

Code Contribution and Authorship

Eva Maxfield Brown

Nicholas Weber

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1 Introduction

Contemporary scientific research fundamentally depends on specialized software tools and computational methods (Edwards et al. 2013; Mayernik et al. 2017; Howison et al. 2015). These include the analysis scripts for data processing and statistical testing, and research tools and computational infrastructure for data collection and storage (Hasselbring et al. 2024), which act as essential components of modern research practices. Software itself additionally plays multiple critical roles: enabling reproducible research and large-scale experiments (Krafczyk et al. 2019; Trisovic et al. 2021), serving as a detailed log of research methodology (Ram 2013), and increasingly being shared alongside research articles as a crucial research artifact (Cao et al. 2023; Trujillo, Hébert-Dufresne, and Bagrow 2022).

Despite the fundamental importance of scientific software, its development and maintenance face persistent challenges in receiving appropriate academic recognition (Muna et al. 2016). Software contributions are often relegated to acknowledgments rather than warranting authorship (Philippe et al. 2019), and this lack of formal credit can significantly impact career advancement in academia (Carver et al. 2022; Biagioli and Galison 2014). The academic community continues to grapple with developing appropriate systems for software citation and credit attribution that adequately reflect these contributions (Merow et al. 2023; Westner et al. 2024; Katz et al. 2020).

Recent initiatives like the Contributor Roles Taxonomy (CRediT) have attempted to address these challenges by expanding academic authorship criteria to include specialized roles (Brand

et al. 2015), but have not fully resolved the issues surrounding software contribution recognition. While previous research has used CRediT and similar systems to understand research labor distribution (Larivière, Pontille, and Sugimoto 2020; Larivière et al. 2016; Sauermann and Haeussler 2017; K. Li, Zhang, and Larivière 2023; Lu et al. 2019), these frameworks remain largely centered on traditional author lists. This traditional focus perpetuates existing problems, including historical and systematic bias, and relies heavily on self-reporting without external verification (Haeussler and Sauermann 2013; Gøtzsche et al. 2007; Ni et al. 2021). The limitations of current contribution frameworks, combined with the growing importance of software in research, highlight the need for a systematic examination of how code contributions actually relate to academic recognition and impact.

These challenges raise several critical questions about the relationship between software development and academic recognition: How does the distribution of coding contributions within research teams relate to article impact? To what extent does sustained engagement in software development influence the likelihood of receiving authorship recognition? How do traditional academic roles align with actual code contributions? What are the career implications for researchers who regularly contribute code to research projects?

To address these questions, we developed a novel predictive model that enables systematic matching between scientific article authors and source code developer accounts. This approach was necessary due to the lack of standardized identifiers (like ORCID) for developers (Haak et al. 2012) and inconsistencies in naming and email conventions across platforms. We leverage recent advances in transformer-based architectures and semantic embeddings to handle subtle variations in identity information (Y. Li et al. 2020; Brunner and Stockinger 2020).

By applying our model across a corpus of 138596 paired research articles and repositories, we provide unique insight into the dynamics of code contribution within research teams. Our analysis moves beyond self-reporting to examine verifiable source-code repository commit histories, providing preliminary quantitative evidence of the exclusion of code contributors from academic authorship. While we find that code-contributing authors are associated with modest increases in article-level impact metrics (~4.5% increase in citations per code-contributing author), these effects are reduced or become statistically non-significant when controlling for factors such as domain, article type, and access status. Across every condition tested, first authors are consistently and significantly more likely to be code contributors compared to other author positions. However, when examining author-level impact metrics, we find a clear negative relationship between coding frequency and scholarly impact - authors who contribute code more frequently show progressively lower h-indices compared to their non-coding peers, a pattern that persists across various controls for author position, domain, and article type.

These findings begin to illuminate the complex and sometimes counterintuitive relationship between code contribution and scientific impact. While our results suggest that software development plays an important role in modern research, they also highlight potential tensions between technical contributions and traditional academic metrics. Our quantitative evidence could help inform ongoing discussions about academic credit systems, though more research

is needed to fully understand how best to recognize and reward software development contributions in research.

2 Background

The relationship between scientific software development and academic credit systems represents a complex intersection of traditional academic practices and modern research requirements. While academic credit has historically focused on analytical, theoretical, and experimental contributions (Larivière et al. 2016; X. Liu, Zhang, and Li 2023), software development has long been viewed as purely technical rather than scholarly work. However, there is growing recognition that research software development requires deep domain expertise (Heroux 2022; Carver et al. 2022), particularly as increased emphasis on large-scale data projects has amplified the need for sophisticated computational methods (Jin et al. 2015; Hampton et al. 2013; Fan, Han, and Liu 2014).

Understanding the dynamics of scientific software development requires examining both team-level patterns and individual contributions. While existing research has extensively studied the impact of general team size and diversity on research outcomes (Franceschet and Costantini 2010; Larivière et al. 2014; AlShebli et al. 2024; Yang, Ding, and Liu 2024; L. Liu et al. 2021; Naik et al. 2023), and software engineering literature has demonstrated correlations between team size and code quality (Wyss, De Carli, and Davidson 2023; Meirelles et al. 2010), we lack systematic evidence of how code contribution patterns affect research impact. Given that technically sophisticated research projects often enable novel scientific insights and methodological advances, **we hypothesize that projects with more code contributors will receive higher citation counts, reflecting the value of technical contributions even when they go uncredited (H1).**

Recent efforts to standardize contribution recognition, such as the CRediT taxonomy, have attempted to establish formal roles for software development (K. Li, Zhang, and Larivière 2023; Ding et al. 2021). However, these frameworks may not capture the full spectrum of technical contributions, particularly the relationship between sustained engagement in software development and authorship recognition. Repository histories provide a unique opportunity to examine this relationship (Ram 2013), leading us to hypothesize that longer project involvement increases the likelihood of authorship recognition. **Specifically, we expect that projects with longer commit durations will be associated with higher percentages of developers who receive authorship (H2).**

Traditional academic authorship conventions carry specific expectations about intellectual contribution and project responsibilities. First authors typically bear primary responsibility for intellectual and experimental contributions (Larivière et al. 2016; Larivière, Pontille, and Sugimoto 2020; Júnior et al. 2016), while corresponding authors maintain research artifacts and serve as primary points of contact. However, expectations regarding technical contributions vary significantly across academic disciplines (E. Smith 2023). **Given these conventions,**

we hypothesize that first authors (H3) and corresponding authors (H4) will show higher rates of code contribution compared to middle and last authors.

The tension between software development and academic advancement is particularly acute given the field’s reliance on traditional impact metrics. **We hypothesize that researchers who frequently contribute code will be associated with lower individual-level impact metrics compared to their non-coding peers (H5).** This may occur through multiple mechanisms: lack of standardized software citation practices reduces recognition of technical contributions, theoretical advances garner more citations than methodological improvements, and version-specific iterative software publications lead to citation-splitting across multiple articles. This pattern could explain the career implications faced by researchers who prioritize coding (Hannay et al. 2009; Heroux 2022; A. M. Smith, Norman, and Cruz 2019; Cosden, McHenry, and Katz 2022).

3 Data and Methods

3.1 Linking Scientific Articles and Source Code Repositories

Modern scientific research increasingly requires the public sharing of research code, creating unique opportunities to study the relationship between academic authorship and software development. Many journals and platforms now require or recommend code and data sharing (Stodden, Guo, and Ma 2013; Sharma et al. 2024), creating traceable links between publications and code that enable systematic study of both article-repository and author-developer relationships (Hata et al. 2021; Kelley and Garijo 2021; Stankovski and Garijo 2024; Milewicz, Pinto, and Rodeghero 2019).

Our data collection process leverages multiple complementary sources of linked scientific articles and code repositories to ensure comprehensive coverage. Our dataset combines multiple sources:

- PLOS: Traditional research articles with code requirements
- JOSS and SoftwareX: Specialized software-focused publications
- Papers with Code (ArXiv): Capturing the pre-print landscape.

To reduce the complexity of dataset processing and enrichment, we filter out any article-source-code-repository pairs which store code somewhere other than GitHub. While this decision prioritizes processing simplicity, we acknowledge that while GitHub is the predominate host of much of scientific software, it is also stored and shared on other platforms (Trujillo, Hébert-Dufresne, and Bagrow 2022).

Through integration of multiple data sources, we extract detailed information about both the academic and software development aspects of each project:

1. Article Resolution and Metadata:

- Use Semantic Scholar API for DOI resolution to find latest versions, particularly important for preprints
- Collect publication metadata via OpenAlex API: open access status, domain, publication date
- Gather author details: name, position, corresponding author status
- Extract impact metrics: citation count, FWCI, h-index

2. Repository Information:

- Use GitHub API to collect repository metadata: name, description, languages, creation date
- Gather contributor details: username, name, email
- Extract engagement metrics: star count, fork count, issue count

While the majority of our data is sourced from Papers with Code, our additional collection from PLOS, JOSS, and SoftwareX as well as the enrichment from GitHub and OpenAlex together form one of the largest and most comprehensively annotated collections of paired scientific articles and associated source-code repositories. In total, we collect and enrich data for 163292 article-repository pairs.

3.2 A Predictive Model for Matching Article Authors and Source Code Contributors

3.2.1 Annotated Dataset Creation

The development of an accurate author-developer matching model requires high-quality labeled training data that captures the complexity of real-world identity matching. Entity matching between authors and developers is non-trivial due to multiple forms of name variation and incomplete information. These variations can include differences in formatting (e.g., “J. Smith” vs “John Smith”), institutional versus personal email addresses, different names across platforms (e.g., academic name vs GitHub username), and incomplete or outdated information.

We developed an annotation process to create a robust training dataset while maximizing efficiency and accuracy. We focused our annotation efforts on JOSS articles to increase positive match density, as these software-focused publications typically have higher overlap between authors and developers. For each JOSS author, we generated three random pairings with developer accounts from the article’s associated repository. From the full set of pairs, we randomly sampled 3,000 for annotation. Two independent annotators labeled each pair as either a match or non-match. After completing all annotations, we systematically resolved any disagreements between the annotators through discussion and additional verification.

The resulting annotated dataset provides a comprehensive foundation for training our predictive model while highlighting common patterns in author-developer identity matching. After resolving all annotated pairs, our final dataset contains:

1. Match Distribution:
 - 451 (15.0%) positive matches
 - 2548 (85.0%) negative matches
2. Unique Individuals:
 - 2027 unique authors
 - 2733 unique developer accounts
3. Developer Profile Completeness:
 - 2191 (80.2%) accounts have associated names
 - 839 (30.7%) accounts have associated emails

This distribution of matches and varying completeness of developer profiles reflects the real-world challenges in identity matching across academic and software development platforms, requiring our model to handle both complete and partial information robustly.

3.2.2 Training and Evaluation

- To optimize our predictive model for author-contributor matching, we evaluate a variety of Transformer-based base models and input features.
 - specifically, we fine-tune from three different base transformer models:
 - * [deberta-v3-base](#) (He, Gao, and Chen 2021; He et al. 2021)
 - * [bert-base-multilingual-cased](#) (Devlin et al. 2018)
 - * [distilbert-base-uncased](#) (Sanh et al. 2019)
 - these three models are all variations or built-upon BERT, and while significant time has passed since BERT was first introduced, BERT-based models remain a strong base for many NLP tasks across a number of domains while being relatively “small” compared to the much larger decoder transformer relatives (GPT, Llama, etc.) (Tran et al. 2024; Yu et al. 2024; Jeong and Kim 2022)
- We employed a systematic evaluation to identify optimal combination of base models and input features.
 - first, to ensure that there was no data leakage, we split our dataset into training and test sets
 - specifically, we created two random sets of 10% of all unique authors and 10% of all unique developers, any pairs containing either the author or developer were placed into the test set
 - in doing so, we ensured that the model was never trained on any author or developer information later used for evaluation

- due to the fact that each author and developer-account can be included in multiple annotated pairs, our final training set contains 2442 (81.4%) and our test set contains 557 (18.6%) author-developer-account pairs
 - we fine-tuned each of our three base models using all combinations of available developer-account features, from including only the developer account username to including the developer’s username, name, and email.
 - to avoid overfitting and ensure generalizability, we fine-tuned each of the base models for only a single training epoch.
 - model evaluation was performed using standard classification metrics, including accuracy, precision, recall, and F1 score
- After extensive model comparison we find that fine-tuning from [Microsoft’s deberta-v3-base](#) and including the developer’s username and name achieves the best performance for author-developer matching.
 - our best model achieves a binary F1 score of 0.944, with an accuracy of 0.984, precision of 0.938, and recall of 0.95
 - analysis of feature importance
 - * note that the addition of developer’s name has a “larger effect” on model performance but that could simply be because of how many more developers have a name available than an email
 - * also note that there is a model that performs just as well as this one using bert-multilingual and includes the developers email however we choose to use the deberta and name only version for its simplicity as well as the fact that deberta is a much more recently developed and released model which was pre-trained on a much larger dataset.
 - * considering that in most cases, deberta out-performs bert-multilingual, we believe that while the overall evaluation metrics between the top two performing models are the same, the deberta based model will generalize to other unseen data better than the bert-multilingual model
 - all model and feature set combination results are available in [Table 4](#)
 - To enable future research, we have made our trained model and supporting application library publicly available.
 - Python library implementation: [sci-soft-models](#)
 - HuggingFace model deployment: [dev-author-em-clf](#)

3.3 Linking Authors and GitHub Developer Accounts

- Our trained entity-matching model enables comprehensive identification of author-developer relationships while accounting for the complex realities of academic software development practices.

- in practice, to fill out our dataset, we apply our trained model to all possible author and developer-account combinations within each article-repository pair
- The presence of multiple developer accounts per individual reflects common practices in academic software development that must be accommodated in our analysis.
- developers often maintain separate accounts for different projects or institutions
- account transitions are common as researchers move between roles
- Further, while our model performs well, we note that there are some limitations to our approach.
 - in most cases predictions are trivial due to minor differences in text (spelling of author name to username)
 - however we do observe a few cases in which our model may not perform as well
 - namely, shorter names, articles and repositories which have contributors with the same last name (i.e. siblings or other relationship), and “organization” accounts (i.e. research lab GitHub accounts used for management, administration, and documentation or a project)
 - TODO: should we take a sample and estimate how widespread these problems are?
 - we include appropriate filtering during analysis to ensure that we do not include author-developer pairs which are unlikely to be the same individual
- Our final dataset provides unprecedented scale and scope for analyzing the relationship between academic authorship and software development contributions.
 - Specifically, our dataset contains 138596 article-repository pairs, 295806 distinct authors, and 152170 distinct developer accounts.
 - From the 295806 distinct authors and 152170 distinct developer accounts we are able to create 108754 annotated author-developer pairs
 - a detailed breakdown of these counts by data source, domain, document type, and open access status is available in Table 1

4 Preliminary Analysis Code Contributor Authorship and Development Dynamics of Research Teams

4.1 Software Development Dynamics Within Research Teams

- We begin by measuring the distributions of different coding and non-coding contributors across all of the article-code-repository pairs within our dataset.
 - individuals in our dataset can fall into three categories:
 - * Code-Contributing Authors (CC-A): authors for which our model predicted a match with at least one developer account which contributed code to the associated repository

Table 1: Counts of Article-Repository Pairs, Authors, and Developers by Data Sources, Domains, Document Types, and Access Status.

Category	Subset	Article-Repository Pairs	Authors	Developers
By Domain	Physical Sciences	116600	240545	130592
	Social Sciences	8838	29269	14043
	Life Sciences	7729	31649	12150
	Health Sciences	5172	25979	7248
By Document Type	preprint	72177	170301	87311
	research article	63528	173183	78935
	software article	2891	9294	12868
By Access Status	Open	132856	286874	147831
	Closed	5740	23668	9352
By Data Source	pwc	129615	262889	134926
	plos	6090	30233	8784
	joss	2336	7105	11362
	softwarex	555	2244	1628
Total		138596	295806	152170

- * Non-Code-Contributing Authors (NCC-A): authors for which our model did not predict any matches with developer accounts which contributed code to the associated repository
- * Code-Contributing Non-Authors (CC-NA): developer accounts which contributed code to the associated repository but were not predicted to be a match with any author
- within our dataset, we find that papers on average have 4.9 ± 1.9 total authors, 3.9 ± 2.0 non-code-contributing authors, 1.0 ± 0.7 code-contributing authors, and 0.4 ± 1.7 code-contributing non-authors (see Figure 1 for a visualization of these distributions).
- Table 2 provides a detailed breakdown of these distributions by domain, article type, and open access status.
- Our finding of, on average, only a single code-contributing author on a paper, is similar to previous work in understanding distributions of labor in knowledge production from Larivière, Pontille, and Sugimoto (2020) which found that the CRediT tasks of “Data Curation”, “Formal Analysis”, “Visualization”, and “Software” were all predominantly performed by a first author.
- However, we do also see that code-contributing non-authors are present in the data, albeit, on average, with less than one code-contributing non-author per article.
- Next we investigate how these distributions have changed over time and how they change by the total number of authors on a paper.

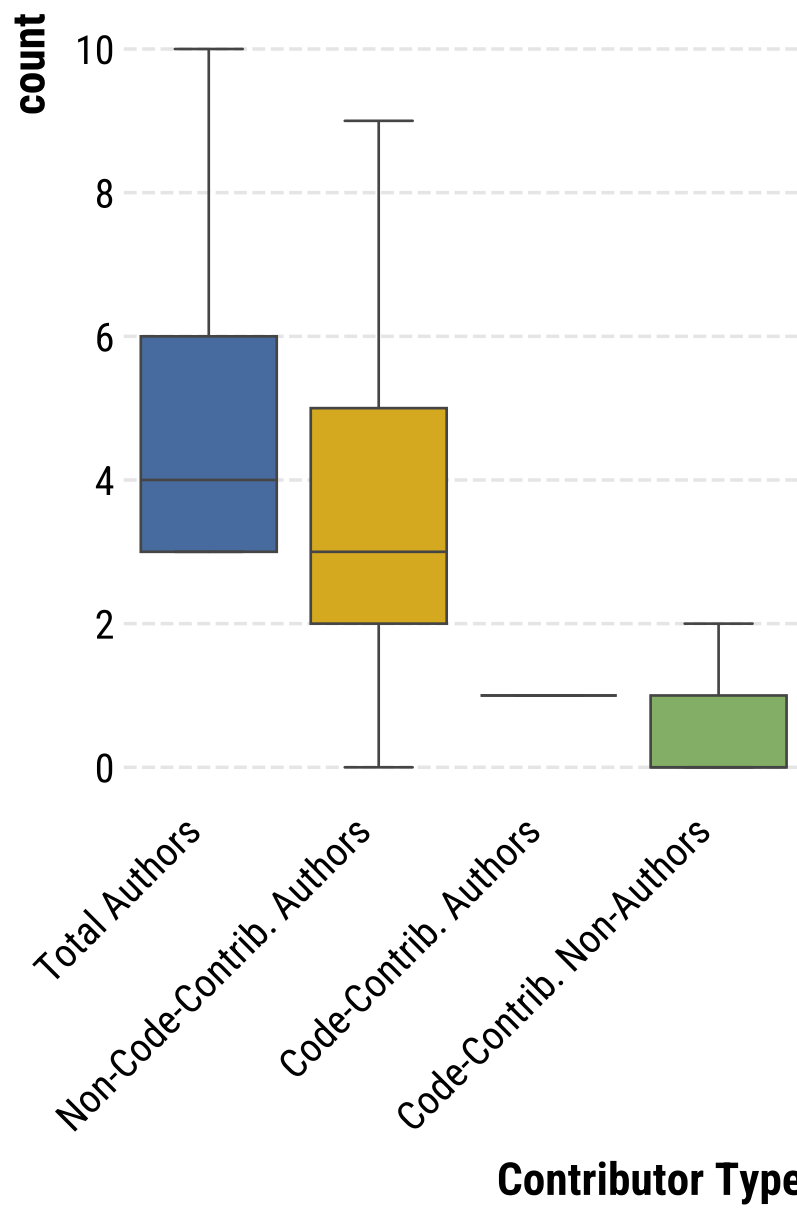


Figure 1: Distribution of the number of Total Authors, Non-Code-Contributing Authors, Code-Contributing Authors, and Code-Contributing Non-Authors across all article-repository pairs. Only includes article-repository pairs with a most recent commit no later than 90 days after publication.

Table 2: Mean and Standard Deviation of Non-Code-Contributing Authors (NCC-A), Code-Contributing Authors (CC-A), and Code-Contributing Non-Authors (CC-NA) Research Team Members by Domain, Article Type, and Open Access Status. Only includes research teams from article-repository pairs with a most recent commit no later than 90 days after publication and excludes research teams which are in the top 3% of total author sizes.

Control	Subset	Total Authors	NCC-A	CC-A	CC-NA
OA Status	Closed	5.1 ± 1.9	4.0 ± 1.9	1.1 ± 0.7	0.5 ± 2.1
	Open	4.9 ± 1.9	3.9 ± 2.0	1.0 ± 0.7	0.4 ± 1.7
Domain	Health Sciences	6.1 ± 2.5	5.1 ± 2.5	1.0 ± 0.6	0.4 ± 1.2
	Life Sciences	5.2 ± 2.1	4.2 ± 2.2	1.0 ± 0.7	0.4 ± 1.2
	Physical Sciences	4.8 ± 1.8	3.8 ± 1.9	1.0 ± 0.7	0.5 ± 1.8
	Social Sciences	4.5 ± 1.7	3.5 ± 1.8	1.1 ± 0.7	0.3 ± 1.1
Article Type	preprint	4.8 ± 1.8	3.8 ± 1.9	1.1 ± 0.7	0.5 ± 2.2
	research article	4.9 ± 1.9	3.9 ± 2.0	1.0 ± 0.7	0.4 ± 1.6
	software article	4.7 ± 1.9	3.2 ± 1.9	1.5 ± 1.4	0.9 ± 1.1
Overall		4.9 ± 1.9	3.9 ± 2.0	1.0 ± 0.7	0.4 ± 1.7

- Figure 2 shows that both, the mean number of code-contributing authors and code-contributing non-authors has remained relatively stable over time and across different total author sizes.
- these descriptive statistics suggest that code-contributing non-authors are an inconsistent feature of research teams, and while their exclusion from authorship is not a recent phenomenon, their exclusion does not appear to be getting worse over time.

4.1.1 Modeling Citations

- Building upon previous work which discuss the effects of team size and team diversity on scientific impact and software quality (see Section 2), we examine how the number of code contributors within a research team may be associated with an article’s research impact.
 - We hypothesize that more code contributors may signal greater technical complexity in research, which may be associated with higher citation counts as the community builds upon more technically sophisticated works.
- However, after analyzing our data, we find few significant associations between the number of code-contributing authors and non-authors, and article citations.
 - Without controlling for any domain, open access, or article type differences (Table 5), we find a positive association between the number of code-contributing

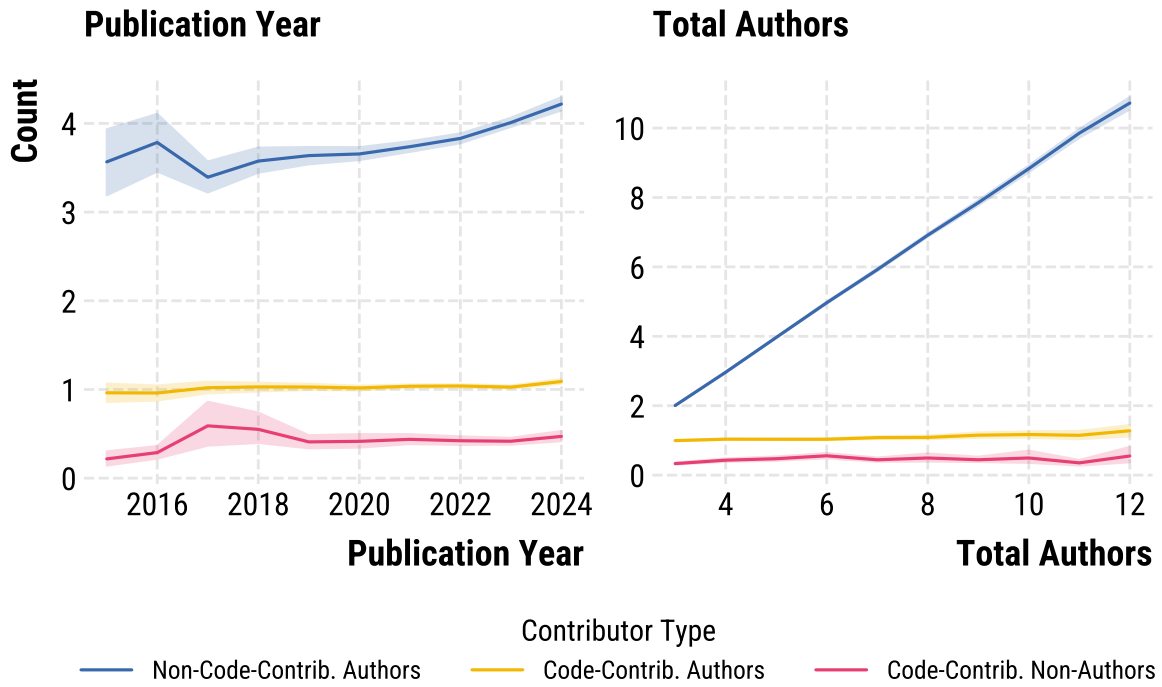


Figure 2: Mean number of Non-Code-Contributing Authors, Code-Contributing Authors, and Code-Contributing Non-Authors by Publication Year and by Total Number of Authors. Only includes article-repository pairs with a most recent commit no later than 90 days after publication and excludes research-teams which are in the top 3% of total author sizes for publication years with 50 or more articles.

authors and article citations with each code-contributing author being associated with a 4.5% increase in article citations ($p < 0.001$).

- Controlling for article type (Table 8), we find that for preprints, each code-contributing non-author is associated with a statistically significant 2.9% decrease in expected citations, holding other variables constant ($p < 0.01$).
- For research articles, we find a significant positive association between the number of code-contributing authors and citations ($p < 0.001$). However, we cannot estimate the precise magnitude of this effect due to the non-significant main effect in the model.
- Finally, we additionally find a statistically significant relationship for code-contributing non-authors for research articles, specifically finding that each code-contributing non-author is associated with a 0.8% increase in expected citations ($p < 0.001$).
- Overall, while we find some statistically significant associations between code contributions and citation counts, these effects are relatively modest in magnitude.

4.1.2 Team Composition and Project Duration

- Building upon previous work examining standardized contribution frameworks and authorship attribution practices (see Section 2), we investigate how project duration may influence the distribution of coding roles between authors and non-authors.
 - We hypothesize that projects with longer development durations may show higher proportions of author-developers compared to non-author developers, as sustained technical engagement could increase the likelihood of receiving authorship recognition.
- However, our analysis finds no evidence to support this hypothesis:
 - When examining the relationship between a repository’s commit duration and the percentage of developers who receive authorship recognition, we find no significant correlation ($r = -0.00$, $p = \text{n.s.}$).
 - This suggests that the length of time a project has been in development has no meaningful relationship with the proportion of developers who are recognized as authors.
 - While this suggests authorship recognition may be driven by other factors, a more precise analysis would examine individual contribution durations rather than overall project length.

4.2 Characteristics of Scientific Code Contributors

4.2.1 Author Positions of Code Contributing Authors

- Building upon previous work examining the relationship between authorship position and research contributions, we investigate how author position may relate to code contribution patterns.
 - We hypothesize that first authors, who traditionally contribute the bulk of intellectual and experimental work, would be most likely to contribute code to a project, while middle and last authors, who often provide oversight and guidance, would be less likely to contribute code.
- Our analysis supports this hypothesis:
 - In every case tested, we find that first authors are statistically significantly more likely to be code contributors than non-code contributors across most scenarios both overall, and regardless of Domain, Article Type, and Open Access Status.
 - Conversely, both middle authors and last authors are statistically significantly more likely to be non-code contributors.
 - These patterns align with traditional academic authorship conventions, where first authors often take primary responsibility for both intellectual and technical aspects of the research.
 - Table 9 provides a detailed breakdown of Bonferroni-correct p-values from post-hoc binomial tests against the counts of author positions of code-contributing authors.

4.2.2 Corresponding Status of Code Contributing Authors

- Building upon our analysis of author position, we next examine how corresponding author status relates to code contribution patterns.
 - We hypothesize that corresponding authors, who traditionally maintain research artifacts and serve as primary points of contact, would be more likely to contribute code compared to non-corresponding authors.
- However, our analysis reveals patterns contrary to this hypothesis:
 - Both corresponding and non-corresponding authors are statistically significantly less likely to be code contributors than would be expected by chance ($p < 0.001$) in most scenarios (Table 10).
 - This pattern holds true across almost all conditions with the only exceptions found in software articles and closed access articles, where corresponding authors show no significant difference in their likelihood to contribute code.

Table 3: Counts of Researcher Coding Status Used in H5

Control	Subset	Any Coding	Majority Coding	Always Coding	Total
Freq. Author Pos.	First	1689	4952	3322	11444
	Middle	12184	3327	701	31911
	Last	2413	568	191	10188
Freq. Domain	Health Sciences	345	202	86	1501
	Life Sciences	369	241	129	1436
	Physical Sciences	15289	8179	3821	49311
	Social Sciences	283	225	178	1295
Freq. Article Type	Preprint	9572	4963	2214	28948
	Research Article	6669	3765	1871	24217
	Software Article	45	119	129	378

- These findings suggest that the responsibility for maintaining research artifacts may not typically extend to direct code contributions, contrary to traditional assumptions about corresponding author roles.

4.2.3 Modeling H-Index

- Building upon previous work examining career implications for researchers who prioritize software development (see Section 2), we investigate how varying levels of code contribution relate to scholarly impact through citation metrics.
 - We hypothesize that researchers who contribute more code may show lower citation counts, potentially due to a lack of formal software citation or more general attitudes towards methodological vs theoretical contributions.
- While we can’t confirm the underlying mechanisms, we do find strong evidence to support our general claim that code-contributing researchers are associated with lower h-indices compared to their non-coding peers.
 - Without any controls (Table 11), we find increasingly negative h-index effects as coding frequency increases compared to non-coding authors with scientists who occasionally make code contributions being associated with ~27.3% lower h-indices than their non-coding counterparts ($p < 0.001$), followed by ~53.5% lower h-indices for majority coders ($p < 0.001$), and ~62.1% lower h-indices for always coding authors ($p < 0.001$).
 - When controlling for author position (Table 12), we find again find a general pattern that more frequent code contribution is associated with reduced h-indices compared to non-coding peers with a few exceptions.

- * Occasional coding first authors are associated with a ~14.9% higher h-index ($p < 0.001$) while always coding first authors are associated with a ~21.6% reduction in h-index compared to non-coding first authors ($p < 0.001$).
- * Occasional coding middle authors are associated with a ~26.6% lower h-index ($p < 0.001$) while always coding middle authors are associated with a ~52.9% lower h-index compared to non-coding middle authors ($p < 0.001$).
- * Occasional coding last authors are associated with a ~13.1% lower h-index ($p < 0.001$) while always coding last authors are associated with a ~45.7% lower h-index compared to non-coding last authors ($p < 0.001$).
- When controlling for domain (Table 13), majority coding scientists in health science domains are associated with a ~76.5% reduction in h-index compared to non-coding peers ($p < 0.001$), while majority coding scientists in life science see a ~47.1% reduction ($p < 0.001$), physical science majority coding scientists see a ~52.3% reduction ($p < 0.001$), and majority coding scientists in social sciences see a ~51.4% reduction in h-index compared to non-coding peers ($p < 0.001$).
- Finally, when controlling for an authors common article type (Table 14), we find that:
 - * for authors whose most common article type is preprints, occasionally coding authors are associated with a ~25.6% lower h-index, majority coding authors are associated with a ~53.5% lower h-index, and always coding authors are associated with a ~62.9% lower h-index compared to their non-coding peers.
 - * while for authors whose most common article type is software articles, majority coding authors are associated with a ~33.1% lower h-index, and always coding authors are associated with a ~33.0% lower h-index compared to their non-coding peers.
- Taken as a whole, these findings tend to indicate that the more frequently an author contributes code, the lower their h-index is likely to be relative to their peers, with one exception being first authors who occasionally contribute code.

5 Discussion

RE: article citations

- Our findings that code contribution and distribution among research teams has only a modest relationship with citations might reflect several underlying factors.
 - First, the relationship between code contributions and research impact may be complex and indirect - while code might enhance research reproducibility or utility, this may not directly translate into increased citations
 - Second, current citation practices may not adequately capture or credit code contributions, as authors might reference papers without explicitly acknowledging the associated code.

- Third, the quality and significance of code contributions likely varies substantially across papers, making it difficult to detect strong aggregate effects.
- Finally, while we have attempted to find and match as many code contributing members of research teams as possible, we must acknowledge that there may be two code-centric reasons why we have not found more code contributors:
 - * First, the code may have been developed by multiple individuals but only a single individual uploaded or committed the code to a repository, thereby removing the opportunity for us to link the code to the other contributors.
 - * Second, we are primarily analyzing “analysis” repositories, which may not contain the full codebase for a project. This is particularly true for projects which separate repositories for tools, libraries, and infrastructures, than the single analysis code repository.

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7 Appendix

Table 4: Comparison of Models for Author-Developer-Account Matching

Optional Feats.	Model	Accuracy	Precision	Recall	F1
name	deberta	0.984	0.938	0.950	0.944
name, email	bert-multilingual	0.984	0.938	0.950	0.944
name, email	deberta	0.982	0.907	0.975	0.940
name	bert-multilingual	0.982	0.938	0.938	0.938
name	distilbert	0.978	0.936	0.912	0.924
name, email	distilbert	0.978	0.936	0.912	0.924
email	deberta	0.957	0.859	0.838	0.848
email	bert-multilingual	0.950	0.894	0.738	0.808
n/a	deberta	0.946	0.847	0.762	0.803
n/a	bert-multilingual	0.941	0.862	0.700	0.772
n/a	distilbert	0.856	0.000	0.000	0.000
email	distilbert	0.856	0.000	0.000	0.000

Table 5: Article Citations by Code Contributorship of Research Team

Variable	coef	P> z	[0.025	0.975]
const ***	0.99	0.00	0.94	1.04
Total Authors ***	0.07	0.00	0.06	0.08
Code-Contrib. Authors ***	0.04	0.00	0.02	0.06
Code-Contrib. Non-Authors	-0.00	0.80	-0.01	0.01
Years Since Publication ***	0.39	0.00	0.38	0.40

Table 6: Article Citations by Code Contributorship of Research Team Controlled by Open Access Status

Variable	coef	P> z	[0.025	0.975]
const ***	0.61	0.00	0.49	0.73
Total Authors ***	0.07	0.00	0.06	0.08
Code-Contrib. Authors	0.05	0.24	-0.04	0.14
Code-Contrib. Non-Authors	0.00	0.96	-0.03	0.03
Years Since Publication ***	0.38	0.00	0.37	0.39
Is Open Access ***	0.42	0.00	0.30	0.54
Code-Contrib. Authors * Is Open Access	-0.01	0.82	-0.10	0.08
Code-Contrib. Non-Authors * Is Open Access	-0.00	0.94	-0.03	0.03

Table 7: Article Citations by Code Contributorship of Research Team Controlled by Domain

Variable	coef	P> z	[0.025	0.975]
const ***	0.88	0.00	0.75	1.01
Total Authors ***	0.07	0.00	0.06	0.08
Code-Contrib. Authors	0.03	0.60	-0.08	0.13
Code-Contrib. Non-Authors	0.01	0.61	-0.04	0.07
Years Since Publication ***	0.40	0.00	0.39	0.41
Domain Life Sciences ***	-0.21	0.01	-0.36	-0.06
Domain Physical Sciences ***	0.14	0.03	0.01	0.26
Domain Social Sciences ***	-0.18	0.03	-0.34	-0.02
Code-Contrib. Authors * Domain Life Sciences	0.07	0.29	-0.06	0.19
Code-Contrib. Authors * Domain Physical Sciences	0.00	0.93	-0.10	0.11
Code-Contrib. Authors * Domain Social Sciences	0.10	0.14	-0.03	0.23
Code-Contrib. Non-Authors * Domain Life Sciences	-0.04	0.23	-0.11	0.03
Code-Contrib. Non-Authors * Domain Physical Sciences	-0.02	0.57	-0.07	0.04
Code-Contrib. Non-Authors * Domain Social Sciences	-0.04	0.31	-0.11	0.03

Table 8: Article Citations by Code Contributorship of Research Team Controlled by Article Type

Variable	coef	P> z	[0.025	0.975]
const ***	0.53	0.00	0.45	0.61
Total Authors ***	0.07	0.00	0.06	0.08
Code-Contrib. Authors	-0.03	0.20	-0.09	0.02
Code-Contrib. Non-Authors ***	-0.03	0.00	-0.05	-0.01
Years Since Publication ***	0.40	0.00	0.39	0.41
Article Type Research Article ***	0.47	0.00	0.40	0.55
Article Type Software Article ***	-0.47	0.00	-0.73	-0.22
Code-Contrib. Authors * Article Type Research Article ***	0.10	0.00	0.05	0.16
Code-Contrib. Authors * Article Type Software Article	-0.06	0.37	-0.19	0.07
Code-Contrib. Non-Authors * Article Type Research Article ***	0.04	0.00	0.02	0.06
Code-Contrib. Non-Authors * Article Type Software Article	0.09	0.24	-0.06	0.24

Table 9: Counts of Code-Contributing Authors ('Coding') as well as Total Authors by Position and Bonferroni Corrected p-values from Post-Hoc Binomial Tests

Control	Subset	Position	Coding	Total	p
Domain	Health Sciences	First	1575	2401	0.000***
		Middle	540	11491	0.000***
		Last	234	2349	0.000***
	Life Sciences	First	2586	3895	0.000***
		Middle	852	11875	0.000***
		Last	491	3784	0.000***
	Physical Sciences	First	28919	41987	0.000***
		Middle	10507	111738	0.000***
		Last	3433	40410	0.000***
	Social Sciences	First	2813	4038	0.000***
		Middle	1021	9249	0.000***
		Last	411	3855	0.000***
Article Type	Preprint	First	13421	19523	0.000***
		Middle	5134	52925	0.000***
		Last	1493	18598	0.000***
	Research Article	First	21940	32081	0.000***
		Middle	7345	89991	0.000***
		Last	2909	31188	0.000***
	Software Article	First	532	717	0.000***
		Middle	441	1437	0.000***
		Last	167	612	0.000***
Open Access Status	Closed Access	First	2637	3742	0.000***
		Middle	918	10965	0.000***
		Last	279	3642	0.000***
	Open Access	First	33256	48579	0.000***
		Middle	12002	133388	0.000***
		Last	4290	46756	0.000***
Overall	Overall	First	35893	52321	0.000***
		Middle	12920	144353	0.000***
		Last	4569	50398	0.000***

Table 10: Counts of Code-Contributing Authors ('Coding') as well as Total Authors by Corresponding Status and Bonferroni Corrected p-values from Post-Hoc Binomial Tests

Control	Subset	Is Corresponding	Coding	Total	p
Domain	Life Sciences	Corresponding	1772	8019	0.000***
		Not Corresponding	2157	11535	0.000***
	Physical Sciences	Corresponding	5248	12487	0.000***
		Not Corresponding	37611	181648	0.000***
	Social Sciences	Corresponding	803	2458	0.000***
		Not Corresponding	3442	14684	0.000***
Article Type	Preprint	Corresponding	772	1036	0.000***
		Not Corresponding	19276	90010	0.000***
	Research Article	Corresponding	7716	27339	0.000***
		Not Corresponding	24478	125921	0.000***
	Software Article	Corresponding	213	438	1.000
		Not Corresponding	927	2328	0.000***
Open Access Status	Closed Access	Corresponding	253	468	0.174
		Not Corresponding	3581	17881	0.000***
	Open Access	Corresponding	8448	28345	0.000***
		Not Corresponding	41100	200378	0.000***
Overall	Overall	Corresponding	8701	28813	0.000***
		Not Corresponding	44681	218259	0.000***

Table 11: Code-Contributing Authors H-Index by Coding Status

Variable	coef	P> z	[0.025 0.975]	
const ***	3.19	0.00	3.18	3.20
Works Count ***	0.00	0.00	0.00	0.00
Any Coding ***	-0.32	0.00	-0.33	-0.30
Majority Coding ***	-0.77	0.00	-0.79	-0.74
Always Coding ***	-0.97	0.00	-1.02	-0.92

Table 12: Researcher H-Index by Coding Status Controlled by Most Freq. Author Position

Variable	coef	P> z	[0.025	0.975]
const ***	2.38	0.00	2.31	2.44
Works Count ***	0.00	0.00	0.00	0.00
Any Coding ***	0.14	0.00	0.05	0.22
Majority Coding	-0.07	0.08	-0.14	0.01
Always Coding ***	-0.24	0.00	-0.33	-0.16
Common Author Position Last ***	1.05	0.00	0.98	1.11
Common Author Position Middle ***	0.74	0.00	0.68	0.81
Any Coding * Common Author Position Last ***	-0.28	0.00	-0.37	-0.19
Any Coding * Common Author Position Middle ***	-0.45	0.00	-0.53	-0.36
Majority Coding * Common Author Position Last ***	-0.35	0.00	-0.45	-0.26
Majority Coding * Common Author Position Middle ***	-0.61	0.00	-0.69	-0.52
Always Coding * Common Author Position Last ***	-0.37	0.00	-0.51	-0.22
Always Coding * Common Author Position Middle ***	-0.51	0.00	-0.64	-0.38

Table 13: Researcher H-Index by Coding Status Controlled by Most Freq. Domain

Variable	coef	P> z	[0.025	0.975]
const ***	3.32	0.00	3.29	3.36
Works Count ***	0.00	0.00	0.00	0.00
Any Coding ***	-0.39	0.00	-0.48	-0.31
Majority Coding ***	-1.45	0.00	-1.65	-1.25
Always Coding ***	-1.22	0.00	-1.58	-0.86
Common Domain Life Sciences ***	0.10	0.00	0.05	0.15
Common Domain Physical Sciences ***	-0.15	0.00	-0.18	-0.11
Common Domain Social Sciences ***	-0.16	0.00	-0.22	-0.10
Any Coding * Common Domain Life Sciences	0.11	0.07	-0.01	0.22
Any Coding * Common Domain Physical Sciences	0.08	0.09	-0.01	0.17
Any Coding * Common Domain Social Sciences	0.01	0.94	-0.14	0.15
Majority Coding * Common Domain Life Sciences ***	0.81	0.00	0.58	1.04
Majority Coding * Common Domain Physical Sciences ***	0.70	0.00	0.50	0.90
Majority Coding * Common Domain Social Sciences ***	0.72	0.00	0.46	0.98
Always Coding * Common Domain Life Sciences	0.33	0.12	-0.08	0.74
Always Coding * Common Domain Physical Sciences	0.25	0.18	-0.12	0.62
Always Coding * Common Domain Social Sciences	0.30	0.17	-0.13	0.73

Table 14: Researcher H-Index by Coding Status Controlled by Most Freq. Article Type

Variable	coef	P> z	[0.025	0.975]
const ***	3.10	0.00	3.09	3.11
Works Count ***	0.00	0.00	0.00	0.00
Any Coding ***	-0.30	0.00	-0.32	-0.27
Majority Coding ***	-0.77	0.00	-0.81	-0.73
Always Coding ***	-0.99	0.00	-1.07	-0.92
Common Article Type Research Article ***	0.18	0.00	0.17	0.20
Common Article Type Software Article ***	0.22	0.00	0.12	0.33
Any Coding * Common Article Type Research Article	-0.02	0.15	-0.05	0.01
Any Coding * Common Article Type Software Article	0.19	0.06	-0.01	0.39
Majority Coding * Common Article Type Research Article	0.01	0.80	-0.05	0.06
Majority Coding * Common Article Type Software Article ***	0.36	0.00	0.19	0.54
Always Coding * Common Article Type Research Article	0.00	0.93	-0.10	0.10
Always Coding * Common Article Type Software Article ***	0.37	0.00	0.15	0.58