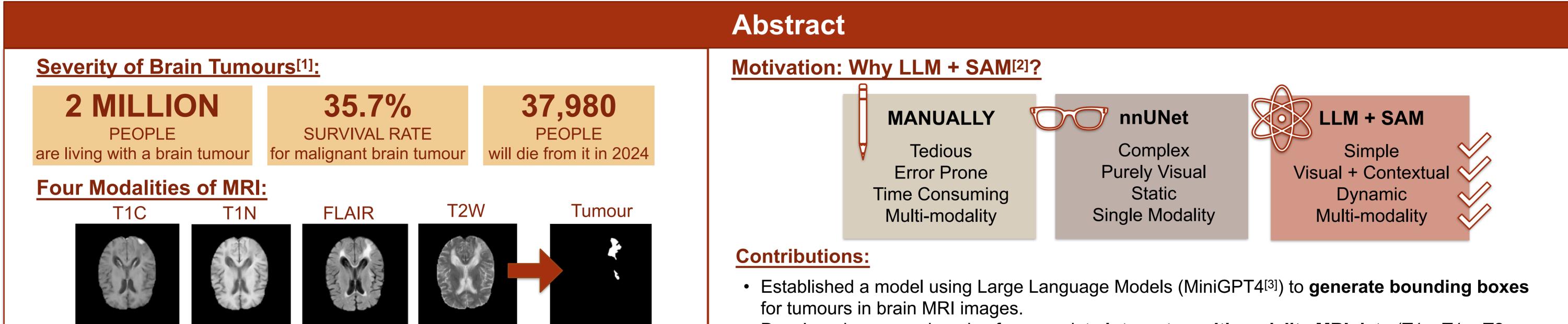


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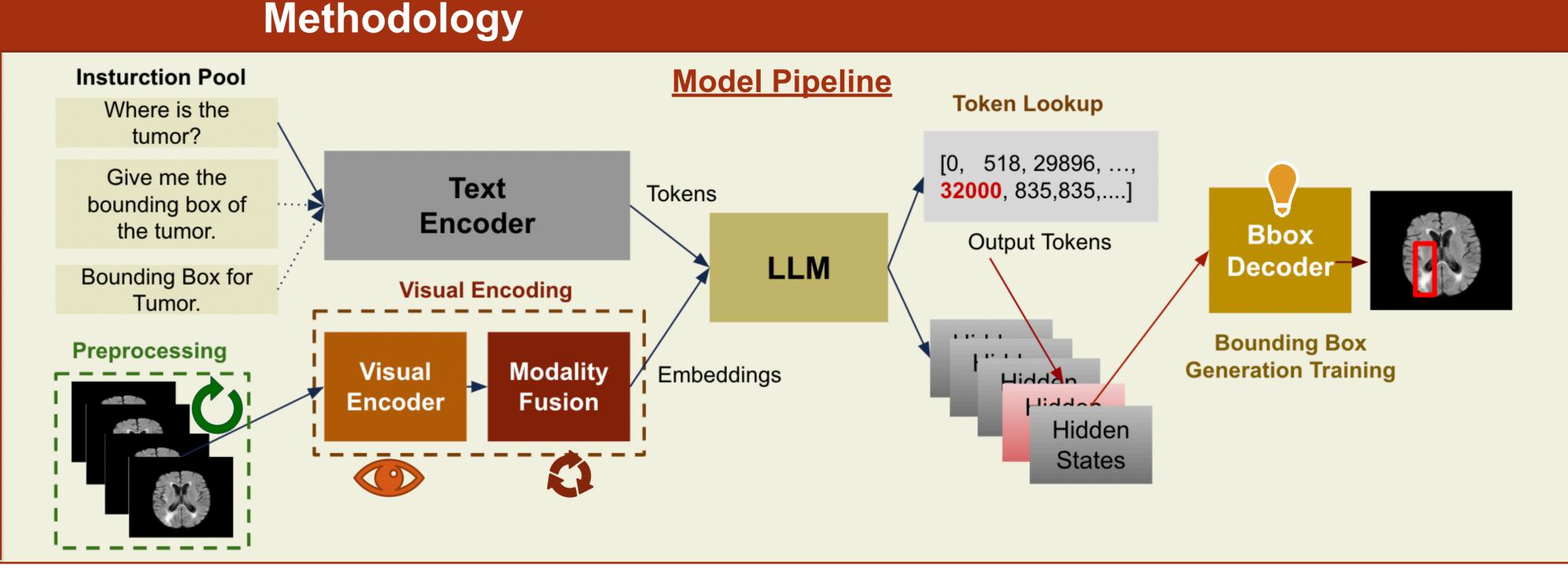
**Objective:** develop a robust, efficient, multimodal framework for accurate brain tumour segmentation by integrating various MRI modalities using LLM + SAM.

• Developed a comprehensive framework to integrate multi-modality MRI data (T1c, T1n, T2, FLAIR) for improved tumour segmentation.

• Established a truly simple, ready to use model for users with zero expertise on machine learning.

## Challenges

- Need to implement a mechanism to train LLM to generate bounding box for images with given instructions.
- LLM is trained on natural images, which struggles with understanding medical images.
- Need to develop an innovative approach to let the LLM to generate a **synchronised bounding box** for four different input pictures.
- Need to increase the correctness (measured by Intersection Over Union) of the LLM predicted bounding boxes.



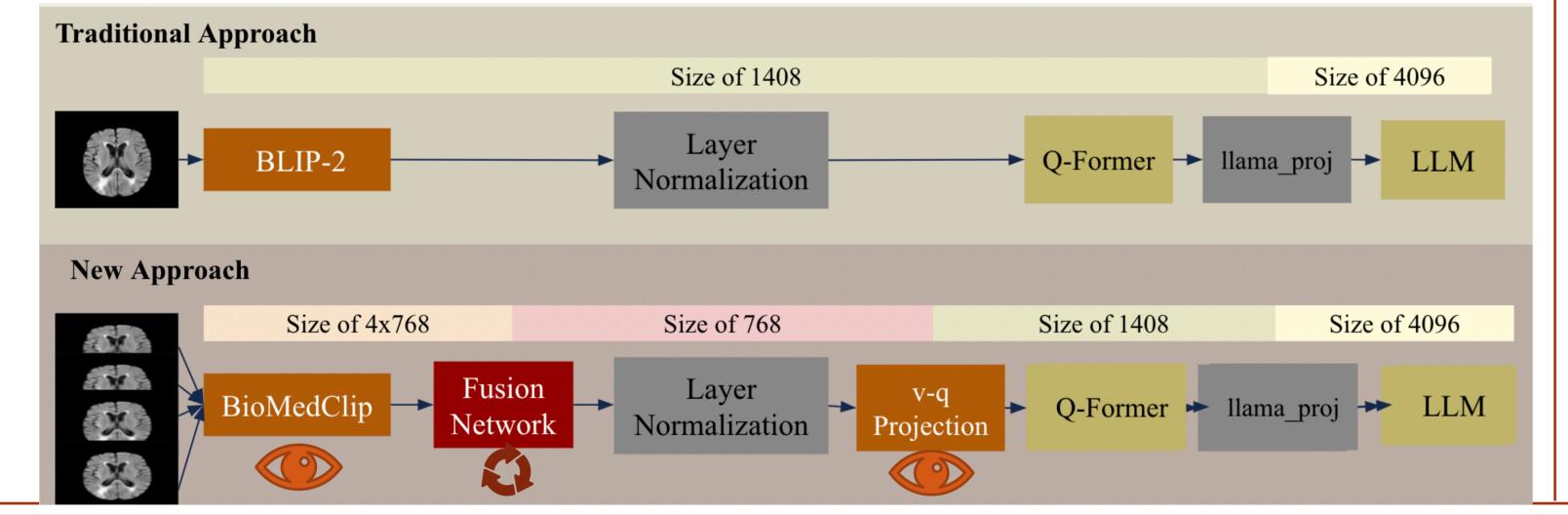
<u>**Preprocessing:**</u> Except for resizing and normalization transformation, add random rotation (p = 0.4) and flipping (p = 0.2) during prepossessing to enhance model understanding.

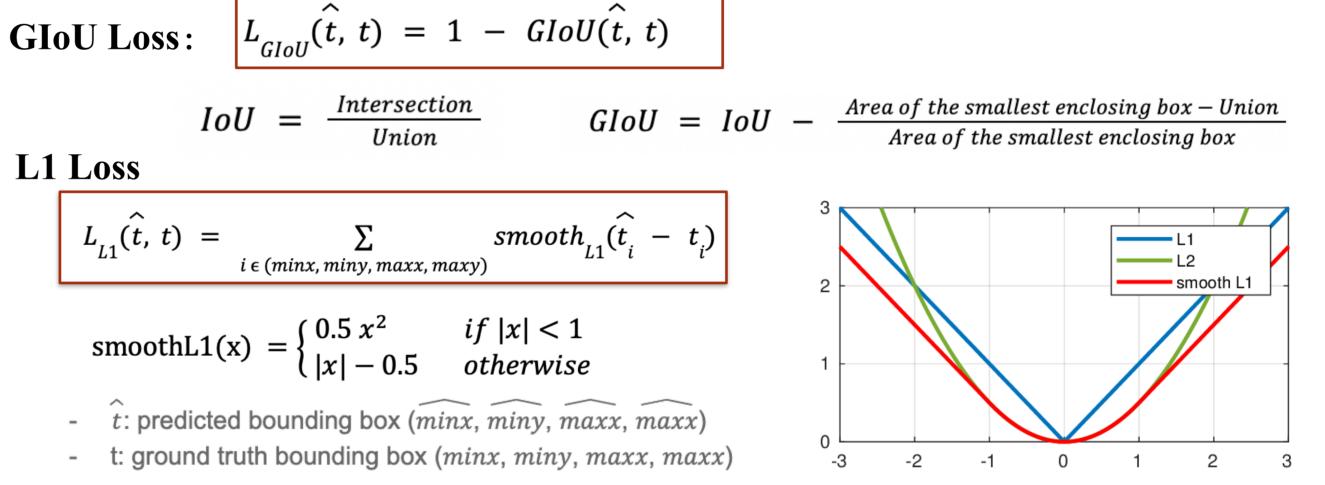
### Visual Encoding

**Specialised Visual Encoder:** Utilized BioMedClip<sup>[4]</sup> to enable LLM to understand medical images. **Modality Fusion**: Established a fusion network that enables LLM to accept multiple images as input.

### **Bounding Box Generation Training**

Used the following loss functions to evaluate the model's performance, guiding it towards the desired direction of achieving higher accuracy in bounding box prediction.





# Results

|             | E      | <u> BLIP-2 Vs. BioMedC</u> | lip <u>Stand</u> | Standard vs. Additional Preprocessing                   |       | Single Modality Vs. Multi-Modality |       |
|-------------|--------|----------------------------|------------------|---|-------|------------------------------------|-------|
| Tumour Type | loU    | Relative Increase          | IoU              | Relative Increase                                       | IoU   | Relative Increase                  | loU   |
| GLI         | 0.208  | + 145.2%                   | 0.582            | + 3.4%  | 0.602 | + 8.4%                             | 0.653 |
| MEN         | 0.232  | + 157.8%                   | 0.598            | + 1.2%  | 0.605 | + 10.7%                            | 0.670 |
| MET         | 0.219  | + 184.0%                   | 0.622            | + 1.5%  | 0.631 | + 3.2%                             | 0.651 |
|             | BLIP-2 |                            | Added BioMedCl   | MedClip Added Additional Preprocessing Added Multi-moda |       |                                    |       |

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Single modality model with standard preprocessing and BLIP-2 as visual encoder



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Single modality model with standard preprocessing and BioMedClip as visual encoder

Single modality model with additional preprocessing and BioMedClip as visual encoder



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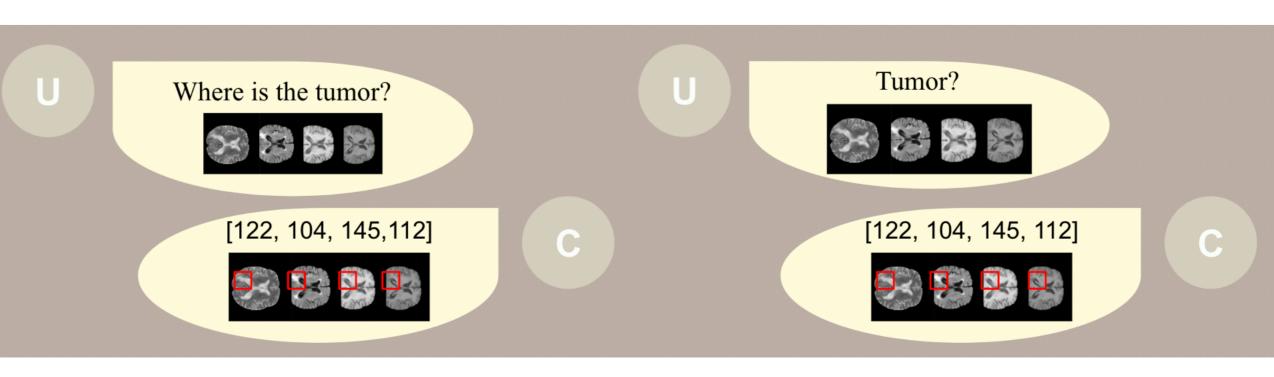
Multi-modality model with BioMedClip as visual encoder, and additional preprocessing

### **Conclusion & Evaluation Overall IoU Overall IoU** + 212.3% 0.211 0.659

Established a LLM for bounding box prediction in Brain Tumour MRI images.

By adding the specialized visual encoder, additional preprocessing, and incorporating MRI data from all modalities, we achieved a 212.3% increase in overall IoU.

### Example Use Case



- Efficient - No expertise on machine learning needed
- Accept various kind of prompt
- Easy to use

### **Future Work**

- 1. Connect to **SAM** for more detailed mask generation.
- 2. Gather **patient data** on contextual information (gender, age) and apply it to the training process.
- 3. Explore potential adaptations to the fusion network.

Reference: [1] Brain Tumor Facts - National Brain Tumor Society. (2024, February 20). National Brain Tumor Society. https://braintumor.org/brain-tumors/about-brain-tumors/brain-tumor-facts/ [2] Kirillov, A., Mintun, E., Ravi, N., Mao, H., Rolland, A., Gustafson, L., ... & Dollár, P. (2023). Segment Anything. arXiv preprint arXiv:2304.02643. Retrieved from https://arxiv.org/abs/2304.02643 [3] Zheng, K., He, X., & Wang, X. E. (2023). MiniGPT-5: Interleaved Vision-and-Language Generation via Generative Vokens. arXiv preprint arXiv:2310.02239.

[4] Zhang, S., et.all . BiomedCLIP: a multimodal biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical foundation model pretrained from fifteen million scientific image-text pairs. arXiv.org. https://arxiv.org/abs/2303.00915biomedical founda 2303.00915