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JoinABLe: Learning Bottom-up Assembly of Parametric CAD Joints

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Abstract

Physical products are often complex assemblies combining a multitude of 3D parts modeled in computer-aided design (CAD) software. CAD designers build up these assemblies by aligning individual parts to one another using constraints called joints. In this paper we introduce JoinABLe, a learning-based method that assembles parts together to form joints. JoinABLe uses the weak supervision available in standard parametric CAD files without the help of object class labels or human guidance. Our results show that by making network predictions over a graph representation of solid models we can outperform multiple baseline methods with an accuracy (79.53%) that approaches human performance (80%). Finally, to support future research we release the Fusion 360 Gallery assembly dataset, containing assemblies with rich information on joints, contact surfaces, holes, and the underlying assembly graph structure.

1. Introduction

The physical products that surround us every day are often complex assemblies combining a multitude of parts modeled using computer-aided design (CAD) software. Well-designed assemblies are critical to ensure that products are cost-efficient, reliable, and easy to physically assemble. CAD designers build up assemblies by aligning pairs of parts together using constraints called *joints*. These joints determine the relative pose and allowed degrees of freedom (DOF) of parts in an assembly [43]. For example, a bolt can be constrained to a hole, then a nut constrained to the bolt, and so on until an entire assembly is designed. Assemblies may contain thousands of parts, represented as solid models in the boundary representation (B-Rep) format [33, 61], and are used for everything from furniture, to vehicles, to electronic devices. Defining individual global positions for each part without using joints quickly becomes cumbersome and prone to error. Joints enable designers to make quick parametric changes to a design while preserving existing part relationships and maintaining design intent.



Figure 1. CAD assemblies contain valuable *joint* information describing how parts are locally constrained and positioned together. We use this weak supervision to learn a bottom-up approach to assembly. *JoinABLe* combines an encoder and joint axis prediction network together with a neurally guided joint pose search to assemble pairs of parts without class labels or human guidance.

However, fully defining joints in assemblies is timeconsuming – roughly one third of time in CAD is spent doing assembly work [27]. As a result many assemblies have missing or partly defined joints. A learning-based approach capable of predicting joints could ease the burden of joint definition and enable other applications such as CAD assembly synthesis [56], robotic assembly [34], optimization of dynamic assemblies [72], part motion prediction [58], assembly-aware similarity search [5] and many more. Although joints for real world assemblies are configured in a bottom-up fashion, recent work largely takes a top-down approach to assembly related tasks [19, 23, 38]. Top-down approaches learn a global arrangement of parts from set object and part classes in carefully annotated data. An open challenge remains to learn to assemble parts without relying on the strong object and part class priors provided in heavily annotated datasets. In this work we ask the following question, illustrated in Figure 1: Given a pair of parts, can we automatically assemble them without prior knowledge of the global design, class labels, or additional human input? Solving this problem is a fundamental building block for leveraging learning-based methods with assemblies. Our long-term motivation is to enable the next generation of assembly aware tools that can increase the reuse of existing components and streamline robotic assembly and disassembly – important steps in reducing the negative impact of physical products [8, 29, 35, 41, 45].

To begin to address this challenge we introduce *JoinABLe* (Joint Assembly Bottom-up Learning), our *bottom-up* approach to assembly that learns how parts connect locally to form parametric CAD joints. *JoinABLe* uses the weak supervision available in parametric CAD files, containing only partial joint labels, to automatically assemble pairs of parts. We make the following contributions:

- We propose a novel learning-based method to automatically assemble pairs of parts using the weak supervision available in parametric CAD files. We do this without the help of object or part class labels, human annotation, or user guidance for the first time.
- We create and release the *Fusion 360 Gallery* assembly dataset, containing CAD assemblies with rich information on joints, contact surfaces, holes, and the underlying assembly graph structure.
- We provide experimental results on both joint axis and joint pose prediction tasks, a human baseline study, and comparisons with multiple other methods.

Our results show that by making network predictions over a graph representation of solid models, we can outperform multiple baseline methods while using fewer network parameters. We demonstrate that our approach performs well with difficult cases where heuristic algorithms can struggle and achieves an accuracy (79.53%) that approaches human performance (80%) on the joint axis prediction task.

2. Related Work

Assemblies have been a critical part of design and engineering for centuries. Since the digitization of CAD in the 1980s, a number of research areas have been explored.

Shape Combination As early as 2004 the power of designing assemblies by combining and reusing existing parts was demonstrated in *Modeling by Example* [13]. Since then a body of work has focused on finding compatible parts to combine together into assemblies [9, 22, 24, 66, 73]. The ability to parametrically assemble parts into novel designs has numerous applications in the media and entertainment

industry, where digital worlds can be populated with novel content. Other lines of work have focused on assemblies that can be physically fabricated [12,32,40,51,52,55,59,60] or conversion to/from assembly instructions [1, 52]. Our work differs in that we automate the pair-wise assembly of real-world CAD parts using a learning based method without class labels or human guidance.

Structure Aware Deep Generative Models 3D shape synthesis has rapidly advanced with the use of structure aware deep generative models [10, 14, 15, 28, 36, 46, 50, 64, 69] that incorporate some notion of assembly structure to describe how the parts of a shape form a whole. Rather than synthesize the parts themselves, we focus instead on assembling existing parts in the industry-standard B-Rep format.

CAD Informed Robotic Assembly Prior knowledge of CAD assemblies has been leveraged for robotic assembly planning [16, 17] and sequencing [11, 26] to constrain the search process and validate assembly sequences. Although not addressed in this work, we envision our approach can aid in improving the sampling efficiency of reinforcement learning based robotic assembly [57] by inferring joint information when it is absent or not fully specified.

Learning to Assemble Learning-based assembly methods from the literature largely follow a top-down approach that predicts the absolute pose of a set of parts to form an assembly [23, 38, 56, 70]. Predicting the absolute pose, however, can lead to noisy results where parts fail to completely align. To deal with this issue several recent works have leveraged supervision from local contact points between parts [18, 19]. We believe a bottom-up approach is a critical part of solving the assembly problem. Rather than rely on contact points, our work uses the joint information found in parametric CAD files as weak supervision. This allows the output of our method to be reconstructed as fully editable parametric CAD files.

Critical to prior work is training on synthetic assemblies [47,65] that belong to set object classes, e.g. chairs, drawers, etc., and are manually segmented, annotated with part class labels, and oriented in a consistent manner. However, semantic segmentation is often incompatible with real-world CAD assemblies that segment parts by manufacturing process [43]. Moreover, while training on set object classes greatly improves within-class performance, generalization to unseen categories is an ongoing area of research [18]. Rather than rely on heavily annotated datasets with strong class priors, our work leverages the weak supervision readily available in standard parametric CAD files, and is trained *without* object classes.

Concurrent to our work, AutoMate [27] leverages similar joint information for use with a learning based recommendation system. Here the user selects an area on each part as guidance, and using those selections, AutoMate recommends to the user multiple joint solutions confined to the



Figure 2. JoinABLe is used to assemble a pair of parts in the B-Rep format (a). We use supervision from parametric CAD files containing user selected B-Rep faces and edges that define joints (cyan). We also identify 'equivalent' faces and edges (pink) sharing the same joint axis for use during evaluation. Graphs for each part G_1 , G_2 are constructed from adjacent B-Rep faces and edges (b), then joint connectivity predictions are made over a graph G_j containing dense connections between all graph vertices. G_j is shown as an $n \times m$ matrix (c) to visualize the prediction space. Finally, the parts are aligned along the predicted joint axes (d), ready for a subsequent search stage.

user-selected input area. Similar to AutoMate, our method enables editable joints to be created in CAD, but we do so in an automated way that does not require user guidance and is not limited to a predefined area. We believe providing an automated solution is critical to enabling advanced assembly applications for CAD and robotics.

Part Mobility Understanding how assembled parts might move, i.e. *part mobility*, is an important problem in both CAD and robotics where the goal is to articulate a given part, such as a hinged door, without knowing the part mobility in advance. Most relevant to our work are systems that automatically predict the relative joint configurations between pairs of parts [37, 58, 68]. Here the input is a point cloud and the output joint axis parameters that define how the parts move in relation to one another. Again, these works rely on strong class priors and heavily annotated synthetic assembly data. We compare our method with adaptions of several part mobility baselines in Section 5.

3. Method

We now present our method, *JoinABLe*, for automatically assembling pairs of parts with joints.

3.1. CAD Joints

Assembly parts are typically represented in the B-Rep format, containing a watertight collection of trimmed parametric surfaces connected together by a well-structured graph [61]. Each face contains a parametric surface, and is bounded by edges that define the trimmed extent of the surface using parametric curves such as lines, arcs, and circles. The B-Rep format is used in all mechanical CAD tools and the selection of B-Rep entities, i.e. faces and edges, is a critical but time-consuming manual task required to set up joints. Our method proposes to learn from these user selections to automate the process of joint creation.

The best practice for CAD assembly is to define relative relationships between pairs of parts, to form joints, also known as *mates*. Joints define the degrees of freedom between the two parts, the parameters for a rest state pose, and the overall motion limits. CAD users select B-Rep entities on each part (highlighted in cyan in Figure 2a) to define a per-part joint axis consisting of an origin point and a direction vector. The joint axes are determined by the type of geometry selection, for a circle the center point becomes the origin point and the normal becomes the direction vector. These two parts can then be aligned along their axes into an assembled state (Figure 2d).

3.2. Joint Prediction Problem Statement

Given a pair of parts (Figure 2a), we aim to create a parametric joint between them, such that the two parts are constrained relative to one another with the same joint axis and pose as defined by the ground truth (Figure 2d). Here the joint axis is defined by two joint origin points and joint direction vectors relative to each part, and the pose is defined by a single rigid transformation in absolute coordinates. We refer to the tasks of predicting these values as *joint axis prediction* and *joint pose prediction*, respectively. We consider only pairs of parts that form rigid joints, and leave full multi-part assembly and non-rigid joints to future

work. We assume that object or part class labels and any form of human guidance are unavailable. We train only using the weak supervision provided by standard parametric CAD files without any manual human annotation such as canonical alignment.

3.3. Input Representation

Our method takes a pair of parts in the B-Rep format (Figure 2a), building upon a line of recent work [25, 31, 62, 67] that utilizes the topology and geometry available within B-Rep CAD data. This approach enables us to make predictions over the exact entities used to define joints, rather than an intermediate representation such as a mesh or point cloud. Importantly, it allows us to frame the problem as a categorical one, by making predictions over the discrete set of B-Rep entities that contain ground truth information about the joint axis. Joints are commonly defined between both B-Rep face and edge entities, e.g. a cylinder (face) can be constrained to another cylinder (face) or a circle (edge). To accommodate this, for each part we build a graph representation, G(V, E), from the B-Rep topology where graph vertices V are either B-Rep faces or edges, and graph edges E are defined by adjacency (Figure 2b).

For graph vertex features we use information about individual B-Rep faces and edges readily available in the B-Rep data structure. For B-Rep faces, we use a one-hot vector for the surface type (plane, cylinder, etc.) and a flag indicating if the surface is reversed with respect to the face. For B-Rep edges, we use a one-hot vector for the curve type (line, circle, etc.), the length of the edge, and a Boolean flag indicating if the curve is reversed with respect to the edge. We evaluate the performance of these input features and others in Section A.2 of the supplementary material.

Finally, given the two graphs G_1, G_2 that we wish to assemble, with n and m vertices respectively, we form a third 'joint connectivity graph' G_j that densely connects the vertices between G_1 and G_2 . G_j has $n \times m$ edges and allows us to formulate a link prediction problem [39], by identifying the connections between G_1 and G_2 that form a joint. G_j can be easily visualized as an $n \times m$ matrix (Figure 2c).

3.4. Weak Supervision from CAD Joints

A pair of parts in the B-Rep format have a finite number of faces and edges that can be paired to form a joint, specifically the $n \times m$ edges in G_j . Each ground truth joint results in a single positive label in the $n \times m$ prediction space and all remaining combinations are negative labels. For complex parts, such as mechanical gears that may contain thousands of discrete B-Rep entities, this results in an *extreme* imbalance between positive and negative labels.

The problem is further compounded by having only weak supervision available in standard parametric CAD files. This is due to several reasons: firstly, specifying joints between parts is time consuming and is often skipped by CAD designers; secondly, each CAD assembly is designed for a specific purpose, rather than to create an exhaustive set of assembly configurations. This weak supervision results in a positive and unlabeled (PU) learning problem [4] where the joints are known positive labels, but the remaining negative labels could be positive (i.e. an unseen but plausible joint) or negative (i.e. an implausible joint). To address the data imbalance and PU learning problem, we organize and augment our data using the following three techniques.

Joint Consolidation To increase the number of positive labels, we consolidate joints between identical pairs of parts into *joint sets*. Figure 4, right shows an example joint set where the same two parts are connected in multiple different ways. This approach allows us to present the network with a *single* data sample, i.e. a joint set, that contains all known joints between a pair of parts. Importantly, joint consolidation avoids presenting the network with multiple contradictory data samples, where a negative label in one sample may be a positive label in another sample. We provide additional implementation details about joint consolidation in Section A.1 of the supplementary material.

Joint Equivalents To further counter the extreme data imbalance, we identify and label 'equivalent' entities that share the same joint axis as the ground truth. For example, if a circle is the labeled entity (highlighted in cyan in Figure 2a), its neighbouring faces, such as the cylinder highlighted in pink, will be labeled as equivalent. These entities represent the same user-selected joint axis and only differ by the origin point that locates the joint axis in 3D space. As we consider a predicted joint axis, we include equivalent labels during *evaluation*. We perform an ablation study in Section A.2 of the supplementary material to evaluate the contribution of equivalent labels.

Unambiguous Evaluation Sets A challenge with PU Learning is establishing a 'clean' test set to accurately measure network performance. Parts that have multiple plausible joints, such as a plate with multiple holes for fasteners, are problematic if only partial joint labels exist, leading to ambiguity at test time. We make a best effort to avoid positive unlabelled samples in the test and validation set by excluding geometrically similar, but unlabeled, 'sibling' entities, e.g. the faces and edges of an unlabeled hole with the same size as a labelled hole. We identify sibling entities by matching the entity type, area or length, and number of connected graph edges to the labeled entities. In Section A.2 of the supplementary material we study the effect of evaluating with sibling entities on a withheld test set that matches the original data distribution.



Figure 3. *JoinABLe* architecture. Given two B-Rep parts in our graph representation, the vertex features from B-Rep faces (green) and edges (orange) pass through separate multi-layer perceptrons (MLP) before being concatenated together and passed through a message passing network (MPN). This yields local vertex embeddings representing each B-Rep entity in the two parts. Our joint axis prediction branch then performs edge convolution *between* the two graphs to estimate the presence of joints over all possible pairs of connections. Finally, the joint parameters are discovered via search, with respect to the predicted joint axis, to complete the assembly.

3.5. JoinABLe Architecture

Our overall architecture is shown in Figure 3 and consists of an encoder module that outputs per-vertex embeddings for each B-Rep face and edge in our graph representation of the input parts. Using these embeddings we can predict a joint axis and then search for joint pose parameters.

3.5.1 Encoder

Our encoder neural network f_{enc} is a Siamese-style network with shared weights for the two parts. It firstly creates graph vertex embeddings by passing the vertex features x_1 and x_2 from the two graphs, through two separate multi-layer perceptrons (MLP). One MLP is used for vertices representing B-Rep faces and another for those representing B-Rep edges; the resulting vertex embeddings are then concatenated together. We next perform message passing *within* each part's graph using a two-layer Graph Attention Network v2 (GATv2) [6] to obtain the per-vertex embeddings h_1 and h_2 for both the graphs.

$$\mathbf{h}_1 = f_{\text{enc}}(\mathbf{x}_1, G_1), \quad \mathbf{h}_2 = f_{\text{enc}}(\mathbf{x}_2, G_2).$$
 (1)

The idea here is to extract local features within each part that consider each B-Rep entity and its neighborhood.

3.5.2 Joint Axis Prediction

When creating a joint, a key piece of design intent is the definition of a joint axis by which two parts can be aligned and constrained to one another. The joint axis forms the basis for the degrees of freedom to be defined and enables downstream tasks such as assembly, part mobility, and animation. We formulate joint axis prediction as a link prediction problem, where the goal is to correctly identify a connection between G_1 and G_2 that aligns the two parts along a ground truth joint axis. This is done by aggregating information *between* parts using an edge convolution along the edges of G_j . The node features \mathbf{x}_1 and \mathbf{x}_2 from graphs G_1 and G_2 , are passed through our shared encoder network f_{enc} to get 384-dimensional embeddings \mathbf{h}_1 and \mathbf{h}_2 (Eq. 1). Then for each edge (u, v) in the graph G_j which densely connects G_1 and G_2 , we predict a logit indicating the presence of a joint:

$$\mathbf{h}_{uv} = \phi(\mathbf{h}_u \oplus \mathbf{h}_v),\tag{2}$$

where $\phi : \mathbb{R}^{768} \mapsto \mathbb{R}$ is a 3-layer MLP, \oplus is the concatenation operator and \mathbf{h}_u and \mathbf{h}_v are gathered from \mathbf{h}_1 and \mathbf{h}_2 based on the source and target vertices for each edge in G_j .

We train the network with a loss function that has two terms. The first term \mathcal{L}_{CE} is the cross-entropy between the edge predictions \mathbf{h}_{uv} and the ground truth edge labels $\mathbf{j}_{uv} \in \{0, 1\}$ normalized into a probability distribution \mathbf{j}_{uv} .

$$\widehat{\mathbf{h}}_{uv} = \text{softmax}_{\text{all}}(\mathbf{h}_{uv}), \mathcal{L}_{\text{CE}} = \text{CE}\left(\widehat{\mathbf{j}}_{uv}, \widehat{\mathbf{h}}_{uv}\right).$$
(3)

Here the subscript in the softmax operation indicates that it is applied over all edges in G_j , and $CE(\mathbf{p}, \mathbf{q}) = -\sum_i \mathbf{p}_i \log \mathbf{q}_i$. This loss encourages true joints to have higher values while simultaneously suppressing non-joints. We observe this is sub-optimal due to the sparsity of positive labels, where \mathcal{L}_{CE} is summed over a large number of terms. To better focus the loss term so that the joints are better contrasted against more likely non-joints, we use a symmetric cross entropy loss \mathcal{L}_{Sym} as the second term in the loss function.

$$\begin{split} & \hat{\mathbf{h}}_{row} = \text{softmax}_{row}(\mathbf{h}_{2D}), \quad \hat{\mathbf{h}}_{col} = \text{softmax}_{col}(\mathbf{h}_{2D}), \\ & \mathcal{L}_{Sym} = \text{CE}(\hat{\mathbf{j}}_{2D}, \hat{\mathbf{h}}_{row}) + \text{CE}(\hat{\mathbf{j}}_{2D}, \hat{\mathbf{h}}_{col}). \end{split}$$
(4)

Here the subscript of the softmax indicates that it is taken over a single axis, and the 2D subscript instead of uv indicates that the predictions and ground truth labels on the edges of G_j are reshaped into $n \times m$ matrices.

3.5.3 Joint Pose Search

The B-Rep entities predicted by our network allow us to query the ground truth B-Rep data to obtain a joint axis prediction for each part. Once these axes are aligned together, three secondary parameters define a rigid joint and can be used for joint pose prediction. An *offset* distance along the joint axis, *rotation* about the joint axis, and a *flip* parameter to reverse the joint axis direction. We find these parameters using a neurally guided search that allows us to enumerate over the top-k joint axis predictions and directly consider interaction between both parts. To evaluate a candidate joint configuration, we propose a cost function $C_{\text{joint}} = C_{\text{overlap}} + \lambda C_{\text{contact}}$ that considers two general criteria for well defined joints: overlap volume and contact area between parts, formulated as:

$$C_{\text{overlap}} = \frac{V_{1\cap 2}}{\min(V_1, V_2)}, \quad C_{\text{contact}} = \frac{A_{1\cap 2}}{\min(A_1, A_2)}.$$
 (5)

Here, V_1 and V_2 are the volume of the two parts, and $V_{1\cap 2}$ represents their overlap volume. Similarly, A_1 and A_2 are the surface area of the two parts, and their contact area is $A_{1\cap 2}$. Intuitively, for the two parts to align closely to each other, minimizing the cost function should encourage a larger contact area while penalizing the overlap volume to prevent penetration. Therefore, we let $\lambda = -10$ if $C_{\text{overlap}} < 0.1$. Otherwise, we set $\lambda = 0$ to increase the overlap penalty. Given this cost function, we search for the optimal joint pose using the Nelder-Mead algorithm [48] as a standard derivative-free optimization.

4. Dataset

To evaluate the performance of our method we create the *Fusion 360 Gallery* assembly dataset, derived from designs created in Autodesk Fusion 360 and submitted to the publicly available Autodesk Online Gallery [3]. The dataset consists of two inter-related sets of data, *Assembly Data*, containing 8,251 assemblies with 154,468 separate parts, and *Joint Data* containing 32,148 joints defined between 23,029 different parts. The data and supporting code are publicly available on GitHub¹ with a license allowing non-commercial research. We now describe the joint data used in our experiments and provide information on the overall dataset in Section A.1 of the supplementary material.

Figure 4, left shows an overview of the joint data in our dataset. We consider a data sample to be a joint set, such as shown in Figure 4, right, containing a pair of parts with one or more joints defined between them. The user-selected B-Rep faces and edges form the ground truth labels together with the joint axis and pose information of each joint. We provide an approximate 70/10/10% data split, for the



Figure 4. An overview of joint data from the *Fusion 360 Gallery* assembly dataset (left). Each sample consists of a unique pair of parts with *one or more* joints defining how they are locally constrained and positioned together (right).

training, validation, test, and original distribution test sets respectively. The validation and test sets do not include samples with potentially ambiguous sibling entities, while the original distribution test set does.

5. Experiments

In this section we perform experiments to qualitatively and quantitatively evaluate our method on two tasks: joint axis prediction and joint pose prediction. We examine how our method compares with a human CAD expert and other methods from the literature. A key criteria for evaluating performance is to gauge how the network performs in scenarios that traditional algorithms find challenging. One such scenario involves designs that do not contain connections between cylindrical shafts and holes, such as a bolt and a hole similar to Figure 2a. Commercial products exist which infer joints of this type by searching for fasteners and holes with similar radii [54]. In our dataset we see that 82% of data samples contain holes and 47.5% of joints constrain circular or cylindrical entities on one part to a hole on the opposing part. In our experiments we report results that gauge the ability of our approach to correctly infer joints, both in the simple Hole case and the more complex No Hole case. Details of experiment procedures are provided in Section A.2 of the supplementary material.

5.1. Human CAD Expert Baseline

Understanding how a human CAD expert performs in a similar setting is important to gauge the efficacy of each method. We conduct a study to establish a human baseline by recruiting a CAD expert, who works on commercial CAD design, and ask them to assemble pairs of parts from

¹https://github.com/AutodeskAILab/Fusion360GalleryDataset

	All Acc.%↑	Hole Acc.% ↑	No Hole Acc.% ↑	Param. #↓
Ours	79.53	80.15	76.59	1.3M
B -Dense	10.59	10.36	10.59	3.2M
B -Discrete	4.28	4.18	4.79	4.0M
B-Grid	65.21	65.09	65.81	3.1M
B -Heuristic	71.39	72.74	64.97	-
B-Random	21.55	21.92	23.29	-
Human	80.00	-	-	-

Table 1. Joint axis prediction accuracy results are shown for all data samples in the test set (All), the subset of data samples with holes (Hole) and without holes (No Hole). The number of network parameters is also shown (Param.). Finally, results from a human CAD expert on 100 test samples are shown.

our dataset with a known ground truth joint. We use 100 data samples picked randomly from a distribution excluding the potentially ambiguous sibling entities. We randomly rotate and translate each part and conduct the study using Fusion 360. We compare the joint axis created by the CAD expert with the ground truth. We find that the CAD expert results match the ground truth 80% of the time. This shows that determining how two isolated parts should be assembled is challenging for CAD experts without the valuable context provided by the object assembly. We provide additional details in Section A.2 of the supplementary material.

5.2. Joint Axis Prediction

Although there are no previous works that address the exact same setting as ours, we adapt several related methods to compare with our approach.

Point Cloud Baselines We adapt two point cloud based methods designed to predict a joint axis for part mobility. For each baseline we use a common architecture, based on a PointNet++ [49] encoder, and adapt the decoder strategy and loss functions from related work. **B-Dense** follows Li et al. [37] to densely regress a joint origin projection vector, projection distance, and joint direction for each point in the point cloud. **B-Discrete** follows Shape2Motion [58] and uses a hybrid of discrete classification and regression to predict the joint origin point and direction.

B-Rep Baselines We compare our method against several baseline methods that take B-Rep graphs as input. **B-Grid** follows UV-Net [25] and uses grid features (points, normals, trimming mask, and tangents) sampled on B-Rep faces and edges together with a CNN encoder. We use the same graph topology, prediction head, and loss as our network. **B-Heuristic** uses a rule-based approach that operates on B-Rep graphs and assigns a score to each B-Rep entity. Higher scores are assigned to entities that are similar, based on the entity type, area, and length information, and that match the training data distribution of entity type

	All CD↓	Hole CD↓	No Hole CD↓	Param. #↓
Ours + Search	0.0580	0.0570	0.0628	1.3M
Ours	0.0627	0.0624	0.0657	1.3M
B-Pose	0.0700	0.0693	0.0730	2.3M

Table 2. Joint pose prediction results using average chamfer distance (CD) where lower is better. We show results for all samples in the test set (All), and the subset of data samples with holes (Hole) and without holes (No Hole). The number of network parameters is also shown (Param.)

pairings. For cylinders and circles, the radius of the entity is also employed. A higher score is given to entity pairs where the radii match to within 5%. **B-Random** makes random predictions over all B-Rep entities and represents the lower bound for B-Rep performance.

Table 1 shows results for the joint axis prediction task on the test set. We report the accuracy of regression based approaches by considering a joint axis prediction to be a 'hit' if it is collinear within a distance and angular threshold of 5%. For classification based approaches we report the top-1 accuracy. We also report accuracy for the subset of data samples that have holes (Hole) and those that do not (No Hole). Recall that traditional algorithms are good at working with the special case of matching fasteners to holes. We observe that the performance gap between our approach and the next highest performing B-Heuristic approach is 8.14%, however this widens to 11.62% for the important No Hole subset where traditional algorithms are known to struggle. We find that the B-Rep based approaches outperform those based on point clouds while also using fewer parameters. Although point cloud approaches perform well with axis aligned parts from the same object class [58], our results show that real world data is significantly more challenging. Finally we note that our approach is within 0.5% of the performance of a human CAD expert. We provide additional details in Section A.2 of the supplementary material.

5.3. Joint Pose Prediction

For the joint pose prediction task we again adapt a baseline method from the literature to our setting. **B-Pose** follows Huang et al. [23] to regress a translation point and rotation quaternion using a combination of L2 and chamfer distance (CD) loss terms. Although a parametric joint is not created, B-Pose represents a common approach used with top-down assembly. We evaluate the performance of our method in two different configurations. **Ours** uses the joint axes derived from network predictions to align the two parts together without an offset, rotation, or flip. **Ours + Search** additionally performs joint pose search over the top 50 predictions to find suitable offset, rotation, and flip parameters.

Table 2 shows results for the joint pose prediction task. We record the minimum CD calculated between the ≥ 1



Figure 5. Qualitative comparison of joint pose prediction results comparing our method, with and without search, with the B-Pose baseline.

ground truth joints from the joint set and the predicted assembly. We then report the average CD across all samples in the test set. We find that using our network predictions alone (Ours) can better match the ground truth when compared with the B-Pose baseline. Introducing search (Ours + Search) can help resolve areas of overlap (Figure 5e) and in some cases resolve incorrect axis predictions (Figure 5b,g). It is important to note that the ground truth data only contains a finite set of discrete states (e.g. door open, door closed) rather than continuous states (e.g. door opening) that may also be valid. For example, our predictions for the belt buckle in Figure 5d do not match the ground truth state but appear plausible. As such, CD should be considered an approximate metric for comparing the relative performance of each method. We provide further qualitative results in Section A.2 of the supplementary material.

6. Discussion

Future Applications Our joint axis prediction network and search approach can serve as fundamental building blocks for a number of applications. One such application is the automated assembly of multiple parts in a design. As a preliminary demonstration we assemble a multi-part



Figure 6. Multi-part assembly demonstration. Parts are aligned sequentially from a given assembly sequence using our joint axis prediction network and pose search.

design given only the individual parts and the sequence of part pairs derived from our assembly dataset. We amend our search strategy to minimize the overlap volume between the new part and the partially assembled design at each assembly step and maximize the contact area between them using a similar cost function. Figure 6 shows an example sequence of parts that are assembled correctly in a bottomup fashion. We provide further details in Section A.3 of the supplementary material.

Limitations A bottom-up approach to assembly may be limited when scaling to large assemblies where global composition is important. Reliance on B-Rep CAD data is another limitation of the current work. Although data availability is improving [27, 30, 63], our method has not been tested beyond mechanical CAD data. Finally, our network does not leverage geometric loss terms that may help with avoiding undesirable overlap between parts and generalize to predicting other joint parameters.

7. Conclusion

Our long-term motivation is to enable assembly-aware design tools, capable of suggesting and automatically placing parts. Such a system could enable greater reuse of existing physical components in new designs and potentially reduce the cost and environmental impact associated with manufacturing and associated supply chains [29]. Understanding how parts are assembled is also critical for robotic assembly and disassembly. CAD-informed robotic disassembly systems may enhance our ability to reuse and recycle components [8, 35, 41, 45]. In this work we have begun the first steps to address these challenges by learning the bottom-up assembly of parametric CAD joints. Our results show the promise of learning-based methods to approach the performance of human CAD experts, and with the publication of our dataset we hope to further aid future research.

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