GeneCIS: A Benchmark for General Conditional Image Similarity

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Abstract

We argue that there are many notions of ‘similarity’ and that models, like humans, should be able to adapt to these dynamically. This contrasts with most representation learning methods, supervised or self-supervised, which learn a fixed embedding function and hence implicitly assume a single notion of similarity. For instance, models trained on ImageNet are biased towards object categories, while a user might prefer the model to focus on colors, textures or specific elements in the scene. In this paper, we propose the GeneCIS (‘genesis’) benchmark, which measures models’ ability to adapt to a range of similarity conditions. Extending prior work, our benchmark is designed for zero-shot evaluation only, and hence considers an open-set of similarity conditions. We find that baselines from powerful CLIP models struggle on GeneCIS and that performance on the benchmark is only weakly correlated with ImageNet accuracy, suggesting that simply scaling existing methods is not fruitful. We further propose a simple, scalable solution based on automatically mining information from existing image-caption datasets. We find our method offers a substantial boost over the baselines on GeneCIS, and further improves zero-shot performance on related image retrieval benchmarks. In fact, though evaluated zero-shot, our model surpasses state-of-the-art supervised models on MIT-States.

We, the architects of the machine, must decide a-priori what constitutes its ‘world’; what things are to be taken as ‘similar’ or ‘equal’ — Karl Popper, 1963

1. Introduction

Humans understand many notions of similarity and choose specific ones depending on the task at hand [21,58]. Consider the task of finding ‘similar’ images illustrated in Figure 1. Which of the rightmost images should be considered ‘most similar’ to the reference? Given different conditions, each image could be a valid answer. For instance, we may be interested in a specific object in the scene, focusing on either the ‘car’ or ‘bridge’. One could even indicate a ‘negative’ similarity condition, specifying a change in the image to identify the bottom image as most similar.

Learning such similarity functions is a central goal in discriminative deep learning [11–13, 34, 63, 68, 75]. Discriminative models, either supervised [30, 75] or self-supervised [9, 10], learn embedding functions such that ‘similar’ images are closer in feature space than ‘dissimilar’ images. However, since there are infinitely many notions of image similarity, how do we allow our models to choose?

Almost all current approaches assume a single notion of similarity, either by explicitly training on a specific concept [68, 75] or through an implicit assumption in the underlying data distribution [9, 12]. Meanwhile, prior works tackling the conditional problem have focused on constrained domains such as fashion [69,73] or birds [46], with a restricted set of similarity conditions. This is because developing and evaluating models that can adapt to generic notions of similarity is extremely challenging. Specifically, curating data to train and evaluate such models is difficult, as collecting annotations for all concepts of similarity is impossible.

\*Work done during an internship at Meta AI Research.
In this work we study the problem of general conditional image similarity, training on an open-set of similarity conditions, and evaluating on diverse similarity notions in a `zero-shot’ manner. We first design a benchmark comprising of four evaluation datasets for conditional image similarity, setting up conditional retrieval tasks. We define these tasks under a unified framework which spans practical use cases, and propose the benchmark as a sparse but broad coverage of the conditional similarity space. We propose these datasets for zero-shot evaluation only, and suggest that models which can perform well without fine-tuning can flexibly adapt to general notions of similarity, as desired. We name this benchmark GeneCIS (`genesis’) for General Conditional Image Similarity. On GeneCIS, we find that baselines built from powerful CLIP backbones struggle and, moreover, that performance on it is only weakly correlated with the backbones’ ImageNet accuracy [17]. This is in contrast to popular vision tasks such as segmentation [39] and detection [45], underlining the benchmark’s utility.

We also propose a solution to training general conditional similarity models, based on parsing large-scale caption datasets [64,66]. Rather than requiring exhaustive similarity annotations, we find that we can automatically mine this information from already abundant image-caption data. We show that training in this way offers substantial gains over the baselines, approaching (and in some cases surpassing) carefully designed specific solutions for each of the GeneCIS tasks. In addition, we demonstrate that our method scales with increasing amounts of caption data, suggesting promising directions for future work. Finally, on related benchmarks from the ‘Composed Image Retrieval’ (CIR) field [44,74], we find our method provides gains over zero-shot baselines. In fact, our model outperforms state-of-the-art on the MIT-States benchmark [28], despite being evaluated zero-shot and never seeing the training data.

**Contributions.** (i) We present a framework for considering conditional image similarity, an important but understudied problem; (ii) We propose the GeneCIS benchmark to test models’ abilities to dynamically adapt to different notions of similarity; (iii) We show that current vision-language models like CLIP struggle on GeneCIS, and that performance on it is only weakly correlated with ImageNet accuracy; (iv) We design a scalable solution to the conditional similarity problem based on automatically parsing large-scale image-caption data; (v) We show our models provide substantial gains over zero-shot CLIP baselines; (vi) We validate our models on related CIR benchmarks, surpassing state-of-the-art on MIT-States despite zero-shot evaluation.

2. Related Work

Our thesis that the similarity between two images should be conditional is generally relevant to the representation learning literature, which aims to learn embedding functions based on a single (often implicit) notion of similarity.

For instance, deep metric learning [30, 34, 63] aims to learn visual representations such that images from the same category are projected nearby in feature space. This idea is used in practical domains such as image retrieval [7, 59, 61], face verification [11, 67, 68] and vehicle re-identification [25, 31, 42]. The key limitation here is that networks are trained to encode a single notion of similarity, namely category-level similarity. While some work considered notions of similarity at different visual granularities [4, 15, 70], we posit that there exist concepts of similarity (e.g. shape and color) which are orthogonal to categories.

Meanwhile, contrastive learning [9, 10, 12, 13] defines notions of similarity by specifying a set of transformations to which the representation should be invariant (e.g. color jitter or random cropping), encouraging augmentations of the same instance to be embedded together. Similarly, vision-language contrastive training [29, 60] learns joint embedding spaces, where images’ representations are aligned with their paired captions. Though the precise notions of similarity are difficult to define in this case, we note that the embeddings are fundamentally unconditional, with a single deterministic embedding of a given image.

Finally, we highlight three relevant sub-fields in the literature: conditional similarity networks (CSNs); compositional learning (CL); and composed image retrieval (CIR). CSNs are networks with multiple subspaces for different notions of similarity [73]. Though their motivation is highly related to our work, CSNs are trained in a supervised manner with pre-defined similarity conditions [41, 46, 73], and/or are evaluated in constrained domains such as fashion [32, 69]. In contrast, we aim to train on an open-set of similarity conditions and evaluate zero-shot on natural images. Meanwhile, our work is related to CL research in that we seek to compose information from images and conditions to establish similarities. However, again, CL models are often assessed on their ability to recognize unseen combinations of a finite set of visual primitives [47, 54, 56]. Lastly, the most similar setup to GeneCIS is proposed in the recent CIR [74]. It tackles the problem of composing an image and text prompt to retrieve relevant images from a gallery [1, 3, 16]. This is typically posed in the context of fashion [23, 76], with the text prompt acting as an image edit instruction (e.g. ‘the same dress but in white’ [1]). As such, CIR tackles a subset of the conditional similarity problem, by presenting models with a ‘negative’ similarity condition.

**Key similarities and differences with prior work:** In this work, we leverage CIRR [44] and MIT-States [28] (natural image CIR datasets) for additional evaluations, and further leverage the ‘Combiner’ architecture [3] to compose text conditions and image features. Broadly speaking, our work differs from CSNs, CL and CIR in that we do not
train on a finite, closed-set of similarity conditions or visual primitives. Instead, we train models on open-world image-caption data, and demonstrate a flexible understanding of conditional similarity through zero-shot evaluation on a range of similarity conditions in natural images.

3. Conditional Similarity

We now describe our setup for the conditional similarity problem and its associated challenges—both with benchmarking models and acquiring data to train them. In § 4 we introduce the GeneCIS benchmark which measures important aspects of the problem. In § 5, we present a scalable solution to automatically acquire training data from widely available image-caption datasets.

Problem Definition: We define the problem of conditional similarity as learning a similarity function between two images given an explicit condition: \( f(I^T; I^R, c) \) yields the scalar similarity between a target image, \( I^T \), and a reference image, \( I^R \), given some external condition, \( c \). We use the scalar \( f(\cdot) \) to find the most conditionally similar image from a target set, i.e., to solve a retrieval task. In this work we consider the condition to be a user-specified text prompt, although other types of condition are possible. We highlight that standard image similarity, framed as \( f(I^T, I^R) \), implicitly assumes a similarity condition, often incorporated into the model or dataset (see § 2). We refer to the case where images are similar under an unspecified condition as the images being implicitly similar.

3.1. Challenges in training and evaluation

Challenges in evaluation: The key difficulty in evaluating conditional similarity is that there are infinitely many possible conditions: from ‘images with the same top-left pixel value are similar’ to ‘the same image but upside down is similar’. Thus, it is impossible to evaluate models’ ability to adapt to every similarity condition. Instead, in § 4, we introduce the GeneCIS benchmark which consists of a subset of such conditions, and covers a broad range of practical use cases. We suggest that models which produce zero-shot gains across GeneCIS, without finetuning, are more capable of flexibly adapting to different notions of similarity.

Challenges in acquiring training data: Since the space and diversity of similarity conditions is huge, acquiring human annotations to train for every type of conditional similarity is not feasible. For instance, to train a function which is sensitive to object category given some conditions (e.g., ‘car’ or ‘bridge’ objects in Figure 1), and ‘color’ given others (e.g., ‘blue’ or ‘black’ car in Figure 1), we need training data containing both features. Prior work addresses this by dramatically restricting the space of conditions and training on human annotations for pre-defined notions of similarity [46, 73]. In § 5, we describe an automatic method which leverages existing large-scale image-text datasets to learn an open-set of similarity conditions. The resulting model can be evaluated in a zero-shot manner across different types of conditional similarity task.

4. The GeneCIS Benchmark

GeneCIS considers two important dimensions of the conditional similarity problem. Firstly, a user may be interested in an object in the scene (‘with the same car’) or an attribute of a given object (‘the same color as the car’). Secondly, the condition could either focus on a particular aspect of the image (‘the same color as the car’) or specify the ‘negative’ space of a similarity condition, by defining a change in the image (‘this car but in black’).
We propose four evaluation tasks in GeneCIS, that covers the combination of the above dimensions and hence a diverse range of conditional similarities. For each of the tasks, we construct retrieval problems with: a reference image, \( I_R \); a text condition, \( c \); and a retrieval gallery of \( M \) target images, \( \{ I^T_i \}_{i=1}^M \), of which only one is ‘correct’ or ‘positive’. The task is to identify which of the target images is most similar to the reference, given the condition. The retrieval tasks, illustrated in Figure 2 with more examples in Appendix G.1, are:

- **Focus on an Attribute:** This task evaluates a model’s ability to focus on a specific attribute type (e.g. ‘color’ or ‘material’). For instance, in Figure 2, we see a white laptop and the condition ‘color’, with the task being to select the laptop with the same color from the gallery.

- **Change an Attribute:** This task contains ‘negative’ similarity conditions, considering target images with a specific attribute changed to be most similar. In Figure 2, the aim is to retrieve the same object (‘train’) but with the color changed from ‘green’ to ‘olive green’.

- **Focus on an Object:** This task considers reference images with many objects, and we refer to the set of objects together as a proxy for the image ‘scene’. The condition selects a single object from the reference as the most important (e.g. ‘refrigerator’ in Figure 2) and the ‘positive’ target contains the condition object as well as the same ‘scene’ (e.g. also contains ‘sky’, ‘chair’ *etc.* in Figure 2).

- **Change an Object:** This task considers ‘negative’ similarity through conditions which specify an object to be added to a scene. For instance, in Figure 2, ‘ceiling’ is specified, with the aim being to retrieve the same scene (a train station) but with a ceiling also present.

The tasks in GeneCIS are designed to be diverse and challenging for a single model while remaining well-posed. In Figure 2, given only the reference image, \( I_R \), and text condition, \( c \), a human can readily identify which of the target images is most ‘similar’. We wish to benchmark vision models’ competency at the same task.

For the benchmark to be challenging, we would want the model to need both the image content and the text condition to solve the problem. Thus, we include different forms of ‘distractor’ images in the galleries. For instance, for tasks with objects in the condition, we include distractors which have a similar ‘scene’ to the reference but do not contain the condition object. Such distractors are likely to affect models which are over-reliant on information from the reference image, without considering the condition. Similarly, we include distractors which contain the object specified in the condition, but not the reference scene, confusing models which solely rely on the condition. Meanwhile, for the attribute-based tasks, we include distractors which contain the reference object category, but not the correct attribute.

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**Table 1. Statistics** of the four tasks in the GeneCIS benchmark.

<table>
<thead>
<tr>
<th>Name</th>
<th>Base Dataset</th>
<th># Templates</th>
<th># Gallery Images</th>
</tr>
</thead>
<tbody>
<tr>
<td>Focus on an Attribute</td>
<td>VAW [56]</td>
<td>2000</td>
<td>10</td>
</tr>
<tr>
<td>Change an Attribute</td>
<td>VAW [56]</td>
<td>2112</td>
<td>15</td>
</tr>
<tr>
<td>Focus on an Object</td>
<td>COCO [36]</td>
<td>1960</td>
<td>15</td>
</tr>
<tr>
<td>Change an Object</td>
<td>COCO [36]</td>
<td>1960</td>
<td>15</td>
</tr>
</tbody>
</table>

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**Figure 3. Distribution of conditions** for attribute- and object-based conditions. For ‘Focus on an Attribute’, we show the distribution of the common attribute between the reference and positive target image (the condition itself is an attribute type, e.g. ‘color’), and vice-versa. As such, many targets are *implicitly similar* to the reference (similar given some condition), but the positive image is the most similar *given the provided condition*.

**Benchmark Details:** We construct all tasks by repurposing existing public datasets. For the two tasks which ‘focus on’ and ‘change’ *attributes*, we leverage the VAW dataset [56], which inherits from Visual Genome [37]. From VAW, we extract crops for individual objects and, for each object, use annotations for: object category; positively labelled attributes (which the object definitely possesses); and negatively labelled attributes (which the object definitely does not possess). For the two tasks which ‘focus on’ or ‘change’ *objects*, we use COCO Panoptic Segmentation data [36, 40] containing dense category annotations for every pixel in the image. We give full details of the template construction process for each task in Appendix A.1.

We show statistics of the evaluations in Table 1, including the number of retrieval templates and number of gallery images. We note that we carefully construct the benchmarks such that there is only one ‘positive’ image among the targets, with gallery sizes of between 10 and 15 images. This is different to many ‘text-to-image’ or ‘image-to-text’ retrieval benchmarks [40, 57], which contain galleries with thousands of targets. Though larger galleries increase the tasks’ difficulty, the galleries inevitably contain some valid targets which are treated as negative. We further show the distribution of objects and attributes specified in the conditions in Figure 3, noting that our space of conditions spans a long-tail of over 400 attributes and 100 objects.
Noise and human verification: Though, in principle, our benchmark should be error free, manual inspection of the templates shows that noise is introduced through underlying inconsistencies in Visual Genome [37], VAW [56] and COCO [36]. We are currently in the process of collecting manual annotations and human verification of the templates, and present the current version as ‘GeneCIS v0’.

5. Method

In § 5.1, we briefly describe preliminaries for our approach to learning general conditional similarity functions. This includes the model architecture and optimization objective which we inherit from prior work [3]. In § 5.2, we describe our main methodological contribution: an automatic and scalable way of mining conditional similarity training data from widely available image-caption datasets.

5.1. Preliminaries

Training data. To learn a conditional similarity function \( f(\cdot) \), we train with triplets \((I^R, I^T, c)\), where \( I^R \) and \( I^T \) are termed reference and target images, and \( c \) is the condition defining a relationship between them.

Model Architecture We parametrize the conditional similarity function \( f(\cdot) \) with deep networks, first encoding features for \((I^R, I^T, c)\) as \((x^R, x^T, e) \in \mathbb{R}^D\). We learn separate encoders, \( \Phi(I) \) and \( \Psi(c) \), for the images and text condition. Next, we train a ‘Combiner’ network [3], which composes the reference image features with the condition text features as \( g(x^R, e) \in \mathbb{R}^D \). Finally, we consider the scalar conditional similarity to be the dot product between the combined feature, and target image feature, as: \( f(I^T; I^R, c) = g(x^R, e) \cdot x^T \). Details of the Combiner architecture can be found in Appendix D and [3].

We initialize our image and text backbones, \( \Phi(\cdot) \) and \( \Psi(\cdot) \), with CLIP [60]. CLIP models are pre-trained on 400M image-text pairs containing a range of visual concepts. Furthermore, the visual and text embeddings from CLIP are aligned, making it easier to learn the composition between reference image and conditioning text features.

Optimisation Objective Given a batch of triplets, \( B = \{(I^R_i, I^T_i, c_i)\}_{i=1}^{|B|} \), we get features as \((x^R_i, x^T_i, e_i)\). Then, given a temperature \( \tau \), we optimise \((\Phi, \Psi, g)\) with a contrastive loss [50], as:

\[
    \mathcal{L} = -\frac{1}{|B|} \sum_{i \in B} \log \frac{\exp\left( g(x^R_i, e_i) \cdot x^T_i / \tau \right)}{\sum_{j \in B} \exp\left( g(x^R_i, e_i) \cdot x^T_j / \tau \right)}
\]

5.2. Scalable training for conditional similarity

To train for general conditional similarity, we wish to curate triplets for training, \( D_{\text{train}} = \{(I^R_i, I^T_i, c_i)\}_{i=1}^N \), with diverse conditions and concepts of similarity. However, as the space of conditions increases, the burden for exhaustively annotating such a dataset increases exponentially. Instead, our method (illustrated in Figure 4) automatically mines training triplets from existing data sources:

**Image-caption Data:** We begin with large-scale image-caption data scraped from the internet, containing images paired with descriptive captions [51, 66]. We hope that the captions contain information about the objects and attributes in the image, which we can utilize for the conditional similarity task. We also hope that such a method can scale with increasing data in the same way that conventional representation learning algorithms do.

**Extract relationships:** We use an off-the-shelf text-to-scene-graph parser [65, 77] to identify ‘Subject’ → ‘Predicate’ → ‘Object’ relationships within the caption [55]. For instance, from the central image in Figure 4, we extract the highlighted relationship ‘Horse’ → ‘on’ → ‘Canvas’. Note that one caption may contain many such relationships.

We find that many of the entities (‘Subjects’ or ‘Objects’) extracted by the parser are not visually grounded in
the image, e.g., pronouns (‘I’, ‘you’) or time-based nouns (‘today’, ‘yesterday’). To address this, we introduce an additional filtering step, where every entity is scored for ‘visual concreteness’ based on a pre-existing database [8]. The database contains human ratings between 1 and 5 for how visually apparent a noun is. For each extracted relationship, we average its ‘Subject’ and ‘Object’ concreteness scores, discarding relationships if their value is below a threshold.

**Construct triplets:** We first randomly select a relationship, taking the image it comes from as the ‘reference’, \( I^R \). Having identified the subject of the relationship (e.g. ‘Horse’ in the rightmost column of Figure 4) we identify all other relationships in the dataset containing the same subject. From this restricted pool of relationships, we randomly sample a ‘target’ relationship and image, \( I^T \), with the same subject but a different object (e.g. a horse on a ‘canvas’ instead of in a ‘meadow’ in Figure 4). Finally, we define the condition of the triplet, \( c \), as the concatenated ‘Predicate’ and ‘Object’ from the target relationship (‘on canvas’ in Figure 4).

**Discussion:** We note that our mined triplets exhibit a bias towards the ‘Change an Object’ GeneCIS task. However, the triplets often involve abstract relationships between reference and target images (e.g. ‘Horse on canvas’ in Figure 4). As such, solving the training task requires the model to use the condition to extract and modify diverse forms of information from the reference, which is the central requirement of the broader conditional similarity problem.

6. Main Experiments

We evaluate baselines, task-specific solutions, and our method on the proposed GeneCIS benchmark. § 6.1 describes the baselines as well as specific solutions which we design for each of the GeneCIS tasks. § 6.3 shows results on GeneCIS and, in § 6.4, we evaluate on related benchmarks from the Composed Image Retrieval (CIR) literature.

6.1. Baselines and Specific Solutions for GeneCIS

**CLIP-Only Baselines:** We provide three simple CLIP-only [60] baselines for GeneCIS. Our Image Only baseline embeds all images with the CLIP image encoder and retrieves the closest gallery image to the reference. The Text Only baseline embeds the text condition with the CLIP text encoder, and the gallery images with the image encoder, and finds the closest gallery image to the text embedding. Finally, our Image + Text baseline averages the reference image with the condition text feature, before using the combined vector to find the closest gallery image.

**CIRR Combiner baseline:** CIRR is a natural image dataset [44] containing 28K curated retrieval templates. All templates contain a human-specified text condition defining the relationship between the reference and ‘positive’ target image. Unlike our automatic and scalable triplet mining method, CIRR is manually constructed with a lengthy annotation process. We include a baseline from [3], which trains a Combiner model with a CLIP backbone on CIRR. For fair comparison with our method, we fine-tune both the image and text backbones on CIRR before evaluating the model zero-shot on GeneCIS, terming it Combiner (CIRR).

**Specific Solutions:** We also design specific solutions for each of the proposed tasks in GeneCIS. These solutions take into account the construction mechanisms of each task and represent sensible approaches to tackling the tasks independently. We design all solutions to respect the zero-shot nature of the evaluations and hence they are all based on ‘open-vocabulary’ models: we use CLIP for the attribute-based tasks and Detic [81] for the object-based ones. For the attribute-based tasks, we use CLIP to predict attributes or categories in the reference image, before using text embeddings of these predictions to search the gallery. For the object-based tasks, we use Detic to detect the object categories present in all images, treating the detected categories as bag-of-word descriptors of the target images. We give full details of the specific solutions in Appendix B.

6.2. Implementation Details

We train our strongest model on 1.6M triplets mined from Conceptual Captions 3 Million (CC3M) [66] which contains 3M image-caption pairs. Each triplet has a visual concreteness of at least 4.8 averaged over the ‘Subject’ and ‘Object’ entities in both the reference and target image. We train the contrastive loss with temperature \( \tau = 0.01 \) and batch size of 256, training for 28K gradient steps. We use early stopping based on the Recall@1 on the CIRR validation set and, for fair comparison with [3], initialize the image and text backbones with the ResNet50×4 CLIP model. Further details are in Appendix E.

6.3. Analysis on GeneCIS

We report results for all methods on the GeneCIS benchmark in Table 2. Our evaluation metric is Recall@K: the frequency with which the model ranks the ‘correct’ gallery image in its top-K predictions. We report results at \( K = \{1, 2, 3\} \) to evaluate under different constraints, and to account for any noise in the benchmark. We also report the Average R@1 over all tasks to measure the overall performance across different forms of conditional similarity.

**Takeaways:** From the baselines we find that both the ‘Image Only’ and ‘Text Only’ models perform poorly as expected, since they only rely on either the reference image content or the text condition. The ‘Image + Text’ and ‘Combiner (CIRR)’ models perform better, validating our claim that both the reference and text condition are required to solve the task. Phrased differently, this suggests the benchmark evaluates conditional similarity, as implicit similarity functions (e.g. the ‘Image Only’ baseline) perform poorly
The CIR task is to retrieve images which best match a concept rather than being specialized to domains such as bird species, insects, or flowers. We turn to the 'Focus Attribute' task, where the baseline performs slightly better than 'Combiner (CIRR)' on GeneCIS. As expected, most per-task specific solutions perform better than our general method. However, the broad zero-shot nature of GeneCIS makes all tasks independently challenging and the specific solutions do not work for all of them. Broadly speaking, we found that CLIP struggles to predict object attributes, and that Detic struggles on the 'stuff' categories in COCO Panoptic. Finally, caveats can be found in 'Image Only' results on the zero-shot evaluation.

### 6.4. Comparisons to Prior Work

GeneCIS uses natural images with general conditions, rather than being specialized to domains such as bird species, insects, or flowers. As such, to find comparable existing benchmarks, we turn to the Composed Image Retrieval (CIR) literature. The CIR task is to retrieve images which best match a composed reference image and editing text condition. This task aligns with the 'Change' dimension of GeneCIS. We evaluate on both the MIT-States benchmark [28] as well as on CIRR [44], with the former precisely reflecting the 'Change Attribute' GeneCIS task.

#### Metrics

On both benchmarks, we evaluate our model zero-shot on the test-sets and compare with prior work trained on the datasets. These datasets are partially labeled and evaluate using a global retrieval setting, i.e., the entire test-set is used as a gallery for each query. Thus, we follow prior work and report Recall@K at multiple $K = \{1, 5, 10\}$ to fully capture the model’s performance.

#### Results

We show results on MIT-States in Table 3. Prior work on this benchmark trains models on the dataset from scratch and thus is not zero-shot. Nonetheless, zero-shot evaluation of our model surpasses state-of-the-art on this task. However, we note that prior methods use smaller models compared to our pre-trained CLIP backbone.

We report on CIRR in Table 4, evaluating through the official test server and again comparing to methods that train for this setting. We report results for the Combiner method from the paper [3] as well as our improved implementation (see § 6.1), which are both trained on CIRR. Our improved implementation is a strong upper bound, surpassing previous fully supervised models. On zero-shot evaluation, our method surpasses the comparable baselines by a significant margin across all the recall metrics. Compared to supervised methods, our model outperforms [16] and [44] zero-shot, though we note [16] trains from scratch. Finally, our method outperforms the comparable zero-shot baselines.

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### Table 2. Evaluation on GeneCIS

<table>
<thead>
<tr>
<th>Specific Solution (Focus Attribute)</th>
<th>Focus Attribute</th>
<th>Change Attribute</th>
<th>Specific Solution (Change Attribute)</th>
<th>Focus Object</th>
<th>Change Object</th>
</tr>
</thead>
<tbody>
<tr>
<td>Image Only</td>
<td>R@1</td>
<td>R@2</td>
<td>R@3</td>
<td>R@1</td>
<td>R@2</td>
</tr>
<tr>
<td>Combiner (CIRR)</td>
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<td>27.7</td>
<td>39.8</td>
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<td>22.8</td>
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<tr>
<td>Combiner (CC3M, Ours)</td>
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<td>31.0</td>
<td>41.5</td>
<td>16.6</td>
<td>27.5</td>
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### Table 3. Results on MIT-States [28]

<table>
<thead>
<tr>
<th>Zero-shot</th>
<th>Recall @ 1</th>
<th>Recall @ 5</th>
<th>Recall @ 10</th>
</tr>
</thead>
<tbody>
<tr>
<td>TIRG [74]</td>
<td>✓</td>
<td>12.2</td>
<td>31.9</td>
</tr>
<tr>
<td>ComposeAE [1]</td>
<td>✓</td>
<td>13.9</td>
<td>35.3</td>
</tr>
<tr>
<td>LBF [27]</td>
<td>✓</td>
<td>14.7</td>
<td>35.3</td>
</tr>
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<td>HCL [79]</td>
<td>✓</td>
<td>15.2</td>
<td>36.0</td>
</tr>
<tr>
<td>MAN [19]</td>
<td>✓</td>
<td>15.6</td>
<td>36.7</td>
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<tr>
<td>Image Only</td>
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<td>3.7</td>
<td>14.1</td>
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<tr>
<td>Text Only</td>
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<td>9.5</td>
<td>22.5</td>
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<td>13.3</td>
<td>31.7</td>
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<td>Combiner (CC3M, Ours)</td>
<td>✓</td>
<td>15.8</td>
<td>37.2</td>
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</table>

### Table 4. Results on CIRR [44]

<table>
<thead>
<tr>
<th>Zero-shot</th>
<th>Recall @ 1</th>
<th>Recall @ 5</th>
<th>Recall @ 10</th>
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<tbody>
<tr>
<td>ARTEMIS [16]</td>
<td>✓</td>
<td>17.0</td>
<td>46.1</td>
</tr>
<tr>
<td>CIRPLANT [44]</td>
<td>✓</td>
<td>19.6</td>
<td>52.6</td>
</tr>
<tr>
<td>Combiner (CIRR, improved)</td>
<td>✓</td>
<td>40.9</td>
<td>73.4</td>
</tr>
<tr>
<td>Image Only</td>
<td>✓</td>
<td>7.5</td>
<td>25.7</td>
</tr>
<tr>
<td>Text Only</td>
<td>✓</td>
<td>20.7</td>
<td>43.9</td>
</tr>
<tr>
<td>Image + Text</td>
<td>✓</td>
<td>21.8</td>
<td>30.9</td>
</tr>
<tr>
<td>Combiner (CC3M, Ours)</td>
<td>✓</td>
<td>27.3</td>
<td>57.0</td>
</tr>
</tbody>
</table>

We do not report on this as we found that the ‘Text Only’ baseline performed comparably with SoTA models on this task, achieving over 60% Recall@1.
Table 5. Ablations of key design choices of our full model with results reported on our GeneCIS benchmark.

<table>
<thead>
<tr>
<th>Ablations</th>
<th>Average Recall @ 1</th>
</tr>
</thead>
<tbody>
<tr>
<td>Full Model</td>
<td>16.8</td>
</tr>
<tr>
<td>No filtering for visual concreteness</td>
<td>15.0</td>
</tr>
<tr>
<td>Freezing CLIP image backbone</td>
<td>14.7</td>
</tr>
<tr>
<td>Freezing CLIP text backbone</td>
<td>15.8</td>
</tr>
<tr>
<td>Freezing entire backbone</td>
<td>15.1</td>
</tr>
<tr>
<td>Training on SBU [51] instead of CC3M [66] caption data</td>
<td>16.5</td>
</tr>
</tbody>
</table>

model reduces the gap between the baselines and specialist Combiner models trained on CIRR.

7. Analysis

Ablations: Table 5 shows the effect of our design choices on the performance on GeneCIS. We find that filtering out relationships which are not visually concrete, and finetuning the entire backbone, both strongly affect the performance. We verify the robustness of our triplet mining procedure by training with SBU Captions [51], a smaller but different source of image-caption data. We find that though the larger CC3M [66] produces slightly better results, different image-caption datasets are also suitable.

Comparing pretrained backbones: In Figure 6, we study the effect of changing the CLIP initialization. We train Combiner models with ResNet [26] and ViT [18] backbones on CC3M, showing their performance as well as the ‘Image + Text’ baseline from § 6.1.

We plot the performance on GeneCIS against the CLIP backbone’s zero-shot ImageNet accuracy [17]. We observe that the performance on GeneCIS is weakly correlated with the ImageNet performance of the backbone: a Top-1 gain of 10% on ImageNet leads to only 1% improvement on GeneCIS. This suggests that improvements on ImageNet do not directly transfer to GeneCIS and that GeneCIS measures a different yet important capability of vision models. In addition, our method offers a substantial boost over the ‘Image + Text’ baseline, and a greater boost than scaling the underlying CLIP model. Both of these results are in stark contrast to trends on popular vision tasks such as segmentation [39] and detection [45], where gains on ImageNet directly transfer to large gains on the downstream task, and often more significantly so than gains from the underlying method.

Scaling the number of triplets: In Figure 5, we investigate the effect of scaling the conditional similarity training data. We successively decrease the number of mined triplets by factors of four (from the 1.6M used to train our strongest models) both with and without concreteness filtering. We find results improve with increasing numbers of triplets and that while our models are trained on a dataset of 3M image-caption pairs [66], open-source caption datasets exist with up to five billion images [64]. We emphasize the utility of this finding, suggesting it is possible to train stronger conditional similarity models by further scaling the training data.

8. Conclusion

In this paper we have proposed the GeneCIS benchmark for General Conditional Image Similarity, an important but understudied problem in computer vision. The benchmark extends prior work and evaluates an open-set of similarity conditions, by being designed for zero-shot testing only. Furthermore, we propose a way forward for scalably training conditional similarity models, which mines information from widely available image-caption datasets.

Our method not only boosts performance over all baselines on GeneCIS, but also provides substantial zero-shot gains on related image retrieval tasks. Moreover, we find that unlike for many popular vision tasks, the performance of our models on GeneCIS is roughly decorrelated from scaling the backbone network’s ImageNet accuracy, motivating further study of the conditional similarity problem.
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