

EXPANDED RESEARCH ON
ATTENTION-GUIDED DISCRIMINATIVE REGION LOCALIZATION
AND LABEL DISTRIBUTION LEARNING FOR

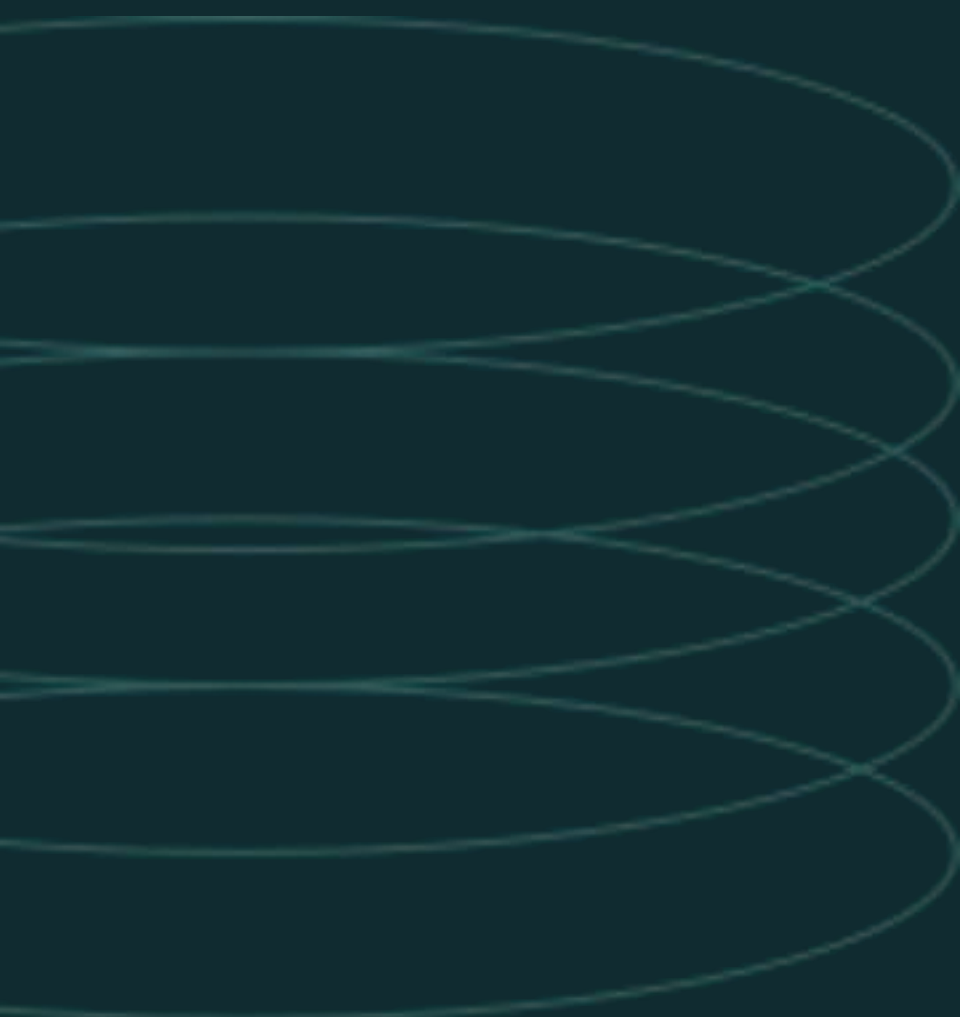
Bone Age Assessment

C&S Bio 185

By Trevor Brokowski, Marlene Lin, Nikhil Patel, Anish Patel, Shiqin Tan

Background

Bone age assessment



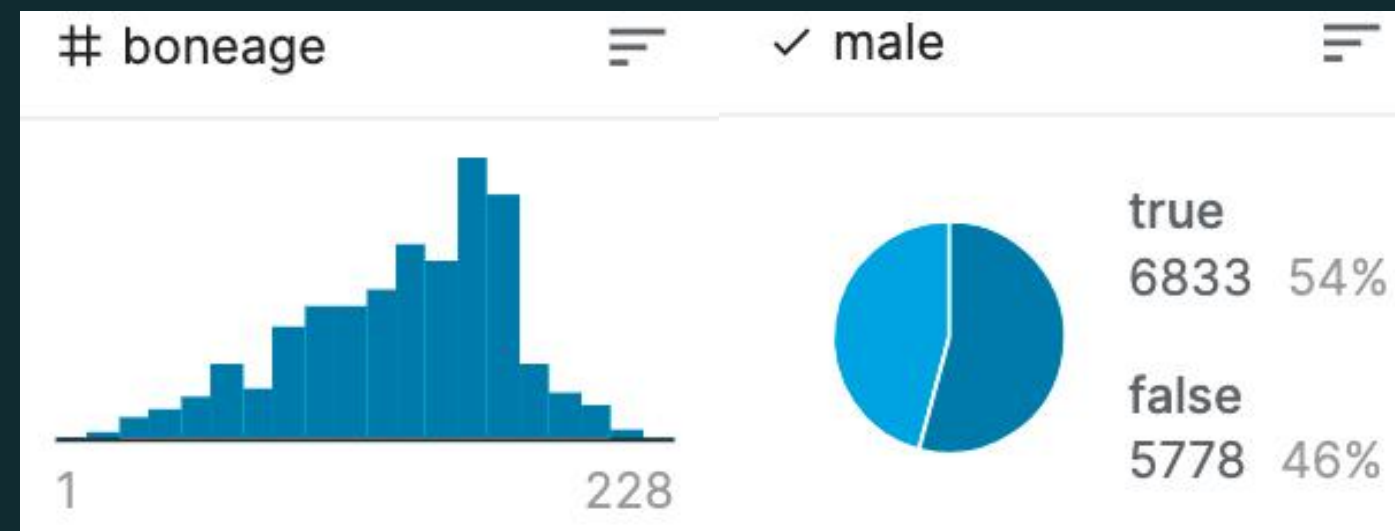
**Attention-Guided Discriminative Region
Localization and Label Distribution
Learning for Bone Age Assessment**

Chao Chen, Zhihong Chen, Xinyu Jin, Lanjuan Li, William Speier, Corey W. Arnold

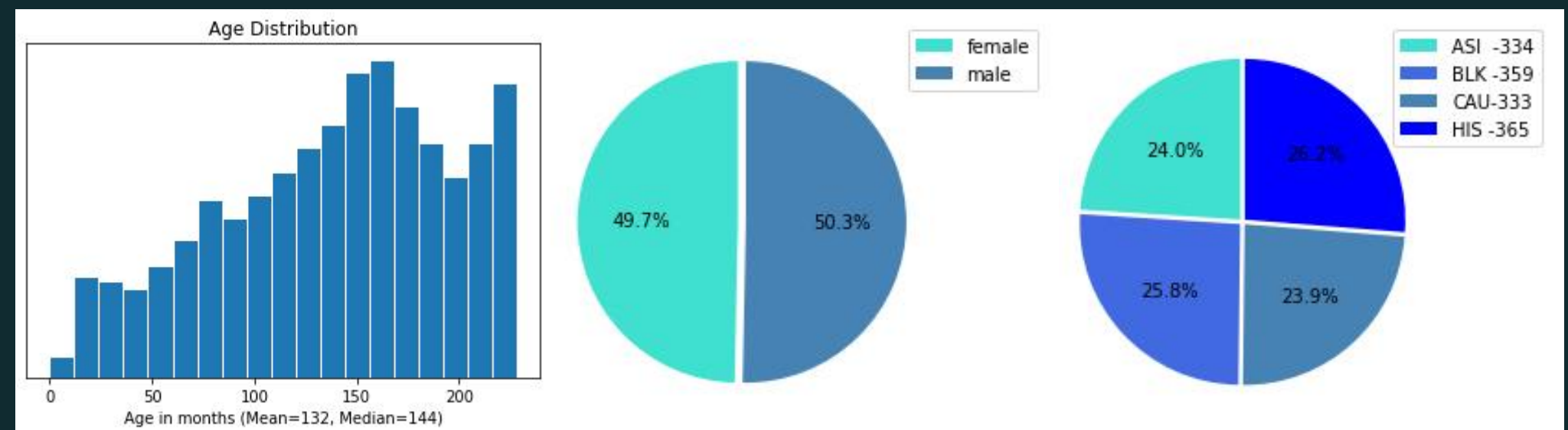
Expanding Research

Transferability to Digital Hand Atlas of USC

RSNA metrics



USC metrics



Our goal

Understand

- ability to transfer to more datasets
- concatenate ethnicity to see if overfit



Selected Paper

- Model: **Xception**
- Accuracy. **More efficient with comparable precision to published BAA methods**

| Methods | Image Size | Extra Labels | Data Augment | Model Ensembling | MAE | | | | |
|---------------------|------------|-----------------|--------------|------------------|------|-------|--------|--------|---------|
| [6] | 750 × 750 | mask & keypoint | Yes | 18 model results | 6.4 | | | | |
| [3] | 224×224 | No | Yes | No | 9.5 | | | | |
| [16] | 512 × 512 | No | Yes | No | 5.99 | | | | |
| | 512 × 512 | No | Yes | 50 model results | 4.26 | | | | |
| [5] | 500×500 | Bbox & keypoint | Yes | No | 4.14 | | | | |
| Ours | O | H | R1 | R2 | H+R1 | R1+R2 | O+H+R1 | H+R1+E | H+R1+R2 |
| l_1 | 7.3 | 6.4 | 6.1 | 7.0 | 5.4 | 5.6 | 5.4 | 4.7 | 4.8 |
| $l_{MAE} + l_{reg}$ | 6.2 | 5.6 | 5.3 | 6.2 | 4.8 | 5.1 | 4.7 | 4.3 | 4.3 |

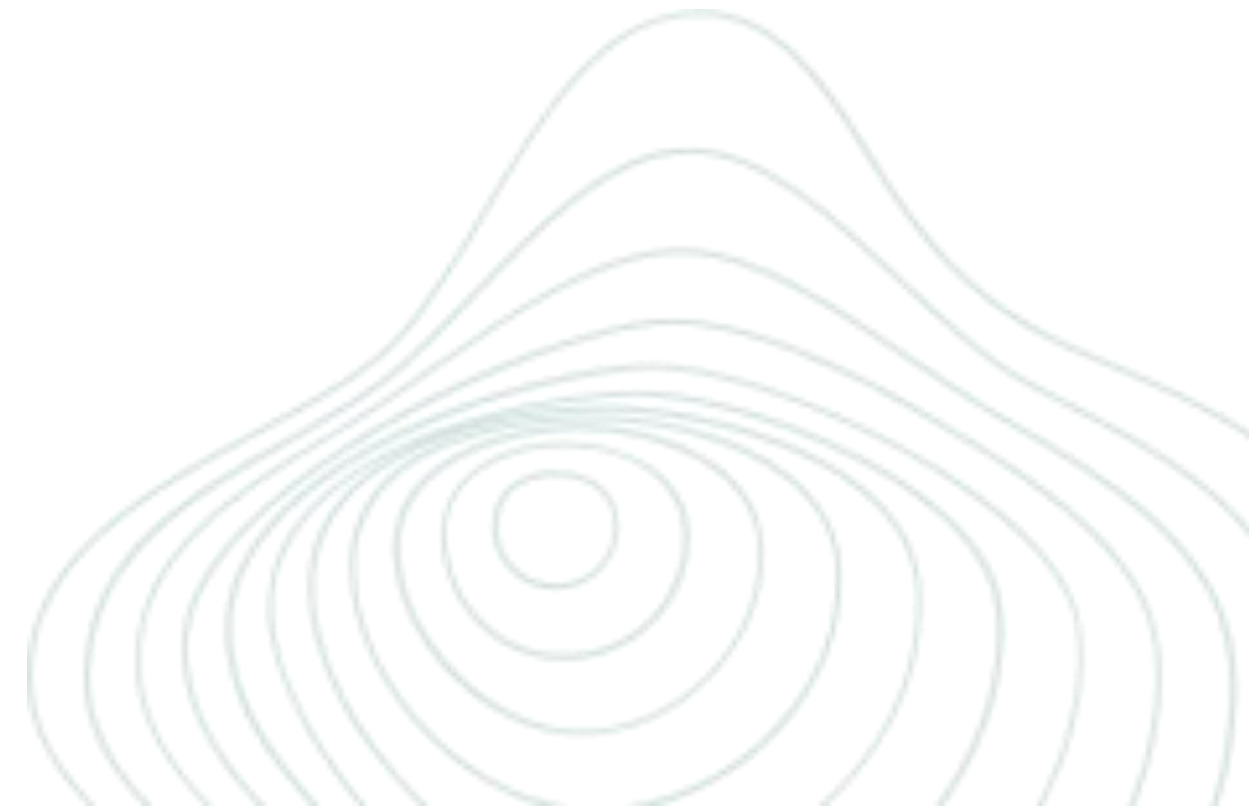
| | Vgg19 | InceptionV3 | ResNet50 | Xception | Xception (with ethnicity) |
|--------------------|-------|-------------|----------|----------|---------------------------|
| Test Results (MAE) | 9.3 | 9.2 | 9.3 | 8.8 | 9.4 |

Our Result

- Model: **Xception**
- Accuracy: **Comparable to conventional BAA methods**
- Training time: **6+ hours**

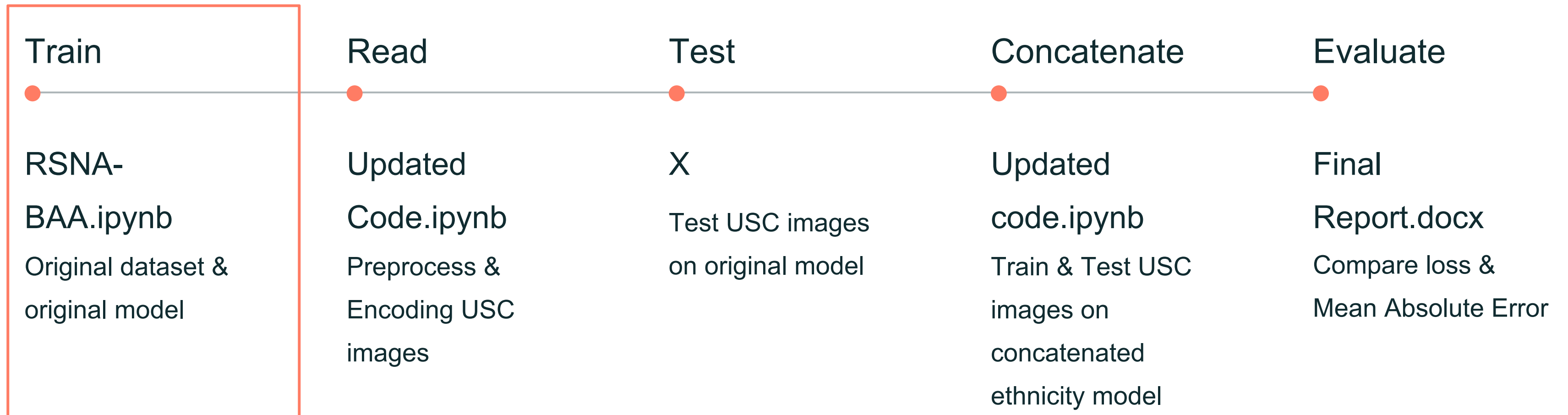
So...How do we get these results?

>>> Ideal Pipeline of Our Implementation



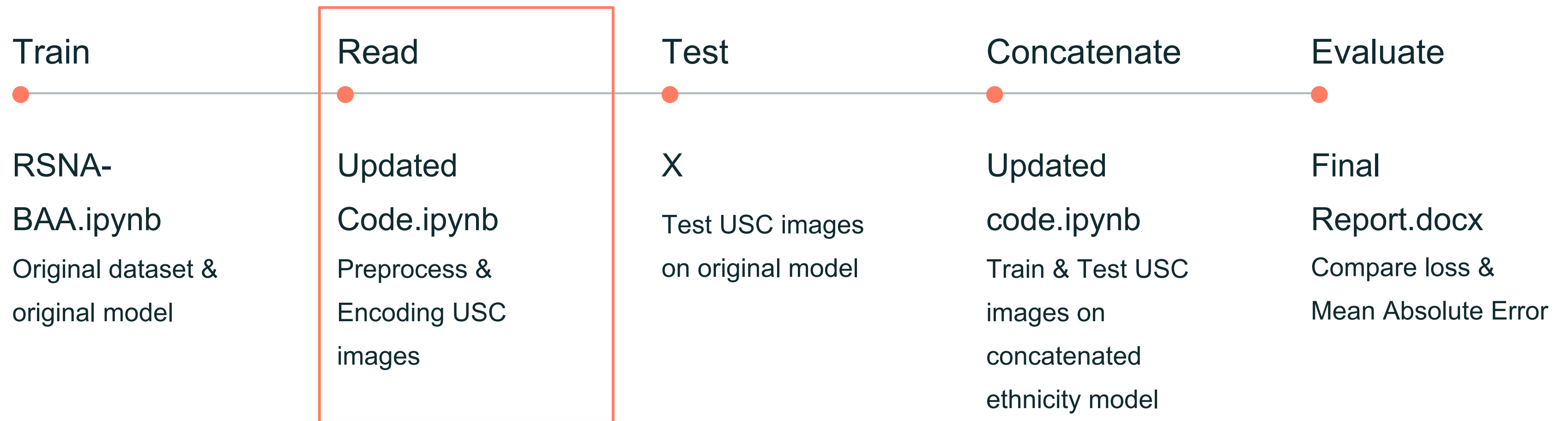
Our Methods

Train, Test, Concatenate



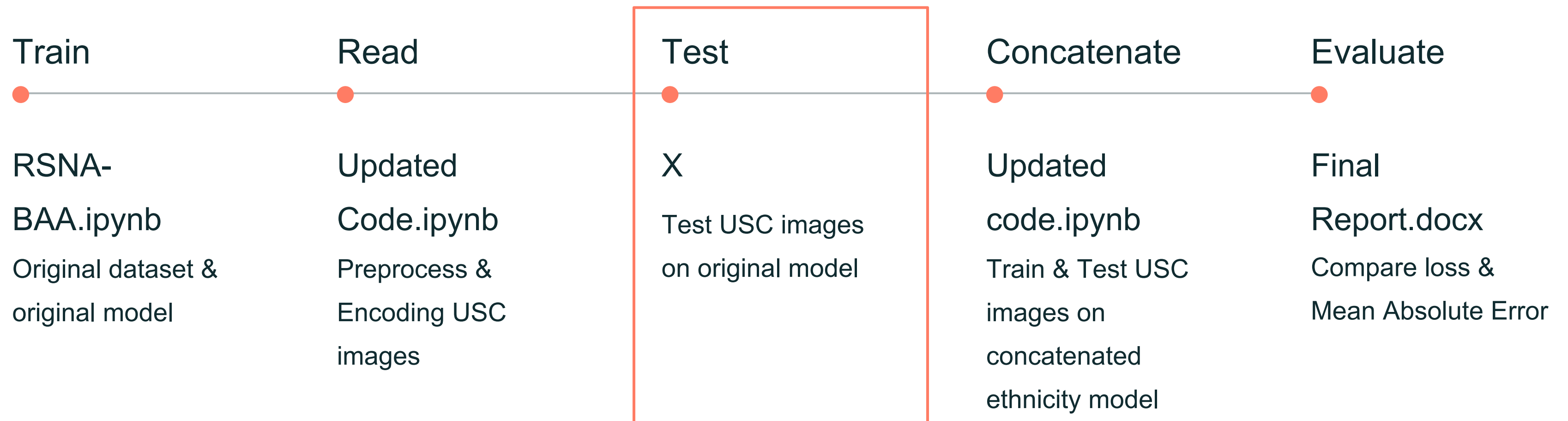
Our Methods

Train, Test, Concatenate



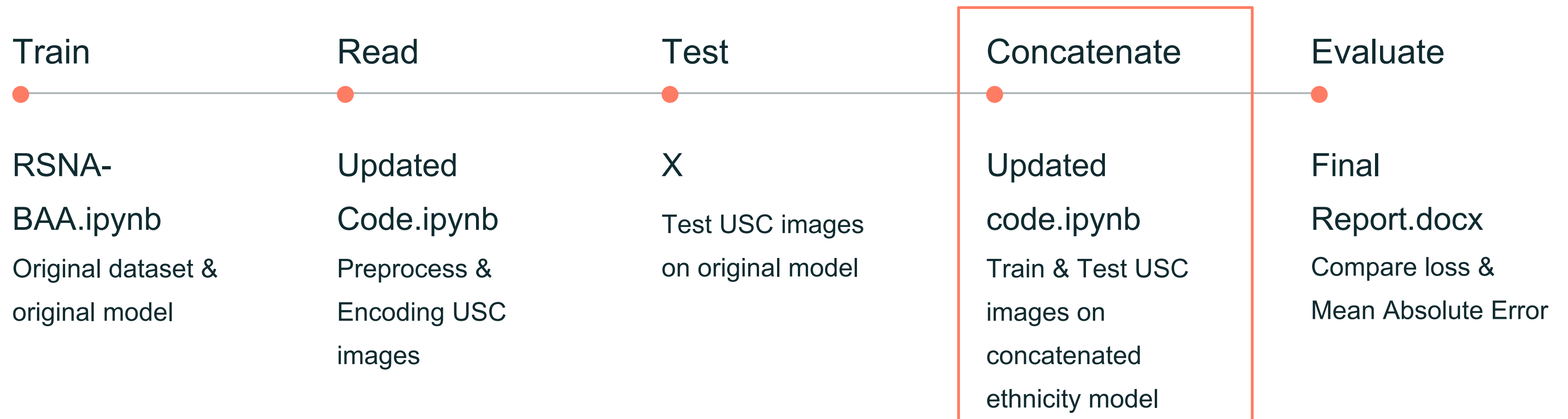
Our Methods

Train, Test, Concatenate



Our Methods

Train, Test, Concatenate



Our Methods

Train, Test, Concatenate

Train



RSNA-

BAA.ipynb

Original dataset &
original model

Read



Updated

Code.ipynb

Preprocess &
Encoding USC
images

Test



X

Test USC images
on original model

Concatenate



Updated

code.ipynb

Train & Test USC
images on
concatenated
ethnicity model

Evaluate



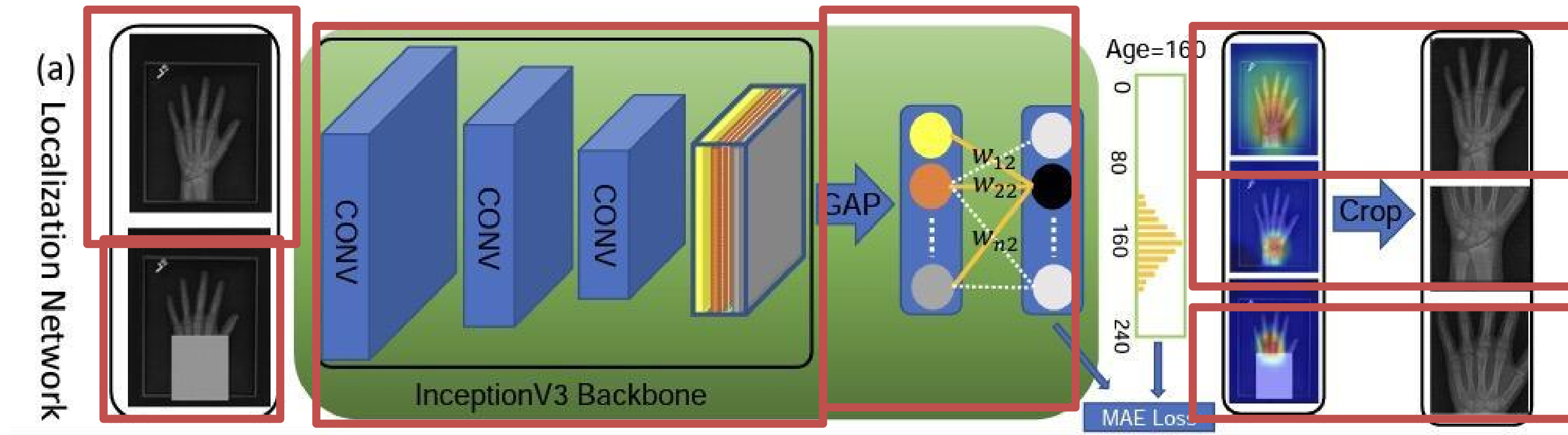
Final

Report.docx

Compare loss &
Mean Absolute Error

Our Methods

Model: Attention-guided localization

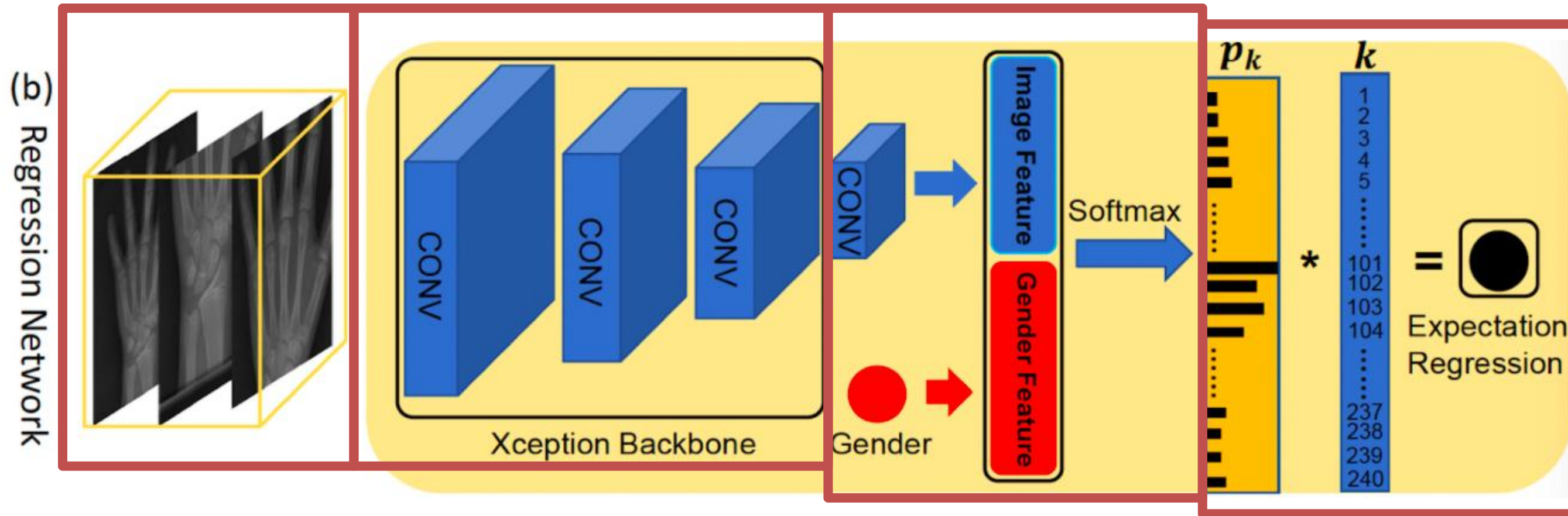


Original x-ray images resized -> localized hand region/localized Region of Interests 1 (carp bones)

Original image with R1 erased -> localized Region of Interests 2

Our Methods

Model: Age regression



Our Methods

Reimplementation (RSNA dataset)

1. Data loading:

RSNA: 12, 611 labeled img -> out of ram -> progressive loading -> out of time -> rand. 1390

2. Localized hand & Regions of Interests 1 & Masked out R1:

learning rate = 0.0003 for the first 60 epochs, learning rate = 0.0001 for the next 30 epochs with a batch size of 32

3. Test results (MAE/Month):

`val_MAE = 8.2505`

learning rate = 0.0003 for the first 40 epochs, learning rate = 0.0001 for the next 20 epochs with a batch size of 16



Our Methods

Reading USC images &
Concatenate ethnicity info

Step 1: Process and extract information from DICOM (.dcm) files

- Extract the patient id, gender, ethnicity, age, and, initially, the image*, from the dicom file and the path name using python string manipulation and the os library.
- Example Path: “/Digital Hand Atlas/DICOM/ASIF/ASIF01/5024.dcm”
- * Due to dimensionality and color channel issues with the Xception base model, the image had to be loaded in using opencv instead of pydicom. (3 color channels vs 1)

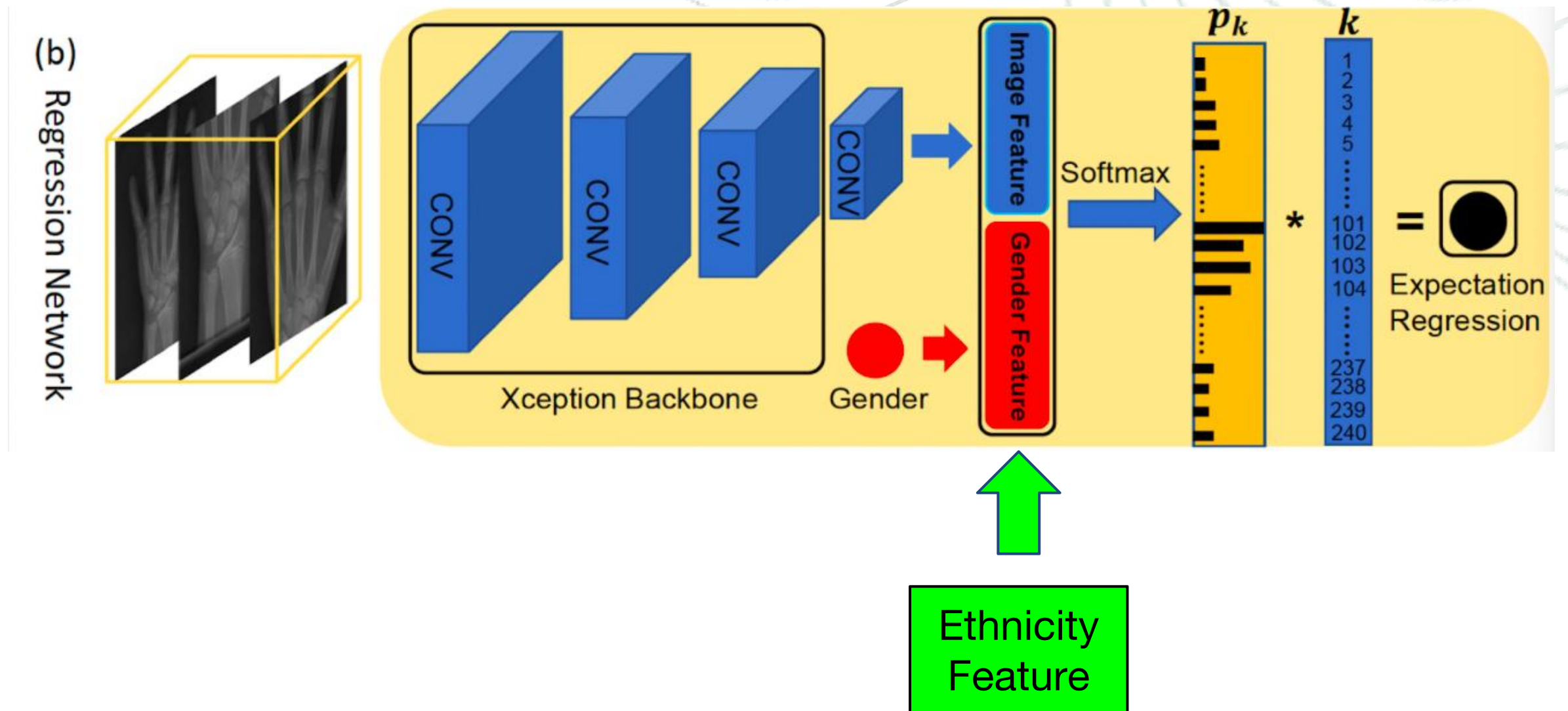
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(0002, 0001) File Meta Information Version      OB: b'\x00\x01'
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(0002, 0003) Media Storage SOP Instance UID     UI: 1.3.6.1.4.1.
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Little Endian
(0002, 0012) Implementation Class UID          UI: 1.3.6.1.4.1.
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.80,10.40,46.99,2.25,2.00'
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(0010, 0020) Patient ID                        LO: '84115024'
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(0018, 1014) Time of Secondary Capture         TM: '151029'
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ration'
(0018, 1018) Secondary Capture Device Manufactur LO: 'IPI LAB'
(0020, 000d) Study Instance UID                 UI: 1.2.392.2005
```


Step 2: Concatenate Ethnicity Information.

- Created a dictionary that performed one-hot-encoding on the variables.

- ASI $\rightarrow [1,0,0,0]$
- CAU $\rightarrow [0,1,0,0]$
- HIS $\rightarrow [0,0,1,0]$
- BLK $\rightarrow [0,0,0,1]$

- Concatenate the list of encoded ethnicities after the convolutions as new input layer for the activation function and the prediction output.





Discussion

Broader Implications

- Concerns: Algorithmic Bias Amplification
- AI Models have proven to:
 - Amplify Healthcare Disparities
 - Exhibit worse results for minority populations
 - Replicate biases in training data



Discussion

Known Biological Differences: Ethnicity

Recent Studies have show that:

1. Significant Differences in Bone Density between ethnicities

1. Rate of bone maturation differs across ethnic lines



Discussion

Inclusion of Ethnicity Data in Models

How to combat algorithmic bias?

No consensus: Proposed including ethnicity data by:

- Entirely separate models
- Including as model input
- Including as model output
- Adjusting input data for known biological differences across ethnic lines



Discussion: Preliminary Results

No significant improvement with Ethnicity.

Important Considerations

- Limited Computing Power
 - Made batching and data set size adjustments to improve training time
 - No time to tune hyperparameters
- Tuning hyperparameters and using larger datasets may improve accuracy beyond baseline
- More research is needed to assess this method of ethnicity inclusion
- Could be a beneficial method to improve algorithmic bias in radiological applications



Thank you
for Listening!

Code Availabilities

- The source code of selected paper is available at their Github page: <https://github.com/chenchao666/Bone-Age-Assessment>
- Our commented code is available at our Google Drive page: <https://drive.google.com/drive/folders/1QGTIIN27TLcWAEKVbjfl8-M8-E6HzAO8?usp=sharing>
- The RSNA dataset is available at Kaggle: <https://www.kaggle.com/kmader/rsna-bone-age>
- The Digital Hand Atlas of University of Southern California is available at their lab page: ipilab.usc.edu/research/baaweb/