

ALTeGraD Data Challenge



# Molecule Retrieval with Natural Language Queries

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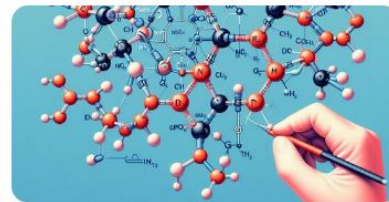
## Context (1/2)

Challenge Goal:

Develop a model capable of accurately retrieving specific molecules through text queries

### **ALTeGraD-2023 Data Challenge**

Molecule Retrieval with Natural Language Queries



## Context (2/2)

### Evaluation metric: **Mean Reciprocal Rank**

Given the rank at which the first relevant item is retrieved, the reciprocal of this rank is calculated, and the average is taken across all queries.

$$\text{MRR} = \frac{1}{|Q|} \sum_{i=1}^{|Q|} \frac{1}{\text{rank}_i}$$

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# **I. Related Work**

# DistilBERT

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- Distilled version of **BERT** model
- **Efficient** and **compact** transformer-based language representation model
- Reduced **computational complexity** and **memory** requirements

This model was used in the baseline architecture upon which we built our final model.

# SciBERT

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- **Specialized** pre-trained language model based on the BERT architecture
- Trained on a vast and diverse corpus of scientific publications, totaling 1.14 million papers
- Training data covers domains like biomedicine (82%) and computer science (12%), making SciBERT adept at processing **molecular properties**

# Graph Convolutional Networks

The diagram illustrates the equation for the  $k$ -th layer embedding of a node  $v$  in a Graph Convolutional Network (GCN). The equation is:

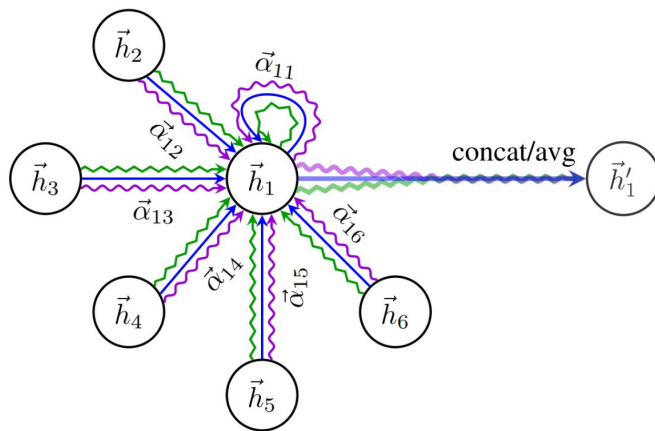
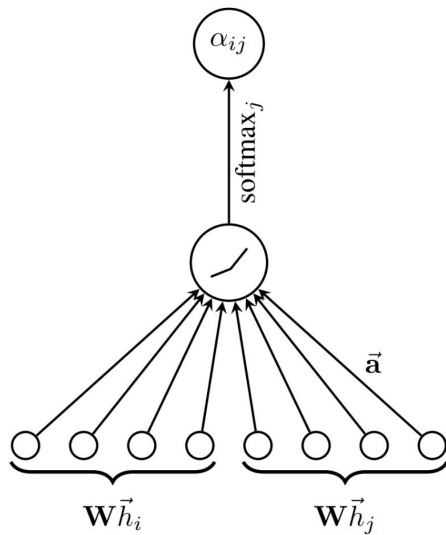
$$\mathbf{h}_v^k = \sigma \left( \mathbf{W}_k \sum_{u \in N(v)} \frac{\mathbf{h}_u^{k-1}}{|N(v)|} + \mathbf{B}_k \mathbf{h}_v^{k-1} \right), \quad \forall k > 0$$

Annotations and components:

- Initial "layer 0" embeddings are equal to node features:** Points to the initial embedding  $\mathbf{h}_v^0 = \mathbf{x}_v$  (purple box).
- previous layer embedding of  $v$ :** Points to the term  $\mathbf{h}_v^{k-1}$  (orange box).
- average of neighbor's previous layer embeddings:** Points to the summation term  $\sum_{u \in N(v)} \frac{\mathbf{h}_u^{k-1}}{|N(v)|}$  (red box).
- non-linearity (e.g., ReLU or tanh):** Points to the activation function  $\sigma$  (blue box).
- $k$ th layer embedding of  $v$ :** Points to the output embedding  $\mathbf{h}_v^k$  (green box).



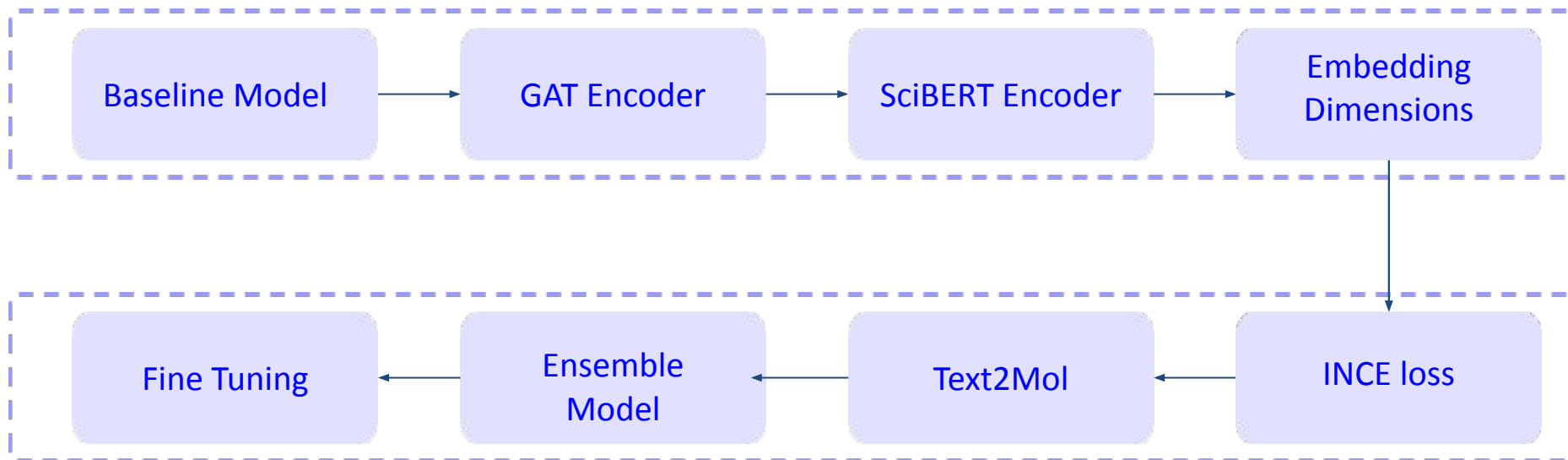
# Graph Attention Networks



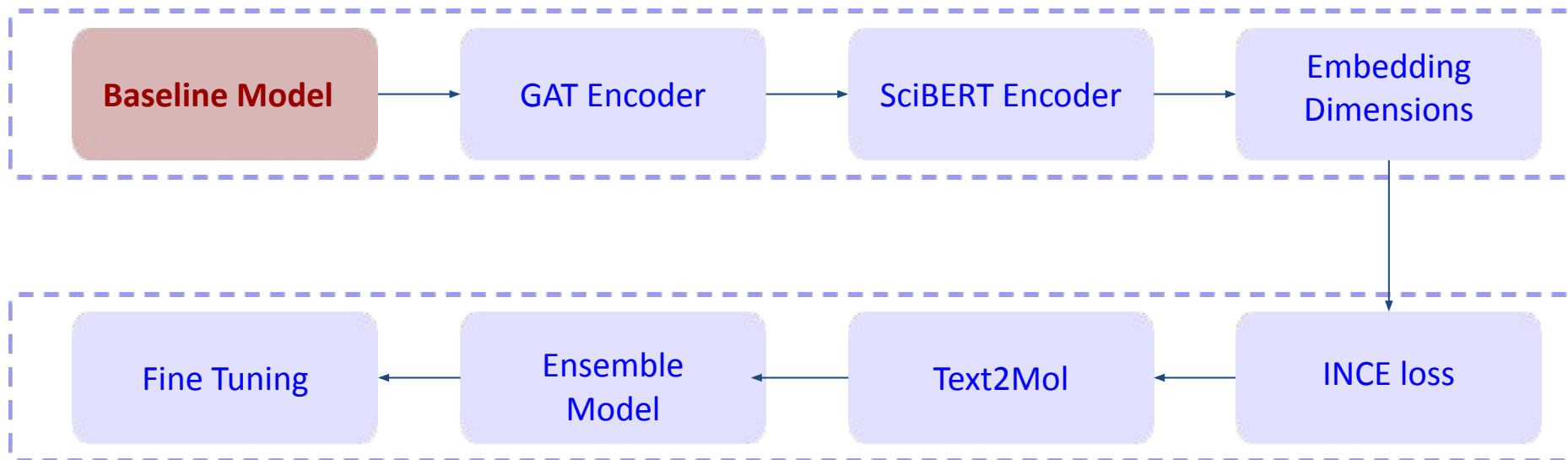
## **II. Models**

# Models

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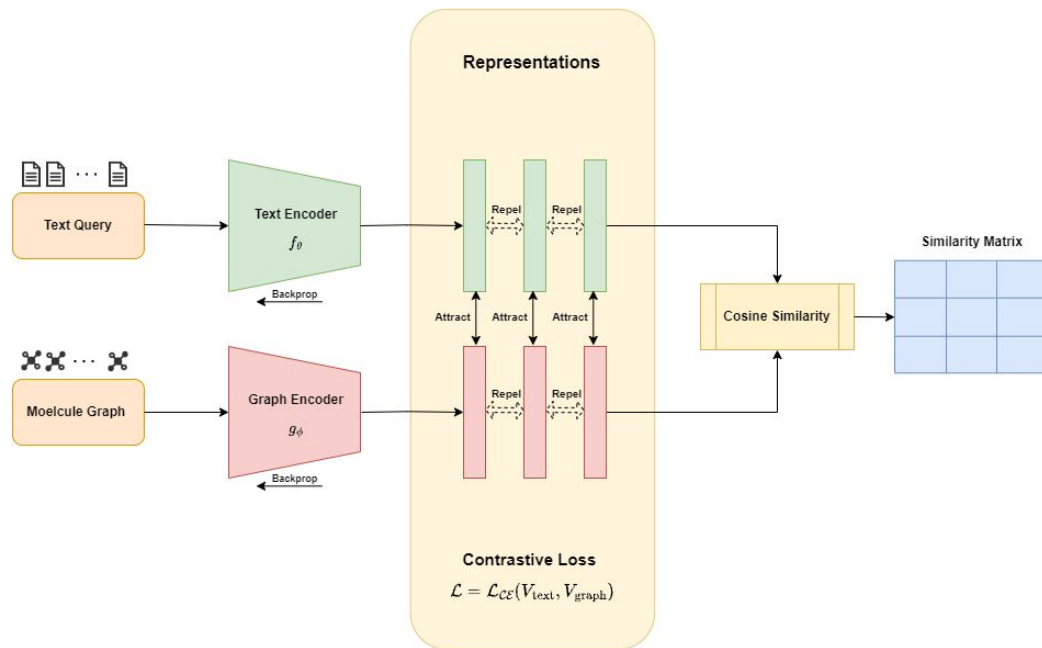


# Models

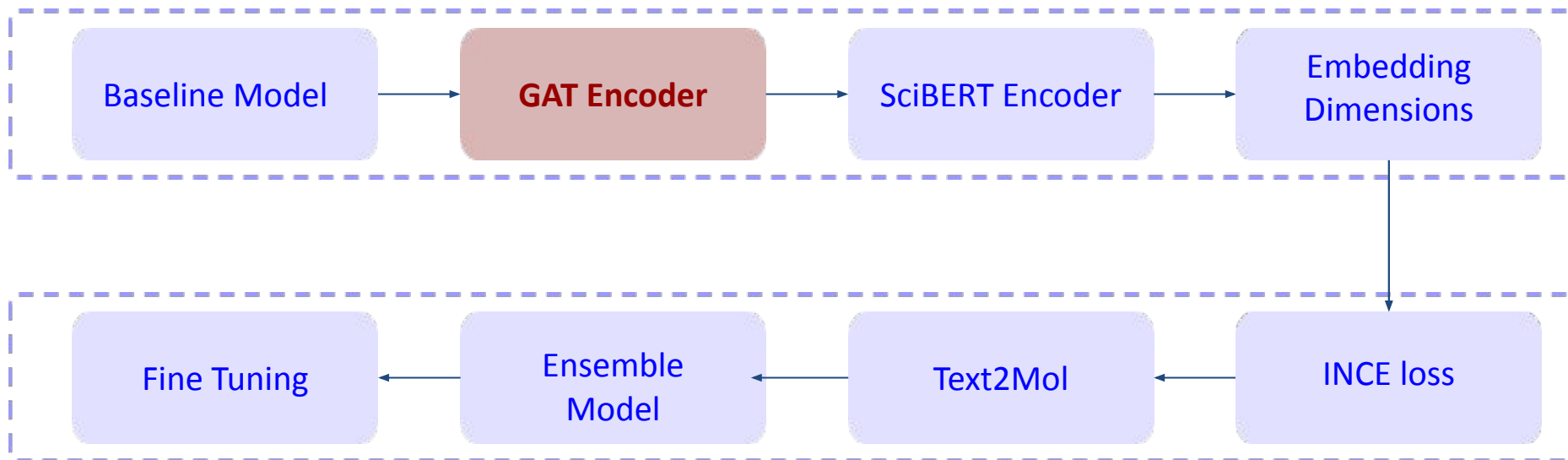


# Baseline model

- Employed **DistilBERT** as the text encoder and **GCN** as the graph encoder
- Trained for **5 epochs**
- Minimized the **contrastive loss** between the representations
- Reaches an MRR score of **0.348**



# Models

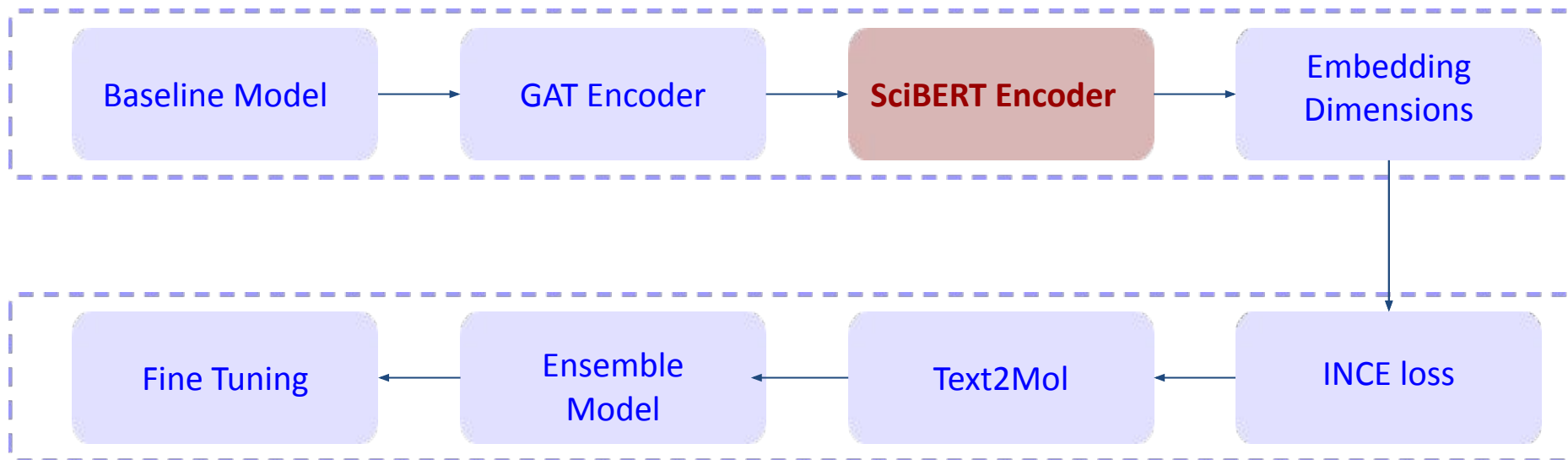


## GAT Encoder

- The substitution of the GCN encoder with a GAT encoder.
- Motivated by the capacity of GATs in **capturing complex relationships** within graph-structured data by applying **self-attention** mechanism on the nodes.
- Chosen GATv2 encoder which used a **dynamic graph attention mechanism** and is more expressive than GAT.
- Trained for **50 epochs** to minimize the **contrastive loss** between the representations.
- Improved the baseline model and achieved an MRR score of **0.5086**.

Model	MRR
Baseline	0.348
GAT	0.5086

# Models



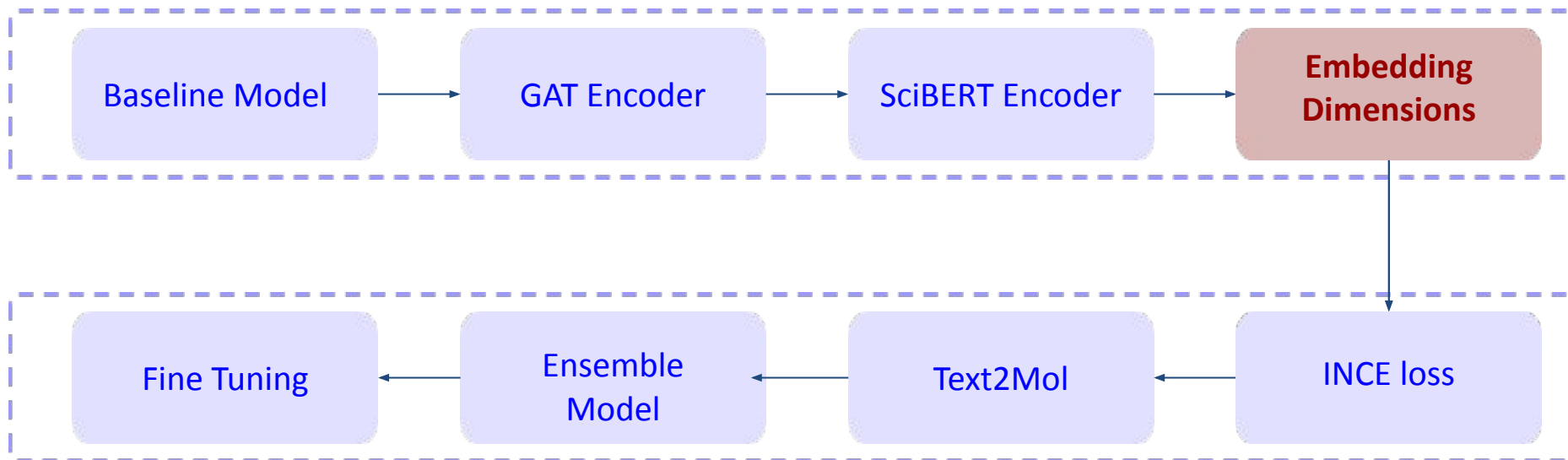


# SciBERT Encoder

- Replaced the DistilBERT text encoder with an **uncased sci vocab SciBERT** encoder.
- Pertained on a data containing 82% **biomedicine** papers -> well-suited for processing texts describing molecular properties
- Trained for **100 epochs** to minimize the contrastive loss between the representations.
- Reaches an MRR score of **0.72**

Model	MRR
Baseline	0.348
GAT	0.5086
GAT and SciBERT	0.72

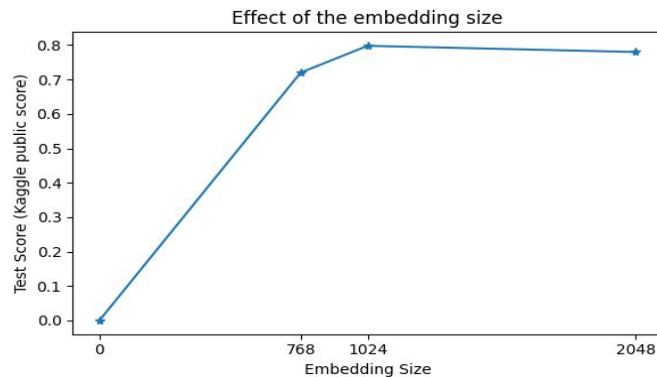
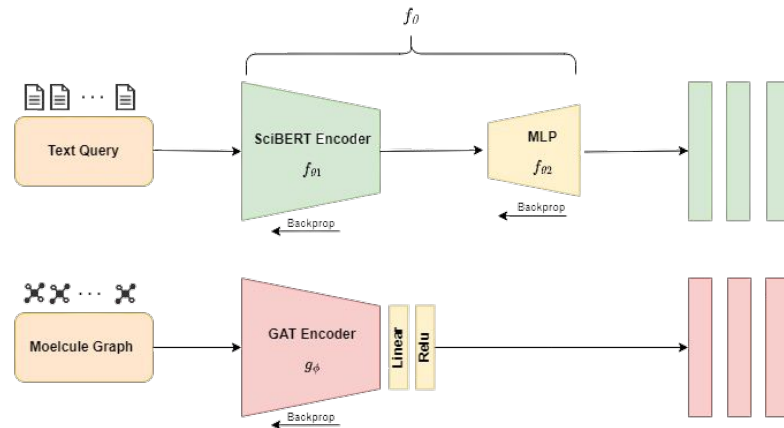
# Models



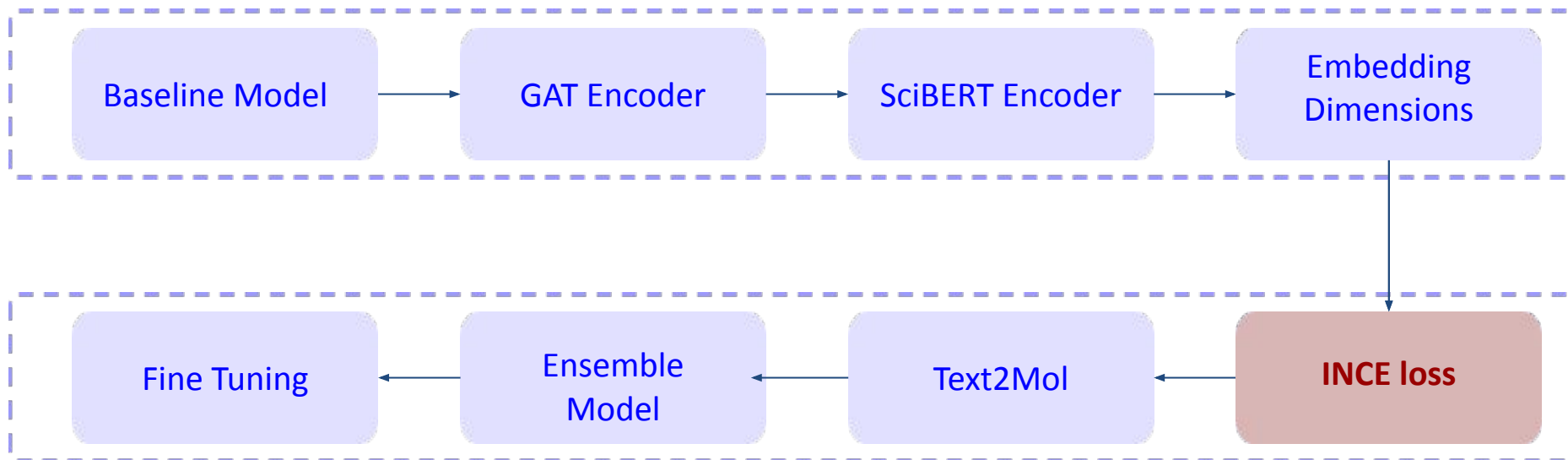
# Higher Dimensional Embeddings

- Used a **richer** embedding space of dimension **1024**
- Appended a Multi-Layer Perceptron (**MLP**) at the end of each encoder
- Achieved an MRR score of **0.798**

Model	MRR
Baseline	0.348
GAT	0.5086
GAT and SciBERT	0.72
High-Dim	0.798



# Models



# Information-Noise-Contrastive Estimation

- Optimizes the negative log probability of classifying the **positive** sample correctly.

$$\mathcal{L}_{\text{INCE}} = -\mathbb{E} \left[ \log \frac{f(\mathbf{x}, \mathbf{c})}{\sum_{\mathbf{x}' \in X} f(\mathbf{x}', \mathbf{c})} \right]$$

- At each step we **incorporated** the embeddings of the **previous batches as negative samples**.
- Memory length =  $2 \times \text{batch\_size}$  to avoid converging to an **uninformative local minimum**
- Enhanced the MRR score to **0.87**.

Model	MRR
Baseline	0.348
High-Dim	0.798
INCE Loss	0.87

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## Algorithm 2 Custom Contrastive Loss

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**Input:**  $v_1$ : Text embeddings,  $v_2$ : Graph embeddings,  $m$ : Memory size

**Initialize:** memory buffers  $m_{\text{text}}$  and  $m_{\text{graph}}$  to None; loss functions: cross-entropy  $ce$  and InfoNCE  $incc$

**Function** RESETMEMORY():

Reset  $m_{\text{text}}$  and  $m_{\text{graph}}$  to None

**Function** FORWARD( $v_1, v_2$ ):

Calculate InfoNCE loss on  $v_1$  and  $v_2$  with  $m_{\text{graph}}$  as negatives

Calculate InfoNCE loss on  $v_2$  and  $v_1$  with  $m_{\text{text}}$  as negatives

Combine losses:

$result = \text{INFONCE}(v_1, v_2, m_{\text{graph}}) + \text{INFONCE}(v_2, v_1, m_{\text{text}})$

Update memory:

If  $m > 0$ :

If  $m_{\text{text}}$  is None:

Set  $m_{\text{text}}$  to  $v_1$  and  $m_{\text{graph}}$  to  $v_2$

Else:

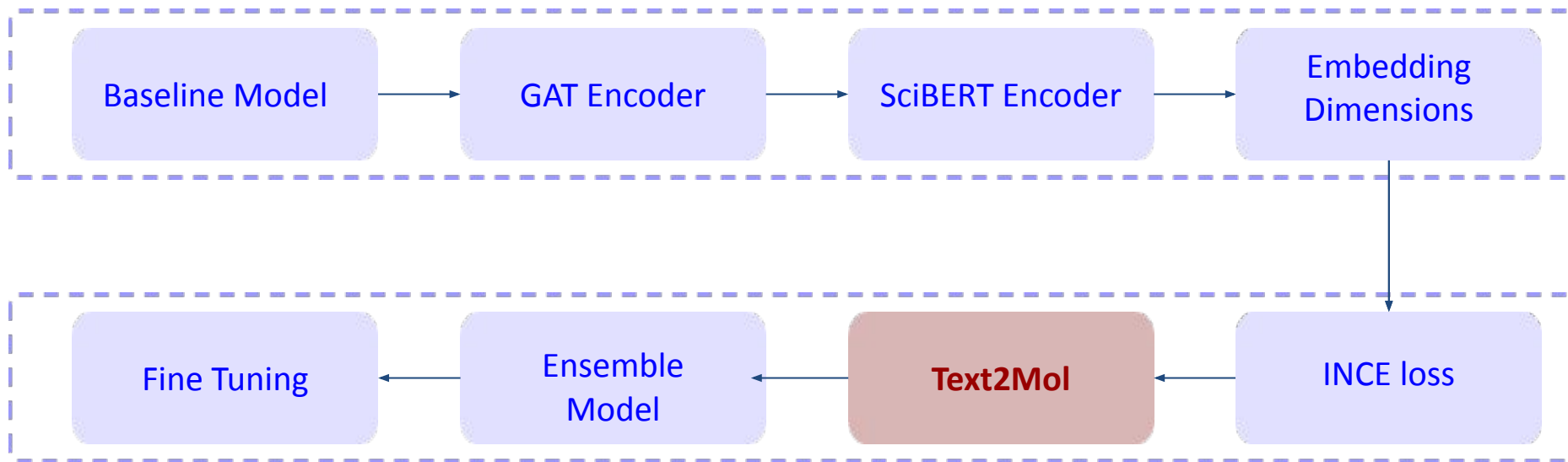
Append  $v_1$  to  $m_{\text{text}}$  and  $v_2$  to  $m_{\text{graph}}$

Keep only last  $m$  elements in  $m_{\text{text}}$  and  $m_{\text{graph}}$  if exceeded

**return**  $result$

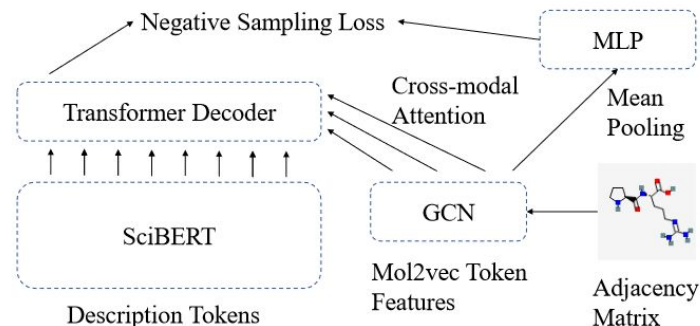
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# Models



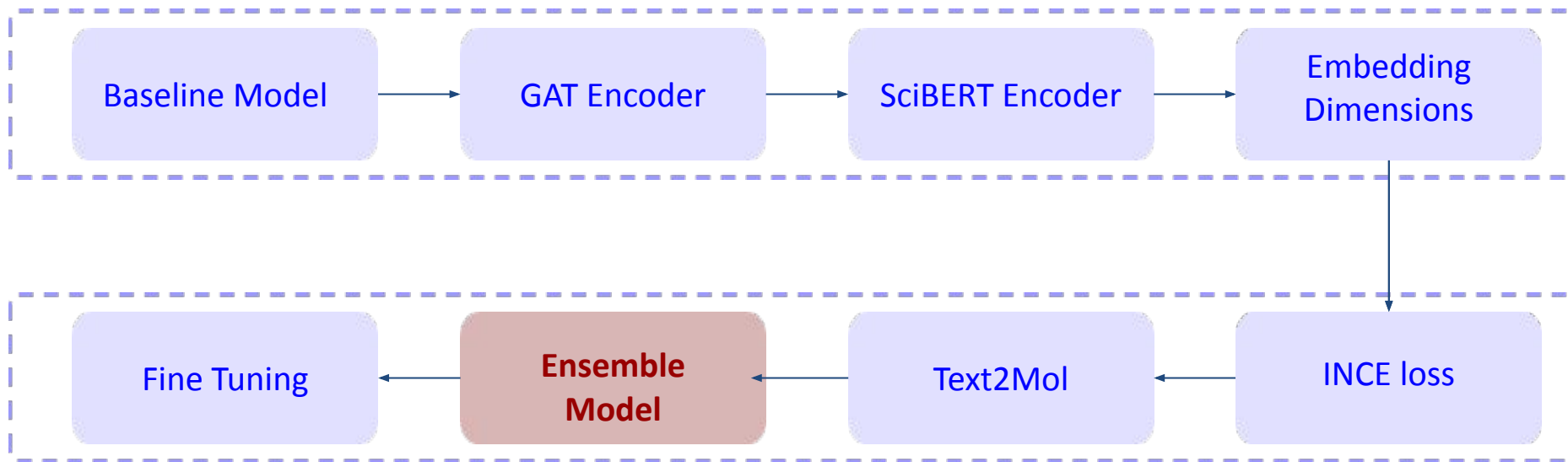
# Text2Mol

- Implemented **Text2Mol** framework using the architecture from Edwards et al. 2021.
- Uses **cross-modal attention mechanism** connecting text and graph embeddings.
- Potential **information leakage** from the graph embeddings which was mitigated through **negative sampling**.
- The model was interesting to explore however we didn't use it for the final results due to:
  - weak performance of **MRR score 0.03**
  - the need to compute the text embedding associated with each individual graph.



The Text2Mol architecture for the cross-modal attention extension and association rules

# Models



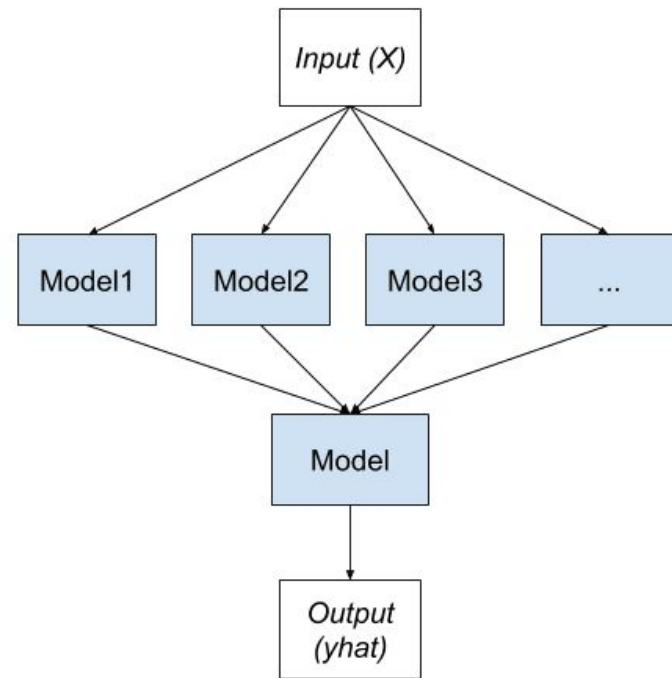


# Ensemble model

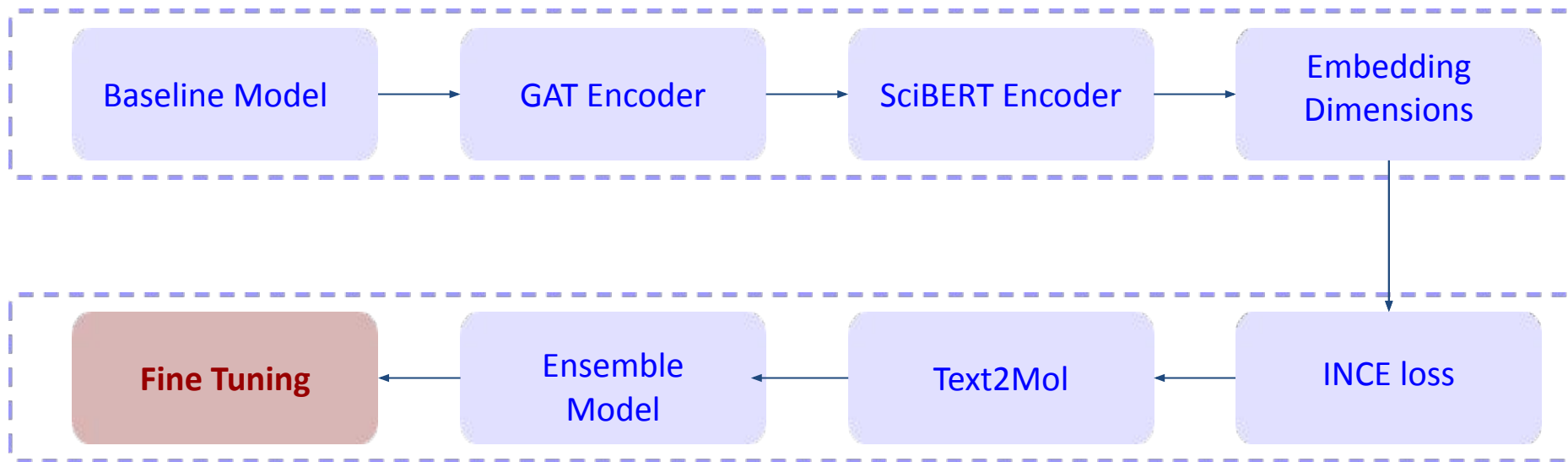
- Noticed that our model weren't overfitting so there is a possibility they can act as "strong learners".
- Implemented naive stacking methods such as the arithmetic and geometric mean of the outputs of different models.
- Improved the result greatly breaking the 0.9 barrier for the **MRR** score 0.92

Model	MRR
Baseline	0.348
High-Dim	0.798
INCE Loss	0.87
Ensemble Model	0.9216 <sub>25</sub>

Stacking Ensemble



# Models



# Fine tuning

For the final training run we:

1. Started from the **best model** obtained so far.
2. **Merged** validation set with the training set to train on the entire dataset.
3. Linear extension of the **memory depth** at intervals of 10 fine-tuning periods, ranging from **2 to 10** times the batch size.
4. Monitoring the performance of the model through **sampled 5000 observations**.
5. Selecting the model with the highest **MRR** score on the sampled validation set : 0.9223

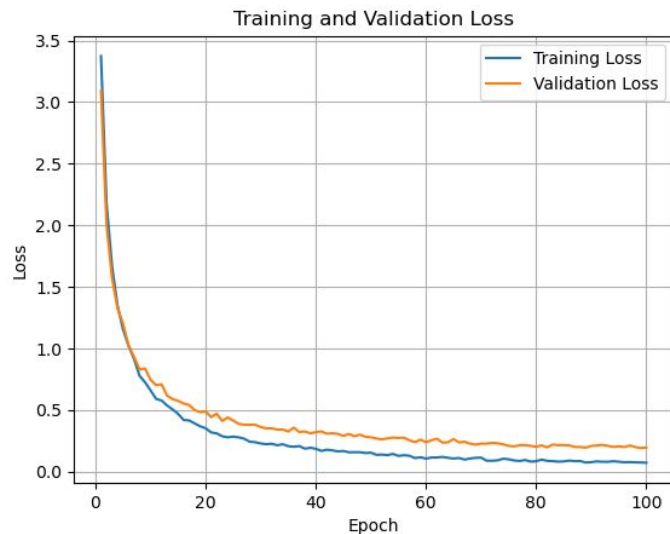
# **III. Results & Discussions**

### III. Results & Discussions:

Model	MRR
Baseline	0.348
GAT	0.5086
GAT and SciBERT	0.72
High-Dim	0.798
INCE Loss	0.87
Text2Mol	0.03
Ensemble Model	0.9216
<b>Fine Tuned</b>	<b>0.9223</b>

### III. Results & Discussions:

- The model didn't overfit :
  1. The model isn't expressive enough
  2. The data is uniformly distributed across training, validation and test



### III. Results & Discussions:

- Training took approximately **12 hours**.
- The training time was pretty much the same for all models.
- The **bottleneck** was the data loader

## **IV. Conclusion & Perspective**



## IV. Conclusion & Perspective:

- The best model was the fine-tuned Scibert + GAT with the modified InfoNCE.
- Exploring more pooling methods for Text and Graphs.
- Results seem to be too promising which might necessitate validating them over other datasets.

## References (1/2)

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