Can you trust your experiments? Generalizability of Experimental Studies

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Abstract

Experimental studies are a cornerstone of Machine Learning (ML) research. A common and often implicit assumption is that the study's results will generalize beyond the study itself, e.g., to new data. That is, repeating the same study under different conditions will likely yield similar results. Existing frameworks to measure generalizability, borrowed from the casual inference literature, cannot capture the complexity of the results and the goals of an ML study. The problem of measuring generalizability in the more general ML setting is thus still open, also due to the lack of a mathematical formalization of experimental studies. In this paper, we propose such a formalization, use it to develop a framework to quantify generalizability, and propose an instantiation based on rankings and the Maximum Mean Discrepancy. We show how this latter offers insights into the desirable number of experiments for a study. Finally, we investigate the generalizability of two recently published experimental studies.

1 Introduction

Experimental studies are a cornerstone of Machine Learning (ML) research. Due to their importance, the community advocates for high methodological standards when performing, evaluating, and sharing studies (Hothorn et al., 2005; Huppler, 2009; Montgomery, 2017).

The quality of an experimental study depends on multiple aspects. First, the experimenter should properly define the *scope* and the *goals* of the study. Particular attention must be given to the choice of benchmarked methods and experimental conditions (Boulesteix et al., 2015; Bouthillier et al., 2021; Dehghani et al., 2021). Second, the study should be *reproducible* by independent parties and hence contain the necessary documentation. This aspect has recently drawn much attention due to the so-called reproducibility crisis (Baker, 2016; Gundersen et al., 2023; Peng, 2011; Raff, 2023; 2021). Third, the results of the study should be sensibly analyzed to draw conclusions regarding, for instance, the *significance* of the findings (Benavoli et al., 2017; Corani et al., 2017; Demsar, 2006). Finally, the *generalizability* of a study concerns how well its results are replicated under unseen experimental conditions, such as datasets not considered in the study (National Academies of Science (2019); Findley et al., 2021; Pineau et al., 2021). The latter two aspects are also known as the internal and external validity of a study. Generalizability and significance, although sometimes confused, are two independent aspects of a study (Findley et al., 2021). On the one hand, significant findings may not generalize to other conditions; on the other hand, results might consistently be not significant.

Generalizability captures how close the results are between two different samples of experiments. Generalizability is, conceptually, closely related to model replicability. A model is ρ -replicable if, given i.i.d. samples from the same data distribution, the trained models are the same with probability $1 - \rho$ (Impagliazzo et al., 2022). An experimental study is generalizable if, when repeated under different experimental conditions, the results are similar with high probability (National Academies of Science (2019)). A quantifiable notion of generalizability thus requires a formalization of experimental studies, of their results, and of similarity between results.

Significance, instead, captures how strong the findings are within the specific sample of experiments performed. Multiple publications have shown how different choices of experimental conditions can lead to very

different results (Benavoli et al., 2017; Boulesteix et al., 2017; Bouthillier et al., 2021; Dehghani et al., 2021; Gundersen et al., 2022; Mechelen et al., 2023). Some recent experimental studies have also reported this phenomenon. Matteucci et al. (2023) discuss how previous studies on categorical encoders disagree on the best-performing ones, even when the results are significant. Similarly, Lu et al. (2023) re-evaluated coreset learning methods and found that all of the methods they considered did not beat a naïve baseline.

Quantifying generalizability can also help determine the appropriate size of experimental studies. While one dataset is intuitively not enough to draw generalizable conclusions (unless all experiments have the same outcome), 10^6 datasets likely are. Of course, such large studies are usually not practical: it is crucial to determine the minimum amount of data needed to achieve generalizability. This principle also applies to other experimental factors, such as initialization seed, task, or quality metric.

Our contributions are the following:

- 1. We introduce a novel measure-theoretic formalization of experimental studies.
- 2. We propose a quantifiable definition of the generalizability of experimental studies.
- 3. We develop an algorithm to estimate the size of a study to obtain generalizable results.
- 4. We analyze two recent experimental studies, Matteucci et al. (2023); Srivastava et al. (2023), and show how well their results generalize.
- 5. We publish the GENEXPY¹² Python module to repeat our analysis in other studies.

Paper outline: Section 2 discusses the related work, Section 3 formalizes experimental studies, Section 4 defines generalizability and provides the algorithm to estimate the required size of a study for generalizability, Section 5 contains the case studies, and Section 6 describes the limitations and concludes.

2 Related work

We first discuss the literature related to the problem we are tackling, i.e., why experimental studies may not generalize. Second, we overview the existing concept of model replicability, closely related to our work. Finally, we show other meanings that these words can assume in other domains.

Non-generalizable results. It is well known that experimental results can significantly vary based on design choices (Lu et al., 2023; Matteucci et al., 2023; Qin et al., 2023; McElfresh et al., 2022). Possible reasons include an insufficient number of datasets (Dehghani et al., 2021; Matteucci et al., 2023; Alvarez et al., 2022; Boulesteix et al., 2015) as well as differences in hyperparameter tuning (Bouthillier et al., 2021; Matteucci et al., 2023), initialization seed (Gundersen et al., 2023), and hardware (Zhuang et al., 2022). As a result, the statistical benchmarking literature advocates for experimenters to motivate their design choices (Bartz-Beielstein et al., 2020; Mechelen et al., 2023; Boulesteix et al., 2017; Bouthillier et al., 2021; Montgomery, 2017) and clearly state the hypotheses they are attempting to test with their study (Bartz-Beielstein et al., 2020; Moran et al., 2023).

Replicability and generalizability in ML. Our work formalizes and extends the definitions of replicability and generalizability given in Pineau et al. (2021) and National Academies of Science (2019). Intuitively, replicable work consists of repeating an experiment on different data, while generalizable work varies other factors as well—e.g., task, seed. In ML, these terms are usually associated to learning algorithms rather than experimental studies. A generalizable model has small generalization error on unseen data (McElfresh et al., 2022), while a replicable model learns the same parameters from different i.i.d. samples (Impagliazzo et al., 2022). Model replicability is also linked to model stability, differential privacy, generalization error, and global stability (Bun et al., 2023; Chase et al., 2023; Ghazi et al., 2023; Moran et al., 2023; Dixon et al., 2023).

External validity. The external validity of a study is a well-studied concept in the context of causal inference, its main applications being in the social and political sciences (Campbell, 1957). In general, the

¹https://anonymous.4open.science/r/genexpy-B94D

²The module will be published on PyPI after acceptance.

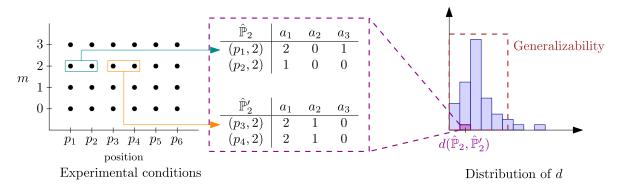


Figure 1: Generalizability of the "checkmate-in-one" task (Example 3.1), as the probability that two substudies yield similar results. A result is a distribution of rankings. Note that the design factor (m) is fixed, while the generalizability factor (position) varies.

external validity of a study performed concerns whether repeating a study on different samples affects the validity of its findings. Generalizability, opposed to transportability, concerns the external validity of results when the samples come from the same population (Findley et al., 2021). Existing methods assess the signand effect-generalization of the treatment on some response variable (Egami & Hartman, 2023). They are thus not applicable to our use-case of ML experimental studies, for which there is—arguably—no treatment and no response variable.

3 Experiments and experimental studies

An experimental study is a collection of experiments comparing the same alternatives under different experimental conditions. An experimental condition is a tuple of levels of experimental factors, the parameters defining the experiments. The study aims at answering a research question, which defines its scope and goals.

Example 3.1. (The "checkmate-in-one" task, cf. Figure 1) An experimenter wants to compare three Large Language Models (LLMs), the alternatives, on the "checkmate-in-one" task (Srivastava et al., 2023; Ammanabrolu et al., 2019; 2020; Dambekodi et al., 2020). The assignment is to find the unique checkmating move from a position of pieces on a chessboard: an LLM succeeds if and only if it outputs the correct move. The experimenter considers two experimental factors: the number of shots, m, and the initial position on the chessboard, p_l . The experimenter wants to find if LLM a_1 ranks consistently (in the same position) against the other two LLMs when changing the initial position, for a fixed number of shots.

3.1 Experiments

An experiment evaluates all the *alternatives* under a valid experimental condition. The result of an experiment is an element of some result space.

Alternatives. An alternative $a \in A$ is an object evaluated in the study, like an LLM in Example 3.1. We call A the finite set of alternatives considered in the study, with cardinality n_a .

Experimental factors. An experimental factor is *anything* that may affect the result of an experiment. We use i to denote a factor, C_i the (possibly infinite) set of *levels* that i can take, $c \in C_i$ a level of i, and I the set of all factors. We adapt Montgomery's classification of experimental factors (Montgomery, 2017, Chapter 1) and distinguish between the *held-constant*, *design*, and *generalizability* factors of a study.

• *Held-constant factors* are presumed not to significantly impact the results, they are hence out of the study's scope and are fixed to a single level; examples are "programming language" or "number of cross-validated folds".

- Design factors are expected to significantly impact the results and have a relatively small set of levels; examples are "quality metric" or "number of shots".
- Generalizability factors (I_{gen}) have a larger number of levels. The experimenter wants to obtain results that generalize to unseen levels of these factors; examples are "dataset" or "chessboard position".

The same factor may play different roles in different studies, according to the studies' objectives (Montgomery, 2017, Chapter 1). For instance, "seed" is a generalizability factor in reinforcement learning studies such as Nauman et al. (2024), as it typically plays a large role in determining the performance of such algorithms. On the other hand, "seed" is a held-constant factor in Matteucci et al. (2023), as the effect of changing the seed is not of interest for the experimenters.

Experimental conditions. An experimental condition \mathbf{c} is a tuple of levels of all experimental factors, $\mathbf{c} = (c_i)_{i \in I} \in C \subseteq \prod_{i \in I} C_i$. We endow C with the product probability measure $\mu^C = \bigotimes_{i \in I} \mu_i$, where μ_i is a probability measure on C_i . For instance, if i is "dataset" and $C_i \subseteq \mathbb{R}^{k \times d}$ for some $k, d \in \mathbb{N}$, then μ_i measures the probability of subsets of $\mathbb{R}^{k \times d}$. If, instead, i is a held-constant factor with $C_i = \{c\}$, then μ_i is $\mu_i : \{c\} \mapsto 1$. The probability space $(C, \mathcal{F}_C, \mu^C)$ is the universe of valid experimental conditions. In our formalization, it plays the role of the sample space for the experiment function E^Q (Section 3.2). We finally assume that one can fix the levels of the experimental factors independently, as long as the resulting experimental condition is in C.

Example 3.1 (Continued). $C = \{(p_l, m)\}_{l,m}$, where p_l is a legal configuration of pieces on a chessboard and m is the non-negative number of shots.

Experimental results. We define the *result space* as $(\mathcal{X}, \mathcal{B}_{\mathcal{X}})$, where \mathcal{X} is a separable topological space and $\mathcal{B}_{\mathcal{X}}$ is the Borel σ -algebra on \mathcal{X} . For instance, if the experimenter is interested in the raw performances of the alternatives on some learning task, $(\mathcal{X}, \mathcal{B}_{\mathcal{X}}) = (\mathbb{R}^{n_a}, \mathcal{B}_{\mathbb{R}^{n_a}})$.

Example 3.1 (Continued). As the goal of the experiment involves ranking the LLMs, the experimenter defines the result of an experiment on (p_l, m) as a ranking of the three LLMs, according to whether or not they output the checkmating move. Suppose that only a_1 and a_2 output the correct move. Then, the result is (0, 0, 1), where a_1 and a_2 are tied best.

3.2 Experimental studies

A study is defined by its research question Q, i.e., its scope and goals. The scope consists of the alternatives A, the valid experimental conditions C, the generalizability factors I_{gen} , and the result space \mathcal{X} . The goal specifies the kind of conclusion one is attempting to draw from the study. We propose a model of the goals in Section 4.2.

Definition 3.1 (Research question). The research question is a tuple $Q = (A, C, I_{gen}, \mathcal{X}, goals)$.

Example 3.1 (Continued). The research question of the "checkmate-in-one" study is as follows. The scope is given by $A = \{a_1, a_2, a_3\}$, $C = \{(p_l, m)\}_{l,m}$, $I_{\text{gen}} = \{\text{"position"}\}$, and \mathcal{X} the set of rankings of 3 alternatives. The goal is "Is a_1 consistently in the same position of the ranking?"

To model the results, we introduce the experiment function $E^Q: (C, \mathcal{F}_C, \mu^C) \to (\mathcal{X}, \mathcal{B}_{\mathcal{X}})$, which evaluates all of the alternatives in A under a valid experimental condition. We also assume that E^Q is a measurable function or, equivalently, a random variable with sample space C. We can thus sample from E^Q . The result of the study on a research question Q is then the distribution of E^Q : since E^Q is not necessarily injective, we assign higher probability to those results that appear more often.

Definition 3.2 (Result of a study). The result of the study on Q is the pushforward probability induced by the experiment function E^Q ,

$$\begin{split} \mathbb{P}^Q \coloneqq E^Q_* \mu^C : \mathcal{B}_{\mathcal{X}} \to [0,1] \\ Y \mapsto \mu^C \left(E^{Q^{-1}}(Y) \right), \end{split}$$

where $E^{Q^{-1}}(Y) = \{\mathbf{c} : E^Q(\mathbf{c}) \in Y\} \in \mathcal{F}_C \text{ is the preimage of } Y.$

In practice, as C might be infinite or too large, one can only run experiments under n experimental conditions and obtain a sample $\hat{E}_n^Q \sim \mathbb{P}^Q$ of size n. We call this an *empirical study of size* n on Q and denote the empirical distribution of \hat{E}_n^Q with $\hat{\mathbb{P}}_n^Q$. To keep the notation clean and unless necessary, we omit Q.

4 Generalizability of experimental studies

The currently accepted definition of generalizability is the property of two independent studies with the same research question to yield similar results, see National Academies of Science (2019) and Pineau et al. (2021). Although intuitive, this notion is not practically useful as it cannot be assessed objectively. We thus propose the following quantifiable definition of generalizability based on our framework (cf. Section 3.1)

Definition 4.1 (Generalizability). Let $Q = (A, C, I_{\text{gen}}, \mathcal{X}, \text{goals})$ be a research question, let \mathbb{P} be the result of the corresponding study, and let d be a distance between probability distributions. The n-generalizability of the study on Q is

$$n\text{-Gen}(Q;\varepsilon) := \mathbb{P}^n \otimes \mathbb{P}^n \left((X_j)_{j=1}^n, (Y_j)_{j=1}^n : d(X,Y) \le \varepsilon \right), \tag{1}$$

where $\varepsilon \in \mathbb{R}^+$ is a dissimilarity threshold.

Intuitively, the n-generalizability is the probability for any two empirical studies of size n on Q to yield "similar" results, as defined by d and ε . We discuss in Section 4.4 how to interpretably choose a value for ε . Definition 4.1 is very flexible and allows for different choices of the result space \mathcal{X} , the goals, and the distance d. The rest of this section proposes a particular instantiation, base on the Maximum Mean Discrepancy (Gretton et al., 2006), which allows for different choices of the experimental space \mathcal{X} and goals.

4.1 Experimental results: Rankings with ties

Experimental results can be formalized in different ways, such as raw performance metrics, time series, or rankings. Among these, rankings are arguably one of the most natural forms:

- (i) Rankings are already widely used for non-parametric tests such as Friedman, Nemenyi, and Conover-Iman (Demsar, 2006; Conover & Iman, 1982).
- (ii) Rankings do not suffer from experimental-condition-fixed effects, such as a dataset being inherently easier to solve than another one. Even though there are multiple ways to deal with these effects, there is no preferred one in the literature. See, for instance, the consensus ranking problem (Matteucci et al., 2023; Nießl et al., 2022).
- (iii) Rankings allow the definition of interpretable kernels to formalize different goals of a study, as we illustrate in Section 4.2.

We define rankings (with ties) in the following way.

Definition 4.2 (Ranking). A ranking r on A is a transitive and reflexive binary endorelation on A. Equivalently, r is a totally ordered partition of A into *tiers* of equivalent alternatives. r(a) denotes the rank of $a \in A$, i.e., the position of the tier of a in the ordering. W.l.o.g., we use \mathcal{R}_{n_a} for the space of rankings of n_a alternatives.

4.2 Goals: Kernels

Goals act as lenses, focusing on specific aspects of the results that are of interest for the experimenter. For instance, consider the goal in Example 3.1: "Is a_1 consistently in the same position of the ranking?". In this case, the goal solely focuses on a_1 's position in the rankings, ignoring the positions of a_2 and a_3 . Changing the goal of a study can thus heavily impact how the results are analyzed. Within our framework, we formalize for goals as kernels on the result space, i.e., positive definite symmetric functions. In the following, we describe three kernels for rankings, covering three representative goals.

Borda kernel. The Borda kernel is suitable for goals in the form "Is alternative a^* consistently ranked the same?". It uses the Borda count, defined as the number of alternatives (weakly) dominated by a given one (Borda, 1781). For a pair of rankings, we compute the Borda counts of a^* and then take their difference.

$$\kappa_b^{a^*,\nu}(r_1,r_2) = e^{-\nu|b_1-b_2|},$$

where $b_l = |\{a \in A : r_l(a) \ge r_l(a^*)\}|$ is the number of alternatives dominated by a^* in r_l and $\nu \in \mathbb{R}^+$. The Borda kernel takes values in $[e^{-\nu n_a}, 1]$. If ν is too large compared to $1/|b_1-b_2|$, the kernel is oversensitive and will heavily penalize even small differences. On the contrary, if ν is too small, the kernel is undersensitive and will not penalize deviations unless they are very large. As $|b_1 - b_2| \in [0, n_a]$, we recommend $\nu = 1/n_a$.

Jaccard kernel. The Jaccard kernel is suitable for goals in the form "Are the best alternatives consistently the same ones?". As it measures the similarity between sets (Gärtner et al., 2006; Bouchard et al., 2013), we use it to compare the top-k tiers of two rankings.

$$\kappa_j^k(r_1, r_2) = \frac{\left|r_1^{-1}([k]) \cap r_2^{-1}([k])\right|}{\left|r_1^{-1}([k]) \cup r_2^{-1}([k])\right|},$$

where $r^{-1}([k]) = \{a \in A : r(a) \le k\}$ is the set of alternatives with rank lower than or equal to k. The Jaccard kernel takes values in [0,1].

Mallows kernel. The Mallows kernel is suitable for goals in the form "Are the alternatives ranked consistently?". It measures the overall similarity between rankings (Jiao & Vert, 2018; Mania et al., 2018; Mallows, 1957). We adapt the original definition in (Mallows, 1957) for ties,

$$\kappa_m^{\nu}(r_1, r_2) = e^{-\nu n_d},$$

where $n_d = \sum_{a_1,a_2 \in A} |\text{sign}(r_1(a_1) - r_1(a_2)) - \text{sign}(r_2(a_1) - r_2(a_2))|$ is the number of discordant pairs and $\nu \in \mathbb{R}^+$. If a pair is tied in one ranking but not in the other, one counts it as half a discordant pair. The Mallows kernel takes values in $\left[\exp\left(-2\nu\binom{n_a}{2}\right), 1\right]$. If ν is too large compared to $1/n_d$, the kernel is oversensitive and it will heavily penalize even small differences. On the contrary, if ν is too small, the kernel is undersensitive and will not penalize deviations unless they are very large. As $n_d \in \left[0, \binom{n_a}{2}\right]$, we recommend $\nu = 1/\binom{n_a}{2}$.

The following example illustrates the three kernels.

Example 4.1. Consider two rankings $\mathbf{r} = (0,0,0)$ and $\mathbf{s} = (0,1,1)$, where x_j is the rank of the j-th alternative. To understand their impact on generalizability, consider a study whose result is a distribution assigning both \mathbf{r} and \mathbf{s} probability $^1/2$. For the goal corresponding to the Borda kernel, \mathbf{r} and \mathbf{s} answer the research question consistently as a_1 weakly dominates all alternatives in both rankings. Hence, the Borda kernel takes a value of 1 and the study is perfectly generalizable. For the Jaccard and Mallows goals, instead, the two rankings are either very different $(\kappa_j^1(r_1, r_2) \approx 0.33)$ or slightly different $(\kappa_m^{1/n_a^2}(r_1, r_2) \approx 0.72)$. Thus, we conclude that the study is more generalizable w.r.t. the Mallows kernel than the Jaccard kernel.

4.3 Distance between experimental results: Maximum Mean Discrepancy

In the previous sections, we have formalized an experimental study, its results, and its goals. The last open point before applying (1) in practice is a definition of d, a distance between probability distributions. Such a distance should satisfy the following requirements. First, it should take into consideration the goal of a study. Second, it should handle sparse distributions well: empirical studies are typically very small compared to the number of all possible rankings, which grows super-exponentially in the number of alternatives.³ Third, it should provide a way to indicate the amount of experiments needed to achieve n-generalizable results. A distance satisfying the above requirements is the Maximum Mean Discrepancy (MMD) (Gretton et al., 2006; 2012).

Definition 4.3 (MMD). Let \mathcal{X} be a set with a kernel κ , and let \mathbb{Q}_1 and \mathbb{Q}_2 be two probability distributions on \mathcal{X} . Let $\mathbf{x} = (x_i)_{i=1}^n \sim \mathbb{Q}_1, \mathbf{y} = (y_i)_{i=1}^m \sim \mathbb{Q}_2$.

$$MMD(\mathbf{x}, \mathbf{y})^{2} := \frac{1}{n^{2}} \sum_{i,j=1}^{n} \kappa(x_{i}, x_{j}) + \frac{1}{m^{2}} \sum_{i,j=1}^{m} \kappa(y_{i}, y_{j}) - \frac{2}{mn} \sum_{\substack{i=1...n\\j=1...m}} \kappa(x_{i}, y_{j}).$$

Finally, there remains the choice of an appropriate ε^* to use in (1), which is hardly interpretable. The following result relates the range of the MMD and the infimum and supremum of the kernel.

Proposition 4.1. The MMD takes values in $\left[0, \sqrt{2 \cdot (\kappa_{sup} - \kappa_{inf})}\right]$, where $\kappa_{sup} = \sup_{x,y \in X} \kappa(x,y)$ and $\kappa_{inf} = \inf_{x,y \in X} \kappa(x,y)$.

Using a similar approach, we propose to replace ε^* with a condition on the desired minimum expected value of the kernel:

$$\varepsilon^*(\delta^*) = \sqrt{2(\kappa_{\sup} - f_{\kappa}(\delta^*))},\tag{2}$$

where f_{κ} is a kernel-specific function and δ^* is an interpretable parameter. We now discuss them for the three kernels discussed in Section 4.2 with their recommended parameters.

- Borda kernel. δ^* is the difference between the fraction of dominated alternatives between two rankings; $f_{\kappa_h}(x) = e^{-x}$.
- Jaccard kernel. δ^* is the Jaccard coefficient between the top-k tiers of two rankings; $f_{\kappa_i}(x) = 1 x$.
- Mallows kernel. δ^* is the fraction of discordant pairs; $f_{\kappa_m}(x) = e^{-x}$.

As a concrete example, achieving ($\alpha^* = 0.90, \delta^* = 0.05$)-generalizable results for the Jaccard kernel means that, w.p. 0.90, the average Jaccard coefficient between two rankings drawn from the results is at least 0.95.

4.4 How many experiments ensure generalizability?

When designing a study, the experimenter has to decide how many experiments to run in order to obtain generalizable results. In other words, they need to choose a (minimum) sample size n^* that achieves the desired generalizability α^* for a given threshold ε^* .

$$n^* = \min \left\{ n \in \mathbb{N} : n\text{-Gen}(Q; \varepsilon^*) \ge \alpha^* \right\}. \tag{3}$$

Example 3.1 (Continued). The experimenter wants to obtain results in which, with probability 0.99, a_1 dominates the same number of alternatives up to a difference of 1. They therefore choose the Borda kernel with $\nu = 1/n_a$, $\delta^* = 1/3$, and $\varepsilon^* = \sqrt{2}\sqrt{1 - e^{-0.33}}$ as in (2). How many experiments are enough?

Let now $\varepsilon_n^{\alpha^*}$ be the α^* -quantile of the MMD. To estimate n^* , we use a linear dependency between $\log(n)$ and $\log(\varepsilon_n^{\alpha^*})$, which we have observed in our experiments; see, for instance, Figure 2. In particular, for any

³Fubini or ordered Bell numbers, https://oeis.org/A000670.

distribution \mathbb{P} and choice of $\alpha^* \in [0,1]$, there exist $\beta_0, \beta_1 \in \mathbb{R}, \beta_i = \beta_i(\alpha^*, \mathbb{P})$, s.t.

$$\log(n) = \beta_1 \log\left(\varepsilon_n^{\alpha^*}\right) + \beta_0. \tag{4}$$

Appendix A.3.1 provides a proof for the distribution-free case. Equation (4) suggests that one can use a small set of N preliminary experiments to estimate n^* . The estimate can then be improved by adding more experiments. On this result is based Algorithm 1, whose working is illustrated in Figure 2.

```
Algorithm 1 Run the necessary amount of experiments
                                                                                                                       Require: \alpha^*
Require: \delta^*
                                                                                                          ▷ similarity threshold on rankings
Require: Q
                                                                                                   \triangleright research question, Q = (A, C, I_{\text{gen}}, \kappa)
Require: N
                                                                                                              > number of initial experiments
Require: N_{\text{step}}
                                                                                                       ▷ number of additional experiments
Require: n_{\rm rep}
                                                             > number of repetitions to estimate the distribution of the MMD
   procedure EstimateNstar(\alpha^*, \delta^*, Q, N, n_{\text{max}}, n_{\text{rep}})
        n_N^* \leftarrow \infty
        while N < n_N^* do
             Run N experiments and get their results \hat{\mathbb{P}}_N
            \varepsilon^* \leftarrow f_{\kappa}(\delta^*)
                                                                                                                                    ▷ cf. Section 4.4
            n_{\text{max}} \leftarrow \lfloor N/2 \rfloor
                                                                               \triangleright we need two disjoint samples of size n_{\max} from \hat{\mathbb{P}}_N
             for n = 1 \dots n_{\text{max}} do
                  mmds \leftarrow empty list
                  for n = 1 \dots n_{\text{rep}} do
                      sample without replacement (x_j)_{j=1}^{2n} \sim \hat{\mathbb{P}}_N
                      \mathbf{x} \leftarrow (x_j)_{j=1}^n
\mathbf{y} \leftarrow (x_j)_{j=n+1}^{2n}
                                                                                                                   ▶ split into disjoint samples
                       append MMD(\mathbf{x}, \mathbf{y}) to mmds
                  end for
                 \varepsilon_n^{\alpha^*} \leftarrow \alpha^*-quantile of mmds
             fit a linear regression \log(n) = \beta_1 \log(\varepsilon_n^{\alpha^*}) + \beta_0
            n_N^* \leftarrow \beta_1 \log(\varepsilon^*) + \beta_0
             N \leftarrow N + N_{\text{step}}
        end while
        return n_N^*
   end procedure
```

5 Case studies

5.1 Case Study 1: A benchmark of categorical encoders

We now evaluate the generalizability of a recent study (Matteucci et al., 2023) that analyzes the performance of encoders for categorical data. The performance of an encoder is approximated by the quality of a model trained on the encoded data. The *design factors* are the model, the tuning strategy for the pipeline, and the quality metric for the model, while the only *generalizability factor* is the dataset. We impute missing values in the results of the study by assigning the worst rank. We evaluate how well the results of the study generalize w.r.t. three goals:

 (g_1) Find out if the one-hot encoder (a popular encoder) ranks consistently amongst its competitors, using the Borda kernel with $\nu = 1/n_a$.

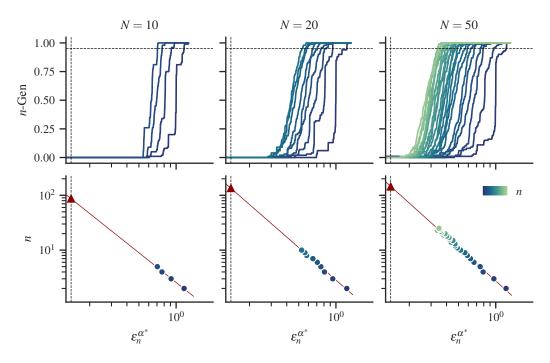


Figure 2: Estimation of the required number of experiments n^* (marked with \blacktriangle) from different amounts N of preliminary experiments. The dashed lines represent $\alpha^* = 0.9$ (horizontal) and $\varepsilon^* = \sqrt{2}/10$ (vertical). We used real data from Matteucci et al. (2023) and the Jaccard kernel (Section 4.2).

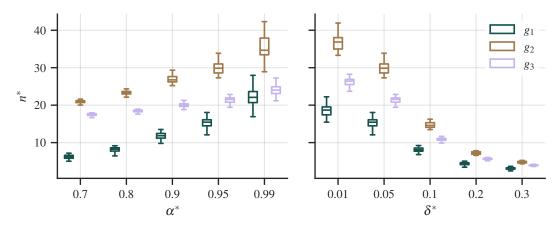


Figure 3: Number of necessary experiments n^* to achieve generalizability for categorical encoders, for different desired generalizability α^* , similarity threshold δ^* , goals g_i . The variation in the plot is due to the combinations of design factors.

- (g_2) Investigate if some encoders outperform all the others using the Jaccard kernel with k=1.
- (g_3) Evaluate whether the encoders are ranked in a similar order, using the Mallows kernel with $\nu = 1/\binom{n_a}{2}$.

Figure 3 shows the predicted n^* for different choices of α^* and δ^* , the other one fixed at 0.95 and 0.05 respectively. The variance in the boxes comes from variance in the design factors. For example, the results for the design factors "decision tree, full tuning, accuracy" have a different (α^*, δ^*) -generalizability than the results for "SVM, no tuning, accuracy". We observe on the left that—as expected—obtaining generalizable results requires more experiments as the desired generalizability α^* increases. We can also see that the variance of the boxes increases with α^* , meaning that the choice of the design factors has a larger influence on the achieved generalizability. This is also possibly due to tail effects in the estimation of the MMD. We

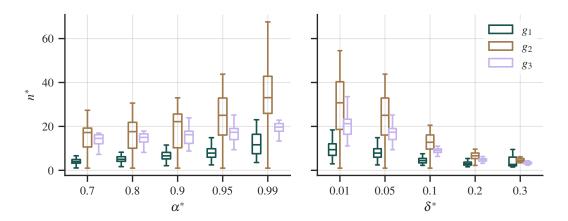


Figure 4: Number of necessary experiments n^* to achieve generalizability for LLMs, for different desired generalizability α^* , similarity threshold δ^* , goals g_i . The variation in the plot is due to the combinations of design factors.

observe the same when decreasing δ^* , as it corresponds to a stricter similarity condition on the rankings. In the rather extreme cases of $\alpha^* = 0.7$ or $\delta^* = 0.3$, even less than 10 datasets are enough to achieve (α^*, δ^*) -generalizability.

Consider now goal g_2 for two different choices of design factors: (A): "decision tree, full tuning, accuracy" and (B): "SVM, full tuning, balanced accuracy". Furthermore, let $(\alpha^*, \delta^*) = (0.95, 0.05)$: we estimate $n^* = 28$ for (A) and $n^* = 34$ for (B), corresponding to the bottom and top whiskers of the corresponding box in Figure 3. As both (A) and (B) were evaluated using n = 30 experiments, we conclude that the results of (A) are (barely) (0.95, 0.05)-generalizable, while those of (B) are not. Hence, one should run more experiments with fixed factors (B) to make the study generalizable.

5.2 Case study 2: BIG-bench — A benchmark of Large Language Models

We now evaluate the generalizability of BIG-bench (Srivastava et al., 2023), a collaborative benchmark of Large Language Models (LLMs). The benchmark compares LLMs on different tasks, such as the "checkmate-in-one" task (cf. Example 3.1), and for different numbers of shots. Task and number of shots are the *design factors*. Every task has a number of subtasks, which is the *generalizability factor*. We stick to the preferred scoring for each subtask. As the results have too many missing values to impute them, we only consider the experimental conditions where at least 80% of the LLMs had results, and to the LLMs whose results cover at least 80% of the conditions.

As before, we define three goals:

- (g_1) Find out if GPT3 (to date, one of the most popular LLMs) ranks consistently amongst its competitors, using the Borda kernel with $\nu = 1/n_a$.
- (g_2) Investigate if some encoders outperform all the others using the Jaccard kernel with k=1.
- (g_3) Evaluate whether the LLMs are ranked in a similar order, using the Mallows kernel with $\nu = 1/\binom{n_2}{2}$.

Figure 4 shows the predicted n^* for different choices of α^* and δ^* , the other one fixed at 0.95 and 0.05 respectively. Again, the variance in the boxes comes from variance in the design factors, i.e., the task and the number of shots. As before, increasing α^* or decreasing δ^* leads to higher n^* . Unlike in the previous section, n^* for g_2 greatly depends on the combination of fixed factors, as we now detail.

Consider now goal g_2 for two different choices of design factors: (A): "conlang_translation, 0 shots", and (B): "arithmetic, 2 shots". Furthermore, let $(\alpha^*, \delta^*) = (0.95, 0.05)$. For this choice of parameters, we estimate $n^* = 44$ for (A), corresponding to the top whisker of the corresponding box in Figure 3. As the study evaluates (A) on 10 subtasks, it is therefore not (0.95, 0.05)-generalizable. In fact, we estimate that this

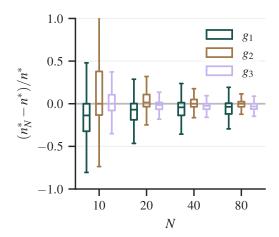


Figure 5: Relative error of the prediction of n^* from N preliminary experiments (n_N^*) for the goals g_i .

would require 34 more subtasks. For (B), on the other hand, we estimate $n^* = 1$: the best 2-shot LLM for the observed subtasks is always PALM 535B. Hence, the result of a single experiment is enough to achieve (0.95, 0.05)-generalizability.

Note that, although we correctly estimated $n^* = 1$ for (B), this estimate relies on 10 preliminary experiments. In other words, our algorithm was able to quantify in hindsight that a single experiment would have been enough to obtain generalizable results. Of course, however, one cannot trust an estimate of n^* based on only one experiment. The next section thus investigates how the number of preliminary experiments influences the estimate of n^* .

5.3 How many preliminary experiments?

We assess the accuracy of our method (cf. Section 4.4) for estimating n^* from N independent experiments. Our procedure is as follows. First, we select a probability distribution \mathbb{P} on the set of rankings \mathcal{R}_{n_a} , representing the true distribution of outcomes. Second, for various values of n, we bootstrap the distribution of MMD(\mathbb{P}^n , \mathbb{P}^n) by repeatedly sampling independently from \mathbb{P} . We compute n^* with its definition in (3). To evaluate our estimate, we generate $N \in \{10, 20, 40, 80\}$ samples and we estimate n^* from \mathbb{P}_N , calling it n_N^* . This procedure is repeated across multiple distributions, with 100 repetitions per distribution.

Figure 5 displays the relative error $\binom{(n_N^*-n^*)}{n^*}$: when greater (lower) than zero, it indicates the overestimation (underestimation) of n^* . Although the specific results vary with the target objective (with goal g_2 , corresponding to the Jaccard kernel, being particularly challenging to estimate), in general, n_N^* approximates n^* within 50% in more than 75% of cases, even when N=10. Consequently, our method provides a reliable estimate of n^* (or at least of its order of magnitude) from as few as 10 preliminary experiments. Appendix B.1 contains additional details regarding the distributions and the results, as well as an application to real data.

6 Conclusion

Limitations. We only investigated the instantiation of our framework based on the MMD, rankings, and kernels for rankings. Keeping the MMD as the cornerstone of an instantation allows different result spaces and goals while potentially maintaining crucial properties such as (4). In this regard, a close formula for the coefficients in (4) would greatly benefit the computational efficiency and the theoretical understanding of our framework.

Future work. First, incorporating information about the experimental conditions into the framework. This would allow to study other aspects of external validity, such as transportability, as well as allow for

active learning to choose the next experiments to run. Second, based on our experiments in Section 5.3, we intend to provide guarantees and confidence intervals on the convergence of n_N^* to n^* . Third, we dealt with missing evaluations by imputing them. Having kernels that can handle missing evaluations might be beneficial. Fourth, rankings, despite their advantages, do not consider the raw performance difference between alternatives. On the other hand, kernels for raw performances (i.e., for vectors in \mathbb{R}^{n_a}) lack an obvious interpretation as the goals of a study. Fuzzy rankings may bridge this gap: performance differences are incorporated into the ranking and the existing kernels for rankings might be adapted to them.

Conclusions. An experimental study is generalizable if, with high probability, its findings will hold under different experimental conditions, e.g., with unseen datasets. Non-generalizable studies might be of limited use or even misleading. This paper is, to our knowledge, the first to develop a quantifiable notion for the generalizability of experimental studies. To achieve this, we formalize experiments, experimental studies, and their results, as well as define an instantiation—rankings and distributions over rankings. Our approach allows us to estimate the number of experiments needed to achieve a desired level of generalizability in new experimental studies. We demonstrate its utility showing generalizable and non-generalizable results in two recent experimental studies.

Acknowledgments

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A Details for Section 4

A.1 Details for Section 4.2

This section contains the proofs to show that the functions introduced in Section 4.2 are kernels, i.e., symmetric and positive definite. As symmetry is a clear property of all of them, we only discuss their positive definiteness. Our proofs for the Borda and Mallows kernels follow Jiao & Vert (2018): we define a distance d on the set of rankings \mathcal{R}_{n_a} and show that (\mathcal{R}_{n_a}, d) is isometric to an L_2 space. This ensures that d is a conditionally positive definite (c.p.d.) function and, thus, that $e^{-\nu d}$ is positive definite (Schoenberg, 1938; Schölkopf, 2000). Our proof for the Jaccard kernel, instead, follows without much effort from previous results. For ease of reading, we restate the definitions as well.

Definition A.1 (Borda kernel).

$$\kappa_h^{a^*,\nu}(r_1, r_2) = e^{-\nu|b_1 - b_2|},$$
(5)

where $b_l = \#\{a \in A : r_l(a) \ge r_l(a^*)\}$ is the number of alternatives dominated by a^* in r_l and $\nu \in \mathbb{R}$.

Proposition A.1. The Borda kernel as defined in (5) is a kernel.

Proof. Define a distance

$$d: \mathcal{R}_{n_a} \times \mathcal{R}_{n_a} \to \mathbb{R}^+$$
$$(r_1, r_2) \mapsto ||b_1 - b_2||,$$

where $b_l = \{a \in A : r_l(a) \ge r_l(a^*)\}$ is the number of alternatives dominated by a^* in r_l . Now, (\mathcal{R}_{n_a}, d) is isometric to $(\mathbb{R}, \|\cdot\|_2)$ via the map $r_l \mapsto b_l$. Hence, d is c.p.d. and κ_b is a kernel.

Definition A.2 (Jaccard kernel).

$$\kappa_j^k(r_1, r_2) = \frac{\left| r_1^{-1}([k]) \cap r_2^{-1}([k]) \right|}{\left| r_1^{-1}([k]) \cup r_2^{-1}([k]) \right|},\tag{6}$$

where $r^{-1}([k]) = \{a \in A : r(a) \le k\}$ is the set of alternatives whose rank is better than or equal to k.

Proposition A.2. The Jaccard kernel as defined in (6) is a kernel.

Proof. It is already know that the Jaccard coefficients for sets is a kernel (Gärtner et al., 2006; Bouchard et al., 2013). As the Jaccard kernel for rankings is equivalent to the Jaccard coefficient for the k-best tiers of said rankings, the former is also a kernel.

Definition A.3 (Mallows kernel).

$$\kappa_m^{\nu}(r_1, r_2) = e^{-\nu n_d},\tag{7}$$

where $n_d = \sum_{a_1,a_2 \in A} |\operatorname{sign}(r_1(a_1) - r_1(a_2)) - \operatorname{sign}(r_2(a_1) - r_2(a_2))|$ is the number of discordant pairs and $\nu \in \mathbb{R}$.

Proposition A.3. The Mallows kernel as defined in (7) is a kernel.

Proof. The number of discordant pairs n_d is a distance on \mathcal{R}_{n_a} (Snell & Kemeny, 1962). Consider now the mapping of a ranking into its adjacency matrix,

$$\begin{split} \Phi: \mathcal{R}_{n_a} &\to \left\{0, 1\right\}^{n_a \times n_a} \\ r &\mapsto \left(\mathbf{1}\left(r(i) \leq r(j)\right)\right)_{i, j = 1}^{n_a}, \end{split}$$

where 1 is the indicator function. Then,

$$n_d = \|\Phi(r_1) - \Phi(r_2)\|_1 = \|\Phi(r_1) - \Phi(r_2)\|_2^2$$

where $\|\cdot\|_p$ indicates the entry-wise matrix p-norm and the equality holds because the entries of the matrices are either 0 or 1. As a consequence, (\mathcal{R}_{n_a}, n_d) is isometric to $(\mathbb{R}^{n_a \times n_a}, \|\cdot\|_2)$ via Φ . Hence, n_d is c.p.d. and κ_m is a kernel.

A.2 Details for Section 4.3

Proposition 4.1. The MMD takes values in $\left[0, \sqrt{2 \cdot (\kappa_{sup} - \kappa_{inf})}\right]$, where $\kappa_{sup} = \sup_{x,y \in X} \kappa(x,y)$ and $\kappa_{inf} = \inf_{x,y \in X} \kappa(x,y)$.

Proof.

$$0 \leq \text{MMD}_{\kappa}(\mathbf{x}, \mathbf{y})^{2} = \frac{1}{n^{2}} \sum_{i,j=1}^{n} \kappa(x_{i}, x_{j}) + \frac{1}{m^{2}} \sum_{i,j=1}^{m} \kappa(y_{i}, y_{j}) - \frac{2}{mn} \sum_{\substack{i=1...n\\j=1...m}} \kappa(x_{i}, y_{j})$$
$$\leq \frac{1}{n^{2}} \sum_{i,j=1}^{n} \kappa_{\text{sup}} + \frac{1}{m^{2}} \sum_{i,j=1}^{n} \kappa_{\text{sup}} - \frac{2}{mn} \sum_{\substack{i=1...n\\j=1...m}} \kappa_{\text{inf}} = 2(\kappa_{\text{sup}} - \kappa_{\text{inf}})$$

A.3 Details for Section 4.4

A.3.1 Distribution-free linear relation

Proposition A.4. Let ε_n^{α} be the α -quantile of the distribution of the MMD for samples of size n. For any $\alpha \in [0,1]$, there exist $\beta_0, \beta_1 \in \mathbb{R}$, $\beta_i = \beta_i(\alpha)$, s.t., for any distribution \mathbb{P} ,

$$\log(n) = \beta_1 \log(\varepsilon_n^{\alpha}) + \beta_0.$$

Proof. We use the distribution-free upper bound on the MMD defined in Gretton et al. (2012, Theorem 8). Let X and Y be samples of size n from \mathbb{P} and define $\mathrm{MMD}_n = \mathrm{MMD}(X,Y)$.

$$\mathbb{P}^{n} \otimes \mathbb{P}^{n} \left(\operatorname{MMD}_{n} - \sqrt{\frac{2\kappa_{\operatorname{sup}}}{n}} > \varepsilon \right) < \exp\left(-\frac{n\varepsilon^{2}}{4\kappa_{\operatorname{sup}}} \right)$$

$$\xrightarrow{(1)} \mathbb{P}^{n} \otimes \mathbb{P}^{n} \left(\operatorname{MMD}_{n} > \varepsilon' \right) < \exp\left(-\frac{n\left(\varepsilon' - \sqrt{\frac{2\kappa_{\operatorname{sup}}}{n}}\right)^{2}}{4\kappa_{\operatorname{sup}}} \right)$$

$$\xrightarrow{(2)} \mathbb{P}^{n} \otimes \mathbb{P}^{n} \left(\operatorname{MMD}_{n} > n^{-\frac{1}{2}} \left(\sqrt{-\log\left(1 - \alpha\right) 4\kappa_{\operatorname{sup}}} \right) + \sqrt{2\kappa_{\operatorname{sup}}} \right) < 1 - \alpha$$

$$\xrightarrow{(3)} \mathbb{P}^{n} \otimes \mathbb{P}^{n} \left(\operatorname{MMD}_{n} \le n^{-\frac{1}{2}} \left(\sqrt{-\log\left(1 - \alpha\right) 4\kappa_{\operatorname{sup}}} \right) + \sqrt{2\kappa_{\operatorname{sup}}} \right) \ge \alpha$$

where:

(1) Define
$$\varepsilon' = \varepsilon + \sqrt{\frac{2\kappa_{\text{sup}}}{n}}$$
.

(2) Define
$$1 - \alpha = \exp\left(-\frac{n\left(\varepsilon' - \left(\frac{2\kappa_{\text{sup}}}{n}\right)\right)^2}{4\kappa_{\text{sup}}}\right)$$
 and $\varepsilon' = n^{-\frac{1}{2}}\left(\sqrt{-\log\left(1 - \alpha\right)4\kappa_{\text{sup}}} + \sqrt{2\kappa_{\text{sup}}}\right)$.

(3) Take the complementary event

Now, let

$$\begin{split} & \varepsilon_n^\alpha = n^{-\frac{1}{2}} \left(\sqrt{-\log\left(1 - \alpha\right) 4\kappa_{\sup}} \right) + \sqrt{2\kappa_{\sup}} \\ \Rightarrow & n = \left(\varepsilon_n^\alpha\right)^{-2} \left(\sqrt{-4\kappa_{\sup} \log\left(1 - \alpha\right)} + \sqrt{2\kappa_{\sup}} \right)^2 \\ \Rightarrow & \log(n) = -2\log(\varepsilon_n^\alpha) + 2\log\left(\sqrt{-4\kappa_{\sup} \log\left(1 - \alpha\right)} + \sqrt{2\kappa_{\sup}}\right). \end{split}$$

The proof is conclude by defining

$$\beta_0 = \log(2\kappa_{\sup}) + \log\left(\sqrt{-2\log(1-\alpha)} + 1\right)$$

$$\beta_1 = -2.$$

Remark. Proposition A.4 leads to overconservative estimates of n^* , usually one or two orders of magnitude greater than the empirical estimate.

B Details for Section 5

B.1 Estimation of n^* from N preliminary experiments

In this section, we discuss how accurately our method estimates n^* from N independent experiments (cf. Section 4.4). Recall that, for a research question Q with corresponding results \mathbb{P} ,

$$n^* = \min \left\{ n \in \mathbb{N} : \mathbb{P}^n \otimes \mathbb{P}^n \left((X_j)_{j=1}^n, (Y_j)_{j=1}^n : \text{MMD}(X, Y) \le \varepsilon \right) \ge \alpha^* \right\},\,$$

where α^* and ε^* are the desired generalizability and dissimilarity respectively. We estimate n^* by running N preliminary experiments and using their results $\hat{\mathbb{P}}_N$ as an approximation for \mathbb{P} (cf. Algorithm 1). We call the obtained estimate n_N^* .

We evaluate our approach with the relative error $(n^* - n_N^*)/n^*$. A relative error greater (lower) than 0 means that we are overestimating (underestimating) n^* .

B.1.1 Synthetic data

We experimented with two families of distributions.

Uniform_{n_a} The uniform distribution assigns the same probability to every ranking in \mathcal{R}_{n_a} . In our experiments, we varied n_a from 2 to 15. Perhaps surprisingly, our estimate improved with the number of alternatives, despite the distribution becoming more "complex".

 $m ext{-}\mathbf{Degenerate}_{n_a}$ The degenerate distribution assigns the same probability to m rankings in \mathcal{R}_{n_a} . As m approaches $|\mathcal{R}_{n_a}|$, the m-degenerate distribution becomes closer to a uniform. In our experiments, we fixed $n_a = 4$ ($|\mathcal{R}_{n_a}| = 75$) and experimented with m ranging from $^{75}/8 \approx 9$ to $^{75}/2 \approx 37$. This parameter doesn't seem to affect the results.

Figure 6 shows that there are no major discrepancies in the behavior of our estimate.

B.1.2 Real data

This section evaluates the influence of the number of preliminary experiments N on n^* . We consider, for both case studies (cf. Section 5), the design factor combinations for which we have at least 50 experiments. This results in 23 out of 48 combinations for the categorical encoders and 9 out of 24 combinations for the

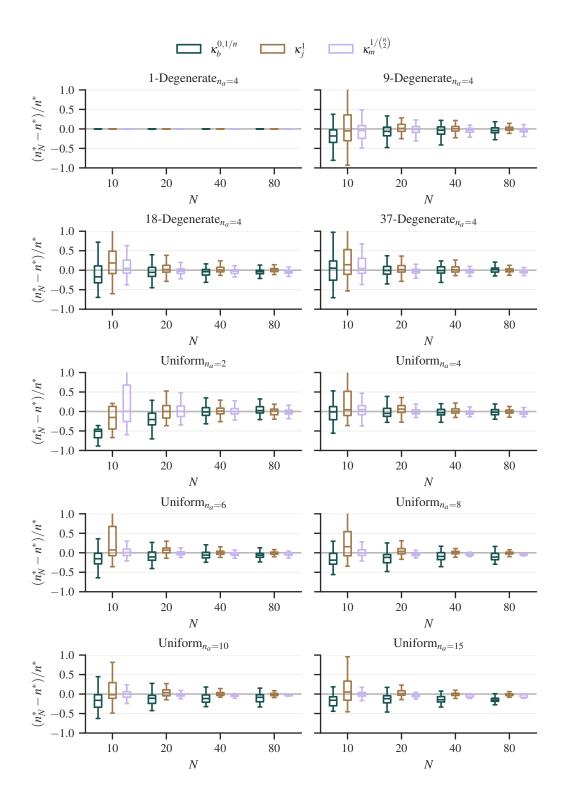


Figure 6: Relative error of the prediction of n^* from N preliminary experiments (n_N^*) for multiple synthetic distributions and the three kernels introduced in Section 4.2.

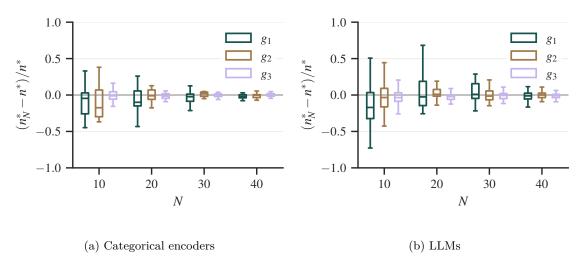


Figure 7: Relative error between the estimate of n^* from N preliminary experiments and n_{50}^* .

LLMs. For each of those combinations and as we do not have access to the true value n^* , we consider the estimate n^*_{50} as the ground truth and observe how the estimates of n^* for N < 50 differ. Figure 7 shows that even N = 10 preliminary experiments provide an estimate within of 50% of n^*_{50} , validating the fact that our method can return reasonably good estimates even for very small sample size.