Abstract

Fact verification systems typically rely on neural network classifiers for veracity prediction, which lack explainability. This paper proposes ProoFVer, which uses a seq2seq model to generate natural logic-based inferences as proofs. These proofs consist of lexical mutations between spans in the claim and the evidence retrieved, each marked with a natural logic operator. Claim veracity is determined solely based on the sequence of these operators. Hence, these proofs are faithful explanations, and this makes ProoFVer faithful by construction. Currently, ProoFVer has the highest label accuracy and the second best score in the FEVER leaderboard. Furthermore, it improves by 13.21% points over the next best model on a dataset with counterfactual instances, demonstrating its robustness. As explanations, the proofs show better overlap with human rationales than attention-based highlights and the proofs help humans predict model decisions correctly more often than using the evidence directly.1

1 Introduction

Fact verification systems typically comprise an evidence retrieval model followed by a textual entailment classifier (Thorne et al., 2018b). Recent high-performing fact verification systems (Zhong et al., 2020; Ye et al., 2020) use neural models for textual entailment whose reasoning is opaque to humans despite advances in interpretability (Han et al., 2020). On the other hand, proof systems like NaturalLI (Angeli and Manning, 2014) provide transparency in their decision making for entailment tasks, by using explicit proofs in the form of natural logic. However, the accuracy of such approaches often does not match that of neural models (Abzianidze, 2017a).

Justifying decisions is central to fact verification (Uscinski and Butler, 2013). While models such as those developed for FEVER (Thorne et al., 2018b) typically substantiate their decisions by presenting the evidence as is, more recent proposals use the evidence to generate explanations. Here, models highlight salient parts of the evidence (Popat et al., 2018; Wu et al., 2020), generate summaries (Kotonya and Toni, 2020b; Atanasova et al., 2020), correct factual errors (Thorne and Vlachos, 2021b; Schuster et al., 2021), answer claim-related questions (Fan et al., 2020), or perform rule discovery (Ahmadi et al., 2019; Gad-Elrab et al., 2019). An explanation is faithful only if it reflects the information that is used for decision making (Lipton, 2018; Jacovi and Goldberg, 2020), which these systems do not guarantee. A possible exception here would be the rule discovery models, although, their performance often suffers due to limited knowledge base coverage and/or the noise in rule extraction from text (Kotonya and Toni, 2020a; Pezeshkpour et al., 2020). Faithful explanations are useful as mechanisms to dispute, debug, or advise (Jacovi and Goldberg, 2021), which may aid a news agency for advice, a user to dispute decisions, and a developer for model debugging in fact verification.

Keeping both accuracy and explainability in mind, we propose ProoFVer—Proof System for Fact Verification—which generates proofs or refutations of the claim given evidence as natural logic-based inference. ProoFVer follows the natural logic based theory of compositional entailment, originally proposed in NatLog (MacCartney and Manning, 2007). In the example of Figure 1 ProoFVer generates the proof shown in Figure 2, for a given claim and evidence. Here, at each step in the proof, a claim span is mutated with a span from the evidence. Each such mutation is marked with an entailment relation, by assigning a natural logic operator (NatOp; Angeli and

1Find our code and data at https://github.com/krishnamrith12/ProoFVer.
Figure 1: The proof generator in ProoFVer, generates the natural logic proofs using a seq2seq model. The natural logic operators from the proof are used as transitions in the DFA to determine the veracity of the claim. The states S, R, and N in the automaton denote the task labels SUPPORTS, REFUTES, and NOT ENOUGH INFO, respectively. The transitions in the automaton are the natural logic operators (NatOPs) defined in Table 1.

Figure 2: Proof steps for the input in Figure 1.

Manning, 2014). A step in the proof can be represented using a triple, consisting of the aligned spans in the mutation and its assigned NatOp. In the example, the mutations in the first and last triples occur with semantically equivalent spans, and hence are assigned with the equivalence NatOp (≡). However, the mutation in the second triple results in a contradiction, as ‘short story’ is replaced with ‘novel’ and an item cannot be both. Hence, the mutation is assigned the alternation NatOp (\(\not\equiv\)). The sequence of NatOPs from the proof become the transitions in the DFA shown in Figure 1, which in this case terminates at the ‘REFUTE (R)’ state, that is, the evidence refutes the claim.

Unlike other natural logic systems (Angeli et al., 2016; Feng et al., 2020), ProoFVer can form a proof by combining spans from multiple evidence sentences, by leveraging the entity mentions linking those sentences. The proof is generated by a seq2seq model trained using a heuristically annotated dataset, obtained by combining information from the publicly available FEVER dataset (Thorne et al., 2018a; Thorne and Vlachos, 2021b) with PPDB (Pavlick et al., 2015), Wordnet (Miller, 1995) and Wikidata (Vrandečić and Krötzsch, 2014). We heuristically generate the training data for the claims in three datasets, namely, FEVER, symmetric FEVER (Schuster et al., 2019), and FEVER 2.0 (Thorne et al., 2019).

ProoFVer is currently the highest scoring system on the FEVER leaderboard in terms of label accuracy and is the second-best system in terms of FEVER score. Additionally, ProoFVer has robustness and explainability as its key strengths. Its veracity predictions are solely determined using the generated proof. Hence by design, ProoFVer’s proofs, when used as explanations, are faithful by construction (Lei et al., 2016; Jain et al., 2020). Similarly, it demonstrates robustness to counterfactual instances from Symmetric FEVER and adversarial instances from FEVER 2.0. In particular, ProoFVer achieved 13.21% higher label accuracy than that of the next best model (Ye et al., 2020) for symmetric FEVER and similarly improves upon the previous best results (Schuster et al., 2021) on Adversarial FEVER.

To evaluate the robustness of fact verification systems against the impact of superfluous information from the retriever, we propose a new metric, Stability Error Rate (SER), which measures the proportion of instances where superfluous information changes the decision of the model. ProoFVer achieves a SER of 5.73%, compared with 9.36% of Stammbach (2021), where a lower SER is preferred. ProoFVer’s proofs as explanations, apart from being faithful, score high in their overlap with human rationales with a token overlap F1-Score of 93.28%, 5.67 percentage points more than attention-based highlights from Ye et al. (2020). Finally, humans, with no knowledge of natural logic, correctly predict ProoFVer’s decisions 81.67% of the times compared with 69.44% when using the retrieved evidence.
Natural Logic Proofs as Explanations

Natural logic operates directly on natural language (Angeli and Manning, 2014; Abzianidze, 2017b). Thus it is appealing for fact verification, as structured knowledge bases like Wikidata typically lag behind text-based encyclopedias such as Wikipedia in terms of coverage (Johnson, 2020). Furthermore, it obviates the need to translate claims and evidence into meaning representations such as lambda calculus (Zettlemoyer and Collins, 2005). While such representations may be more expressive, they require the development of semantic parsers, introducing another source of potential errors in the verification process.

Natural Logic has been previously used in several information extraction and NLU tasks such as Natural Language Inference (NLI, Abzianidze, 2017a; Feng et al., 2020), question answering (Angeli et al., 2016), and open information extraction (Angeli, 2016, Chapter 5). NatLog (MacCartney and Manning, 2007), building on earlier theoretical work on natural logic and monotonicity calculus (Van Benthem, 1986; Valencia, 1991), uses natural logic for textual inference.

NaturalLI (Angeli and Manning, 2014) extended NatLog by adopting the formal semantics of Icard III and Moss (2014), and it is a proof system formulated for the NLI task. It determines the entailment of a hypothesis by searching over a database of premises. The proofs are in the form of a natural logic based logical inference, which results in a sequence of mutations between a premise and a hypothesis. Each mutation is marked with a natural logic relation, and is realized as a lexical substitution, forming a step in the inference. Each mutation results in a new sentence, and the natural logic relation assigned to it identifies the type of entailment that holds between the sentences before and after the mutation. NaturalLI adopts a set of seven natural logic operators, as shown in Table 1. The operators were originally proposed in NatLog (MacCartney, 2009, p. 79). We henceforth refer to these operators as NatOps.

To determine whether a hypothesis is entailed by a premise, NaturalLI uses a deterministic finite state automaton (DFA). Here, each state is an entailment label, and the transitions are the NatOps (Figure 1). The sequence of NatOps in the inference is used to traverse the DFA, and the state where it terminates decides the label of the hypothesis-premise pair. The decision making process relies solely on the steps in the logical inference, and thus form faithful explanations.

Other proof systems that apply mutations between text sequences have been previously explored. Stern et al. (2012) explored how to transform a premise into a hypothesis using mutations, however their approach was limited to two-way entailment instead of three-way that is handled by NaturalLI. Similar proof systems have used mutations in the form of tree-edit operations (Mehdad, 2009), transformations over syntactic parses (Heilman and Smith, 2010; Harmeling, 2009), knowledge-based transformations in the form of lexical mutations, entailment rules, rewrite rules, or their combinations (Bar-Haim et al. 2007; Szpektor et al., 2004).

3 ProofVer

ProofVer uses a seq2seq generator that generates a proof in the form of natural-logic based logical inference, which becomes the input to a DFA for predicting the veracity of the claim. We elaborate on the proof generation process in Section 3.1, and on the veracity prediction in Section 3.2.

3.1 Proof Generation

The proof generator, as shown in Figures 1 and 3, takes as input a claim along with one or more retrieved evidence sentences. It generates the steps of the proof as a sequence of triples, each consisting of a span from the claim, a span from the evidence and a NatOp. The claim span being substituted and the evidence span replacing it form a mutation, and each mutation is assigned a NatOp. In a proof, we start with the claim, and the mutations are iteratively applied from left to right. Figure 1 shows a proof containing a sequence

| NatOp | Name | Definition |
|-------|------|------------|
| ∨: Alternation | $x \cap y = \emptyset \land x \cup y \neq U$ |
| ⊇: Cover | $x \cap y \neq \emptyset \land x \cup y = U$ |
| ≡: Equivalence | $x = y$ |
| ⊑: Forward Entailment | $x \subset y$ |
| ⌠: Negation | $x \cap y = \emptyset \land x \cup y = U$ |
| ⌡: Reverse Entailment | $x \supset y$ |
| #: Independence | All other cases |

Table 1: Natural logic relations (NatOps) and their set theoretic definitions.
of three triples. The corresponding mutated statements at each step of the proof, along with the assigned NatOps, are shown in Figure 2.

We use a seq2seq model following an autoregressive formulation for the proof generation. In the proof, successive spans of the claim form part of the successive triples. However, the corresponding evidence spans in the successive triples need not follow any order. As shown in Figure 3, the evidence spans may come from multiple sentences, and may not all end up being used. Finally, the NatOps, as shown in Table 1, are represented using a predetermined set of tokens.

To obtain valid proofs during prediction, we need to lexically constrain the inference process by switching between three different search spaces depending on which element of the triple is being predicted. To achieve this, we use dynamically constrained markup decoding (De Cao et al., 2021), a modified form of lexically constrained decoding (Post and Vilar, 2018). This decoding uses markups to switch between the search spaces, and we use the delimiters ‘{’, ‘}’, ‘[’, and ‘]’ as the markups. Using these markups, we constrain the tokens predicted between a ‘{‘ and ‘}’ to be from the claim, between a ‘[‘ and ‘]’ to be from the evidence, and the token after ‘]’ to be a NatOp token. The prediction of a triple begins with predicting a ‘{‘, and it proceeds by generating a claim span where the tokens are monotonically copied from the claim in the input, until a ‘}’ is predicted. The prediction then continues by generating a ‘[‘, which initiates the evidence span prediction in the triple. The evidence span can begin with any word from the evidence, and is then expanded by predicting subsequent tokens, until ‘]’ is predicted. Finally, the NatOp token is predicted. In the next triple, copying resumes from the next token in the claim. All triples until the one with the last token in the claim are generated in this manner.

3.2 Veracity Prediction

The DFA shown in Figure 1 uses the sequence of NatOps predicted by the proof generator as transitions to arrive at the outcome. Figure 2 shows the corresponding sequence of transitions for the claim and evidence from Figure 1. Based on this, the DFA in Figure 1 determines that the evidence refutes the claim, that is, it terminates in state $R$. NaturalLI (Angeli and Manning, 2014) designed the DFA for the three classes in the NLI classification task, namely, entail, contradict, and neutral. Here, we replace them with SUPPORT (S), REFUTE (R), and NOT ENOUGH INFO (N), respectively for fact verification. Angeli and Manning (2014) chose not to distinguish between negation ($\nabla$) and alternation ($\forall$) relations for NLI, and assign $\forall$ for both. However, there is a clear distinction between cases where each of these NatOps is applicable in fact verification, and thus we treat them as different NatOps. For instance, in the second mutation for the claim in Figure 1, an evidence span “is not a short story” would be assigned negation ($\nabla$), and not the currently assigned alternation ($\forall$) for the mutation with the evidence span “is a novel”. However, we follow Angeli and Manning (2014) in not using the cover ($\nabla$) NatOp.

In rare occasions where this NatOp would be applicable, say in a mutation with the spans “not a novel” and “fiction”, we currently assign the independence NatOp ($\forall$).

4 Generating Proofs for Training

Training datasets for evidence-based fact verification consist of instances containing a claim, a label indicating its veracity, and the evidence, typically a set of sentences (Thorne et al., 2018a; Hanselowski et al., 2019; Wadden et al., 2020). However, we need sequences of triples to train the proof generator of Section 3.1. Manually
annotating them would be laborious; thus, we heuristically generate them from existing resources. As shown in Figure 4, we perform a two-step annotation process: chunking and alignment, followed by the NatOp assignment.

4.1 Chunking and Alignment

Chunking the claim into spans is conducted using the chunker of Akbik et al. (2019), and any span that does not contain any content words is merged with its subsequent span. Next, as shown in Figure 4, a word aligner (Jalili Sabet et al., 2020) aligns each evidence sentence in the input separately with the claim. For each claim span, each evidence sentence provides an aligned span by grouping together words that are aligned to it, including any words in between to ensure contiguity. However, if the aggregated similarity score from the aligner for a given pair of claim and evidence spans falls below an empirically set threshold, then it is ignored and instead the claim span is aligned with the string “DEL”. In Figure 4, “DEL” appears once in each of the evidence sentences.

Next, we convert the alignments into a sequence of mutations, which requires no additional effort in instances with only one evidence sentence. However, a claim span may have multiple evidence spans aligned with it in cases with multiple evidence sentences, as shown in Figure 4. Here, for a claim span, we generally select the evidence span with the highest cosine similarity with it. Such spans are marked with solid red borders in Figure 4. Further, we assume that the evidence sentences are linked via entity mentions, such as “Spanish Empire” the only hyperlinked mention (from Evidence-1 to 2) in Figure 3. These hyperlinked mentions must always be added as a mutation, as they provide the context for switching the source of the evidence from one sentence to another. In Figure 3, “Spanish Empire” is not selected as an alignment based on the similarity scores with the claim spans. Hence, it is inserted as the third mutation, at the juncture at which the switch from Evidence-1 to 2 happens. It is aligned with the string ‘‘Ins’’ in the place of a claim span. Use of hyperlink structure in Wikipedia or performing entity linking to establish hyperlinked mentions, similar to our approach here, has been previously explored in multi-hop open domain question answering (Asai et al., 2020; Nie et al., 2019). Mutations with a “DEL” instead of an evidence span, and an “INS” instead of a claim span, are treated as deletions and insertions of claim and evidence spans, respectively.

4.2 NatOp Assignment

As shown in Figure 4, the NatOp assignment step produces a sequence of NatOps, one for each mutation. Here, the search space becomes exponentially large (i.e., $6^n$ possible NatOp sequences for $n$ mutations). First, we assign NatOps to individual mutations relying on hand-crafted rules and external resources, without considering the other mutations in the sequence (§ 4.2.1). With this partially filled NatOp sequence, we perform two filtering steps to further reduce the search space. We describe these steps below: one using veracity label information from training data in FEVER (Thorne et al., 2018a) and another using some additional manual annotation information from annotation logs of FEVER (§ 4.2.2).
4.2.1 Initial Assignment

The initial assignment of NatOps considers each mutation in the sequence in isolation. Here, mutations that fully match lexically are assigned with the equivalence NatOp ($\equiv$), like the mutations 1, 4, and 5 in Figure 4. Similarly, mutations where the claim or evidence span has an extra negation word but lexically match otherwise, are assigned the negation NatOp ($\not\equiv$). Further, insertions and deletions, that is, mutations with Ins and Del, respectively (§4.1), containing negation words are also assigned the negation NatOp. To obtain these words, we identify a set of common negation words from the list of stop words in Honnibal et al. (2020), and combine them with the list of negative sentiment polarity words from Hu and Liu (2004). Remaining cases of insertions (deletions) are treated as making the existing claim more specific (general), and hence assigned the forward (reverse) entailment NatOp, like mutation 3 in Figure 4. Furthermore, as every paraphrase pair present in Paraphrase Database (PPDB Ganitkevitch et al., 2013; Pavlick et al., 2015) is marked with an entailment relation, we identify mutations which are present in it as paraphrases and assign the corresponding NatOp.

In several cases, the NatOp information need not be readily available at the span level. Here, we retain the word-level alignments from the aligner and perform lexical level NatOp assignment with the help of Wordnet (Miller, 1995) and Wikidata (Vrandečić and Krötzsch, 2014). We follow MacCartney (2009, Chapter 6) for NatOp assignment of open-class terms using Wordnet. Additionally, we define rules to assign a NatOp for named entities using Wikidata. Here, aliases of an entity are marked with an equivalence NatOp ($\equiv$), as shown in third triple in Figure 1. Further, we manually assign NatOps to the 500 most frequently occurring Wikidata relations in the aligned training data. For instance, as shown in Figure 5, the entities ‘The Trial’ and ‘novel’ have the relation ‘genre’. A claim span containing ‘The Trial’, when substituted with an evidence span containing ‘novel’, would result in a generalisation of the claim, and hence will be assigned the reverse entailment NatOp ($\not\equiv$). A substitution in the reverse direction would be assigned a forward entailment NatOp ($\equiv$), indicating specialization.

The KB relations we annotated occur between the entities linked in Wikidata, and they do not capture hierarchical multihop relations between the entities in the KB. We create such a hierarchy by combining the “instance of”, “part of”, and “subclass of” relations in Wikidata. Thus, a pair of entities connected via a directed path of length $k \leq 3$, such as “Work of art” and “Rashomon” in Figure 5, is considered to have a parent-child relation, and assigned the forward or reverse entailment NatOp, depending on which span appears in the claim and the evidence. Similarly, two entities (e.g., “Rashomon” and “Inception”) are considered to be siblings if they have a common parent, and are assigned the alternation NatOp (§). However, two connected entities that do not satisfy the aforementioned distance criterion (e.g., “novel” and “Rashomon”) are assigned with the independence NatOp ($\not\equiv$), signifying that they are unrelated.

4.2.2 Filtering the Search Space

While in Section 4.2.1 we assigned a NatOp to each mutation in isolation, there can still be unfilled NatOps. For instance, the unfilled NatOp in the second mutation of Figure 4 leads to six possible NatOp sequences as candidates, one per available NatOp. Recall that these NatOp sequences act as a transition sequence in the DFA (§ 3.2). Thus we make use of the partially filled NatOp sequence and the veracity label from the training data to filter out NatOp sequences that do not terminate at the same state as the veracity label according to the DFA. The instance in Figure 4 has the SUPPORT label, and among the six possible candidate sequences only two terminate in this label. Hence, we retain those two sequences.

For the final filtering step we use the additional manual annotation that was produced during the construction of the claims in FEVER. There, the annotators constructed each claim by manipulating a factoid extracted from Wikipedia using...
Table 2: NatOp assignment based on transformations and veracity label information.

| Transformation                          | S | R | N |
|----------------------------------------|---|---|---|
| substitute with similar info.          | □ | □ | □ |
| substitute with dissimilar info.       | □ | □ | □ |
| paraphrasing                           | ★ | ★ | # |
| negation                               | ★ | ★ | # |
| transform to specific                  | ❌ | ❌ | ❌ |
| transform to general                   | ❌ | ❌ | ❌ |

one of the six transformations listed in Table 2. Our proofs can be viewed as an attempt at reconstructing the factoid from a claim in multiple mutations, whereas these transformations can be considered claim-level mutations that transition directly from the last step (reconstructed factoid) in the proof to the first step (claim). This factoid is treated as the corrected claim in Thorne and Vlachos (2021b), who released this annotation. For each veracity label we define the mapping of each transformation to a NatOp, as described in Table 2. The assumption is that if a transformation has resulted in a particular veracity label, then the corresponding NatOp is likely to occur in the proof. To identify the mutation to assign it, we obtain the text portions in the claim manipulated by the annotators to construct it, by comparing the claim and the original Wikipedia factoid. In the example of Figure 4, this transformed text span happens to be part of the second mutation, and as per Table 2 forward entailment is the corresponding NatOp given the veracity label, resulting in the selection of the first NatOp sequence. In rare occasions (2.55% claims in FEVER), we manually performed NatOp assignment, as the filtering steps led to zero candidates in those cases. As the heuristic annotation requires manual effort, we explore how it can be obtained using a supervised classifier (see §5.5).

5 Experimental Methodology

5.1 Data

ProoFVer is trained using heuristically annotated proofs (§4) obtained from FEVER (Thorne et al., 2018a), which has a train/test/development split of 145,449/19,998/19,998 claims. Further, the heuristic proof annotation involves the use of additional information from the manual annotation logs of FEVER, recently released by Thorne and Vlachos (2021b). Finally, claims with the label NOT ENOUGH INFO (NEI) require retrieved evidence for obtaining their proofs for training, as no ground truth evidence exists for such cases. Here, we use the same retriever that would be used during the prediction time as well.

In addition to FEVER, we train and evaluate ProoFVer on two other related datasets. First, we use Symmetric FEVER (Schuster et al., 2019), a dataset designed to assess the robustness of fact verification systems against the claim-only bias present in FEVER. The dataset consists of 1,420 counterfactual instances, split into development and test sets of 708 and 712 instances, respectively. Here, we heuristically generate the ground truth proofs for the dataset’s development data and use it to fine tune ProoFVer, before evaluating it on the dataset’s test data. Similarly, we also evaluate ProoFVer on the FEVER 2.0 adversarial examples (Thorne et al., 2019). Specifically, we use the same evaluation subset of 766 claims that was used by Schuster et al. (2021). To fine-tune ProoFVer on this dataset, we generate the ground truth proofs for 2,100 additional adversarial claims, separate from the evaluation set, which were curated by the organisers and participants of the FEVER 2.0 shared task.

Finally, we also use the manual annotation logs of FEVER (Thorne and Vlachos, 2021b) to obtain rationales for claims in the development data. In particular, we obtain the rationale for a claim by extracting from its corresponding Wikipedia factoid the words which were removed by the annotators during its creation. If these words are part of an evidence sentence, then they become the rationale for veracity label of the claim given the evidence. Further, we require that the words extracted as rationale form a contiguous phrase. We identified 300 claims that satisfy all these criteria.

5.2 Evaluation Metrics

The evaluation metrics for FEVER are label accuracy (LA, i.e., veracity accuracy) and FEVER Score (Thorne et al., 2018b), which rewards only those predictions which are accompanied by at least one correct set of evidence sentences. We report mean LA and standard deviation for experiments with Symmetric FEVER, where we use its development data for training and train with five random initialisations due to its limited size. We further introduce a new evaluation metric, to
assess model robustness, called Stability Error Rate (SER). Neural models, especially with a retriever component, have shown to be vulnerable to model overstability (Jia and Liang, 2017). Overstability is the inability of a model to distinguish superfluous information that merely has lexical similarity with the input, from the information truly relevant to arrive at the correct decision. In the context of fact verification, it is expected that an ideal model should always predict NOT ENOUGH INFO, whenever it lacks sufficient evidence to make a decision otherwise. Further, it should arrive at a REFUTE or SUPPORT decision only when the model possesses sufficient evidence to do so, and any additional evidence should not alter its decision. To assess the model overstability in fact verification, we define SER as the percentage of claims where additional evidence alters the SUPPORT or REFUTE decision of a model.

5.3 Baseline Systems

**KGAT (Liu et al., 2020)** uses a graph attention network, where each evidence sentence, concatenated with the claim, forms a node in the graph. We use their best configuration, where the node representations are initialized using RoBERTA (Large). The relative importance of each node is computed with node kernels, and information propagation is performed using edge kernels. They also propose a new evidence sentence retriever, a BERT model trained with a pairwise ranking loss, though they rely on past work for document retrieval (Hanselowski et al., 2018).

**CorefBERT (Ye et al., 2020)** follows KGAT and differs only in terms of the LM used for the node initialisation. Here, they further pretrain the LM on a task that involves prediction of referents of a masked mention to capture co-referential relations in context. We use CorefRoBERTA, their best-performing configuration.

**DominikS (Stammbach, 2021)** focuses primarily on sentence-level evidence retrieval, scoring individual tokens from a given Wikipedia document, and then selecting the highest scoring sentences by averaging token scores. It uses a fine-tuned document level BigBird model (Zaheer et al., 2020) for this purpose. For claim verification it uses a DeBERTa (He et al., 2021) based classifier.

5.4 ProoFVer: Implementation Details

We follow most previous works on FEVER which model the task in three steps, namely, document retrieval, retrieval of evidence sentences from them, and finally veracity prediction based on the evidence. ProoFVer’s novelty lies in the proof generation in the third step. Hence, for better comparability, we follow two popular, well-performing retrieval approaches, Liu et al. (2020) and Stammbach (2021). Liu et al.’s (2020) sentence retriever, also used in Ye et al. (2020), is a sentence level pairwise ranking model, whereas that of Stammbach (2021) is a document level token score aggregation model. ProoFVer’s configuration which uses the former is our default configuration, referred to as ProoFVer, and the configuration using the latter will henceforth be referred to as ProoFVer-SB. We retrieve five sentences for each claim as required in the FEVER evaluation.

For the proof generator, we use the pretrained BART (Large) model (Lewis et al., 2020) and fine-tune it using the heuristically annotated data from Section 4. During prediction, the search spaces for the claim and evidence are populated using two separate tries. We add all possible subsequences of the claim and evidence, each with one to seven words, into the respective tries. The default configuration takes the concatenation of a claim and all the retrieved evidence together as a single input, separated by a delimiter.

We consider three additional configurations which differ in the way the retrieved evidence is handled. In ProoFVer-MV, a claim is concatenated with one evidence sentence at a time; this produces five proofs and five decisions per claim, and the final label is decided based on majority voting (MV). Both ProoFVer-A and -AR are designed to restrict the proof generator’s flexibility in inferring the textual spans in the mutations, and thus assess the gains obtained by allowing it in ProoFVer. ProoFVer-A (aligned) considers during prediction only the subsequences from each evidence sentence aligned with the claim using word-level alignment, which are then concatenated with the claim as its input during training and prediction. Thus, the evidence search space becomes narrower, as the unaligned portions in the evidence are not considered. ProoFVer-AR (aligned-restricted) further restricts the search space of both the claim and evidence, by predetermining the number of
mutations, the claim spans in these mutations and five candidate evidence spans for each mutation (one per evidence sentence). It obtains this information using the chunker and aligner used in the heuristic annotation (§4).

5.5 Heuristic Annotation Using Kepler

To reduce the reliance on manual annotation from Thorne and Vlachos (2021b) during the annotation in Section 4, we experiment with replacing the ground truth transformations with predicted ones using a classifier. We use KEPLER (Wang et al., 2021), a RoBERTA-based pretrained LM enhanced with KB relations and entity pairs from WikiData for the classification. KEPLER covers 97.5% of the entities present in FEVER. We first train it with the FEVER training dataset for the fact verification task. Then we fine-tune it for the six-class classification task of predicting the transformations, given a claim, evidence sentence and veracity label as input from the FEVER training data. We train it with varying training dataset sizes ranging from 1.24% (1,800; 300 per class) to 41.24% (60,000; 10,000 per class) of the FEVER training data. We consider two configurations: ProoFVer-K, which uses gold data to identify the transformed span for applying the predicted transformation, and ProoFVer-K-NoS, which instead only ensures that the predicted transformation occurs at least once in the final NatOp sequence.

6 Results

6.1 Fact Verification

Table 3 reports the fact verification results for ProoFVer and the baselines. Overall, ProoFVer-SB, our configuration using Stammbach’s (2021) retriever, is the best performing model in our experiments. ProoFVer-SB, which outperforms Stammbach (2021) itself, is currently the highest scoring model in terms of label accuracy in the FEVER leaderboard. It also is the second best model in terms of FEVER Score, second only to the currently unpublished model titled “mitchell.dehaven”, in the leaderboard.

ProoFVer, our default configuration using the retriever from Liu et al. (2020), differs from ProoFVer-SB only in terms of the retriever they use. ProoFVer is the best performing model among all the baselines and other ProoFVer configurations (-MV, -A, and -AR) that use Liu et al.’s (2020) retriever. As compared to ProoFVer-MV, ProoFVer’s gains come primarily from its ability to handle multiple evidence sentences together, as opposed to handling each separately and then aggregating the predictions. 9.8% (1,960) of the claims in the FEVER development set require multiple evidence sentences for verification. While ProoFVer-MV predicts 60.1% of these instances correctly, ProoFVer correctly predicts 67.45% of these. Further, around 80.73% (of 18,038) of the single evidence instances are correctly predicted by ProoFVer-MV, in comparison to 81.62% instances for ProoFVer. Allowing the proof generator to infer the mutations dynamically, instead of having them predefined, benefits the overall performance of the model. The increasingly restricted variants with narrower search spaces (i.e., ProoFVer-A and ProoFVer-AR) lead to decreasing performances as shown in Table 3. ProoFVer-AR, the most restricted version, performs worse than all the other models.

| System              | Dev LA | Dev Fever Score | Test LA | Test Fever Score |
|---------------------|--------|-----------------|---------|------------------|
| ProoFVer            | 80.23  | 78.17           | 79.25   | 74.37            |
| ProoFVer-MV         | 78.71  | 74.62           | 74.18   | 70.09            |
| ProoFVer-A          | 79.83  | 76.33           | 77.16   | 72.47            |
| ProoFVer-AR         | 77.42  | 75.27           | –       | –                |
| KGAT                | 78.29  | 76.11           | 74.07   | 70.38            |
| CorefBERT           | 79.12  | 77.46           | 75.96   | 72.30            |

Table 3: Fact verification results on FEVER.

| KEPLER               | ProoFVer                               |
|----------------------|----------------------------------------|
| Training Data Size   | Classifier Accuracy                   |
| 1,800                | 69.07                                  |
| 6,000                | 74.02                                  |
| 18,000               | 79.67                                  |
| 30,000               | 80.61                                  |
| 45,000               | 82.76                                  |
| 60,000               | 84.85                                  |
| -K-NoS (LA)          | 64.65                                  |
| -K (LA)              | 66.73                                  |
| -                   | 68.86                                  |
| -                   | 72.41                                  |
| -                   | 74.25                                  |
| -                   | 76.23                                  |
| -                   | 75.39                                  |
| -                   | 77.76                                  |
| -                   | 77.62                                  |
| -                   | 78.84                                  |
| -                   | 78.61                                  |
| -                   | 79.67                                  |

Table 4: LA of ProoFVer-K and -NoS using predictions from KEPLER. Training data size used for KEPLER and its classifier accuracy is also provided.
Table 5: Label accuracy of models on FEVER-development (DEV) and Symmetric FEVER with and without fine tuning. All results marked with * and # are statistically significant (unpaired t-test) with \( p < 0.05 \) against their FT and Original variants respectively. FEVER-DEV predictions are using gold standard evidence.

**Impact of Additional Manual Annotation**
Because the final filtering step in NatOp assignment (§4.2.2) requires additional manual annotation, we experimented with a proof set obtained without this step. Here, we arbitrarily select a NatOp sequence from the candidates remaining after the veracity label based filtering. The latter reduced the search space to just two possible NatOp sequences in 93.59% of the claims. However, training ProoFVer with these proofs resulted in a LA of 58.29% on the FEVER development set. In comparison, ProoFVer-K-NoS achieves a LA of 64.65%, even when using predictions from a KEPLER configuration trained on as little as 1,800 instances. Table 4 shows the LA for ProoFVer-K-NoS and ProoFVer-K when using KEPLER predictions, with varying training data sizes for KEPLER; the largest KEPLER configuration is trained on only 41.24% of claims in FEVER. Using this amount of training data, ProoFVer-K and ProoFVer-K-NoS achieve a LA of 79.67% and 78.61%, respectively. Here, ProoFVer-K outperforms all the baseline models, including CorefBERT, which also uses additional annotation for pretraining.

**6.2 Robustness**

**Symmetric FEVER** As shown in Table 5, ProoFVer shows better robustness with a mean accuracy of 81.70% on the Symmetric FEVER test dataset, an improvement of 13.21% over CorefBERT, the next best model. All models improve their accuracy and are comparable on the test set when we fine-tune them on its development set. However, this results in more than 9% reduction on the original FEVER-DEV data for both the classifier based models, KGAT and CorefBERT. This catastrophic forgetting (French, 1999) occurs primarily due to the shift in label distribution during fine-tuning, as Symmetric FEVER contains only claims with SUPPORT and REFUTE labels. ProoFVer accuracy drops by only less than 3%, as it is trained with a seq2seq objective. To mitigate the effect of catastrophic forgetting, we apply L2 regularization (Thorne and Vlachos, 2021a), which improves all models on the FEVER development set. Nevertheless, ProoFVer has the highest accuracy on both FEVER and Symmetric FEVER among the competing models after regularization.

**Generalizing to FEVER 2.0** ProoFVer when evaluated on FEVER 2.0 adversarial data, reports a LA of 82.79%, outperforming the previously best reported LA of 82.51% by Schuster et al. (2021). ProoFVer, after training on FEVER, is further fine-tuned (with L2 regularization) on heuristically generated proofs from the data contributed by the participants of the FEVER 2.0 shared task (disjoint from the evaluation set), and the proofs generated from the FEVER Symmetric data. On the other hand, Schuster et al. (2021) was trained on the VitaminC training data. When they further fine-tune their default model with FEVER, their performance drops to 80.94%.

**Stability Error Rate (SER):** SER quantifies the rate of instances where a system alters its decision due additional evidence in the input, passed on by the retriever component. KGAT, CorefBERT, and DominikS have a SER of 12.35%, 10.27%, and 9.36% respectively. ProoFVer has an SER of only 6.21%, which is further reduced to 5.73% for ProoFVer-SB. The SER results confirm that the baselines change their predictions from SUPPORT or REFUTE after providing them with additional information more often than ProoFVer.
6.3 ProoFVer Proofs as Explanations

6.3.1 Rationale Extraction
Rationales extracted based on attention are often used as means to highlight the reasoning involved in the decision making process of various models (DeYoung et al., 2020). For this evaluation, we compare using token-level F-score of the predicted rationales with human-provided rationales for 300 claims from the FEVER development data, as elaborated in Section 5.1. We ensure that all the systems are provided with the same set of evidence sentences, and consider only those words from the evidence as rationales that do not occur in the claim. For ProoFVer, we additionally remove evidence spans which are part of mutations with an equivalence NatOp. For KGAT and CorefBERT, we obtain the rationales by sorting the eligible words in descending order of their attention scores, and for each instance we find the set of words with the highest token overlap F-score with the rationale. Here, we consider the words in the top 1% of attention scores, and also those ranging from 5% to 50% of the words in step sizes of 5%. We find that ProoFVer achieves a token level F-score of 93.28, compared to 87.61 and 86.42, the best F-Scores for CorefBERT and KGAT.

Figure 6 shows the rationales for 3 instances extracted from ProoFVer, one for each label. All the three proofs result in correct decisions. While for the first two claims there is a perfect overlap with the human rationale, the third claim in Figure 6 has some extraneous information in the predicted proof.

6.3.2 Human Evaluation
We use forward prediction (Doshi-Velez and Kim, 2017) here, where humans are asked to predict the system output based on the explanations. For assessing ProoFVer, we provide the claim, the proof as the explanation, and those evidence sentences from which the evidence spans in the proof were extracted. Since we are interested in evaluating the applicability of our proofs as natural language explanations, we ensure that none of our subjects are aware of the deterministic nature of determining the label from natural logic proofs. Moreover, we replaced the NatOps in the proof with plain English phrases for better comprehension by the subjects, as shown in Table 6. As the baseline setup for comparison, we provide the claim with all five retrieved evidence sentences. We form a set of 24 different claims, 12 each from ProoFVer and baseline, and 3 individual subjects independently annotate the same set. Finally, we altogether obtain annotations for 5 sets, resulting in 60 claims, 120 explanations, and a total of 360 annotations from 15 subjects. Although 19 subjects volunteered, one of them annotated a set that did not receive any other annotations. In another set, two of them had prior knowledge in natural logic, leading to disqualification of these 3 annotations from the set.

| NatOps | Human Rationale |
|--------|-----------------|
| ≡ | Equivalent Spans |
| % | Evidence span contradicts the claim span |
| ⊒ | Claim span follows from evidence span |
| ⊢ | (Insert) New information from evidence |
| ⊥ | Incomplete Evidence |
| ⊥ | Evidence span refutes claim span |
| ⊥ | Claim span negated (Deletion) |
| # | Unrelated claim span and evidence span |
| ⊥ | No related evidence found (Deletion) |

Table 6: NatOps and the corresponding paraphrases.
claims, ProoFVer, CorefBERT, and KGAT predicted the same labels, though not necessarily the correct ones (the subjects were not aware of this). All the subjects were pursuing a PhD or postdocs in fields related to computer science and computational linguistics, or industry researchers/data scientists.

With ProoFVer’s proofs, subjects are able to predict the model decisions correctly in 81.67% of the cases as against 69.44% of the cases with only the evidence. In both setups, subjects were often confused on instances with a NOT ENOUGH INFO label, and the forward predictions were comparable, with 66.67% (ProoFVer) and 65% (baseline). In many such cases, subjects subconsciously filled in their own world knowledge that is not found in the evidence to arrive at a SUPPORT or REFUTE decision. Further, for instances with both REFUTE and SUPPORT labels, subjects correctly predicted ProoFVer’s decisions 86.67% and 91.67% times, respectively, against only 70% and 73.33% for the baseline. The inter-annotator agreement for ProoFVer’s explanations is 0.7074 in Fleiss $\kappa$ (Fleiss, 1971), and 0.6612 for the baseline.

7 Limitations

Figure 7 shows three instances of incorrect proof generation from ProoFVer. The claim and evidence spans are enclosed within ‘{ }’ and ‘[ ]’, respectively, with numbered superscripts showing the correspondence between the spans.

Figure 7: Cases of incorrect proof generation from ProoFVer. The claim and evidence spans are enclosed within ‘{ }’ and ‘[ ]’, respectively, with numbered superscripts showing the correspondence between the spans.

...
decoding. Nevertheless, in the future we plan to experiment with alternative decoding approaches, including some of the recent developments in non-autoregressive conditional language models (Xu and Carpuat, 2021) and transformer-based proof generators (Saha et al., 2021).

8 Conclusion

We presented ProoFVer, a natural logic-based proof system for fact verification. Currently, we report the best results in terms of label accuracy, and the second best results in FEVER Score in the FEVER leaderboard. Moreover, ProoFVer is more robust in handling superfluous information from the retriever, and handling counterfactual instances. Finally, ProoFVer’s proofs are faithful explanations by construction, and improve the understanding of the decision making process of the models by humans.

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