YOLO-SG: An Efficient Framework for Scene Graph Generation

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Abstract

Scene graph generation (SGG) is the task of detecting object pairs and their relations in a visual medium, widely used for captioning, generation, and visual question answering. 2D scene graph generation is a subtask that focuses on generating a 2D graph given an image. While the development of models capable of performing 2D SGG has improved in both accuracy and speed, the computational complexity of the problem and the inherently long-tailed distribution of large, available datasets have led to generation speed and accuracy less than ideal for real-time use. Mainstream approaches focus on two-stage generation, where object detection is performed first, followed by a series of comparisons for relation inference. However, these have an inherent drawback where the computational complexity of detecting nobjects and their relationships is n^2 . More recent models have utilized encoderdecoder structures to reduce generation into a 1-stage problem. Unfortunately, the computation required for these architectures is still too high. Additionally, the model bias caused by long-tailed data distributions remains a key problem in both approaches. In this work, we propose YOLO-SG, a novel SGG framework capable of operating in real-time by decoupling object detection and relation detection and by performing relationship inference with multiple detection models in parallel. Our proposal seeks to both alleviate the effects of the long-tailed distribution problem and perform high-speed inference. Preliminary experiments on the Visual Genome 1.2 dataset demonstrate that YOLO-SG can achieve competitive performance with state-of-the-art models while maintaining high inference speed.

1 Introduction

In recent decades, object detection problems have become an increasingly popular subject in research literature[32] as the rise of deep learning[13] has led to many breakthroughs in the field. In this work, we leverage this rapid advancement in object detection to improve upon a downstream task: scene graph generation (SGG)[10]. SGG encompasses the set of tasks that focus on detecting object pairs and their relationships in visual media, creating scene graphs that can then be used for applications such as image captioning[7, 6], image generation[26], and visual question answering[31]. Two significant barriers stand in the way of higher performances for SGG tasks: The long-tailed distribution problem [15, 3] and the quadratic increase in candidate triplets [15, 3, 29]. When collecting data regarding a subject in an image and its relation to other objects, a small number of predicate terms often appear significantly more than all other terms. For example, 'on' can be used to

describe the relation between almost any subject-object pair where the subject is 'above' the object, and hence will appear significantly more than predicates such as 'mounted' or 'reaching'. Despite many attempts to alleviate the effect this uneven distribution has on the final results [14, 5, 18, 8], the prediction bias that arises from the uneven data distribution remains an important problem to be solved. The second barrier is the quadratic increase in computation required proportional to the number of detected objects. Mainstream approaches first obtain a list of objects in an image, and then infer the predicate (if one exists) for every possible subject-object pair [29, 10]. Unfortunately, the number of parameters required in the second stage scales quadratically with the number of objects found, as n objects would have n^2 potential predicates. Recent works have proposed a one-stage approach that leverages transformers to generate a set of triplets using object information[3, 16]. However, the large number of parameters required in a transformer architecture also slows down the model inference times, preventing the use of these approaches in real-time applications. This paper seeks to address the aforementioned long-tailed data distribution by leveraging the recent advancements in object detection accuracy. To do this, we first reformulate the SGG task into an object detection and classification problem, where each relationship triplet is represented as three objects: the subject, the object, and the relationship predicate. By formulating the problem in this way, we can divide the training data into balanced subsets, where every subset consists of predicate classes with a similar number of instances. This allows us to train multiple detection models, each specialized in detecting a subset of predicates. More importantly, we propose this approach as a solution to the long-tailed distribution problem, as these models will have significantly less bias than a single model trained on all predicate instances. Specifically, we propose a new framework, YOLO-SG, that leverages the YOLOv11 object detection model[21] to detect objects and predicates in parallel. YOLO-SG consists of four main components: object detection, predicate detection, object clustering, and predicate inference. The object and predicate detection module is composed of two sets of YOLOv11 models[21]. The first set is trained to detect and label objects, while the second set is trained to detect and label predicates. Additionally, we divide the training data such that every subset consists of 4 predicate classes, where the number of instances of each class in the subset is balanced. By doing so, we obtain multiple unbiased models that detect and label their own set of predicate classes instead of a single heavily biased model that predicts all predicate classes at once. The predicate inference module takes the output of the first stage and obtains a list of objects for each predicate using a simple clustering algorithm. It then passes every possible pair of objects in this list through a lightweight multilayer perceptron which is trained on the same subset of predicate classes. This then returns confidence scores for that subset of predicates. We then return the triplet with the highest confidence score of the detected predicate as the final output. Since we only use the object labels and positions as inputs as opposed to visual features, we can significantly reduce the number of trained parameters compared to methods that rely on convolutional layers or transformer architectures. We evaluate our model on the Visual Genome 1.2 dataset[12] and demonstrate that our model can achieve competitive performance with state-of-the-art models while maintaining real-time inference speeds.

2 Team Members

Shui Jie

As the overall project manager of the team, I am responsible for the overall work of the projects, including different iterations of goals and work distribution. I am responsible for designing the overall pipeline, training the YOLO model, and data analysis on individual model performance on different sets of objects and relationships. I also took part in coding the APIs to integrate my models into the complete pipeline.

Aleksei Lobanov

My name is Aleksei Lobanov, I am 23 years old, and I did my undergrad at Shenzhen MSU-BIT University, studying Applied Mathematics, where I did research on Formal Languages. At PKU, I study at the School of CS, NEEC Lab under Professor Luo Guojie. My research interests include programming languages, compiler development, and other systems programming-related topics.

This project was the first time I had to do anything ML-related, it was a big learning experience for me. I had two big tasks: pre-processing the dataset to try to improve the class imbalance within

the predicates, and developing a small neural network that tries to assemble the final output of object-predicate-subject triplets. I found working on the neural network particularly challenging, since I had to learn everything from scratch, and I have to thank my amazing teammates for helping me throughout the semester.

Lawrence Leroy Chieng Tze Yao

Charles Young

My name's Charles Young, I'm a 23-year-old who went to undergrad at UC San Diego for a bachelor's in Mathematics and Computer Science. I study under Professor Tao Xie, and my research interests include mutation testing, metaheuristics, and developer tool use. I was responsible for writing the report and researching related works to improve our framework. I was able to learn a lot about the current state of the art in scene graph generation and object detection, and I'm grateful to my teammates for their hard work and dedication to the project.

3 Background and Related Work

Scene Graph Generation

Since beginning in 2015 with Johnson et als. work[10], many works have been published improving upon and expanding the scope of SGG[3, 4, 29, 17, 11, 16, 20, 9, 14, 5, 8, 26, 18, 24, 2, 28]. Mainstream works often focus on the development of a two-stage scene graph generation approach composed of an object detection stage and a contextual reasoning stage [29?, 11]. This two-stage approach the is foundation for most modern SGG methods[29, 4, 17, 5, 8]. The first stage is often completed using pre-trained detection models[1], while contextual reasoning is done using a variety of methods, from convolutional neural networks[29] to transformers[3]. To address the quadratic growth of computation relative to object count common in two-stage models, recent works have proposed several one-stage approaches to the problem[3, 16]. These models leverage transformer architectures to generate a set of triplets using object information and have been shown to be competitive with two-stage models in terms of performance. However, the large number of parameters required in a transformer architecture also slows down the model inference times, preventing the use of these approaches in real-time applications[3, 18].

Long-tailed Distribution Problem Several works have been proposed to mitigate the long-tail bias introduced by the data[24, 30, 2, 9]. These works can be split into two categories: approaches that used biased training data paired with extra learning techniques or approaches that attempt to remove bias during training. In the first category, TDE[24] leverages a causal graph to infer the effect of long-tail bias. CogTree[30] builds a cognitive structure that distinguishes course-to-fine relationships based on biased predictions. DLFE[2] uses Dynamic Label Frequency Estimation to recover unbiased probabilities and reduce reporting bias. In the second category, The second category of works often approaches the problem by preprocessing training data to remove bias. PCPI[28] utilizes the correlation between predicate classes to adjust the loss weights. BGNN[17] uses a bi-level data resampling strategy with a confidence-aware bitartite graph to reduce bias. One particular work that heavily motivated our approach is the Context Knowledge Network (CKN) proposed by Jin et al. [9], which only uses the object labels and bounds as an input to a multilayer perceptron with the number of possible predicates as the output. Despite the lack of visual information, the authors were still able to achieve competitive high-speed performance with state-of-the-art models, whilst also reducing model bias. This work motivated our approach with the use of a lightweight MLP similar to CKN, as Jin et al. demonstrated potential in the ability of machine learning models to extract relationship information from non-visual contexts[9]. However, our approach differs as we use more than one model, with each model specializing in obtaining confidence scores for only a subset of predicates.

You Only Look Once (YOLO)

YOLO is a popular object detection model that has been widely used in many computer vision tasks due to its ability to perform single-pass, real-time inferencing. The first version of YOLO proposed by Redmon et al. in 2016 allowed for end-to-end training, as well as object bounding box proposal and

object classification in a single forward pass[21]. More importantly, the model was able to achieve this at a rate of 45 frames per second, more than enough for real-time applications. Since then, multiple versions of YOLO have been introduced by various different authors, each of which attempt to improve upon the original whilst following its core design philosophy: open source, end-to-end, and one-shot[25]. These newer versions often apply novel techniques to achieve greater overall performance, measured by inference speed, training cost, and inference accuracy[25, 22, 23], For our framework, we chose to use the YOLOv11 model due to its higher inference speed compared to YOLOv8 due to the lower number of trained parameters[27]. However, it is still possible that performance could be improved using an older version, but this will be left as potential future work.

4 Method

5 Object Detection

Our proposed framework, YOLO-SG, decouples the process of object and predicate detection by framing both subproblems as standard object detection tasks. Rather than inferring predicates between every pair of detected objects in a computationally expensive second stage, we directly detect "predicate entities" in parallel alongside object detection. By modeling predicates as distinct objects, we leverage YOLOv11[27] for both object and predicate recognition. This approach reduces the complexity of pairwise comparisons from n^2 operations to a more manageable parallel detection task.

6 Clustering Algorithm

We chose to use an Intersection-Over-Union (IoU) based clustering algorithm to associate detected objects with their corresponding predicates due to its simplicity and low time complexity. Given n objects, the algorithm has a worst-case complexity of $O(n\log n)$. For each predicate entity detected, we calculate the IoU of objects that overlap with the predicate bounds, grouping all objects that have an IOU larger than 0.5 with the predicate bounding box. Due to time constraints, we did not perform experiments to determine the optimal IOU threshold, and this is left as potential future work.

6.1 Preprocssing

Since we need the neural network to interpret words and letters meaningful, we first convert object and and predicate labels into vectors using GLOVE[19], a word-to-vector library commonly used by other works on SGG[14].

We use Visual Genome 1.2, a dataset composed of more than 108 thousand images and 2.3 million relationship predicates [12] as our training and test set. To treat predicates as objects, we assign each annotated predicate to a bounding box that encapsulates both the subject and object bounding boxes. For example, given a pair of objects $A, B \in \mathbb{R}^2$ with predicate r in Visual Genome 1.2, we create a new bounding box R that minimally encloses both R and R such that Our dataset are separated into wo In order to represent relationships and objects rather than edges between objects, we must add a corresponding predicate bounding box for each relationship triplet. We define the predicate bounding box R as the minimal bounding box that encapsulates both the subject and object bounding boxes. Given a pair of object bounding positions R0 with predicate R1, we define the predicate bounding box R2 as

$$R = (\min(A[1], B[1]), \max(A[2], B[2]))$$

This ensures that each predicate is represented as a single entity in the image which encompasses the subject and object. For example, given a 'man' with bounding box A = [0] We use the original scene graph annotations to produce a secondary dataset of predicate bounding boxes, each labeled with its corresponding predicate. We now have two datasets which are composed of the same images but different bounding boxes and labels: one has the bounds and labels for objects, while the other has the bounds and labels of predicates.

Data Splitting We split the dataset as follows. We first sort the predicates found in the dataset by the number of instances associated with each predicate. Next, we separate a number of predicates

into their own subset. We determine how many predicates to place into the set by calculating the ratio between the number of a instances containing the most frequently occurring predicate with the number of instances containing the least frequently occurring predicate. It follows that the larger the resulting number, the more unbalanced the dataset. For every subset, we select the largest number of predicates ordered by frequency such that the ratio does not exceed a chosen threshold.

Metric

Cleaning One significant issue with the Visual Genome dataset is that it contains a large number of object and predicate labels which are semantically equivalent to one another (for example, 'under' and 'below'). In a recent release, the maintainers of the dataset has provided us with a collection of synonym sets that group similar objects and predicates into the same class, with one term assigned by group. After converting all synonymous terms to their respective group representative term, we remove objects which rarely appear along with any associated triplets.

6.2 Model training

Object detection models Here we arrive at the long-tailed distribution problem. For objects, we train a YOLO model on objects Object YOLO: Trained on the object dataset, this model detects all objects in a given image. predicate YOLO: Trained on the predicate dataset, this model detects all predicate entities R in the same image. Each model is trained using conventional object detection procedures and loss functions, taking advantage of YOLO's efficient architecture. By isolating objects and predicates into their own detection tasks, we allow each model to specialize and minimize biases that often arise when both tasks are intertwined.

6.3 Inference

We start with weights pro the YOLO v11 model on the COCO dataset and evaluate it on the CoCo Dataset on 5 NVIDIA 4090Ti graphics cards. Training took 25 hours, with a maximum epoch of 300 and a batch size of 8. We evaluate our model using mean Recall@50.

Parallel detection For a given test image, we pass it through both the Object YOLO and the predicate YOLO models independently and in parallel. The Object YOLO output gives us a set of detected objects with corresponding bounding boxes and class labels. The predicate YOLO output provides a set of detected "predicate boxes," each with a predicted predicate label.

Clustering and association we need to determine which objects are involved in each detected predicate. We apply a clustering algorithm to group each predicate bounding box R with candidate object bounding boxes that it encloses or overlaps. This gives us one or more candidate subject-object pairs for each predicate detection.

Triplet selection using MLP scoring After obtaining two sets of detections—one for objects and one for predicates—we need the determine the subject and object associated with each triplet. For every predicate bounding box and label, we collect a list of detected objects have have a large overlap within the predicate box. To select the most like subject-object pair associated with the label, we use a lightweight MLP classifier trained to score the plausibility of a triplet (subject, predicate, object). For each predicate detection, we run the MLP once with every potential object pair that can be formed by objects in the list, selecting the triplet which returns the highest confidence score.

7 Evaluation

7.1 Evaluation Metrics

The task of scene graph generation can be divided into several subtasks, each with varying levels of difficulty. Predicate classification (PredCLS)[3, 9, 5] predicts predicates given the subject and object labels and bounds. Scene graph classification (SGCLS)[3, 5] predicts the triplet labels given the bounding boxes. Scene graph detection (SGDET)[3?], also referred to simply as Scene graph generation (SGGen)[9, 5], predicts the labels and bounding boxes of all three elements. To evaluate

our model, we follow the common standard set by previous works[] and adopt the evaluation metrics recall@K (R@K) and mean recall@K (mR@K). Given a single image, Recall@k is defined as the fraction of ground truth triplets which can be found in the top k triplet predictions. Two triplets are considered equivalent when all three elements (subject, predicate, and object) have been labelled correctly and both object and subject bounding boxes have IoU > 0.5 compared to the ground truth bounding boxes.

- 7.2 dataset
- 7.3 setting
- 7.4 implementation
- 7.5 results
- 8 Conclusion

9 Submission of papers to NeurIPS 2024

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9.1 Style

Papers to be submitted to NeurIPS 2024 must be prepared according to the instructions presented here. Papers may only be up to **nine** pages long, including figures. Additional pages *containing only acknowledgments and references* are allowed. Papers that exceed the page limit will not be reviewed, or in any other way considered for presentation at the conference.

The margins in 2024 are the same as those in previous years.

Authors are required to use the NeurIPS LaTeX style files obtainable at the NeurIPS website as indicated below. Please make sure you use the current files and not previous versions. Tweaking the style files may be grounds for rejection.

9.2 Retrieval of style files

The style files for NeurIPS and other conference information are available on the website at

The file neurips_2024.pdf contains these instructions and illustrates the various formatting requirements your NeurIPS paper must satisfy.

The only supported style file for NeurIPS 2024 is neurips_2024.sty, rewritten for LATEX 2ε . Previous style files for LATEX 2.09, Microsoft Word, and RTF are no longer supported!

The LATEX style file contains three optional arguments: final, which creates a camera-ready copy, preprint, which creates a preprint for submission to, e.g., arXiv, and nonatbib, which will not load the natbib package for you in case of package clash.

Preprint option If you wish to post a preprint of your work online, e.g., on arXiv, using the NeurIPS style, please use the preprint option. This will create a nonanonymized version of your work with the text "Preprint. Work in progress." in the footer. This version may be distributed as you see fit, as long as you do not say which conference it was submitted to. Please **do not** use the final option, which should **only** be used for papers accepted to NeurIPS.

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The file neurips_2024.tex may be used as a "shell" for writing your paper. All you have to do is replace the author, title, abstract, and text of the paper with your own.

The formatting instructions contained in these style files are summarized in Sections 10, 11, and 12 below.

10 General formatting instructions

The text must be confined within a rectangle 5.5 inches (33 picas) wide and 9 inches (54 picas) long. The left margin is 1.5 inch (9 picas). Use 10 point type with a vertical spacing (leading) of 11 points. Times New Roman is the preferred typeface throughout, and will be selected for you by default. Paragraphs are separated by $\frac{1}{2}$ line space (5.5 points), with no indentation.

The paper title should be 17 point, initial caps/lower case, bold, centered between two horizontal rules. The top rule should be 4 points thick and the bottom rule should be 1 point thick. Allow 1/4 inch space above and below the title to rules. All pages should start at 1 inch (6 picas) from the top of the page.

For the final version, authors' names are set in boldface, and each name is centered above the corresponding address. The lead author's name is to be listed first (left-most), and the co-authors' names (if different address) are set to follow. If there is only one co-author, list both author and co-author side by side.

Please pay special attention to the instructions in Section 12 regarding figures, tables, acknowledgments, and references.

11 Headings: first level

All headings should be lower case (except for first word and proper nouns), flush left, and bold.

First-level headings should be in 12-point type.

11.1 Headings: second level

Second-level headings should be in 10-point type.

11.1.1 Headings: third level

Third-level headings should be in 10-point type.

Paragraphs There is also a \paragraph command available, which sets the heading in bold, flush left, and inline with the text, with the heading followed by 1 em of space.

12 Citations, figures, tables, references

These instructions apply to everyone.

12.1 Citations within the text

The natbib package will be loaded for you by default. Citations may be author/year or numeric, as long as you maintain internal consistency. As to the format of the references themselves, any style is acceptable as long as it is used consistently.

The documentation for natbib may be found at

http://mirrors.ctan.org/macros/latex/contrib/natbib/natnotes.pdf

Of note is the command \citet, which produces citations appropriate for use in inline text. For example,

\citet{hasselmo} investigated\dots

produces



Figure 1: Sample figure caption.

Hasselmo, et al. (1995) investigated...

If you wish to load the natbib package with options, you may add the following before loading the neurips_2024 package:

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\usepackage[nonatbib] {neurips_2024}

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12.2 Footnotes

Footnotes should be used sparingly. If you do require a footnote, indicate footnotes with a number in the text. Place the footnotes at the bottom of the page on which they appear. Precede the footnote with a horizontal rule of 2 inches (12 picas).

Note that footnotes are properly typeset after punctuation marks.²

12.3 Figures

All artwork must be neat, clean, and legible. Lines should be dark enough for purposes of reproduction. The figure number and caption always appear after the figure. Place one line space before the figure caption and one line space after the figure. The figure caption should be lower case (except for first word and proper nouns); figures are numbered consecutively.

You may use color figures. However, it is best for the figure captions and the paper body to be legible if the paper is printed in either black/white or in color.

12.4 Tables

All tables must be centered, neat, clean and legible. The table number and title always appear before the table. See Table 1.

Place one line space before the table title, one line space after the table title, and one line space after the table. The table title must be lower case (except for first word and proper nouns); tables are numbered consecutively.

¹Sample of the first footnote.

²As in this example.

Table 1: Sample table title

	Part	
Name	Description	Size (μ m)
Dendrite Axon Soma	Input terminal Output terminal Cell body	$\begin{array}{c} \sim \! 100 \\ \sim \! 10 \\ \text{up to } 10^6 \end{array}$

Note that publication-quality tables *do not contain vertical rules*. We strongly suggest the use of the booktabs package, which allows for typesetting high-quality, professional tables:

https://www.ctan.org/pkg/booktabs

This package was used to typeset Table 1.

12.5 Math

Note that display math in bare TeX commands will not create correct line numbers for submission. Please use LaTeX (or AMSTeX) commands for unnumbered display math. (You really shouldn't be using \$\$ anyway; see https://tex.stackexchange.com/questions/503/why-is-preferable-to and https://tex.stackexchange.com/questions/40492/what-are-the-differences-between-align-equation-and-displaymath for more information.)

12.6 Final instructions

Do not change any aspects of the formatting parameters in the style files. In particular, do not modify the width or length of the rectangle the text should fit into, and do not change font sizes (except perhaps in the **References** section; see below). Please note that pages should be numbered.

13 Preparing PDF files

Please prepare submission files with paper size "US Letter," and not, for example, "A4."

Fonts were the main cause of problems in the past years. Your PDF file must only contain Type 1 or Embedded TrueType fonts. Here are a few instructions to achieve this.

- You should directly generate PDF files using pdflatex.
- You can check which fonts a PDF files uses. In Acrobat Reader, select the menu Files>Document Properties>Fonts and select Show All Fonts. You can also use the program pdffonts which comes with xpdf and is available out-of-the-box on most Linux machines.
- xfig "patterned" shapes are implemented with bitmap fonts. Use "solid" shapes instead.
- The \bbold package almost always uses bitmap fonts. You should use the equivalent AMS Fonts:

\usepackage{amsfonts}

followed by, e.g., \mathbb{R} , \mathbb{R} , \mathbb{R} , or \mathbb{R} , \mathbb{R} or \mathbb{R} . You can also use the following workaround for reals, natural and complex:

```
\newcommand{\RR}{I\!\!R} %real numbers
\newcommand{\Nat}{I\!\!N} %natural numbers
\newcommand{\CC}{I\!\!\!C} %complex numbers
```

Note that amsfonts is automatically loaded by the amssymb package.

If your file contains type 3 fonts or non embedded TrueType fonts, we will ask you to fix it.

13.1 Margins in LATEX

Most of the margin problems come from figures positioned by hand using \special or other commands. We suggest using the command \includegraphics from the graphicx package. Always specify the figure width as a multiple of the line width as in the example below:

```
\usepackage[pdftex]{graphicx} ... \includegraphics[width=0.8\linewidth]{myfile.pdf}
```

See Section 4.4 in the graphics bundle documentation (http://mirrors.ctan.org/macros/latex/required/graphics/grfguide.pdf)

A number of width problems arise when L^ATEX cannot properly hyphenate a line. Please give LaTeX hyphenation hints using the \- command when necessary.

Acknowledgments and Disclosure of Funding

Use unnumbered first level headings for the acknowledgments. All acknowledgments go at the end of the paper before the list of references. Moreover, you are required to declare funding (financial activities supporting the submitted work) and competing interests (related financial activities outside the submitted work). More information about this disclosure can be found at: https://neurips.cc/Conferences/2024/PaperInformation/FundingDisclosure.

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References

References follow the acknowledgments in the camera-ready paper. Use unnumbered first-level heading for the references. Any choice of citation style is acceptable as long as you are consistent. It is permissible to reduce the font size to small (9 point) when listing the references. Note that the Reference section does not count towards the page limit.

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A Appendix / supplemental material

Optionally include supplemental material (complete proofs, additional experiments and plots) in appendix. All such materials **SHOULD be included in the main submission.**

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