# arXiVeri: Automatic table verification with GPT

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**NEURAL INFORMATION** PROCESSING SYSTEMS

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# Automatic table verification Task

Researchers often manually transfer performance metrics between academic papers, a practical but error-prone process. To meet this challenge, automatic table verification aims to verify the numerical data in tables by cross-referencing cited sources.

target paper					
target table					
model	metric 1	metric 2	metric 3		
A [1]	0.12	0.56	0.41		
B [2]	0.34	0.78	0.50		
Ours	0.41	0.80	0.56		

Table 1. Comparison to A [1] and B [2] in terms

[A placeholder paper text for illustrative purposes] Lorem ipsum

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of metric 1, 2, and 3.

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#### source paper (B [2])

metric 4 (%)

metric 5 (%)

model

D [27] 33.0 55.0 44.0 66.0 Ours Table 1. We compare our model to D [27]. source table metric 1 (%) model metric 2 (%) A' [29] 10.2 41.2 A [29] 12.1 56.3 Ours 34.1 77.8 Table 2. Quantitative results of our model with regards to metric 1 and 2.



I. Table matching

### II. Cell matching

### arXiVeri benchmark

#### Metrics



# Text prompt for cell matching

Target-source cell matching			
Input	a target table (target_table), a source table (source_table)		
System User	You are a helpful assistant. Compare the following target and source tables and identify cells that contain floating point numbers with the same meaning present in both tables. Return the matched cells in a Python dictionary with the following format: { (target_table_row_index, target_table_column_index): (source_table_row_index, source_table_column_index),  } Use 0-based indexing, including headers, rowspan, and colspan at- tributes. Locate as many matching cell pairs as possible. If no matches are found, return an empty dictionary ({}). The target table and its caption: {target_table} The source table and its caption: {source_table}		

- Table matching accuracy evaluates the verifier's ability to accurately identify a source table that matches a given target table, or to determine that no such source table exists in the cited document.
- Cell matching recall quantifies the percentage of targetsource cell matches that are accurately identified (i.e., true positives) among a ground-truth set of cell matches across a source table and a target table.
- Cell matching precision measures how many targetsource cell pairs are true positives among all the detected target-source cell pairs.

# Qualitative example



Table 1. For each method, we report supervised metrics (i.e., ones requiring human references): B@1 = BLEU-1, M = METEOR, C = CIDEr, S = SPICE. We also report diversity metrics, which measures the vocabulary size (Vocab), and the number of novel sentences w.r.t the training set (%Novel). Finally, we report semantic relatedness to the image (CLIP-S), and to the human references (CLIP-SRef) based on

Cells marked in green denote accurate correspondences, while those highlighted in orange indicate mismatches.

### Links







# Code & benchmark

## Conclusion

• We address the critical task of ensuring numerical data accuracy in academic documents by introducing a novel task **-automatic table verification**—leveraging the capabilities of large language models.

• We presented **arXiVeri**, a benchmark comprising tabular data from arXiv papers, and proposed metrics for evaluating verification performance.

Despite the sophistication of advanced models like GPT-4, our findings underline the inherent complexity of the task, underscoring the necessity for further research in this field.