

Previous datasets for evaluating PDF content extraction rely on machine-generated labels of imperfect quality, and comprise papers from a limited range of scientific disciplines. To better evaluate our proposed methods, we design a new benchmark suite, Semantic Scholar Visual Layout-enhanced Scientific Text Understanding Evaluation (**S2-VLUE**). The benchmark extends two existing resources (Tkaczyk et al., 2015; Li et al., 2020) and introduces a newly curated dataset, S2-VL, which contains high-quality human annotations for papers across 19 disciplines.

Our contributions are as follows:

1. We introduce a new strategy for PDF content extraction that uses VILA structures to inject layout information into language models, and show that this improves accuracy *without* the expensive pretraining required by existing methods, and generalizes to different language models.
2. We design two models that incorporate VILA features differently. The I-VILA model injects layout indicator tokens into the input texts and improves prediction accuracy (up to +1.9% Macro F1) and consistency compared to the previous layout-augmented language model LayoutLM (Xu et al., 2020). The H-VILA model performs group-level predictions and can reduce model inference time by 47% with less than 0.8% loss in Macro F1.
3. We construct a unified benchmark suite S2-VLUE which enhances existing datasets with VILA structures, and introduce a novel dataset S2-VL that addresses gaps in existing resources. S2-VL contains hand-annotated gold labels for 15 token categories on papers spanning 19 disciplines.

The benchmark datasets, modeling code, and trained weights are available at <https://github.com/allenai/VILA>.

2 Related Work

2.1 Structured Content Extraction for Scientific Documents

Prior work on structured content extraction for scientific documents usually relies on textual or visual features. Text-based methods like ScienceParse (Ammar et al., 2018), GROBID (GRO, 2008–2021) or Corpus Conversion Service (Staar

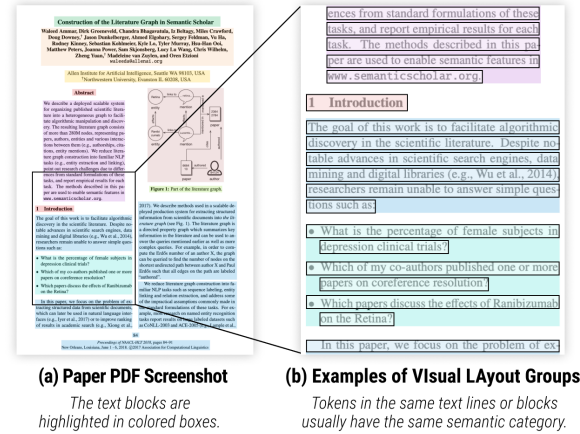


Figure 1: (a) Real-world scientific documents often have intricate layout structures, so analyzing only flattened raw text forfeits valuable information, yielding sub-optimal results. (b) The complex structures can be broken down into groups (text blocks or lines) that are composed of tokens with the same semantic category.

et al., 2018) combine PDF-to-text parsing engines like CERMINE (Tkaczyk et al., 2015) or pdfalto,¹ which output a sequence of tokens extracted from a PDF, with machine learning models like RNN (Hochreiter and Schmidhuber, 1997), CRF (Lafferty et al., 2001), or Random Forest (Breiman, 2001) trained to classify the token categories of the sequence. Though these models are practical and fairly efficient, they fall short in prediction accuracy or generalize poorly to out-of-domain documents. Vision-based Approaches (Zhong et al., 2019; He et al., 2017; Siegel et al., 2018), on the other hand, treat the parsing task as an image object detection problem: given document images, the models predict rectangular bounding boxes, segmenting the page into individual components of different categories. These models excel at capturing complex visual layout structures like figures or tables, but because they operate only on visual signals without textual information, they cannot accurately predict fine-grained semantic categories like title, author, or abstract, which are of central importance for parsing scientific documents.

2.2 Layout-aware Language Models

Recent methods on layout-aware language models improve prediction accuracy by jointly modeling documents’ textual and visual signals. LayoutLM (Xu et al., 2020) learns a set of novel posi-

¹<https://github.com/kermitt2/pdfalto> (last accessed Jan. 1, 2022).

