

Code Contribution and Authorship

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1 Random Citations

- may be interesting / useful: <https://doi.org/10.1145/1772690.1772766>

2 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).
 - define scientific software (analysis scripts, research tools, computational infrastructure) (Hasselbring et al. 2024)
 - software enables reproducible research and large-scale experiments (Krafczyk et al. 2019; Trisovic et al. 2021)
 - code serves as a detailed log of research methodology (Ram 2013)
 - due to all of the above, code is increasingly being shared alongside research articles (Cao et al. 2023; Trujillo, Hébert-Dufresne, and Bagrow 2022)
- The development and maintenance of scientific software requires substantial contribution, yet faces persistent challenges in receiving academic recognition (Muna et al. 2016).
 - software contributions often receive only acknowledgments rather than authorship (Philippe et al. 2019)

- lack of formal credit affects career advancement in academia (Carver et al. 2022; Biagioli and Galison 2014)
 - other general discussion of software citations and credit systems (Merow et al. 2023; Westner et al. 2024; Katz et al. 2020)
- Recent initiatives to expand academic credit systems, while promising, have not fully addressed the challenges of recognizing software contributions.
 - describe the Contributor Roles Taxonomy (CRediT) (Brand et al. 2015)
 - previous research using CRediT (and prior 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)
 - CRediT research is still centered on traditional author lists (historic and systematic bias, self-reporting without verification, etc.) (Haeussler and Sauermann 2013; Gøtzsche et al. 2007; Ni et al. 2021)
- Our novel predictive model addresses these challenges by enabling systematic matching between scientific article authors and source code developer accounts.
 - we use predictive modeling due to the lack of standardized identifiers (i.e. ORCID) for developers (Haak et al. 2012)
 - further, lack consistency in naming and email overlap [GET CITATION FOR SORTING HAT FROM GOGGINS??]
 - semantic models handle subtle variations in identity information (general entity matching has moved to transformers and semantic embeddings) (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, the impact of code contribution on research outcomes, and an understanding of the authors who are and who aren't code contributors.
 - move from self-reporting to verifiable source code repository commit histories
 - provide preliminary quantitative evidence of exclusion of code contributors from academic authorship
 - model article level impact metrics as a function of software development dynamics to show the benefit code contributors have on research
 - find that first authors are more likely to be code contributors than not
 - find that code-contributing authors have reduced individual level impact metrics compared to their non-coding counterparts
- These findings not only illuminate the relationship between code contribution and scientific impact but also provide an empirical foundation for reforming academic credit systems to better recognize software development contributions in research.

3 Background

- The relationship between scientific software development and academic credit systems represents a complex intersection of traditional academic practices and modern research requirements.
 - academic credit traditionally focuses on analytical, theoretical, and experimental contributions (Larivière et al. 2016; X. Liu, Zhang, and Li 2023)
 - software development historically viewed as technical rather than scholarly work
 - growing recognition that research software development requires deep domain expertise (Heroux 2022; Carver et al. 2022)
 - increased emphasis on large scale (big data) projects has resulted in larger need for software development (Jin et al. 2015; Hampton et al. 2013; Fan, Han, and Liu 2014)
 - understanding this relationship requires examining both team-level dynamics and individual contributions
- (H1) Modern research increasingly depends on collaborative software development, yet we lack systematic evidence of how code contribution patterns affect research outcomes.
 - existing research focuses primarily on general team size and diversity (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)
 - software engineering literature shows correlation between team size and code quality (many eyes make all bugs shallow) (Wyss, De Carli, and Davidson 2023; Meirelles et al. 2010)
 - limited understanding of how code contribution is associated to research impact
 - need to understand relationship between code contributors and citation metrics to understand the value of these technical, potentially uncredited, contributions
 - we believe that more code contributors may signal a more technical research project and that technical complexity may be rewarded with more citations
- (H2) Despite formal taxonomies like CRediT attempting to standardize contribution recognition, the criteria for granting authorship to technical contributors remain inconsistent and poorly understood across research communities.
 - existing contribution frameworks provide definitions for software development roles (K. Li, Zhang, and Larivière 2023; Ding et al. 2021)
 - however, these frameworks may not capture the full spectrum of technical contributions
 - repository histories allow us to examine how sustained technical engagement relates to authorship status (Ram 2013)
 - we believe that longer project involvement increases likelihood of authorship recognition

- specifically, we hypothesize that projects with longer durations will show higher proportions of author-developers compared to non-author developers
- (H3 and H4) Academic authorship conventions signal both intellectual contribution and project responsibilities, yet their relationship to software development remains poorly understood.
 - first authors traditionally responsible for primary intellectual and experimental contributions (Júnior et al. 2016)
 - corresponding authors serve as primary points of contact and often maintain research artifacts
 - varying expectations across academic disciplines regarding technical contributions
 - limited research examining how these authorship roles relate to code contributions
 - potential insights into how software development responsibilities are distributed within research teams
 - we believe that first authors and corresponding authors will have higher proportions of code contribution than not.
 - conversely, middle and last authors and non-corresponding authors will have lower proportions of code contribution than not.
- (H5) Academic career advancement has historically depended on traditional impact metrics, creating potential tension for researchers who dedicate significant time to software development.
 - time invested in code development may reduce traditional scholarly output
 - potential career implications for researchers who prioritize coding
 - need to understand relationship between code contributions and academic impact
 - we believe that code contributing researchers will have lower individual level impact metrics than non-coding researchers, likely due to a combination of lack of recognition and authorship credit as well as reduced time for traditional scholarly output
- Understanding these relationships is crucial for developing equitable academic credit systems that recognize the full spectrum of research contributions.
 - findings will inform policy making around academic credit
 - importance of large-scale quantitative evidence for understanding current credit systems
 - implications for academic hiring and promotion decisions
 - potential to develop new impact metrics that capture software contributions

4 Data and Methods

4.1 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
 - this requirement creates traceable links between publications and code
 - these links enable systematic study of both article-repository and author-developer relationships
- Our data collection process leverages multiple complementary sources of linked scientific articles and code repositories to ensure comprehensive coverage.
 - PLOS: Traditional research articles with code requirements
 - JOSS and SoftwareX: Specialized software-focused publications
 - Papers with Code / ArXiv: Capturing 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
- Through integration of multiple data sources, we extract detailed information about both the academic and software development aspects of each project.
 - specifically we utilize the Semantic Scholar API for article DOI resolution to ensure that we find the latest version for each article.
 - this is particularly important for working with preprints as they may have been published in a journal since their inclusion in the Papers with Code dataset
 - we then utilize the OpenAlex API to gather publication metadata (i.e. open access status, domain, publication date), author details (i.e. name, author position, corresponding author status), and article- and individual-level metrics (i.e. citation count, FWCI, h-index).
 - the GitHub API provides similar information for source code repositories, including repository metadata (i.e. name, description, languages, creation date), contributor details (i.e. username, name, email), and repository-level metrics (i.e. 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 collections of linked, metadata enriched, datasets of paired scientific articles and associated source code repositories.
 - in total, we collect and enrich data for 163292 article-repository pairs

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

4.2.1 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
 - multiple forms of name variation and incomplete information
 - need to expand on specific matching challenges
 - add figure showing example matches/non-matches
- We developed an annotation process to create a robust training dataset while maximizing efficiency and accuracy.
 - focus on JOSS articles to increase positive match density
 - we create author-developer pairs for annotation by creating all possible combinations of authors and developers within a single JOSS article-repository pair
 - we take a random sample of 3000 pairs from the full set and have two independent annotators label each
 - structured conflict resolution process
 - need to add details about annotation guidelines/criteria
- The resulting annotated dataset provides a comprehensive foundation for training our predictive model while highlighting common patterns in author-developer identity matching.
 - after resolution of all annotated pairs, our annotated dataset contains 451 (15.0%) positive and 2548 (85.0%) negative author-developer-account pairs
 - there are 2027 unique authors and 2733 unique developer accounts within this annotated set
 - however, not all developer accounts contain complete information, in our set 2191 (80.2%) have associated names and 839 (30.7%) have associated emails

4.2.2 Training and Evaluation

- To optimize our predictive model for author-contributor matching, we evaluate a variety of Transformer-based base models and input features.
 - multiple transformer base models available
 - various potential feature combinations
- 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 used three different transformer models as our fine-tuning bases and fine-tuned each 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 only 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 (see Figure 1 for a confusion matrix of model predictions on the test set).
 - 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 (including multilingual data).
 - * 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 11
 - To enable future research, we have made our trained model and supporting application library publicly available.

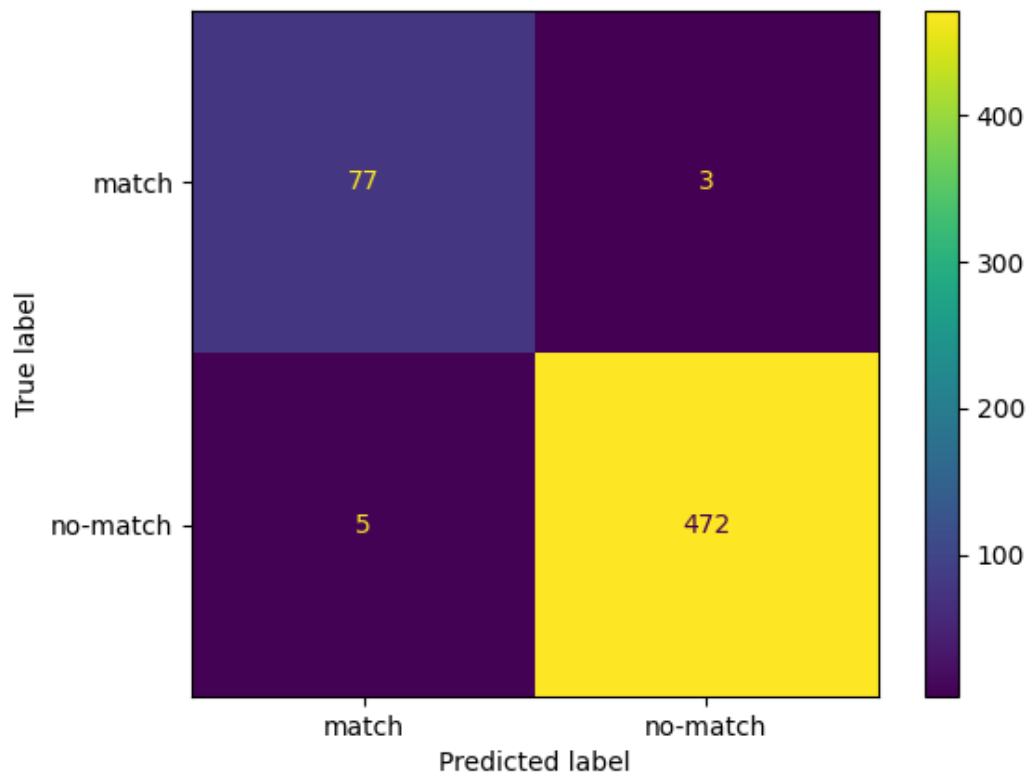


Figure 1: Confusion Matrix Produced From Evaluation of Best Performing Model (deberta-v3 with developer username, developer name, and author name).

- Python library implementation
- HuggingFace model deployment

4.3 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.
 - model applied to all possible author and developer-account combinations within each article-repository pair
 - recognition that one-to-one matching may not reflect reality
 - deliberate choice to allow many-to-many relationships in matching
- 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
 - CHAOSS project provides precedent for this type of identity resolution
- Further, while our model performs well overall, 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.
 - a detailed breakdown of these counts by data source, domain, document type, and open access status is available in Table 1

Table 1: Counts of Article-Repository Pairs, Authors, and Developers broken out 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

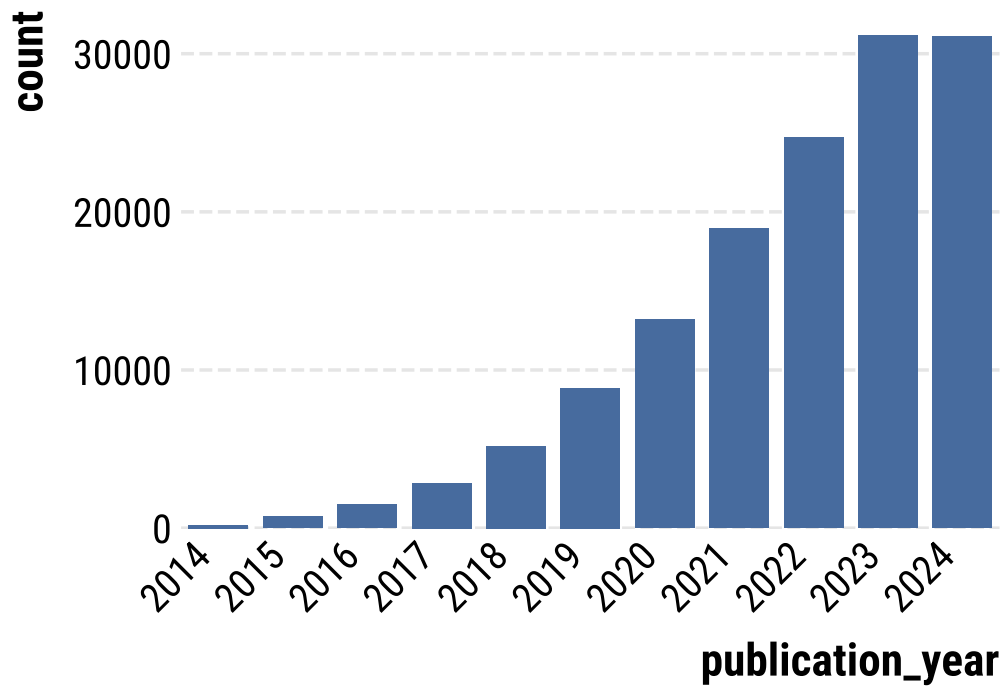


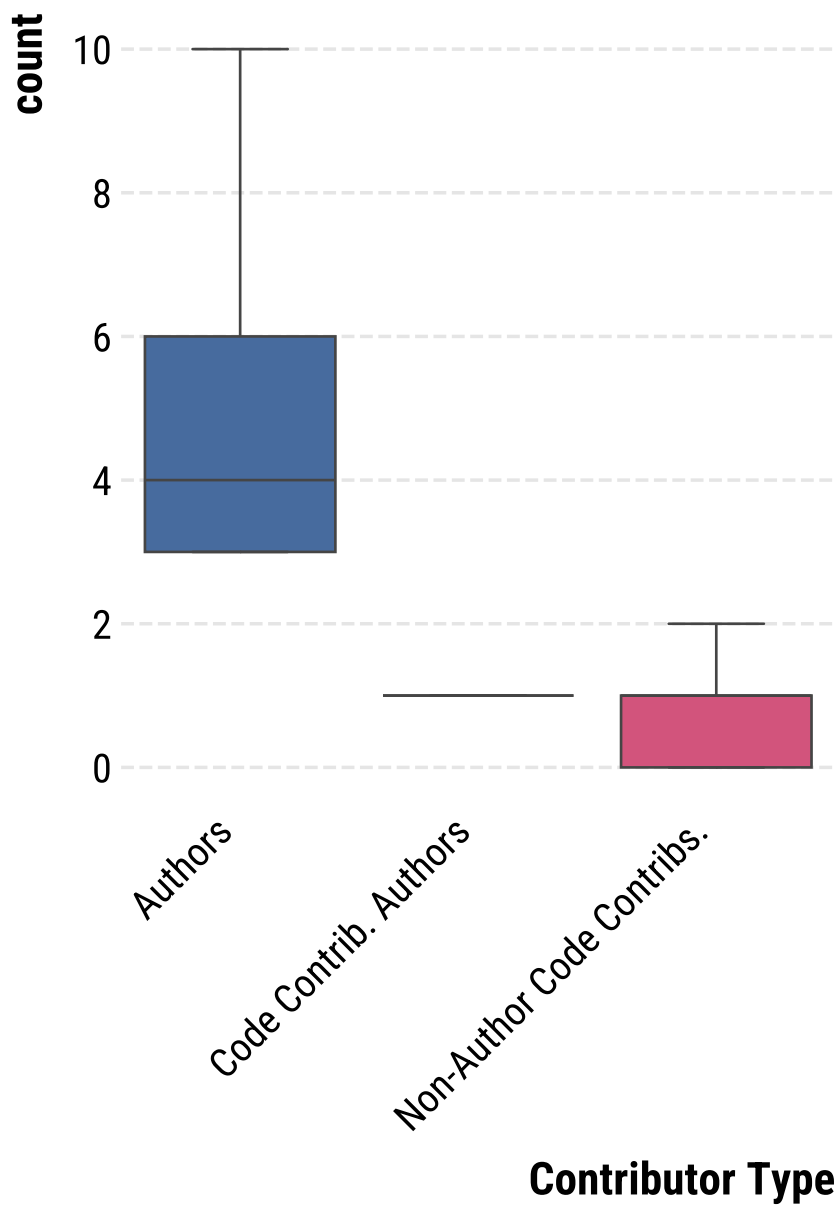
Figure 2: Number of articles by publication year. Only publication years with 100 or more articles are included.

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

- To enrich our pre-existing dataset, we apply our trained predictive model across pairs of authors and developer accounts.
 - again, these pairs are all combinations of author and developer account within an individual paper
 - specifics, how many unique author-developer account pairs are we able to find
 - table of author-developer account pairs for by data source / by field
 - we next use this enriched dataset to understand software development dynamics within research teams, and characterize the authors who are and who aren't code contributors.

5.1 Software Development Dynamics Within Research Teams

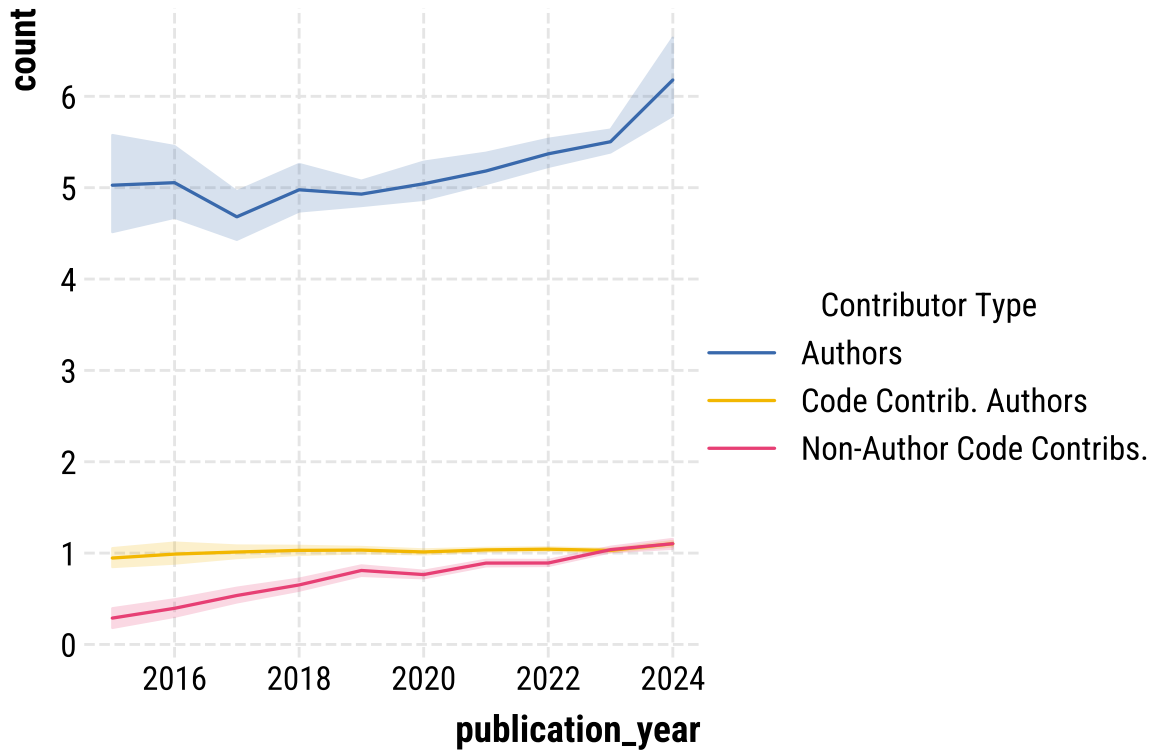
- We begin by measuring the distributions of different coding and non-coding contributors across all of the article-code-repository pairs within our dataset.
 - explain more, what are the different types of contributions? (coding contributor, coding-with-authorship contributor, non-coding-author, etc.)
 - what are the basics / what do we see across the board? What are the distributions of each of these contributor types
 - compare against analysis built on CRediT statements?
- Next we investigate if these distributions change over time, or, by “research team size”.
 - define research team size, in our case this is the total number of author-developers + non-coding authors + non-credited developers
 - plot the medians of the contributor type distributions over time (by publication year)
 - results in summary



Contributor Type	mean	std	min	25%	50%	75%	max
Authors	5.4	6.3	3	3	4	6	428
Code Contrib. Authors	1.0	0.7	0	1	1	1	19
Non-Author Code Contribs.	0.9	1.2	0	0	1	1	28

Table 3: Team Composition and Coding Status Counts Used in H1

Control	Subset	Authors	Code Cntrb. Auth.	Non-Auth. Code Cntrb.
OA Status	Closed	5.3 ± 3.4	1.1 ± 0.7	1.1 ± 1.2
	Open	5.4 ± 6.5	1.0 ± 0.7	0.9 ± 1.2
Domain	Health Sciences	7.7 ± 7.4	1.0 ± 0.6	0.7 ± 1.3
	Life Sciences	5.9 ± 9.8	1.0 ± 0.8	0.5 ± 1.0
	Physical Sciences	5.2 ± 5.9	1.0 ± 0.7	1.0 ± 1.3
	Social Sciences	4.7 ± 2.1	1.1 ± 0.7	0.6 ± 1.0
Article Type	preprint	5.1 ± 4.6	1.1 ± 0.8	1.1 ± 1.4
	research article	5.4 ± 6.6	1.0 ± 0.7	0.9 ± 1.2
	software article	5.0 ± 2.9	1.6 ± 1.5	0.2 ± 0.5



5.1.1 Modeling Citations

- We model an article's total citations by the coding contributorship of the research team and controlled by a number of different factors.
- Each control variable is modeled separately and the results are presented in the tables below:

Table 4: Article Citations by Code Contributorship of Research Team Controlled by Open Access Status

Variable	coef	P> z	[0.025	0.975]
const ***	0.71	0.00	0.58	0.84
n authors ***	0.01	0.00	0.00	0.01
n author devs ***	0.10	0.04	0.01	0.19
n non author devs ***	0.15	0.00	0.10	0.20
years since publication ***	0.39	0.00	0.38	0.39
is open access ***	0.48	0.00	0.35	0.61
n author devs * is open access	-0.05	0.32	-0.14	0.04
n non author devs * is open access	-0.01	0.83	-0.06	0.05

- Controlling for article open access status: Table 4
- Controlling for article domain: Table 5
- Controlling for article type: Table 6

5.1.1.1 Open Access Status

- open access articles
 - n_author_devs is not significant
 - gain 1.1621828452888712 more citations per non-author code contributor
- closed access articles
 - n_author_devs is not significant
 - gain 1.1681251177146603 more citations per non-author code contributor

5.1.1.2 Domain

- health sciences
 - gain 1.0989992577120393 more citations per author code contributor compared to no author code contributor health science papers
 - gain 1.0733665310933769 more citations per non-author code contributor compared to no non-author code contributor health science papers
- life sciences
 - n_author_devs is not significant
 - n_non_author_devs is not significant
- physical sciences

Table 5: Article Citations by Code Contributorship of Research Team Controlled by Domain

Variable	coef	P> z	[0.025	0.975]
const ***	1.20	0.00	1.08	1.31
n authors ***	0.01	0.00	0.00	0.01
n author devs	0.07	0.13	-0.02	0.16
n non author devs ***	0.07	0.00	0.03	0.12
years since publication ***	0.40	0.00	0.39	0.41
domain Life Sciences ***	-0.22	0.00	-0.35	-0.09
domain Physical Sciences	-0.03	0.65	-0.14	0.09
domain Social Sciences ***	-0.28	0.00	-0.43	-0.13
n author devs * domain Life Sciences	0.04	0.46	-0.07	0.15
n author devs * domain Physical Sciences	-0.03	0.51	-0.12	0.06
n author devs * domain Social Sciences	0.07	0.28	-0.05	0.18
n non author devs * domain Life Sciences	-0.04	0.18	-0.11	0.02
n non author devs * domain Physical Sciences ***	0.07	0.00	0.03	0.12
n non author devs * domain Social Sciences	0.02	0.67	-0.06	0.09

- n_author_devs is not significant
- gain 1.0836121025480543 more citations per non-author code contributor compared to no non-author code contributor physical science papers
- social sciences
 - n_author_devs is not significant
 - n_non_author_devs is not significant

5.1.1.3 Article Type

- preprint
 - n_author_devs is not significant
 - gain 1.132242296815202 more citations per non-author code contributor compared to no non-author code contributor preprints
- research article
 - gain 1.0934086297167769 more citations per author code contributor compared to no author code contributor research articles
 - gain 1.0381081810730037 more citations per non-author code contributor compared to no non-author code contributor research articles
- software article
 - n_author_devs is not significant

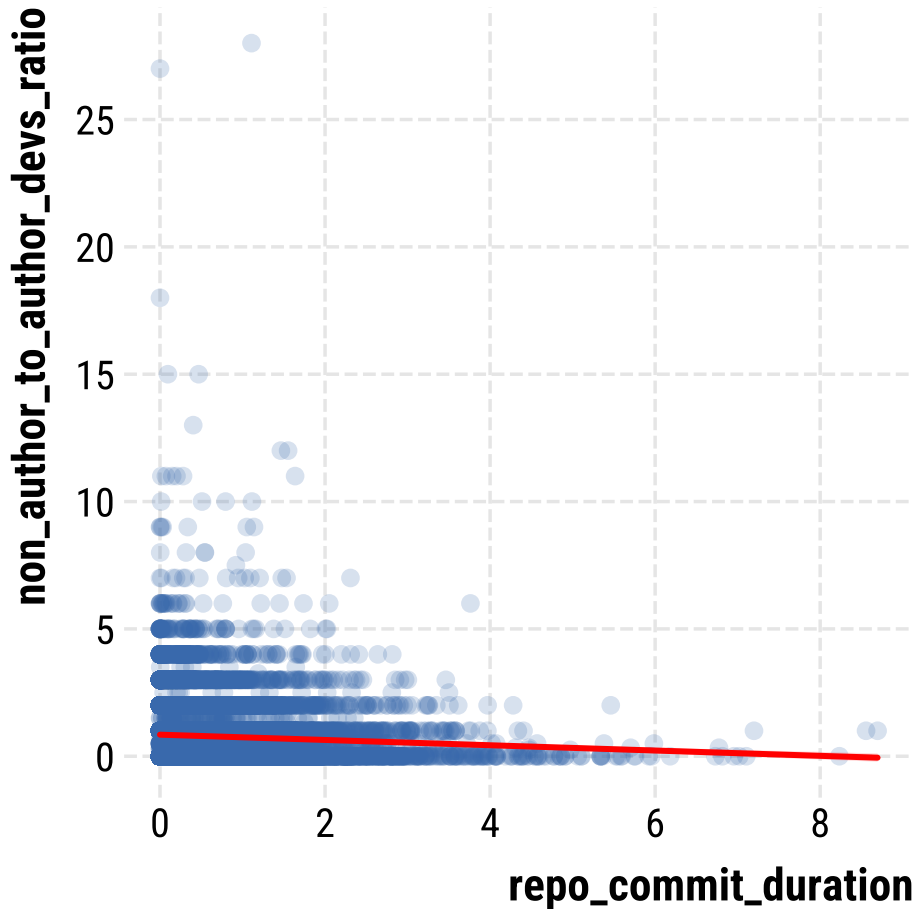
Table 6: Article Citations by Code Contributorship of Research Team Controlled by Article Type

Variable	coef	P> z	[0.025	0.975]
const ***	0.64	0.00	0.57	0.72
n authors ***	0.00	0.00	0.00	0.01
n author devs	-0.02	0.51	-0.06	0.03
n non author devs ***	0.13	0.00	0.10	0.16
years since publication ***	0.41	0.00	0.40	0.41
article type research article ***	0.52	0.00	0.44	0.59
article type software article ***	-0.41	0.00	-0.65	-0.17
n author devs * article type research article ***	0.09	0.00	0.04	0.14
n author devs * article type software article	0.06	0.31	-0.05	0.17
n non author devs * article type research article	0.03	0.07	-0.00	0.06
n non author devs * article type software article	0.11	0.45	-0.17	0.39

– n_non_author_devs is not significant

5.1.2 Team Composition and Project Duration

- We next investigate the relationship between the duration of a project and the coding contributorship of the research team.



PearsonRResult(statistic=-0.06420365596401129, pvalue=1.8808669891907014e-16)

5.2 Characteristics of Scientific Code Contributors

- Next we investigate the differences between coding and non-coding article authors.
 - specifics, author position in authorship list is a commonly used tool in scientometrics
 - similarly, metrics of “scientific impact” such as h-index, i10 index, and two-year mean citedness are also available to us.
 - plot / table of the distributions between coding and non-coding authors
 - ANOVA / Chi2 tests to see if these differences are significant
 - results in summary
- Just as before, we next investigate if these results are affected by article type and research domain.

- subplot + stats tests for differences by each article type
- subplot + stats tests for differences by each domain
- results in summary

5.2.1 Author Positions of Code Contributing Authors

Overall

is_code_contributor	False	True
position		
first	16428	35893
last	45829	4569
middle	131433	12920

Chi2: 86552.98658551606, p: 0.0, n: 247072

position: first, p: 0.0, statistic: 0.6860151755509261, n: 52321

position: middle, p: 0.0, statistic: 0.0895028160135224, n: 144353

position: last, p: 0.0, statistic: 0.09065835945870868, n: 50398

5.2.1.1 Domain

Physical Sciences

is_code_contributor	False	True
position		
first	13068	28919
last	36977	3433
middle	101231	10507

Chi2: 68220.51167362447, p: 0.0, n: 194135

position: first, p: 0.0, statistic: 0.6887608069164265, n: 41987

position: middle, p: 0.0, statistic: 0.09403246881096852, n: 111738

position: last, p: 0.0, statistic: 0.08495421925266024, n: 40410

Life Sciences

is_code_contributor	False	True
position		
first	1309	2586
last	3293	491
middle	11023	852

Chi2: 6554.038088460125, p: 0.0, n: 19554

position: first, p: 1.3228240814067576e-94, statistic: 0.6639281129653402, n: 3895

position: middle, p: 0.0, statistic: 0.07174736842105263, n: 11875

position: last, p: 0.0, statistic: 0.1297568710359408, n: 3784

Social Sciences

is_code_contributor	False	True
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position
first          1225   2813
last           3444    411
middle         8228   1021
Chi2: 5715.82059242655, p: 0.0, n: 17142
position: first, p: 2.2055993260095607e-141, statistic: 0.6966319960376424, n: 4038
position: last, p: 0.0, statistic: 0.1066147859922179, n: 3855
position: middle, p: 0.0, statistic: 0.11039031246621256, n: 9249
Health Sciences
is_code_contributor  False  True
position
first              826   1575
last              2115    234
middle            10951    540
Chi2: 5998.504729922407, p: 0.0, n: 16241
position: first, p: 1.8308248282746384e-53, statistic: 0.6559766763848397, n: 2401
position: middle, p: 0.0, statistic: 0.04699329910364633, n: 11491
position: last, p: 0.0, statistic: 0.09961685823754789, n: 2349

```

5.2.1.2 Article Type

```

research article
is_code_contributor  False  True
position
first              10141  21940
last              28279   2909
middle            82646   7345
Chi2: 54917.35337329746, p: 0.0, n: 153260
position: first, p: 0.0, statistic: 0.6838938935818709, n: 32081
position: middle, p: 0.0, statistic: 0.08161927303841496, n: 89991
position: last, p: 0.0, statistic: 0.09327305373861741, n: 31188
preprint
is_code_contributor  False  True
position
first              6102  13421
last              17105   1493
middle            47791   5134
Chi2: 31620.70287233051, p: 0.0, n: 91046
position: first, p: 0.0, statistic: 0.6874455770117297, n: 19523
position: last, p: 0.0, statistic: 0.08027744918808474, n: 18598
position: middle, p: 0.0, statistic: 0.09700519603212092, n: 52925

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```

software article
is_code_contributor  False  True
position
first                185    532
last                 445    167
middle               996    441
Chi2: 436.6562463251282, p: 1.5181016986238546e-95, n: 2766
position: first, p: 2.8268609500046396e-39, statistic: 0.7419804741980475, n: 717
position: last, p: 4.070740833733961e-30, statistic: 0.272875816993464, n: 612
position: middle, p: 4.079797536800063e-49, statistic: 0.3068893528183716, n: 1437

```

5.2.1.3 Open Access Status

```

Open Access
is_code_contributor  False  True
position
first                15323  33256
last                 42466  4290
middle               121386  12002
Chi2: 79588.66090114709, p: 0.0, n: 228723
position: first, p: 0.0, statistic: 0.684575639679697, n: 48579
position: middle, p: 0.0, statistic: 0.08997810897532012, n: 133388
position: last, p: 0.0, statistic: 0.09175293010522714, n: 46756
Closed Access
is_code_contributor  False  True
position
first                1105    2637
last                 3363    279
middle               10047    918
Chi2: 6990.309755750654, p: 0.0, n: 18349
position: first, p: 2.5647584282503314e-142, statistic: 0.7047033671833244, n: 3742
position: last, p: 0.0, statistic: 0.07660626029654036, n: 3642
position: middle, p: 0.0, statistic: 0.08372093023255814, n: 10965

```

5.2.2 Corresponding Status of Code Contributing Authors

```

Overall
is_code_contributor  False  True
is_corresponding

```

```

Corresponding          20112    8701
Not Corresponding       173578   44681
Chi2: 1421.1208069996246, p: 5.40576546314e-311, n: 247072
is_corresponding: Not Corresponding, p: 0.0, statistic: 0.2047154985590514, n: 218259
is_corresponding: Corresponding, p: 0.0, statistic: 0.301981744351508, n: 28813

```

5.2.2.1 Domain

```

Physical Sciences
is_code_contributor    False   True
is_corresponding
Corresponding          7239    5248
Not Corresponding       144037   37611
Chi2: 3086.5504905422213, p: 0.0, n: 194135
is_corresponding: Not Corresponding, p: 0.0, statistic: 0.2070543028274465, n: 181648
is_corresponding: Corresponding, p: 3.0892901003685015e-71, statistic: 0.42027708817169857, n: 218259
Life Sciences
is_code_contributor    False   True
is_corresponding
Corresponding          6247    1772
Not Corresponding       9378    2157
Chi2: 33.80559720544868, p: 6.090338640251236e-09, n: 19554
is_corresponding: Not Corresponding, p: 0.0, statistic: 0.1869960988296489, n: 11535
is_corresponding: Corresponding, p: 0.0, statistic: 0.2209751839381469, n: 8019
Social Sciences
is_code_contributor    False   True
is_corresponding
Corresponding          1655     803
Not Corresponding      11242   3442
Chi2: 95.74853878212862, p: 1.304427435637673e-22, n: 17142
is_corresponding: Corresponding, p: 2.2084447510143967e-67, statistic: 0.32668836452400324, n: 14684
is_corresponding: Not Corresponding, p: 0.0, statistic: 0.23440479433396894, n: 14684
Health Sciences
is_code_contributor    False   True
is_corresponding
Corresponding          4971     878
Not Corresponding      8921   1471
Coding by is_corresponding not significant

```

5.2.2.2 Article Type

```

research article
is_code_contributor  False  True
is_corresponding
Corresponding          19623   7716
Not Corresponding       101443  24478
Chi2: 1044.0017267689143, p: 4.898689779565618e-229, n: 153260
is_corresponding: Not Corresponding, p: 0.0, statistic: 0.1943917217938231, n: 125921
is_corresponding: Corresponding, p: 0.0, statistic: 0.2822341709645561, n: 27339
preprint
is_code_contributor  False  True
is_corresponding
Corresponding          264     772
Not Corresponding       70734  19276
Chi2: 1678.8685858544789, p: 0.0, n: 91046
is_corresponding: Not Corresponding, p: 0.0, statistic: 0.2141539828907899, n: 90010
is_corresponding: Corresponding, p: 2.737621889235552e-58, statistic: 0.7451737451737451, n: 43
software article
is_code_contributor  False  True
is_corresponding
Corresponding          225     213
Not Corresponding       1401     927
Chi2: 11.450236853781284, p: 0.0007148488745836787, n: 2766
is_corresponding: Corresponding, p: 0.5992164561363355, statistic: 0.4863013698630137, n: 43
is_corresponding: Not Corresponding, p: 1.5581829692815375e-22, statistic: 0.398195876288659

```

5.2.2.3 Open Access Status

```

Open Access
is_code_contributor  False  True
is_corresponding
Corresponding          19897   8448
Not Corresponding       159278  41100
Chi2: 1263.1447877833145, p: 1.1546882926708494e-276, n: 228723
is_corresponding: Not Corresponding, p: 0.0, statistic: 0.20511233768178144, n: 200378
is_corresponding: Corresponding, p: 0.0, statistic: 0.29804198271300053, n: 28345
Closed Access
is_code_contributor  False  True
is_corresponding
Corresponding          215     253
Not Corresponding       14300  3581
Chi2: 317.52534860502976, p: 5.011183853047046e-71, n: 18349

```

Table 7: 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
	last	2413	568	191	10188
	middle	12184	3327	701	31911
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

is_corresponding: Not Corresponding, p: 0.0, statistic: 0.20026844136233993, n: 17881

is_corresponding: Corresponding, p: 0.08709725881534564, statistic: 0.5405982905982906, n: 4

5.2.3 Modeling H-Index

- We model an authors total citations by their coding status and controlled by a number of different factors.
- Each control variable is modeled separately and the results are presented in the tables below:
 - Controlling for an authors most frequent author position: Table 8
 - Controlling for an authors most frequent domain: Table 9
 - Controlling for an authors most frequent article type: Table 10

5.2.3.1 Author Position

- first authors
 - any coding has a positive association with citations (1.2636444922077779)
 - majority coding has a negative association with citations (0.8860339595928756)
 - always coding has a negative association with citations (0.764143255648199)
- middle authors
 - any coding has a negative association with citations (0.7482635675785653)
 - majority coding has a negative association with citations (0.9039330328858641)
 - always coding is not significant

Table 8: 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

- last authors
 - any coding has a negative association with citations (0.8737159116880344)
 - majority coding is not significant
 - always coding is not significant

In general, any coding has a positive association while majority and always coding have negative associations with citations. “The more you code the less you are cited” – granted that we don’t have a lot of data for always coding authors (which itself backs up qual lit).

in general, coding is associated with about a ~10 - 30% decrease in citations for a number of conditions when compared to non-coding first authors.

5.2.3.2 Domain

- health sciences
 - any coding is not significant
 - majority coding has a negative association with citations (0.7527666447061963)
 - always coding has a negative association with citations (0.6120140740013499)
- life sciences
 - any coding is not significant
 - majority coding is not significant
 - always coding has a positive association with citations (1.5983949987546404)

Table 9: 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

- physical sciences
 - any coding is not significant
 - majority coding is not significant
 - always coding is not significant
- social sciences
 - any coding has a negative associate with citations (0.8033217181536265)
 - majority coding is not significant
 - always coding has a positive association with citations (1.587245303225596)

We couldn't get significant results for a majority of the groupings so I don't think that we can say anything too strongly, however, health sciences doesn't seem to favor coding (which I think is inline most with "RSE" dynamics). Social sciences is split, and social science is broad so this may be a "qual" vs "quant" split, if you include mixed methods researchers in there its hard to parse.

Physical sciences being entirely non-significant is interesting because a majority of our data comes from Physical sciences. this could indicate that coding is just a part of the culture and doesn't have a significant impact on citations (which is inline with CS being the bulk of our physical sciences data).

Table 10: 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

5.2.3.3 Article Type

- preprint
 - any coding is not significant
 - majority coding has a negative association with citations (0.6736800392488677)
 - always coding has a negative association with citations (0.6306526773980542)
- research article
 - any coding is not significant
 - majority coding has a positive association with citations (1.0565406146754943)
 - always coding has a positive association with citations (1.0843708965667604)
- software article
 - any coding has a positive association with citations (1.7471746543074462)
 - majority coding is not significant
 - always coding has a positive association with citations (1.4681454416819895)

Preprints (from arXiv) have a negative association with coding generally. Research articles have a very slim positive association with coding. Software articles have a strongly positive association with coding (unsuprising).

Table 11: 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

6 Appendix

6.1 Full Comparison of Models and Optional Features for Author-Developer-Account Matching

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