Summarizing Biomedical Texts with Encoder Decoder Models

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Problem



- Doctors need to summarize complex findings efficiently
- Make quick decisions on urgent patients cases.

Background

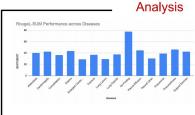
Radiologists write reports to report observation on chest X-rays and synthesize those findings into impressions to hand off to other specialists. We introduce a Biomedical-BERT2BERT model to help radiologists generate concise impressions from reports, with state of the art performance.

Methods

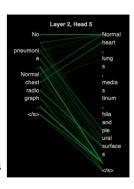
Model centric methods

Data centric methods:

Text-to-Text: T5 (small / large)Encoder-Decoder: BERT-to-BERTLinear Attention: BigBird

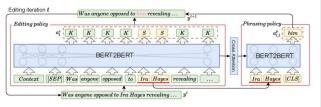


- Model Robustness: All models learned to detect No Findings very well
- Learned disease level fairly well
- Learned several radiology facts such as pneumonia corresponds with "pleural surfaces"



Results

Model	Epochs	time/ epoch	ROUGE-1	ROUGE-2	ROUGE-L	baseline
Finetuned T5-Small	3	1.5 h	48.71	37.98	47.42	•
DistillBART	3	2.5 h	30.81	19.51	26.88	0
Finetuned T5-base-long	12	0.63 h	55.68	45.52	54.74	©
Finetuned BERT2BERT	6	0.65 h	59.61	48.99	58.75	0
Finetuned PubMed BigBird	7	1.55 h	57.83	47.12	56.66	2



Conclusions

- New state-of-art ROUGE-L score of 58.75
- Previous knowledge on similar tasks and vocabulary showed advantageous
- Very efficient model, compared to 2nd best performing model
- **Data-centric approach** enabled new opportunities for improvement in performance
 - Data augmentation techniques with input fields shuffling: "paraphrasing"
- Analysis on diseases shows that may be still room for improvement if class imbalance is addressed

References

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Acknowledgments

Kathy Yu (project Mentor)

