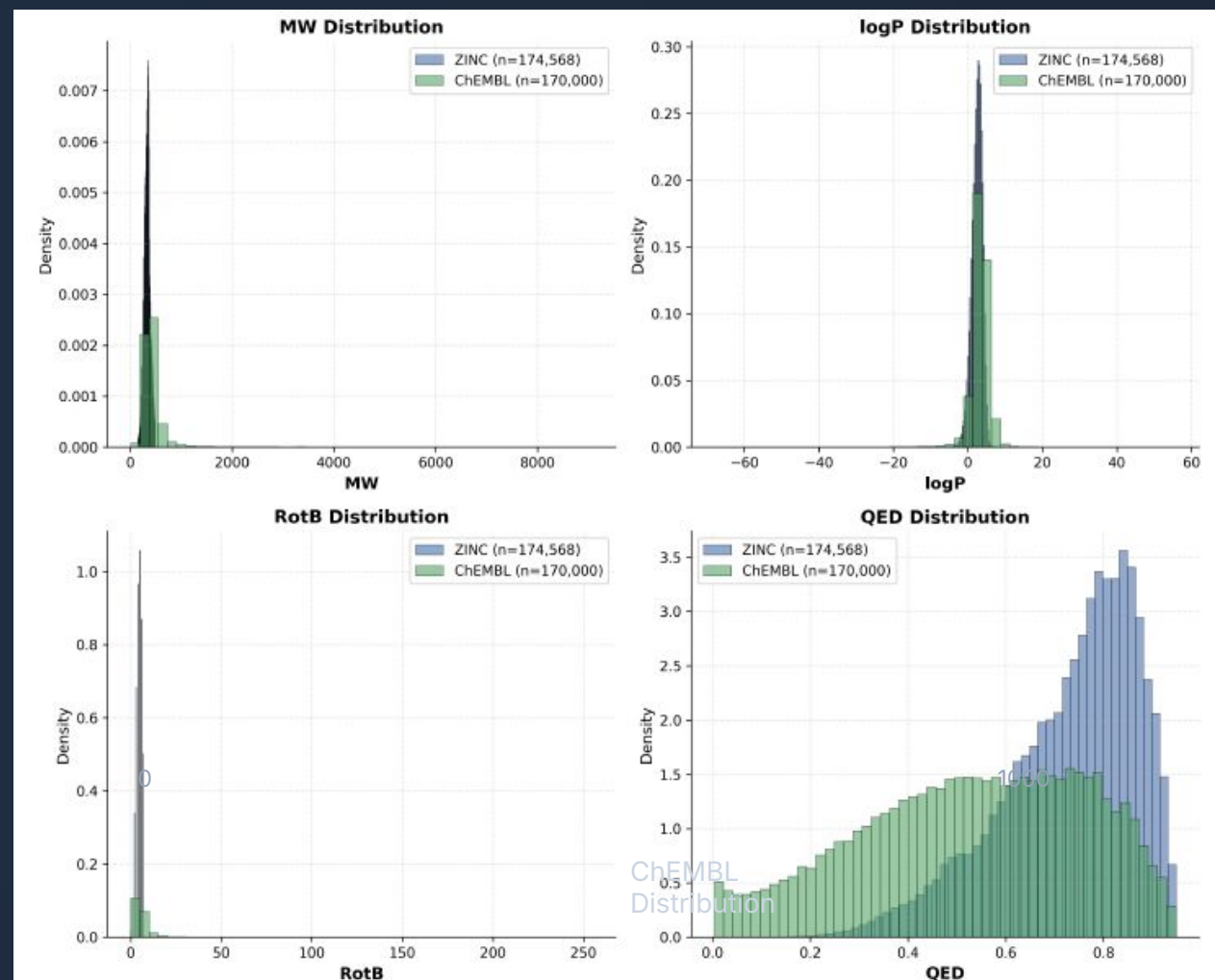
SMILES String:  
O=C(Nc1ccc(OC(F)(F)F)cc1)N1CCCCC1

# Data Distribution Analysis

## Property Overlap

Despite the immense difference in dataset size (ZINC: ~480M vs ChEMBL: ~2M), the distributions of key properties are remarkably similar.

- ✓ **ZINC** : High density in drug-like space.
- ✓ **ChEMBL** : Broad pharmacological coverage.
- ✓ **Conclusion**: The chosen properties (MW, logP, RotB) follow similar manifolds in both datasets, ensuring a fair comparison of generative capabilities.



# Constraint Paradigms

## Range-Based

Testing strict dual-bound control derived from data percentiles.

- ✓ **Loose:** 5th-95th percentile
- ✓ **Tight:** 25th-75th percentile
- ✓ **Ultra-tight:** 40th-60th percentile

## Gradual (Upper-Bound)

Matching instruction-following formats with increasing difficulty.

- ✓ **Loosen:**  $MW \leq 500$
- ✓ **Tight:**  $MW \leq 400, \log P \leq 4$
- ✓ **Ultra-tight:**  $MW \leq 350, \log P \leq 3.5, \text{RotB} \leq 8$

# Reward Design

We shift from binary filtering to continuous optimization, shaping the reward surface to guide particles toward valid regions.



## Symmetric (Range)

Centers the optimizer in the middle of the range for maximum robustness.



## Asymmetric (Bound)

Penalizes violations while rewarding a "Safety Margin" buffer zone.

# Initialization Prefixes

To prevent mode collapse and ensure structural diversity, we initialize SMC particles with 20 distinct chemical prefixes ranging from aromatic rings to aliphatic chains.

C1=CC=

Aromatic Core

CCN

Amino  
Aliphatic

COC

Polar Ether

CC(=O)

Carbonyl

C1CN

Heterocycle

CCS

Sulfur  
Motif

CCOCON

Extended Polar

CCC

Generic Aliphatic

C1CCCCC1

Aliphatic  
Ring

c1ccccc1

Benzene

n1ccccc1

Pyridine

N1CCCCC1

Piperidine

O1CCNCC1

Morpholin  
e

NC(=O)

Amid  
e

NS(=O)(=O)

Sulfonamide

NC(=O)N

Urea

C#N

Nitrile

Clc1ccccc1

Aryl Chloride

P(=O)(O)O

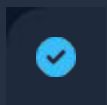
Phosphonat  
e

COC

Ether Builder



# Results



## 100%

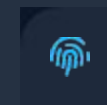
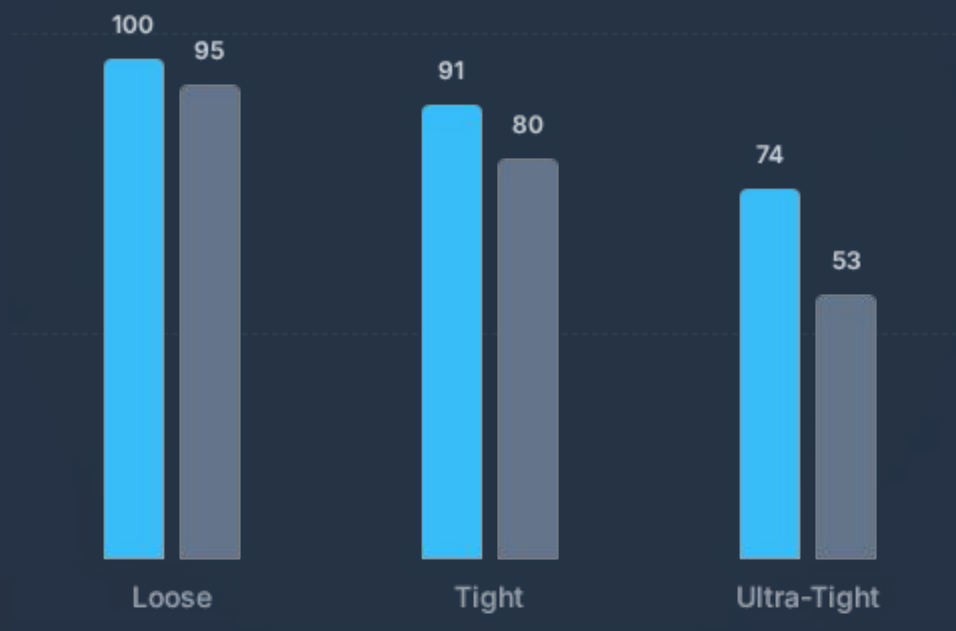
### Validity

Perfect structural validity achieved across all models and constraint settings.

### Adherence %

■ GPT2+SMC ■ SmileyLlama

Comparison under "Gradual" constraints.



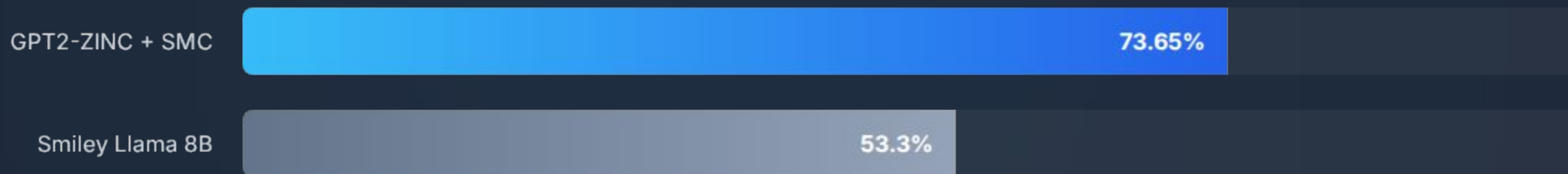
## 0.91

### Diversity

High chemical diversity maintained via distinct prefix initialization.

# Highlight: Ultra-Tight Constraints

Comparing the percentage of generated molecules that strictly adhere to the most difficult "Gradual" constraints.



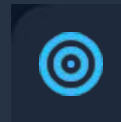
💡 **Insight:** Despite being ~100x smaller, the SMC-guided model outperforms the 8B parameter model by over 20 percentage points.

# Key Takeaways



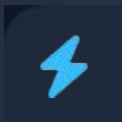
## Small Models Could Win

A specialized small model (87M) with SMC can rival or surpass a larger fine-tuned model (8B).



## SMC Effectiveness

Potential-based rewards steer generation towards desired properties without fine-tuning.



## Computational Efficiency (Not 100% confirmed)

This approach operates at a fraction of the cost of large-scale inference.



## Current Limitations

Extremely narrow, dual-bound ranges remain a difficult open problem.

# Future Work



## Non-Linear Restrictions

Refining SMC potentials to handle more complex chemical constraints beyond simple ranges.



## Multi-Property Balancing

Developing sophisticated weighting mechanisms to balance conflicting objectives.



## Core-Driven Synthesis

Exploring scaffold-based generation where the model builds around a fixed core.