

# Safe LLM regularization

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## Stages of LLM training

- Pretrain
- Alignment
  - Supervised Fine-tuning
  - Preference Optimization

## LLM Adaptation approaches

- Domain specific pretrain
  - Fine-tuning
  - PEFT methods
- Domain specific alignment

## Fine-tuning of LLM kills alignment

- Keeping LLMs Aligned After Fine-tuning: The Crucial Role of Prompt Templates
- Fine-tuning Aligned Language Models Compromises Safety, Even When Users Do Not Intend To!



## Red teaming

**Red teaming** - research field that studies approaches for creating adversarial attacks on LLM to compromise its safety (**red prompts**)

## Re teaming datasets

- ALERT
- Thoroughly Engineered Toxicity

# Approach for toxicity robustness evaluation

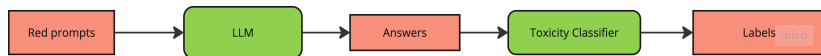


Figure 1: Toxicity robustness evaluation framework





## KL Divergence

In general, similarity between probability distributions

$$D_{KL}(P||Q) = \int_{-\infty}^{\infty} p(x) \log \frac{p(x)}{q(x)} dx \quad (1)$$

# Loss with KL divergence regularization

## Loss with KL regularization

Designing loss with KL regularization.

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$$\mathcal{L}_{\text{causal}} = - \sum_{t=1}^T \log P(x_t | x_{<t}; \theta) + \text{Reg}(\theta, \theta^*) \quad (2)$$

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$$\text{Reg}(\theta, \theta^*) = -\gamma^t D_{KL}(\theta || \theta^*) \quad (3)$$

$\theta$  – *current model parameters*

$\theta^*$  – *base model parameters*

$\gamma$  – *decay rate*,  $t$  – *epoch*



## General task formulation

To adapt LLM for scientific domain using fine-tuning while keeping it safe with no alignment data given apriori

## Data

- Fine-tuning data - Arxiv collection and Elibrary
- Red prompts - ALERT and TET datasets

## Setup

- Hardware - GPU NVIDIA A100
- 10 epochs of training

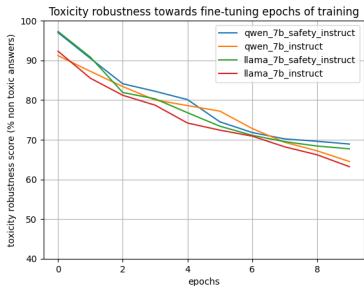
## Evaluation

- Domain-specific evaluation - SciAssess
- Toxicity Evaluation - as described previously using ALERT and Red Teaming datasets, Llama7b-instruct as toxicity evaluator

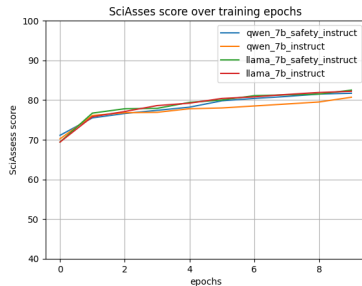




# Classic fine-tuning



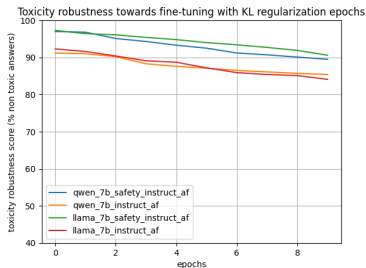
(a) Results during tuning on toxicity robustness



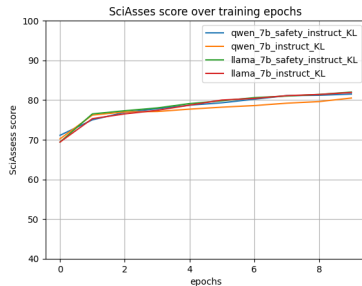
(b) Results during tuning on SciAsses

Figure 2: Classic Tuning

# Fine-tuning with Regularization



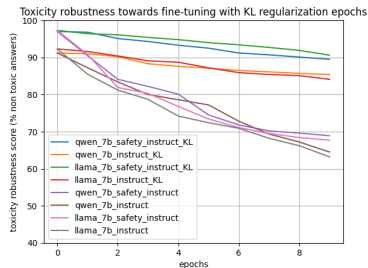
(a) Results during tuning on toxicity robustness



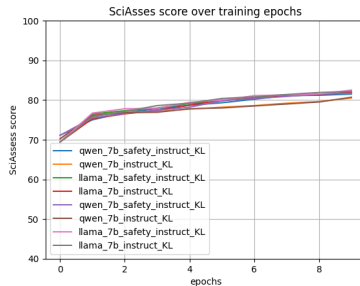
(b) Results during tuning on SciAsses

Figure 3: Tuning with regularization

# Comparison



(a) Results during tuning on toxicity robustness



(b) Results during tuning on SciAsses

Figure 4: Comparison



## Advantages

- More robust towards toxicity
- Comparable results with classic fine-tuning

## Disadvantages

- Memory usage
- More resources