

Methods for measuring social and conceptual dimensions of convergence science

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Abstract

Convergence science is an intrepid form of interdisciplinarity defined by the US National Research Council as ‘the coming together of insights and approaches from originally distinct fields’ to strategically address grand challenges. Despite its increasing relevance to science policy and institutional design, there is still no practical framework for measuring convergence. We address this gap by developing a measure of disciplinary distance based upon disciplinary boundaries delineated by hierarchical ontologies. We apply this approach using two widely used ontologies—the Classification of Instructional Programs and the Medical Subject Headings—each comprised of thousands of entities that facilitate classifying two distinct research dimensions, respectively. The social dimension codifies the disciplinary pedigree of individual scholars, connoting core expertise associated with traditional modes of mono-disciplinary graduate education. The conceptual dimension codifies the knowledge, methods, and equipment fundamental to a given target problem, which together may exceed the researchers’ core expertise. Considered in tandem, this decomposition facilitates measuring social-conceptual alignment and optimizing team assembly around domain-spanning problems—a key aspect that eludes other approaches. We demonstrate the utility of this framework in a case study of the human brain science (HBS) ecosystem, a relevant convergence nexus that highlights several practical considerations for designing, evaluating, institutionalizing, and accelerating convergence. Econometric analysis of 655,386 publications derived from 9,121 distinct HBS scholars reveals a 11.4% article-level citation premium attributable to research featuring full topical convergence, and an additional 2.7% citation premium if the social (disciplinary) configuration of scholars is maximally aligned with the conceptual (topical) configuration of the research.

Keywords: convergence; team science; team assembly; ontology; interdisciplinary distance; alignment.

The scientific frontier is increasingly characterized by domain-spanning problems calling for the strategic integration of disparate domains of expertise to strategically address high-stakes challenges faced by society (Helbing 2012, 2013; Petersen, Ahmed and Pavlidis 2021). In response, the convergence science paradigm—defined by its originators as ‘the coming together of insights and approaches from originally distinct fields’ (National Research Council 2014)—has emerged as an organizational model constructed around a mission-oriented agenda that promotes social-engineering to fortify existing interdisciplinary approaches to addressing boundary-spanning grand challenges (NSF, accessed February 2021). With team science becoming the predominant mode of knowledge production (Wuchty, Jones and Uzzi 2007; Börner et al. 2010; Pavlidis, Petersen and Semendeferi 2014; Petersen, Pavlidis and Semendeferi 2014), convergence represents a holistic strategy for harnessing social and conceptual diversity, and for accelerating action on multi-dimensional problems (Page 2008; Linkov, Wood and Bates 2014; Pavlidis, Akleman and Petersen 2022). Specific examples include deforestation and illicit wildlife trade (Di Minin et al. 2018; Arroyave et al. 2020, 2021), two wicked problems that span sociocultural, technological, political, and environmental dimensions (Orsatti, Quatraro and Pezzoni 2020).

Even in the best-case scenario, where traditional mono-domain approaches exist that address certain facets of the target

problem, convergence is needed to address the multi-dimensionality of such problems, as partial solutions are likely to be fragmented and all together incomplete (Linkov, Wood and Bates 2014). As such, designing and assembling a complete and feasible composite solution is a principal barrier to addressing grand challenges. Another reason multi-dimensional problems call for convergence is due to the intrepid interdisciplinary distances commonly entailed, which can alter the required assumptions and generalizability of mono-domain approaches. All together, the integration of disparate disciplines and their specialized capabilities is unlikely to be straightforward or clear. However, by extending principles of recombinant innovation (Weitzman 1998; Fleming 2001; Orsatti, Quatraro and Pezzoni 2020) to social-engineering contexts, effective multidisciplinary integration can be achieved by repurposing and reconfiguring of disparate elements—such as scholars of varying expertise, and conceptual theories and methods—into a configuration that represents a specific strategy (a key) that sufficiently satisfies the constraints associated with all facets of the domain-spanning problem (the lock). For this reason, exploiting diversity also serves a valuable hedge against the uncertainty inherent in exploring the space of relevant and accessible social and conceptual recombinations (Fleming 2004; Orsatti, Quatraro and Pezzoni 2020; Petersen 2022).

Owing to these considerations, the application of convergence science to domain-spanning problems can clearly be

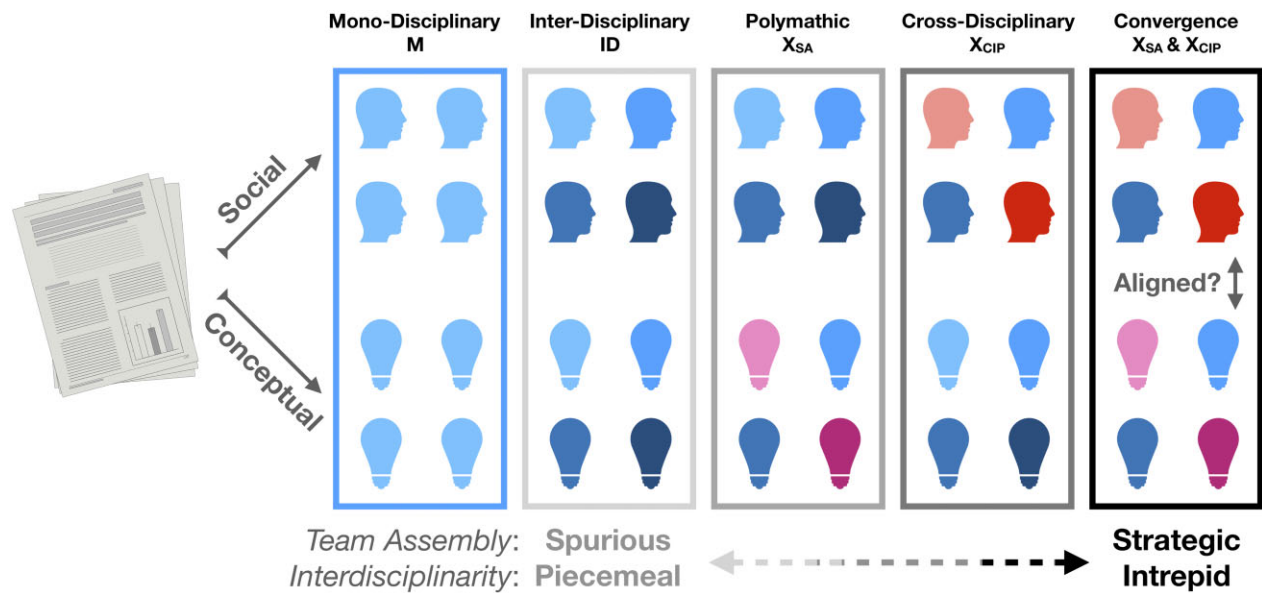


Figure 1. A definition of convergence by way of its deconstruction along social and conceptual dimensions. Convergence is achieved by way of strategic team assembly that leverages synergies among originally distinct domains to address specific target problems defined by a mission-oriented agenda. According to this definition, convergence is an intrepid form of interdisciplinarity, distinguished from more happenstance manifestations of piecemeal integration, and instead requiring a consistent and robust measurement framework that facilitates evaluating (mis)alignment across both social and conceptual dimensions (Wagner et al. 2011), which is an essential task of research project selection, evaluation, and assessment. Accordingly, in this schematic, the different shades of a common color base indicate neighboring domains characterizing piecemeal diversity, whereas different color bases (e.g. red, blue, magenta) indicate more intrepid configurations spanning distinct social and conceptual domains. Two partial modes of convergence can thus be codified and identified (Petersen et al. 2018; Petersen, Ahmed and Pavlidis 2021; Pavlidis, Akleman and Petersen 2022; Yang, Pavlidis and Petersen 2023): (1) *polymathic* research (represented as X_{SA}) integrates distant concepts and methods by way of expansive learning by a team featuring more narrow disciplinary diversity and (2) conversely, *cross-disciplinary* research (X_{CIP}) features multi-disciplinary teams focusing on problems spanning a relatively narrow conceptual scope. According to this framework, complete convergence ($X_{SA\&CIP}$) incorporates both modes of cross-domain integration, with the additional requirement of evaluating the quality of alignment between the social and conceptual configurations. Conceptualized as such, distinguishing $X_{SA\&CIP}$ from IDR requires operationalizing a distance between social and conceptual entities defining a given research agendas and its output, which is the main methodological contribution of this work.

science, and a myriad of other sciences (Dzau and Balatbat 2018). And we conclude with outlook and policy recommendations addressing several practical considerations associated with designing, evaluating, institutionalizing and accelerating convergence.

Background and motivation

Convergence science—a mission-oriented paradigm for addressing transdisciplinary grand challenges

While the structure of convergence is multi-dimensional (social and conceptual), the institutional scope of convergence is multi-level. At the highest level of aggregation are national innovation systems characterized as configurations of industry, university, and government that leverage cross-sectoral (e.g. triple-helix) synergies, whereby a common agenda that respects individual prerogatives and practices can be established around a common objective to promote economic growth (Leydesdorff and Etzkowitz 1996; Etzkowitz and Leydesdorff 2000; Stephan 2012). Such strategic synergies are also important to endeavors less appreciated as innovation and growth oriented, yet still relying on cross-sectoral knowledge co-production, such as protected area land management for preserving critical ecosystems (Arroyave et al. 2022). This model of integration-mediated innovation is readily extended to other domains of knowledge production that organize around the principle triple-helix components—namely, demand for solutions (applications), supply of knowledge

(theory), and techno-socio-political capabilities (catalysts). For example, the biomedical health sector can be cast as a triple-helix forming around a disease, drug, and techno-informatic capabilities (Petersen, Rotolo and Leydesdorff 2016; Yang, Pavlidis and Petersen 2023), yielding breakthrough successes in the last two decades ranging from the map of the human genome (Petersen et al. 2018) to rapid development of COVID-19 mRNA vaccines; more prospective examples include the coming era of bio-mechanics and human-machine systems (Kose and Sakata 2019; Pavlidis, Akleman and Petersen 2022). Principles and practices of convergence science can be applied to ongoing efforts to integrate non-STEM fields, such as humanities and arts, as demonstrated by STEAM and digital humanities initiatives that exemplify the integration of ‘soft’ and ‘hard’ methodologies (Pedersen 2016).

It is also notable that disciplinary convergence has a long-standing role as the counter-balance to divergence (Roco et al. 2013; Ballelli, Mäs and Helbing 2015; Watson 2017; Pavlidis, Akleman and Petersen 2022). Yet the transition toward convergent problem solving has become integral to national innovation systems charged with developing mission-oriented agendas and policy (Fealing 2011; Wanzenböck et al. 2020). Convergence has been championed in the last decade by the US NSF, specifically the Office of Integrative Activities (OIA), which aims to accelerate problem-solving around specific target areas characterized by grand societal challenges (Helbing 2012) by calling for strategic collaboration across disciplines and sectors (National

target molecules in the sample') represent altogether different approaches to operationalizing biological research—the former representing *in silico* and the latter being *in vitro* approaches. One also encounters various approaches in the IDR literature for measuring disciplinary diversity as either variety, balance and/or disparity, or as a combination (Harrison and Klein 2007; Stirling 2007; Rafols and Meyer 2010).

Another issue is that the underlying data used to quantify variety, balance and disparity are typically derived from ‘flat’ (i.e., non-hierarchical) classification systems that were originally designed for altogether distinct library science objectives, namely cataloguing journals. The main advantage of flat classification systems is they are simple. And because there is no structure associated with the categories, the category systems can be readily extended without having to also amend the relationships between categories.¹ As such, these flat classification systems are conveniently available in large publication indices such as Clarivate Web of Science (WOS) and Scopus (namely, the WC and SU fields in the former index, and the ‘Subject Areas’ field in the latter). And while the research area (SU) classification does feature entities grouped according to five broad categories (Arts and Humanities; Life Sciences and Biomedicine; Physical Sciences; Social Sciences; Technology), these categories lack the granularity needed to establish distances in-between the two extremes of heterotype and homotype. Because most classification systems lack the requisite resolution for delineating more nuanced disciplinary boundaries, they are not appropriate for measuring convergence. One manifestation of this inadequacy is the ‘multidisciplinary’ category applied when a type does not fit neatly into an existing category, which is increasingly common, and speaks to the relevance of identifying a more robust and objective approach to identifying boundary-spanning configurations. In order to measure the similarity or distance between categories, metrics such as the Stirling Index and other variants call for ad hoc assignment of a distance d_{ij} between any two categories i and j (Stirling 2007; Leydesdorff, Wagner and Bornmann 2018, 2019). Nevertheless, without subjective manual grouping of journal categories into disciplinary clusters, there is no objective rubric for identifying whether a given journal category combination represents convergence.

This may be the most critical disadvantage of commonly used classification systems—i.e., their intended design for classifying journals, and not individual research articles (Boyack and Klavans 2011). In addition to lacking information regarding the distance between different categories, flat classifications tend to over-generalize disciplinary content. This issue was demonstrated in a study by Leydesdorff and Opthof (2013) showing that research published on a very narrow conceptual topic ‘Brugada Syndrome’ nevertheless maps onto 24 different WC. This example shows how WC lack information specifying relationships between categories that could be used to counter the tendency for category diversification, which is an advantage of relational ontologies. Moreover, the vast majority of journals, and hence all articles published by that journal, are classified by a single category (see [Supplementary Appendix](#) for specifics on WOS), and the assignment of which has been criticized as being subjective (Boyack, Klavans and Börner 2005; Rafols and Leydesdorff 2009; Rafols and Meyer 2010; Leydesdorff, Wagner and Bornmann 2018). Consequently, these systems lack sufficient

resolution to distinguish piecemeal interdisciplinary combinations from more intrepid cross-domain combinations.²

Indeed, if the technical objective is to classify and compare the content of individual research articles, then article-level keywords are more appropriate. Extant methods to define a keyword concept space include externally defined dictionaries (Leahey and Moody 2014) and clustering title and abstract words using natural language processing (Mane and Börner 2004). Ideally, article-level keywords are assigned based upon the article content only and are not conditioned by other information such as the journal; see Shu et al. (2019) on the differences between journal and article-level classifications. Another necessity is that keyword dictionaries be standardized—such as the ‘Keywords Plus’ (ID) recorded in WOS annotations, as opposed to alternative author-defined keywords (DE)—so that they are not subject to assignment idiosyncrasies associated with author, discipline, language, and international context. And as above with journal categories, if the organizational structure of the standardized keywords is flat then they also offer limited ability to measure cross-domain integration. Bibliographic coupling and keyword clustering approaches may provide a step in the right direction to develop measures of (dis)similarity by identifying topical or social groups based upon co-occurrence statistics (Velden et al. 2017).

A final limitation regarding the scope of approaches used in extant IDR literature is the relatively narrow focus on conceptual components (commonly identified by way of keywords or journal classifications) (Wagner et al. 2011)—as opposed to its social components. In what follows, we develop a parallel classification of social dimensions as informed by authors' departmental affiliations, which connotes scholars' particular domains of core training and expertise. Alternative approaches might involve classifying individuals according to their PhD field. Yet, because hiring culture in traditional academic settings has reinforced longstanding disciplinary identities connoted by departments, an author's departmental affiliation is likely to highly correlate with their PhD field, and so these two approaches are likely to yield the same insights. Such intra-disciplinary hiring bias is consistent with the strong role of prestige-oriented in-group sorting in faculty hiring (Wapman et al. 2022). Independent of the methodology for classifying social components, there are relatively few studies that systematically construct disciplinary categories based upon author attributes (see for example Qiu 1992; Qin, Lancaster and Allen 1997; Schummer 2004; Abramo, D'Angelo and Di Costa 2017; Petersen et al. 2021, 2018). One reason for this is the difficulty in obtaining and classifying unstandardized author affiliation metadata. As a result, many studies that take this approach are limited in data sample size (Schummer 2004; Wagner et al. 2011).

Advantages of hierarchical ontologies for defining distinct disciplinary domains

We address the aforementioned issues associated with defining distinct disciplinary domains by leveraging two existing hierarchical ontologies used to classify the social and conceptual dimensions of research, respectively: (1) the Classification of Instructional Programs (CIP) ontology comprised of 2,100+ educational program types, useful for classifying authors' departmental affiliations; and (2) the Medical Subject Heading (MeSH) ontology comprised of 30,000+

individual keywords spanning a wide range of biological and medical concepts. Both ontologies offer varying depth resolution due to their hierarchical design, and are sufficiently broad to support the evaluation of nearly all the 10 challenge areas listed above. For example, the CIP ontology includes a branch dedicated to ‘Multi/Interdisciplinary Studies’, including but limited to ‘Science, Technology and Society’, and ‘Data Science’. Similarly, in addition to core biomedical and health concepts, MeSH also includes equipment, technology, methods, and other far-reaching entities representing intersections with other domains, such as ‘sustainable development’ and ‘algorithms’. See the [Supplementary Appendix](#) regarding the scope and limitations of these ontologies.

Another advantage of thesaurus and entity-oriented ontologies, examples including MeSH and PhySH, is they can readily be combined by way of advanced alignment techniques ([Wang et al. 2018](#)), since they are comprised of objective entities as opposed to subjectively defined and broad categories. Notably, the PhySH ontology has replaced the longstanding PACS system used for decades to classify physics research ([Smith 2019](#)). Hence, the foresight of ontological design supports the generalizability and extendibility of our framework beyond the ontologies developed in what follows. As such, building on recent efforts ([Petersen et al. 2021](#); [Yang, Pavlidis and Petersen 2023](#)), we use this convergence framework to develop useful methods for representing, visualizing, and quantifying convergence as cross-domain integration—corresponding to multidisciplinary integration if the domains being considered are disciplines; or epistemological integration if the domains correspond to research concepts.

Methods

Hierarchical CIP and MeSH ontologies for representing social and conceptual dimensions of research

The measurement of convergence requires a measure of a distance between any two given entities. As such, the first methodological imperative is to be able to identify whether two given entities are neighboring variants or sufficiently distant to qualify as belonging to ‘originally distinct’ domains ([National Research Council 2014](#)).

[Figure 2A](#) is a schematic that illustrates our method that leverages existing hierarchical ontologies to differentiate whether two concepts (alternatively departments) belong to the same or to distinct subject areas (respectively, disciplines). Neighboring and distinct domains are clearly delineated by the hierarchical structure of the ontology, and depend on the selection of an aggregation level parameterized by a level cut. The schematic shows a L_2 level cut, which thereby defines members of subgroups and establishes a first approximation of a metric distance according to the ontological lineage. Level cuts at higher levels of the hierarchy yield more distinct domains. The choice of the level cut facilitates variable domain resolution scales, e.g. see [Yang, Pavlidis and Petersen \(2023\)](#) for comprehensive historical MeSH co-occurrence analysis at both L_1 and L_2 levels.

The hierarchical structure facilitates aggregating counts for entities located above the level cut into the counts for their parent entity: for example, the hypothetical keyword 1.1.2.3 and 1.1.2 would both be counted as entity 1.1 for a level cut at L_2 . Similarly, 1.1.2.3 and 1.1.3.3 would also be aggregated

for a L_2 cut, but would be counted separately for a L_3 cut. This ability to merge entity counts facilitates establishing a weighted content representation, which is another advantage of our method.³

Based upon their locations in the ontology, the distance between any two entities can be objectively defined as neighboring (mono-domain) or distant (cross-domain). By way of example, our schematic illustrates how classifying all types within the hypothetical ontology according to the L_2 level produces six distinct sub-domains: category types 1.1 and 1.1.2 and 1.1.2.3 would all be classified as type 1.1, and all these types would be considered different than entities belonging to 1.3 (e.g. 1.3.1, 1.3.2, and so on). In this way, a hierarchical ontology yields a flexible basis set that offers a weighted vector representation of the conceptual (or alternatively, social) dimensions of a research article, as illustrated in [Figure 2B](#).

[Figure 3](#) illustrates two existing ontologies, one social and one conceptual, applied in recent research ([Petersen et al. 2021, 2018, 2016](#); [Yang, Pavlidis and Petersen 2023](#)). In the case of the social dimension, we use the CIP ontology maintained by the US National Center for Education Statistics ([National Center for Education Statistics 2022](#)), which was developed for classifying instructional degree-granting programs for programmatic certification and assessment. Because faculty departments are typically strongly aligned with the degree-granting educational programs they offer, the CIP ontology is useful for measuring distances between disciplines as proxied by scholars’ departmental affiliations, which can be inferred by information listed in an article byline or on their faculty home page or departmental home page. And with thousands of categories, the CIP ontology is sufficiently comprehensive to span the entire space of author affiliations. Also, a researcher could in principle have multiple primary affiliations, which could map onto two or more CIP categories. Such instances are likely to be exceptional corner cases where the individual represents a cross-disciplinary or hybrid scholar. However, if we restrict our annotation to primary affiliations (e.g. excluding courtesy appointments and external institutional affiliations), then such cases are likely to also be exceptional. Regardless, for each research article p (or any other research output, such as a patent or grant proposal) one obtains a count vector $v_{CIP,p}$ by aggregating the CIP counts across the set of coauthors.

In the case of the conceptual dimension, we use the *Medical Subject Heading (MeSH)* ontology maintained by the US National Library of Medicine ([US National Library of Medicine 2022](#)), which is presently comprised of more than 30,000 MeSH terms organized in a 13-level hierarchical knowledge network that spans a number of distinct conceptual domains that are concentrated upon, but not limited to, biological, health, and medical subject areas ([Yang, Pavlidis and Petersen 2023](#)). Individual MeSH are assigned to articles indexed within the PubMed index by professional annotators using algorithmic assistance, which connote the principal entities entailed by the research, such as diseases, chemicals, syndromes, methods, equipment, etc. From 2022 onwards, annotation has become increasingly automated by way of the Medical Text Indexer algorithm (MTIA). MeSH is constructed according to a thesaurus, such that different terms map onto a single MeSH according to the variant ‘Entry Terms’ specified in each MeSH’s description page. Hence, the MeSH ontology corrects for multiple and ambiguous

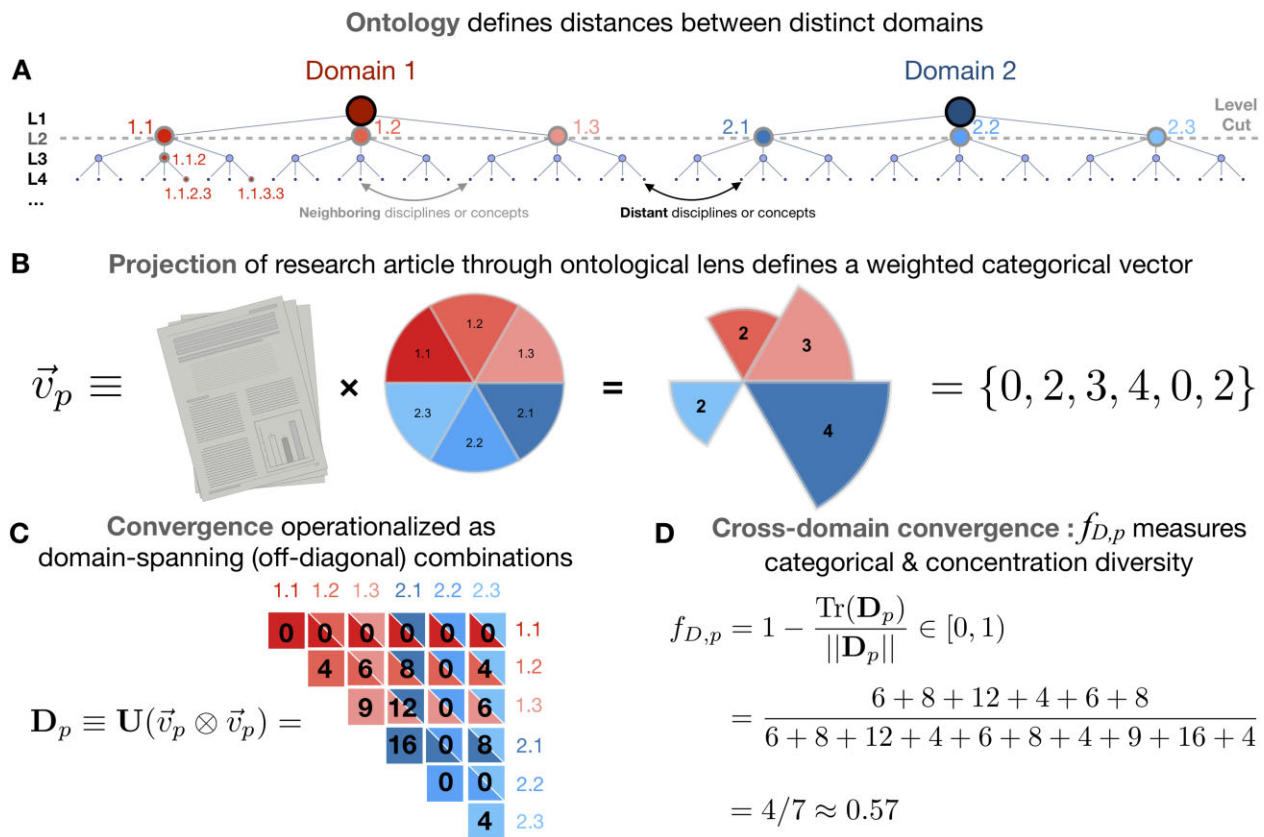


Figure 2. Measuring convergence as cross-domain integration. (A) Identifying boundaries that define distinct domains is an important step toward developing a metric distance between entity types, e.g. concepts or disciplines in the present case. To this end, the boundaries explicitly delineated by hierarchical ontologies are useful for classifying entities as neighboring variants, or alternatively, sufficiently distant so as to be considered ‘originally distinct’—a dichotomy required for evaluating convergence (National Research Council 2014). (B) Projecting research outputs (e.g. a publication or patent or grant proposal, generically denoted by the index p), against a consistent framework yields the type-count vector v_p . (C) As a combinatorial construct, we systematically measure convergence by way of the outer-product matrix that tabulates the proportion of each cross-domain combination (corresponding to off-diagonal matrix elements), relative to mono-domain concentrations (diagonal elements). (D) Illustration of simple steps to quantify convergence according to $f_{p,p}$ defined in Equation (1), which accounts for both categorical variation and concentration disparity (Harrison and Klein 2007).

meanings of individual descriptors. As in the case of multiple affiliations above, when a single MeSH maps onto multiple L_1 MeSH branches (a relatively infrequent case, corresponding to 6% of all MeSH (Yang, Pavlidis and Petersen 2023)), the ontology provides a systematic way for identifying and managing these edge cases. As such, each individual MeSH keyword is classified according to a given topical domain, which we call a subject area (SA). We then combine all the MeSH counts into a count vector $v_{SA,p}$ for each article.

These examples highlight one of the advantages of hierarchical ontologies, namely they support identifying particular cross-domain configurations to be evaluated, which can be specified by the manual merging of distinct domains into a ‘super-group’. By way of example, the color scheme in Figure 3 illustrates a manual merging of L_1 categories into three ‘super-group’ or L_0 domains: Health (purple); Science Technology Engineering and Mathematics (STEM; green); and Social Sciences, Humanities and Arts (SSHA; orange).

Measuring convergence according to boundary-spanning configurations

Because convergence is a fundamentally combinatorial construct, we operationalize it by tabulating all pairwise cross-domain combinations occurring in a given p . [Figure 2C](#) illustrates how all pairwise combinations can be represented

by the tensor-product matrix $D_p = U(v_p \otimes v_p)$, where U represents an operator that selects the upper-triangular matrix elements, since convergence is operationalized as categorical combinations as opposed to permutations. Each matrix element $D_{ij} = v_i \times v_j$ corresponds to a simple Hadamard product of the corresponding vector elements (for $j \geq i$; conversely, $D_{ij} = 0$ for $j < i$, according to the arbitrary choice of U to correspond to upper- as opposed to lower-triangular elements). Rather intuitively, elements along the diagonal of D_p capture the relative weight of intra-domain combinations, whereas the off-diagonal elements capture cross-domain combinations. Additional higher-level organization, e.g. L_1 information encoded in the schematic as red and blue color schemes, can be inferred according to the location of each domain within the ontology.

Figure 2D illustrates a straightforward and intuitive measure of convergence, calculated as the relative contribution to D_p by off-diagonal elements, given by the fraction

$$f_{D,p} = 1 - \text{Tr}(\mathbf{D}_p) / \|\mathbf{D}_p\| \quad (1)$$

where $\text{Tr}(D_p)$ indicates the matrix trace corresponding to the sum of the diagonal elements, and $\|\dots\|$ indicates the matrix total calculated by summing across all matrix elements. This measure is standardized in that its upper and lower limits are

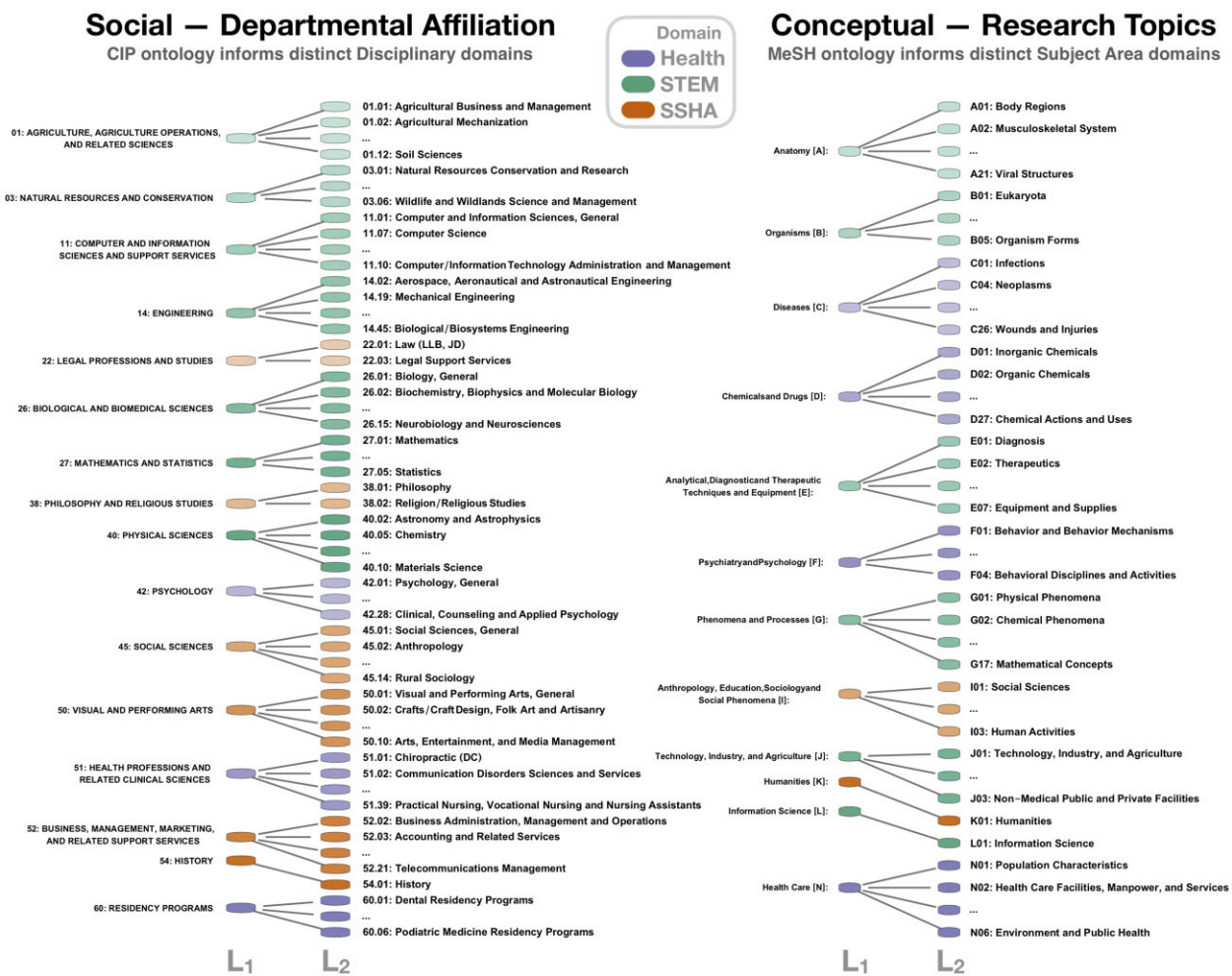


Figure 3. Social and conceptual ontologies for measuring cross-domain convergence. Shown is only a small portion of the (left) CIP ontology maintained by the US National Center for Education Statistics (National Center for Education Statistics 2022); and (right) Medical Subject Heading (MeSH) ontology maintained by the US National Library of Medicine (U.S. National Library of Medicine 2022). Ellipses indicate the L_2 categories that are too numerous to show (connoting 1,000 s of CIP and MeSH not shown), and extend each ontology across a broad scope that covers nearly all social and conceptual domains. The entire CIP ontology can be explored here: <https://nces.ed.gov/ipeds/cipcode/browse.aspx?y=56>. See the [Supplementary Appendix](#) regarding the limitations and scope of these ontologies. For each shown entity (ellipses connote 1,000 s of CIP and 30,000 s MeSH not shown), we manually classified the parent (L_1) category according to three distinct ‘super-group’ domains: Health science (purple); STEM (green); and Social Sciences, Humanities and Arts (orange); variable color tones are provided as visual aid for distinguishing distinct categories at higher branch level cuts. Depending on the convergence resolution being considered, the choice of level cut separates the ontology into various domains. For example, a partition according to the tripartite super-group (L_0) implies that Materials Science and Statistics are neighboring disciplines; however defining boundaries according to L_1 means these two disciplines are considered distinct; and for both L_0 and L_1 partitions, Aerospace and Biological/Biosystems are neighboring Engineering sub-disciplines.

bounded, $0 \leq f_{D,p} < 1$. Mono-domain publications, yielding v_p with just a single non-zero element located on the matrix diagonal, correspond to $f_{D,p} = 0$. Contrariwise, in the case of a uniform distribution, when all vector elements having the same value, the measure records the maximum value, $f_{D,p} = (d - 1)/(d + 1) \approx 1$, where d is the number of distinct domains within the ontology and thus the dimensionality of v_p .

As formulated, $f_{D,p}$ is a Blau-like measure of both categorical variety and concentration disparity (Harrison and Klein 2007), since by construction the number of distinct domains is fixed by the level cut applied to the hierarchical ontology. As such, $f_{D,p}$ increases as the number of distinct domains represented by p increases; it also increases as the parity in weight values encoded in v_p increases. There are various alternative diversity measures employed in the scientometrics of IDR

(Stirling 2007; Leydesdorff, Wagner and Bornmann 2018, 2019), the most similar being the ‘Stirling Index’ Δ_p (Stirling 2007), which requires an ad hoc prescription of a distance d_{ij} between any two categories i and j . By comparison, our method uses the hierarchical ontology to organically define all d_{ij} . While it is neither our objective nor our interest in comparing or establishing the superiority of these various diversity measures, this definition opens the possibility for exploring the utility of these other diversity measures by leveraging the d_{ij} encoded in the ontology.

In summary, we chose the tensor-product formulation of D_p to capture the combinatorial features of convergence. We anticipate that higher-order matrix decomposition methods and measures, such as the distribution of eigenvalues of D_p , will reveal new insights into the structure and dynamics of convergence. For example, recent work analyzing $f_{D,p}$

calculated across the entire 21.6 million research articles indexed within PubMed over the period 1970–2018 shows a steady increase in conceptual convergence over the last half century, with wide levels of variation across individual journals likely attributable to the propensity for different scholarly communities to support convergence science (Yang, Pavlidis and Petersen 2023).

HBS research corpus

The broad frontier of HBS is an appropriate testbed for developing this convergence measurement framework given that it represents domain-spanning research in the core biological sciences (physiology of structure, function, and evolution), the behavior and public health sciences, and also relies of advanced medical imaging technologies, as well the cognitive science of intelligence, artificial, and natural. It is also a relevant area to study given several ongoing national funding initiatives such as BRAIN in the USA and the Human Brain Projects in Europe, which are on the order of a billion US\$ in total funding size.

To demonstrate the practical application of this framework, we constructed a comprehensive scholar-centric representation of the HBS ecosystem by collecting and merging publication data from Web of Science (WOS), Scopus, and PubMed. The former dataset was used to identify articles associated with the topic field query ‘Human Brain’ from the WOS Core Collection over the period 1955–2016. This initial search returned 224,201 publication records. From this set we identified the full first and last names of all authors with ≥ 5 publications, including their most recent affiliation. To address the name disambiguation problem, we then used the Scopus Author API to identify 9,121 distinct HBS profiles over the period 1945–2018. We manually classified each Scopus Author’s affiliation according to 9 CIP groups: (1) Neurosciences, (2) Biology, (3) Psychology, (4) Biotech and Genetics, (5) Medical Specialty, (6) Health Sciences, (7) Pathology and Pharmacology, (8) Engineering and Informatics, and (9) Chemistry, Physics, and Math. The collection of CIP categories across all coauthors thereby define the social dimensions of a given article. Similarly, in order to define the conceptual dimension of HBS research, we matched each Scopus record to its PubMed entry in order to obtain the set of MeSH for each article. The final dataset is comprised of 655,386 research articles systematically classified according to the MeSH and CIP ontologies. For more dataset construction details, including the open dataset (see [Petersen et al. 2021](#); [Pavlidis and Zhukov 2022](#)).

To measure each article’s research impact through late 2019, we obtained the citation count $c_{p,t}$ for each article p published in year t using the Scopus API. Because nominal citation counts suffer from systematic temporal bias (Petersen et al. 2018), in what follows we use a normalized citation measure denoted by

$$z_{p,t} = (\ln(c_{p,t} + 1) - \mu_t) / \sigma_t, \quad (2)$$

where $\mu_t \equiv \langle \ln(c_t + 1) \rangle$ is the mean and $\sigma_t \equiv \sigma[\ln(c_t + 1)]$ is the SD of the citation distribution for a given t ; we add 1 to $c_{p,t}$ to avoid the divergence of $\ln 0$ associated with uncited publications—a common method which does not alter the interpretation of results. Consequently, the normalized citation measure z_p is a robust measure that is well-fit by the Normal

$N(0, 1)$ distribution, independent of t ; see [Petersen et al. \(2021\)](#) for a demonstration of this statistical stationarity. Publications with $z_{p,t} > 0$ (respectively, $z_{p,t} \leq 0$) can be collected by year into above-average (respectively, below-average) article groups. The scale of the logarithmic citation distribution σ_t is also relatively stable over the 49-year period 1970–2018 (average and SD value are $\langle \sigma \rangle \pm \text{SD} = 1.24 \pm 0.09$). Hence, in our regression model that follows, we model the dependent variable $Y \equiv z_{p,t}$ and estimate the regression coefficient β_x associated with an independent variable x that ranges between 0 and 1. As elaborated in [Petersen et al. \(2021\)](#), the percent change in citations $c_{p,t}$ associated with the variable x shifting from 0 to 1 is $100\Delta c_p / c_p \approx 100\langle \sigma \rangle \beta_x$.

Results

Evaluating the prevalence of different convergence science modes

This convergence framework facilitates a variety of novel perspectives in the science of science and science policy (Fealing 2011; Fortunato et al. 2018) for understanding how and when convergence emerges, and for also detailing the structural properties of nascent integration interfaces. Consider, for example, patterns of cross-disciplinary collaboration of scholars from two or more originally distinct domains. From this perspective, social convergence is highly dynamic, involving both the entry and exit of scholars into the interface between domains. Such a convergence nexus is seeded by individuals and their social interactions. An example of the former is the cross-disciplinary mobility of a scholar from one domain to another; an example of the latter is the formation of a potent cross-disciplinary collaboration between scholars. Either scenario can give rise to a persistent interface that accelerates the cross-pollination of theory, methods, and culture—a scenario that typifies the emergence of cross-disciplinary collaboration in the Human Genome Project and the role of cross-disciplinary mobility embodied by Dr Eric Lander and several other subsequent bioinformatics leaders (Petersen et al. 2018). Both individual and group-level social convergence modes are critical for addressing complex multidimensional problems calling on systems-thinking approaches (Orsatti, Quatraro and Pezzoni 2020; Wanzenböck et al. 2020).

Whereas some interfaces involve just two domains, as in the genomics revolution (Petersen et al. 2018), others may involve multiple domains, as is typical of environmental problems (Petersen, Vincent and Westerling 2019; Arroyave et al. 2021). In particular, Figure 4A shows the emergence of a triple-domain nexus—the convergence of the neuro-biological ↔ health ↔ techno-informatic domains—that characterizes the HBS frontier. Structural comparison of cross-disciplinary collaboration networks constructed across two decades indicates the increasing densification at this human brain (HB) science frontier coinciding with the emergence of massive flagship funding programs in the USA, Europe, and Australasia occurring in the period 2009–18 (Petersen et al. 2021). Discrepancies in the configurations of social and conceptual dimensions identify science policy pathways to adjust, incentivize, institutionalize—and in the long run—to accelerate convergence.



Figure 4B juxtaposes the average domain-spanning diversity $\langle f_D(t) \rangle$, a measure of convergence within the social and conceptual dimensions, individually. Comparison of historical trends points to different drivers of convergence in each dimension. To emphasize the distinct trends observed for each dimension, we also show $\langle f_D(t) \rangle$ values reported in units of percent difference from the mean value calculated across the entire period. Two distinct patterns emerge, suggesting different challenges associated with effecting cross-domain integration of each type. Whereas disciplinary (SA) convergence has fluctuated around its mean value (with no statistically significant trend), conceptual (CIP) convergence has steadily increased $\sim 30\%$ over the three-decade period 1990–18 (P-value < 0.0001). This result suggests that it is relatively easier to integrate cross-domain knowledge than to integrate cross-disciplinary expertise, likely owing to coordination costs and other constraints associated with crossing disciplinary and organizational boundaries (Cummings and Kiesler 2005, 2008; Feller 2006; Van Rijnsoever and Hessels 2011; Bromham, Dinnage and Hua 2016). For more on this disparity in social and conceptual integration, and econometric

Naturally, the question arises as to which of three convergence modes—research characterized as polymathic convergence only (X_{SA}), cross-discipline only (X_{CIP}), or full convergence ($X_{SA\&CIP}$)—prevails in practice and impact. To address this question, we partitioned the publication data depending on $f_{D,p}^{SA}$, $f_{D,p}^{CIP}$, and z_p and show in Figure 4C their joint frequency distributions. While there is little variation in the joint distribution $P(f_{D,p}^{SA}, f_{D,p}^{CIP})$, aside from the marginal growth of $f_{D,p}^{SA}$ indicated in panel A, comparing above-average ($z > 0$) relative to below-average cited research ($z \leq 0$) shows that higher joint convergence values correlate with highly cited research. For more robust econometric analysis (see Petersen et al. 2021), which shows that research featuring full convergence ($X_{SA\&CIP}$, corresponding to research with $f_{D,p}^{SA} > 0$ and $f_{D,p}^{CIP} > 0$) features a 6% citation premium relative to polymathic research (X_{SA}). This differential reflects the additional quality and rigor of research that passes the thresholds of multi-disciplinary evaluation and communication.

Complementary analysis of this HBS dataset provides insights into the mechanism giving rise to this disparity, finding an increasing propensity for teams to pursue conceptual convergence without appropriate social convergence—i.e., a *convergence shortcut* (Petersen et al. 2021). This result points to the increasing prevalence of a strategy for rapidly and efficiently competing for large flagship funding opportunities that foregoes the more timely and costly efforts associated with cross-disciplinary team assembly. While expanding beyond one's core expertise, as endowed by the traditionally mono-disciplinary channels of graduate education, largely reflects the innate curiosity at the foundation of scholarship, it also signals the dawn of a new era of autodidactic education by way of open science data, code and tutorials. The theory of expansive learning (Engeström and Sannino 2010) provides understanding for this innate autodidactic propensity, which arises naturally in highly interconnected social systems, and is further supported by cross-pollinating team science.

Evaluating the alignment of social and conceptual dimensions

Convergence is optimally operational when multidisciplinary teams are appropriately aligned with the boundary-spanning target problem, as illustrated in Figure 5A. In effect this means

that the strategic assembly of teams should follow *form follows function* design principles. The function of the team should be tailored to the focal subject area of the target problem. However, most challenges are grand in that they are inextricably embedded in a network of contingencies and risk (Helbing 2013), a complex state of dependency that support chain reactions that extend well beyond the source domain, and thereby requires deep understanding pertaining to each of those domains as well. Further exacerbating this issue, different communities of expertise, which in the most ideal case are in agreement as to what is the core problem, may not necessarily be in (or reach in a timely manner) a working level of consensus around which pathway to take, thereby giving rise to 'wicked problems' (Stirling 2010; Arroyave et al. 2021).

The process by which a sustainable convergence nexus emerges is likely akin to the nucleation and growth of stable surfaces in disordered media developed in statistical physics (Krapivsky, Redner and Ben-Naim 2010). Nucleation commences when problem-solving expertise identifies an emerging target problem, forms a local community around some partial dimension of the problem, and then coalesce with other stakeholders at subsequent stages of cross-disciplinary integration in order to avoid eventual disassociation. Analysis of 'wicked' target problems in the environmental sciences (Arroyave et al.

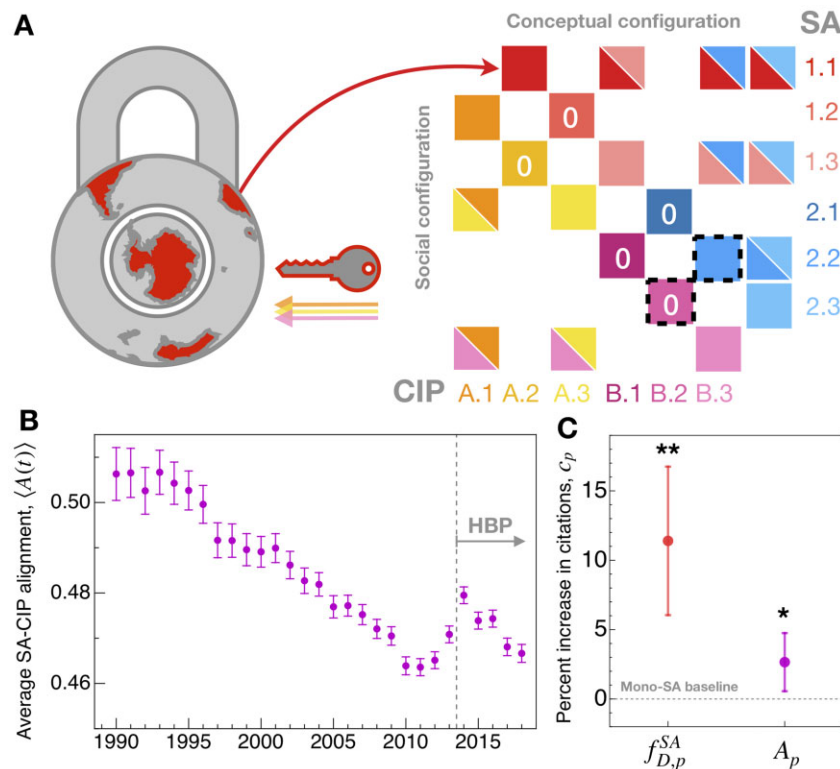


Figure 5. Evaluation and impact of socio-conceptual (CIP-SA) alignment. (A) Schematic of a target problem, e.g. melting polar ice-caps deriving from global warming, which maps onto a particular configuration of SA hypothetically spanning two 'super-group' domains, each featuring three L_2 domains. A particular research team tackling this particular problem can be represented by a social configuration codified by six corresponding CIP (A.1 through B.3). As such, this approach to measuring and evaluating convergence can aid policymakers and principal investigators in designing teams that represent strategic configurations of expertise aimed at specific target problems, which are themselves conceptualized as configurations of core concepts (e.g. established knowledge, methods, and tools). Consistent codification of social and conceptual configurations can help identify candidate pathways for unlocking solutions to grand challenges. The relative composition of the conceptual and social configurations can also identify when teams are not sufficiently aligned with the problem, as indicated by the cells with dashed black borders suggesting that the social configuration is misaligned since there is no expertise ('0' weight in CIP B.2) to match the corresponding dimension SA 2.2 of the target problem. (B) Average social-conceptual alignment $\langle A(t) \rangle$ calculated by year, with error bars indicating the standard error of the mean. (C) Regression coefficients for the two main model variables reported in terms of the percent increase in citations attributable to research featuring full SA convergence ($f_{D,p}^{SA} = 1$) and full alignment of CIP with the SA ($A_p = 1$), measured relative to the mono-disciplinary baseline with $f_{D,p}^{SA} = 0$.

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