Learning Counterfactually Invariant Predictors

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Abstract

Notions of counterfactual invariance (CI) have proven essential for predictors that are fair, robust, and generalizable in the real world. We propose graphical criteria that yield a sufficient condition for a predictor to be counterfactually invariant in terms of a conditional independence in the observational distribution. In order to learn such predictors, we propose a model-agnostic framework, called Counterfactually Invariant Prediction (CIP), building on the Hilbert-Schmidt Conditional Independence Criterion (HSCIC), a kernel-based conditional dependence measure. Our experimental results demonstrate the effectiveness of CIP in enforcing counterfactual invariance across various simulated and real-world datasets including scalar and multi-variate settings.

1 Introduction

Invariance, or equivariance to certain data transformations, has proven essential in numerous applications of machine learning (ML), since it can lead to better generalization capabilities \cite{Arjovsky2019, Bloem-Reddy2020, Chen2020}. For instance, in image recognition, predictions ought to remain unchanged under scaling, translation, or rotation of the input image. Data augmentation, an early heuristic to promote such invariances, has become indispensable for successfully training deep neural networks (DNNs) \cite{Shorten2019, Xie2020}. Well-known examples of “invariance by design” include convolutional neural networks (CNNs) for translation invariance \cite{Krizhevsky2012}, group equivariant NNs for general group transformations \cite{Cohen2016}, recurrent neural networks (RNNs) and transformers for sequential data \cite{Vaswani2017}, DeepSet \cite{Zaheer2017} for sets, and graph neural networks (GNNs) for different types of geometric structures \cite{Battaglia2018}.

Many applications in modern ML, however, call for arguably stronger notions of invariance based on causality. This case has been made for image classification, algorithmic fairness \cite{Hardt2016, Mitchell2021}, robustness \cite{Buehlmann2020}, and out-of-distribution generalization \cite{Lu2021}. The goal is invariance with respect to hypothetical manipulations of the data generating process (DGP). Various works develop methods that assume observational distributions (across environments or between training and test) to be governed by shared causal mechanisms, but differ due to various types of distribution shifts encoded by the causal model \cite{Arjovsky2019, Buehlmann2020, Heinze-Deml2018, Makar2022, *Part of this work was done while Francesco Quinzan visited the Max Planck Institute for Intelligent Systems, Tübingen, Germany.
While we share the broader motivation, these works are orthogonal to ours, because even though counterfactual distributions are also generated from the same causal model, they are fundamentally different from such shifts [Peters et al., 2017]. Intuitively, counterfactuals are about events that did not, but could have happened had circumstances been different in a controlled way. A formal discussion of what we mean by counterfactuals is required to properly position our work in the existing literature and describe our contributions.

2 Problem setting and related work

2.1 Preliminaries and terminology

Definition 2.1 (Structural causal model (SCM)). A structural causal model is a tuple $S = (U, V, F, P_U)$ such that $U$ is a set of background variables that are exogenous to the model; $V$ is a set of observable (endogenous) variables; $F = \{f_v \}_{v \in V}$ is a set of functions from (the domains of) $pa(V) \cup U_V$ to (the domain of) $V$, where $U_V \subseteq U$ and $pa(V) \subseteq V \setminus \{V\}$ such that $V = f_v(pa(V), U_V)$; $P_U$ is a probability distribution over the domain of $U$. Further, the subsets $pa(V) \subseteq V \setminus \{V\}$ are chosen such that the graph $G$ over $V$ where the edge $V' \to V$ is in $G$ if and only if $V' \in pa(V)$ is a directed acyclic graph (DAG).

Observational distribution. An SCM implies a unique observational distribution over $V$, which can be thought of as being generated by transforming the distribution over $P_U$ via the deterministic functions in $F$ iteratively to obtain a distribution over $V$.

Interventions. Given a variable $A \in V$, an intervention $A \leftarrow a$ amounts to replacing $f_A$ in $F$ with the constant function setting $A$ to $a$. This yields a new SCM, which induces the \textit{interventional distribution} under intervention $A \leftarrow a$. Similarly, we can intervene on multiple variables $V \supseteq A \leftarrow a$. For an outcome (or prediction target) variable $Y \subseteq V$, we then write $Y_a$ for the outcome in the intervened SCM, also called \textit{potential outcome}. Note that the interventional distribution $P_{Y_a}(y)$ differs in general from the conditional distribution $P_{Y|A}(y \mid a)$. This is typically the case when $Y$ and $A$ have a shared ancestor, i.e., they are confounded. In interventional distributions, potential outcomes are random variables via the exogenous variables $u$, i.e., $Y_a(u)$ where $u \sim P_U$. Hence, interventions capture “population level” properties, i.e., the action is performed for all units $u$.

Counterfactuals. Counterfactuals capture what happens under interventions for a “subset” of possible units $u$ that are compatible with observations $W = w$ for a subset of observed variables $W \subseteq V$. This can be described in a three step procedure. (i) \textit{Abduction}: We restrict our attention to units compatible with the observations, i.e., consider the new SCM $S^w = (U, V, F, P_{U|W = w})$. (ii) \textit{Intervention}: Within $S^w$, perform an intervention $A \leftarrow a$ on some variables $A$ (which need not be disjoint from $W$). (iii) \textit{Prediction}: Finally, we are typically interested in the outcome $Y$ in an interventional distribution of $S^w$, which we denote by $P_{Y_a|W = w}(y)$ and call a \textit{counterfactual distribution}: “Given that we have observed $W = w$,

\footnote{Note that all randomness stems from $P_U$. The observational distribution is well-defined and unique, essentially because every DAG allows for a topological order.}

\footnote{The observational distribution in an intervened SCM is called interventional distribution of the base SCM.}

\footnote{We use $P$ for distributions (common in the kernel literature) and $Y_a$ instead of the do notation.}
what would \( Y \) have been had we set \( A \leftarrow a \), instead of the value \( A \) has actually taken?". In this notation the conditioning set \( W \) is conditioned on in the pre-interventional world. Counterfactuals capture properties of a “subpopulation” \( u \sim P_{U \mid W=w} \) compatible with the observations.\(^4\) Even for fine-grained \( W \), there may be multiple units \( u \) in the support of this distribution. In contrast, “unit level counterfactuals” often considered in philosophy contrast \( Y_u^a(u) \) with \( Y_u^a(u) \) for a single unit \( u \). Such unit level counterfactuals are too fine-grained in our setting. Hence, our used definition of counterfactual invariance is:

**Definition 2.2 (Counterfactual invariance).** Let \( A, W \) be (not necessarily disjoint) sets of nodes in a given SCM. Then, \( Y \) is **counterfactually invariant** in \( A \) w.r.t. \( W \) if \( P_{Y_u^a \mid W=w}(y) = P_{Y_u^a \mid W=w} \) almost surely, for all \( a, a' \) in the domain of \( A \) and all \( w \) in the domain of \( W \).\(^5\)

**Predictors in SCMs.** Ultimately, we aim at learning a predictor \( \hat{Y} \) for the outcome \( Y \). Originally, the predictor \( \hat{Y} \) is not part of the DGP, because we get to learn \( f_Y \) from data. Using supervised learning, the predictor \( f_Y \) depends both on the chosen inputs \( X \subset V \) as well as the target \( Y \). However, once \( f_Y \) is fixed, it is a deterministic function with arguments \( X \subset V \), so \( (U, V \cup \{Y\}, F \cup \{f_Y\}, P_U) \) is a valid SCM and we can consider \( \hat{Y} \) an observed variable with incoming arrows from only \( X \). Hence, the definition of counterfactual invariance can be applied to the predictor \( \hat{Y} \).

**Kernel mean embeddings (KME).** Our method relies on kernel mean embeddings (KMEs). We describe the main concepts pertaining KMEs and refer the reader to Berlinet and Thomas-Agnan [2011], Muandet et al. [2017], Schölkopf et al. [2002], Smola et al. [2007] for details. Fix a measurable space \( \mathcal{Y} \) with respect to a \( \sigma \)-algebra \( F_\mathcal{Y} \), and consider a probability measure \( P \) on the space \( (\mathcal{Y}, F_\mathcal{Y}) \). Let \( \mathcal{H} \) be a reproducing kernel Hilbert space (RKHS) with a bounded kernel \( k_Y : \mathcal{Y} \times \mathcal{Y} \to \mathbb{R} \), i.e., \( k_Y \) is such that \( \sup_{y \in \mathcal{Y}} k(y, y) < \infty \). The kernel mean embedding \( \mu_P \) of \( P \) is defined as the expected value of the function \( k(\cdot, y) \) with respect to \( y \), i.e., \( \mu_P := \mathbb{E}[k(\cdot, y)] \). The definition of KMEs can be extended to conditional distributions [Fukumizu et al., 2013, Grünewälder et al., 2012, Song et al., 2013, 2009]. Consider two random variables \( Y, S \), and denote with \( (\Omega_Y, F_Y) \) and \( (\Omega_S, F_S) \) the respective measurable spaces. These random variables induce a probability measure \( P_{Y \mid S} \) in the product space \( \Omega_Y \times \Omega_S \). Let \( H_Y \) be a RKHS with a bounded kernel \( k_Y(\cdot, \cdot) \) on \( \Omega_Y \). We define the KME of a conditional distribution \( P_{Y \mid S}(\cdot \mid s) \) via \( \mu_{P_{Y \mid S}(\cdot \mid s)} := \mathbb{E}[k_Y(\cdot, y) \mid S = s] \). Here, the expected value is taken over \( y \). KMEs of conditional measures can be estimated from samples [Grünewälder et al., 2012]. Pogodin et al. [2022] recently proposed an efficient kernel-based regularizer for learning features of input data that allow for estimating a target while being conditionally independent of a distractor given the target. Since CIP ultimately enforces conditional independence (see Theorem 3.2), we believe it could further benefit from leveraging the efficiency and convergence properties of their technique, which we leave for future work.

2.2 Related work and contributions

While we focus on counterfactuals in the SCM framework [Pearl, 2000, Peters et al., 2016], there are different incompatible frameworks to describe counterfactuals [Dorr, 2016, von Kügelgen et al., 2022, Woodward, 2021], which may give rise to orthogonal notions of counterfactual invariance.

Research on algorithmic fairness has explored a plethora of causal “invariance” notions with

\(^4\)Note that conditioning in an interventional distribution is different from a counterfactual and our notation is quite subtle here \( P_{Y_u^a} \mid W=w \neq P_{Y_u^a \mid W=w} \).

\(^5\)With an abuse of notation, if \( W = \emptyset \) then the requirement of conditional counterfactual invariance becomes \( P_{Y_u^a} = P_{Y_u^a} \) almost surely, for all \( a, a' \) in the domain of \( A \). The “almost surely” part in our definition merely refers to the type of equality of distributions and is not related to the “almost sure” in a.s.-CI defined by Fawkes and Evans [2023].
the goal of achieving fair predictors [Carey and Wu, 2022, Loftus et al., 2018, Plecko and Bareinboim, 2022]. Kilbertus et al. [2017] conceptually introduce a notion based on group-level interventional, which has been refined to take into account more fine-grained context by Galhotra et al. [2022], Salimi et al. [2019], who then obtain fair predictors by viewing it as a database repair problem or a causal feature selection problem, respectively. A counterfactual-level definition was proposed by Kusner et al. [2017] and followed up by path-specific counterfactual notions [Chiappa, 2019, Nabi and Shpitser, 2018], where the protected attribute may take different values along different paths to the outcome. Recently, Dutta et al. [2021] developed an information theoretic framework to decompose the overall causal influence allowing for exempted variables and properly dealing with synergies across different paths.

Our focus is on counterfactuals because they are fundamentally more expressive than mere interventions [Bareinboim et al., 2022, Pearl, 2000], but do not require a fine-grained path- or variable-level judgment of “allowed” and “disallowed” paths or variables, which may be challenging to devise in practice. Since CI already requires strong assumptions, we leave path-specific counterfactuals—even more challenging in terms of identifiability [Avin et al., 2005]—for future work. While our Definition 2.2 requires equality in distribution, Veitch et al. [2021] suggest a definition of a counterfactually invariant predictor \( \hat{f}_Y \) which requires almost sure equality of \( \hat{Y}^*_a \) and \( \hat{Y}^*_{a'} \), where we view \( \hat{Y} \) as an observed variable in the SCM as described above. Fawkes and Evans [2023] recently carefully formulated various precise technical definitions of what counterfactual invariance may mean in different contexts, such as “almost sure CI” (a.s.-CI in Definition A.1), “distributional CI” (D-CI in Definition A.3), and “CI of predictors” (F-CI in Definition A.2). They provide various connections between them such as the fact that \( f_Y \) being F-CI is equivalent to \( \hat{Y} \) being D-CI conditioned on \( X \), yielding an equivalence to the definition of counterfactual fairness [Kusner et al., 2017]. The notion of counterfactual invariance in Definition 2.2 is most closely related to D-CI from Fawkes and Evans [2023], but we do not enforce conditioning on the intervening variable.

Inspired by problems in natural language processing (NLP), Veitch et al. [2021] aim at “stress-testing” models for spurious correlations. It differs from our work in that they (i) focus only on two specific graphs, and (ii) provide a necessary but not sufficient criterion for CI in terms of a conditional independence. Their method enforces the conditional independence via maximum mean discrepancy (MMD) (in discrete settings only). However, enforcing a consequence of CI, does not necessarily improve CI. Indeed, Fawkes and Evans [2023, Prop. 4.4] show that while a.s.-CI implies certain conditional independencies, no set of conditional independencies implies any bounds on the difference in counterfactuals. On the contrary, distributional notions of CI such as D-CI or our Definition 2.2 are weaker than a.s.-CI (see [Fawkes and Evans, 2023, Lem. 2.4]). Therefore, these weaker notions can indeed be written equivalently as conditional independencies in the observational distribution under additional assumptions about the data generating mechanism [Fawkes and Evans, 2023, Lem. A.3]. For example, Fawkes and Evans [2023, Lem. A.3] shows that D-CI can be implied by conditional independence in special settings where the counterfactual distribution is identified from the observational one. Similarly, our reduction of CI (as in Definition 2.2 ) to conditional independence in Theorem 3.2 requires a strong injectivity assumption that essentially amounts to being able to remove exogenous uncertainty.

**Contributions.** We provide such a sufficient graphical criterion for D-CI under an injectivity condition of a structural equation. Depending on the assumed causal graph, this can also come at the cost of requiring certain variables to be observed. As our main contribution, we propose a model-agnostic learning framework, called Counterfactually Invariant Prediction (CIP), using a kernel-based conditional dependence measure that also works for mixed categorical and continuous, multivariate variables. We evaluate CIP extensively in (semi-)synthetic settings and demonstrate its efficacy in enforcing counterfactual invariance even when the strict assumptions may be violated.
We now state the sufficient graphical criterion that renders CI equivalent to conditional independence. The proof builds on known results from the literature such as the full characterization of valid adjustment sets in fully observed structural causal models and the backdoor criteria, but the final statement requires a substantial original theoretical contribution, which we detail in Section B.

**Theorem 3.2.** Let $\mathcal{G}$ be a causal graph, $A$, $W$ be two (not necessarily disjoint) sets of nodes in $\mathcal{G}$, such that $(A \cup W) \cap Y = \emptyset$, let $S$ be a valid adjustment set for $(A \cup W, Y)$. Further, for any random variable $X \in W \setminus A$ denote with $g_X(\text{pa}(X), U_X)$ its structural equation, and suppose that $\text{pa}(X) \subseteq A \cup W$. Suppose that $g_X$ is injective in the variable $u$. Then, in all

$$X \not\perp Y \mid S \quad \text{for any random variable } X \in A \cup W \setminus \text{pa}(X).$$
SCMs compatible with $G$, if a predictor $\hat{Y}$ satisfies $\hat{Y} \perp \perp A \cup W \mid S$, then $\hat{Y}$ is counterfactually invariant in $A$ with respect to $W$.

**Assumptions.** First, we do not assume the SCM (or any structural equations) to be known. We do assume the causal graph to be known. This is a standard assumption widely made in the causality literature, even though it is a strong one [Cartwright, 2007]. The additional assumption of Theorem 3.2, namely injectivity of $g_X$ is satisfied—but more general than—a wide variety of commonly considered models in causality such as the widely used Additive Noise Models [Peters et al., 2017]. More broadly, Fawkes and Evans [2023, Lem. A.3] show that to achieve CI from the observational distribution and the causal graph, additional assumptions are always required.

### 3.2 Example use-cases of counterfactually invariant prediction

Fig. 1(a) shows an exemplary graph in which the outcome $Y$ is affected by (disjoint) sets $A$ (in which we want to be CI), $L$ and $S$ (inputs to $f$). We consider $W = L \cup A \cup S$. Here we aim to achieve $\hat{Y} \perp \perp A \cup L \mid S$ to obtain CI in $A$ w.r.t. $W$. In our synthetic experiments, we also allow $S$ to affect $A$, see Fig. 1(d). Let us further illustrate concrete potential applications of CI, which we later also study in our experiments.

**Counterfactual fairness.** Counterfactual fairness [Kusner et al., 2017] informally challenges a consequential decision: “Would I have gotten the same outcome had my gender, race, or age been different with all else being equal?”. Here $Y \subset V$ denotes the outcome and $A \subset V \setminus Y$ the protected attributes such as gender, race, or age—protected under anti-discrimination laws [Barocas and Selbst, 2016]—by $A \subset V \setminus Y$. Collecting all remaining observed covariates into $W := V \setminus Y$ counterfactual fairness reduces to counterfactual invariance. In experiments, we build a semi-synthetic DGP assuming the graph in Fig. 1(e) for the UCI adult dataset [Kohavi and Becker, 1996].

**Robustness.** CI serves as a strong notion of robustness in settings such as image classification: “Would the truck have been classified correctly had it been winter in this situation instead of summer?” For concrete demonstration, we use the dSprites dataset [Matthey et al., 2017] consisting of simple black and white images of different shapes (squares, ellipses, ...), sizes, orientations, and locations. We devise a DGP for this dataset with the graph depicted in Fig. 1(f).

**Text classification.** Veitch et al. [2021] motivate the importance of counterfactual invariance in text classification tasks. Specifically, they consider the causal and anti-causal structures depicted in Veitch et al. [2021, Fig. 1], which we replicate in Fig. 1(b,c). Both diagrams consist of protected attributes $A$, observed covariates $X$, and outcomes $Y$. To apply our sufficient criterion to their settings, we must assume that $A$ and $Y$ are unconfounded. We show in Section H.1 that CIP still performs on par with Veitch et al. [2021] even when this assumption is violated. Theorem 3.2 provides a sufficient condition for CI (Definition 2.2) in terms of the conditional independence $\hat{Y} \perp \perp A \cup W \mid S$. We next develop an operator $\text{HSCIC}(\hat{Y}, A \cup W \mid S)$ that is (a) efficiently estimable from data, (b) differentiable, (c) a monotonic measure of conditional dependence, and (d) is zero if and only if $\hat{Y} \perp \perp A \cup W \mid S$. Hence, it is a practical objective to enforce CI.

Consider $\text{pa}(X)$ and $u, u'$ in the support of $U_X$. Suppose that $P_{\text{pa}(X),U_X}(p, u) \neq 0$ and $P_{\text{pa}(X),U_X}(p, u') \neq 0$. Then, it holds $g(p, u) = g(p, u')$ if and only if $u = u'$.
3.3 HSCIC for conditional independence

Consider two sets of random variables \( \mathbf{Y} \) and \( \mathbf{A} \cup \mathbf{W} \), and denote with \((\Omega_{\mathbf{Y}}, \mathcal{F}_\mathbf{Y})\) and \((\Omega_{\mathbf{A} \cup \mathbf{W}}, \mathcal{F}_{\mathbf{A} \cup \mathbf{W}})\) the respective measurable spaces. Suppose that we are given two RKHSs \( \mathcal{H}_\mathbf{Y} \), \( \mathcal{H}_{\mathbf{A} \cup \mathbf{W}} \) over the support of \( \mathbf{Y} \) and \( \mathbf{A} \cup \mathbf{W} \) respectively. The tensor product space \( \mathcal{H}_\mathbf{Y} \otimes \mathcal{H}_{\mathbf{A} \cup \mathbf{W}} \) is defined as the space of functions of the form \( f \otimes g(y, [a, w]) := f(y)g([a, w]) \), for all \( f \in \mathcal{H}_\mathbf{Y} \) and \( g \in \mathcal{H}_{\mathbf{A} \cup \mathbf{W}} \). The tensor product space yields a natural RKHS structure, with kernel \( k \) defined by \( k(y \otimes [a, w], y' \otimes [a', w']) := k(y, y')k_{\mathbf{A} \cup \mathbf{W}}([a, w], [a', w']) \). We refer the reader to Szabó and Sriperumbudur [2017] for more details on tensor product spaces.

**Definition 3.3 (HSCIC).** For (sets of) random variables \( \mathbf{Y}, \mathbf{A} \cup \mathbf{W}, \mathbf{S} \), the HSCIC between \( \mathbf{Y} \) and \( \mathbf{A} \cup \mathbf{W} \) given \( \mathbf{S} \) is defined as the real-valued random variable \( \text{HSCIC}(\mathbf{Y}, \mathbf{A} \cup \mathbf{W} | \mathbf{S}) = \mathcal{H}_{\mathbf{Y}, \mathbf{A} \cup \mathbf{W}|\mathbf{S}} \circ \mathbf{S} \) where \( \mathcal{H}_{\mathbf{Y}, \mathbf{A} \cup \mathbf{W}|\mathbf{S}} \) is a real-valued deterministic function, defined as \( \mathcal{H}_{\mathbf{Y}, \mathbf{A} \cup \mathbf{W}|\mathbf{S}}(s) := \| \mathcal{H}_{\mathbf{Y}, \mathbf{A} \cup \mathbf{W}|\mathbf{S}=s} - \mathcal{H}_{\mathbf{Y}|\mathbf{S}=s} \otimes \mathcal{H}_{\mathbf{A} \cup \mathbf{W}|\mathbf{S}=s} \| \) with \( \| \cdot \| \) the norm induced by the inner product of the tensor product space \( \mathcal{H}_\mathbf{X} \otimes \mathcal{H}_{\mathbf{A} \cup \mathbf{W}} \).

Our Definition 3.3 is motivated by, but differs slightly from Park and Muandet [2020, Def. 5.3], which relies on the Bochner conditional expected value. While it is functionally equivalent (with the same implementation, see Eq. (2)), ours has the benefit of bypassing some technical assumptions required by Park and Muandet [2020] (see Sections D and E for details). The HSCIC has the following important property, proved in Section C.

**Theorem 3.4** (Theorem 5.4 by Park and Muandet [2020]). If the kernel \( k \) of \( \mathcal{H}_\mathbf{X} \otimes \mathcal{H}_{\mathbf{A} \cup \mathbf{W}} \) is characteristic\(^5\), \( \text{HSCIC}(\mathbf{Y}, \mathbf{A} \cup \mathbf{W} | \mathbf{S}) = 0 \) almost surely if and only if \( \mathbf{Y} \perp \mathbf{A} \cup \mathbf{W} | \mathbf{S} \).

Because “most interesting” kernels such as the Gaussian and Laplacian kernels are characteristic, and the tensor product of translation-invariant characteristic kernels is characteristic again [Szabó and Sriperumbudur, 2017], this natural assumption is non-restrictive in practice. Combining Theorems 3.2 and 3.4, we can now use HSCIC to reliably achieve counterfactual invariance.

**Corollary 3.5.** Under the assumptions of Theorem 3.2, if \( \text{HSCIC}(\hat{\mathbf{Y}}, \mathbf{A} \cup \mathbf{W} | \mathbf{S}) = 0 \) almost surely, then \( \hat{\mathbf{Y}} \) is counterfactually invariant in \( \mathbf{A} \) with respect to \( \mathbf{W} \).

In addition, since HSCIC is defined in terms of the MMD (Definition 3.3 and Park and Muandet [2020, Def. 5.3]), it inherits the weak convergence property, i.e., if \( \text{HSCIC}(\hat{\mathbf{Y}}, \mathbf{A} \cup \mathbf{W} | \mathbf{S}) \) converges to zero, then the counterfactual distributions (for different intervention values \( \mathbf{a} \)) weakly converge to the same distribution. We refer to Simon-Gabriel et al. [2020], Simon-Gabriel and Schölkopf [2018] for a precise characterization. Hence, as HSCIC decreases, the predictor approaches counterfactual invariance and we need not drive HSCIC all the way to zero to obtain meaningful results.

3.4 Learning counterfactually invariant predictors (CIP)

Corollary 3.5 justifies our proposed objective, namely to minimize the following loss

\[
\mathcal{L}_\text{CIP}(\hat{\mathbf{Y}}) = \mathcal{L}(\hat{\mathbf{Y}}) + \gamma \cdot \text{HSCIC}(\hat{\mathbf{Y}}, \mathbf{A} \cup \mathbf{W} | \mathbf{S}),
\]

where \( \mathcal{L}(\hat{\mathbf{Y}}) \) is a task-dependent loss function (e.g., cross-entropy for classification, or mean squared error for regression) and \( \gamma \geq 0 \) is a parameter that regulates the trade-off between predictive performance and counterfactual invariance.

**The meaning of \( \gamma \) and how to choose it.** The second term in Eq. (1) amounts to the additional objective of CI, which is typically at odds with predictive performance within the observational distribution \( \mathcal{L} \). In practice, driving HSCIC to zero, i.e., viewing our task as a constrained optimization problem, typically deteriorates predictive performance too much.

\(^5\)The tensor product kernel \( k \) is characteristic if \( \mathbb{P}_{\mathbf{Y}, \mathbf{A} \cup \mathbf{W}} \rightarrow \mathbb{E}_{\mathbf{Y}|[a, w]}[k(\cdot, y \otimes [a, w])] \) is injective.
to be useful for prediction—especially in small data settings.\textsuperscript{8} As the choice of $\gamma$ amounts to choosing an operating point between predictive performance and CI, it cannot be selected in a data-driven fashion. As different settings call for different tradeoffs, we advocate for employing the following procedure: (i) Train an unconstrained predictor for a base predictive performance (e.g., 92% accuracy or 0.21 MSE). (ii) Fix a tolerance level $\alpha$, indicating the maximally tolerable loss in predictive performance (e.g., at most 5% drop in accuracy or at most 10% increase in MSE). (iii) Perform a log-spaced binary search on $\gamma$ (e.g., on $[10^{-4}, 10^{4}]$) to find the largest $\gamma$ such that the predictive performance of the resulting predictor achieves predictive performance within the tolerance $\alpha$—see Section G.6 for an illustration. A similar search for the optimal value of $\gamma$ can be conducted, when there is a fixed requirement for a maximum tolerance of counterfactual invariance as measured by HSCIC.

**Estimating the HSCIC from samples.** The key benefit of HSCIC as a conditional independence measure is that it does not require parametric assumptions on the underlying probability distributions, and it is applicable for any mixed, multi-dimensional data modalities, as long as we can define positive definite kernels on them. Given $n$ samples $\{(\hat{y}_i, a_i, w_i, s_i)\}_{i=1}^n$, denote with $\hat{K}_Y$ the kernel matrix with entries $[\hat{K}_Y]_{i,j} := k_Y(\hat{y}_i, \hat{y}_j)$, and let $\hat{K}_{A \cup W}$ be the kernel matrix for $A \cup W$. We estimate the $H_{\hat{Y}, A \cup W \mid S} \equiv H_{\hat{Y}, A \cup W \mid S}(\cdot)$ as

$$
\hat{H}^2_{\hat{Y}, A \cup W \mid S} = \hat{\omega}^T_{\hat{Y}, A \cup W \mid S} \left( \hat{K}_Y \circ \hat{K}_{A \cup W} \right) \hat{\omega}_{\hat{Y}, A \cup W \mid S} - 2 \left( \hat{\omega}^T_{\hat{Y} \mid S} \hat{K}_Y \hat{\omega}_{\hat{Y}, A \cup W \mid S} \right) \left( \hat{\omega}^T_{A \cup W \mid S} \hat{K}_{A \cup W} \hat{\omega}_{\hat{Y}, A \cup W \mid S} \right) + \left( \hat{\omega}^T_{\hat{Y} \mid S} \hat{K}_Y \hat{\omega}_{Y \mid S} \right) \left( \hat{\omega}^T_{A \cup W \mid S} \hat{K}_{A \cup W} \hat{\omega}_{A \cup W \mid S} \right),
$$

(2)

where $\circ$ is element-wise multiplication. The functions $\hat{\omega}_{Y \mid S} \equiv \hat{\omega}_{Y \mid S}(\cdot)$, $\hat{\omega}_{A \cup W \mid S} \equiv \hat{\omega}_{A \cup W \mid S}(\cdot)$, and $\hat{\omega}_{\hat{Y}, A \cup W \mid S} \equiv \hat{\omega}_{\hat{Y}, A \cup W \mid S}(\cdot)$ are found via kernel ridge regression. Caponnetto and Vito [2007] provide the convergence rates of the estimand $\hat{H}^2_{\hat{Y}, A \cup W \mid S}$ under mild conditions. In practice, computing the HSCIC approximation by the formula in Eq. (2) can be computationally expensive. We provide the runtime for some of our experimental setting showing that including the HSCIC term can slow down training anywhere between 2x and several 100x in Section G.7. To speed it up, we can use random Fourier features to approximate $\hat{K}_Y$ and $\hat{K}_{A \cup W}$ [Avron et al., 2017, Rahimi and Recht, 2007]. The details on how to substantially speed up the computations with such approximations are described in Section F and we describe the improvements in terms of computational complexity in Section G.7. We emphasize that Eq. (2) allows us to consistently estimate the HSCIC from observational i.i.d. samples, without prior knowledge of the counterfactual distributions.

### 3.5 Measuring counterfactual invariance.

Besides predictive performance, e.g., mean squared error (MSE) for regression or accuracy for classification, our key metric of interest is the level of counterfactual invariance achieved by the predictor $\hat{Y}$. First, we emphasize again that counterfactual distributions are generally not identified from the observational distribution (i.e., from available data) meaning that CI is generally untestable in practice from observational data. We can thus only evaluate CI in (semi-)synthetic settings where we have access to the full SCM and thus all counterfactual distributions.

A measure for CI must capture how the distribution of $\hat{Y}_{a'}^s$ changes for different values of $a'$ across all conditioning values $w$ (which may include an observed value $A = a$). We propose

\textsuperscript{8}In particular, HSCIC does not regularize an ill-posed problem, i.e., it does not merely break ties between predictors with equal $L(\hat{Y})$. Hence it also need not decay to zero as the sample size increases.
The variance of CounterFactuals (VCF) as a metric of CI

\[ \text{VCF}(\hat{Y}) := \mathbb{E}_{\mathbf{w} \sim P_{\mathbf{w}}} \left[ \text{var}_{\mathbf{a}' \sim P_{\mathbf{A}}} \left( \mathbb{E}_{\mathbf{Y}_{\mathbf{a}'}|\mathbf{W}=\mathbf{w}}[\hat{Y}] \right) \right]. \] (3)

That is, we quantify how the average outcome varies with the interventional value \( \mathbf{a}' \) at conditioning value \( \mathbf{w} \) and average this variance over \( \mathbf{w} \). For deterministic predictors (point estimators), which we use in all our experiments, the prediction is a fixed value for each input \( \mathbb{E}_{\mathbf{Y}_{\mathbf{a}'}|\mathbf{W}=\mathbf{w}}[\hat{Y}] = \hat{y} \) and we can drop the inner expectation of Eq. (3). In this case, the variance term in Eq. (3) is zero if and only if \( P_{\mathbf{Y}_{\mathbf{a}'}|\mathbf{W}=\mathbf{w}}(\hat{y}) = P_{\mathbf{Y}_{\mathbf{a}'}|\mathbf{W}=\mathbf{w}}(\hat{y}) \) almost surely. Since the variance is non-negative, the outer expectation is zero if and only if the variance term is zero almost surely, yielding the following result.

**Corollary 3.6.** For point-estimators, \( \hat{Y} \) is counterfactually invariant in \( \mathbf{A} \) w.r.t. \( \mathbf{W} \) if and only if \( \text{VCF}(\hat{Y}) = 0 \) almost surely.

Estimating VCF in practice requires access to the DGP to generate counterfactuals. Given \( d \) i.i.d. examples \( (\mathbf{w}_i)^d_{i=1} \) from a fixed observational dataset we sample \( k \) intervention values from the marginal \( P_{\mathbf{A}} \) and compute corresponding predictions. The inner expectation is simply the deterministic predictor output, and we compute the empirical expectation over the \( d \) observed \( \mathbf{w} \) values and empirical variances over the \( k \) sampled interventional values (for each \( \mathbf{w} \)). Since the required counterfactuals are by their very nature unavailable in practice, our analysis of VCF is limited to (semi-)synthetic settings. Notably, the proposed procedure for choosing \( \gamma \) does not require VCF. Our experiments corroborate the weak convergence property of HSCIC—small HSCIC implies small VCF. Hence, HSCIC may serve as a strong proxy for VCF and thus CI in practice.

In practical terms, HSCIC is considered the relevant metric, which is also estimable from data. Theoretically, however, counterfactual invariance is only implied by an exact HSCIC value of zero (Corollary 3.5). Even if HSCIC exhibits continuity, meaning it approaches zero as distributions converge weakly, VCF may still be preferable for directly assessing counterfactual invariance because it provides a more interpretable metric. Our experimental investigations into VCF aim to establish the practical utility of HSCIC as a measure of counterfactual invariance.

### 4 Experiments

The code for the experiments is available at: [https://github.com/ceciliacasolo/CIP](https://github.com/ceciliacasolo/CIP).
Baselines. As many existing methods focus on cruder purely observational or interventional invariances (see Section 2.2), our choice of baselines for true counterfactual invariance is highly limited. First, we compare CIP to Veitch et al. [2021] in their two limited settings (Fig. 6(b-c)) in Section H.1, showing that our method performs on par with theirs. Next, we compare to two methods proposed by Kusner et al. [2017] in settings where they apply. CF1 (their ‘Level 1’) consists of only using non-descendants of $A$ as inputs to $\hat{f}_Y$. CF2 (their ‘Level 2’) assumes an additive noise model and uses the residuals of descendants of $A$ after regression on $A$ together with non-descendants of $A$ as inputs to $\hat{f}_Y$. We refer to these two baselines as CF1 and CF2 respectively. We also compare CIP to the ‘naive baseline’ which consists in training a predictor ignoring $A$. In settings where $A$ is binary, we also compare to Chiappa [2019], devised for path-wise counterfactual fairness. Finally, we develop heuristics based on data augmentation as further possible baselines in Section H.2.

4.1 Synthetic experiments

First, we generate various synthetic datasets following the causal graph in Fig. 1(d). They contain (i) the prediction targets $Y$, (ii) variable(s) we want to be CI in $A$, (iii) covariates $L$ mediating effects from $A$ on $Y$, and (iv) confounding variables $S$. The goal is to learn a predictor $\hat{Y}$ that is CI in $A$ w.r.t. $W := A \cup L \cup S$. The datasets cover different dimensions for the observed variables and their correlations and are described in detail in Section G.

Model choices and parameters. For all synthetic experiments, we train fully connected neural networks (MLPs) with MSE loss $L_{\text{mse}}(\hat{Y})$ as the predictive loss $L$ in Eq. (1) for continuous outcomes $Y$. We generate 10k samples from the observational distribution in each setting and use an 80 to 20 train-test split. All metrics reported are on the test set. We perform hyper-parameter tuning for MLP hyperparameters based on a random strategy (see Section G for details). The $\text{HSCIC}(\hat{Y}, A \cup W | S)$ term is computed as in Eq. (2) using a Gaussian kernel with amplitude $1.0$ and length scale $0.1$. The regularization parameter $\lambda$ for the ridge regression coefficients is set to $\lambda = 0.01$. We set $d = 1000$ and $k = 500$ in the estimation of VCF.

Model performance. We first study the effect of the HSCIC on accuracy and counterfactual invariance on the simulated dataset in Section G.1. Fig. 2 (left) depicts the expected trade-off between MSE and VCF for varying $\gamma$, whereas Fig. 2 (middle) highlights that HSCIC (estimable from observational data) is a strong proxy of counterfactual invariance measured by VCF (see discussion after Eq. (3)). Figure 2 (right) compares CIP to baselines for a simulated non-additive noise model in Section G.2. For a suitable choice of $\gamma$, CIP outperforms the baseline CF2 and the naive baseline in both MSE and VCF simultaneously. While CF1 achieves perfect CI by construction (VCF = 0), its MSE is higher than CIP at almost perfect CI (VCF...
near zero). To conclude, our method flexibly and reliably trades predictive performance for counterfactual invariance via a single parameter $\gamma$ and Pareto-dominates existing methods. In Section G.2 we present extensive results on further simulated settings and compare CIP to other heuristic methods in Section H.2.

**Effect of dimensionality of $A$.** A key advantage of CIP is that it can deal with multi-dimensional $A$. We consider simulated datasets described in Section G.3, where we gradually increase the dimension of $A$. The results in Fig. 3 for different trade-off parameters $\gamma$ and different dimensions of $A$ demonstrate that CIP effectively enforces CI also for multi-dimensional $A$.\footnote{In all boxplots, boxes represent the interquartile range, the horizontal line is the median, and whiskers show minimum and maximum values, excluding outliers (dots).} Further results are shown in Section G.3.

### 4.2 Image experiments

We consider an image classification task on the dSprites dataset [Matthey et al., 2017], with a causal model as depicted in Fig. 1(f). The full structural equations are provided in Section G.4. This experiment is particularly challenging due to the mixed categorical and continuous variables in $C$ (shape, y-pos) and $L$ (color, orientation), with continuous $A$ (x-pos). We seek a predictor $\hat{Y}$ that is CI in the x-position w.r.t. all other observed variables. Following Theorem 3.2, we achieve $\hat{Y} \perp \{\text{x-pos}, \text{scale}, \text{color}, \text{orientation}\} \mid \{\text{shape}, \text{y-pos}\}$ via the HSCIC operator. To accommodate the mixed input types, we first extract features from the images via a CNN and from other inputs via an MLP. We then use an MLP on all concatenated features for $\hat{Y}$. Fig. 4 shows that CIP gradually enforces CI as $\gamma$ increases and illustrates the inevitable increase of MSE.

### 4.3 Fairness with continuous protected attributes

Finally, we apply CIP to the widely-used UCI Adult dataset [Kohavi and Becker, 1996] and compare it against a ‘naive baseline’ which simply ignores $A$, CF1, CF2, and path-specific counterfactual fairness (PSCF) [Chiappa, 2019]. Like CF1, PSCF always achieves $VCF = 0$ by design. We explicitly acknowledge the shortcomings of the UCI Adult dataset to reason about social justice [Ding et al., 2021]. Instead, we chose it due to previous investigations into plausible causal structures based on domain knowledge [Zhang et al., 2017]. The task is to predict whether an individual’s income is above a threshold based on demographic information, including protected attributes. We follow Chiappa [2019], Nabi and Shpitser [2018], where a causal structure is assumed for a subset of the variables as in Fig. 1(e) (see Section G.5 and Fig. 8 for details). We choose gender and age as the protected attributes $A$, collect marital status, level of education, occupation, working hours per week, and income.

![Figure 4: On the dSprites image dataset, CIP trades off MSE for VCF and achieves almost full CI as $\gamma$ increases. Boxes are for 8 random seeds.](image)

![Figure 5: Accuracy and VCF on the Adult dataset. CIP achieves better VCF than CF2 and the naive baseline (NB), improved in accuracy compared to PSCF and is on par with CF1 in accuracy at $VCF \approx 0$.](image)
and work class into $L$, and combine the remaining observed attributes in $C$. Our aim is to learn a predictor $\hat{Y}$ that is CI w.r.t. $W = C \cup L$. Achieving (causal) fairness for (mixed categorical and) continuous protected attributes is under active investigation [Chiappa and Pacchiano, 2021, Mary et al., 2019], but directly supported by CIP.

We use an MLP with binary cross-entropy loss for $\hat{Y}$. Since this experiment is based on real data, the true counterfactual distribution cannot be known. Following Chiappa and Pacchiano [2021] we estimate a possible SCM by inferring the posterior distribution over the unobserved variables using variational autoencoders [Kingma and Welling, 2014]. Even though CF1, PSCF always achieves $VCF = 0$ by design, Figure 5 shows that CIP gradually achieves CI and even manages to keep a constant accuracy after an initial drop. It outperforms PSCF in accuracy and reaches comparable accuracy to CF1 when reaching $VCF \approx 0$. It Pareto-dominates CF2 and can achieve better VCF than the naive baseline, even though NB is more accurate for $\gamma \geq 5$ than CIP.

5 Discussion and future work

We developed CIP, a method to learn counterfactually invariant predictors $\hat{Y}$. First, we presented a sufficient graphical criterion to characterize counterfactual invariance and reduced it to conditional independence in the observational distribution under an injectivity assumption of a causal mechanism. We then built on kernel mean embeddings and the Hilbert-Schmidt Conditional Independence Criterion to devise an efficiently estimable, differentiable, model-agnostic objective to train CI predictors for mixed continuous/categorical, multi-dimensional variables. We demonstrated the efficacy of CIP in extensive empirical evaluations on various regression and classification tasks.

A limitation of this work is the computational cost of estimating HSCIC, which may limit the scalability of CIP to very high-dimensional settings even when using efficient random Fourier features. While the increased robustness of counterfactually invariant predictors are certainly desirable in many contexts, this presupposes the validity of our assumptions. Thus, an important direction for future work is to assess the sensitivity of CIP to misspecifications of the causal graph or insufficient knowledge of the required adjustment set. Lastly, we envision our graphical criterion and KME-based objective to be useful also for causal representation learning to isolate causally relevant, autonomous factors underlying the data.

Another key limitation of our work, shared by all studies in this domain, is the assumption that the causal graph is known. Assessing the validity of a causal DAG in real-world settings is challenging, since one can often conjecture plausible confounding mechanisms for any missing edge. For this reason, in this work we found it challenging to provide convincing examples that corroborate the “potential validity” of our assumptions. However, our empirical results demonstrate that CIP is effective even when these assumptions are violated. Similarly, our real-world experiments show a comparable invariance and accuracy trade-off trend as the simulated experiments. We believe this to be a better indicator of validity and robustness than isolated toy examples where assumptions may be satisfied. Practitioners using our proposed framework are recommended to take these considerations into account before they apply the method to their own data, and combine causal discovery methods with domain knowledge to assess the validity of the underlying DGP.

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Appendix

A Discussion on related work

In this section, we give an overview of the definitions used in the related work Fawkes and Evans [2023].

Definition A.1 (Def. 2.1 by Fawkes et al. [2022]). A variable $Y$ satisfies almost sure counterfactual invariance (a.s.-CI) with respect to $A$ if:

$$Y_a = Y_{a'} \text{ a.s. for all } a, a'$$

Definition A.2 (Def. 2.3 by Fawkes and Evans [2023] and Def. 1.1 Veitch et al. [2021]). A function $f : \mathcal{X} \rightarrow \mathcal{Y}$ is counterfactually invariant ($\mathcal{F}$-CI) if the variable $\hat{Y} := f(X)$ satisfies almost sure counterfactual invariance. That is:

$$f(X_a) = f(X_{a'}) \text{ a.s.}$$

Definition A.3 (Def. 2.2 by Fawkes and Evans [2023]). A variable $Y$ satisfies distributional counterfactual invariance ($\mathcal{D}$-CI) conditional on some set of variables $W$, with respect to $A$, if:

$$P(Y_a = y | W = w, A = a) = P(Y_{a'} = y | W = w, A = a)$$

for all $a, a'$ and for almost all $y, w$.

Definition A.3 is similar to the one provided in the main paper Definition 2.2, however enforces conditioning on the intervening variable $A$. As it can be noticed from the definitions, distributional notions of Counterfactual Invariance (CI), such as Definition A.3 and Definition 2.2, are less stringent than Definition A.1, as indicated in the following result in Fawkes and Evans [2023, Lem. 2.4]:

Lemma A.4 (Lem. 2.4 by Fawkes and Evans [2023]). We have that a.s.-CI implies $\mathcal{D}$-CI conditional on any set of variables, but $\mathcal{D}$-CI implies a.s.-CI only if the conditioning set contains the outcome $Y$.

B Proof of Theorem 3.2

To provide a mostly self-contained proof, we start by stating the necessary definitions and repeat required known results from the literature. Sections B.1 to B.3 contain these preliminary definitions and results, whereas the final statement and proof provided in Section B.4 is our original contribution (making use of the preliminaries stated before).

B.1 Overview of the proof techniques

We restate the main theorem for completeness.

Theorem 3.2. Let $G$ be a causal graph, $A, W$ be two (not necessarily disjoint) sets of nodes in $G$, such that $(A \cup W) \cap Y = \emptyset$, let $S$ be a valid adjustment set for $(A \cup W, Y)$. Further, for any random variable $X \in W \setminus A$ denote with $g_X(pa(X), U_X)$ its structural equation, and suppose that $pa(X) \subseteq A \cup W$. Suppose that $g_X$ is injective in the variable $u$.\(^\text{10}\) Then, in all SCMs compatible with $G$, if a predictor $\hat{Y}$ satisfies $\hat{Y} \perp \perp A \cup W | S$, then $\hat{Y}$ is counterfactually invariant in $A$ with respect to $W$.

\(^\text{10}\)The injectivity of $g_X$ is defined as follows. Consider two pairs $\{p, u\}$ and $\{p, u'\}$ with $p$ in the support of $pa(X)$ and $u, u'$ in the support of $U_X$. Suppose that $P_{pa(X), U_X}(p, u) \neq 0$ and $P_{pa(X), U_X}(p, u') \neq 0$. Then, it holds $g(p, u) = g(p, u')$ if and only if $u = u'$. 

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Our proof technique generalizes the work of Shpitser and Pearl [2009]. To understand the proof technique, note that conditional counterfactual distributions of the form \( P_{Y_a^w | Y} (y | w) \) involve quantities from two different worlds. The variables \( W \) belong to the pre-interventional world, and the interventional variable \( Y_a \) belongs to the world after performing the intervention \( A \leftarrow a \). The variable \( Y_a^w \) refers to the counterfactual of \( Y_a \) after conditioning in the pre-interventional world on \( W \).

### B.2 Identifiability of counterfactual distributions

In this section, we discuss the well-known backdoor criterion for the identifiability of conditional distributions, which we will then use to prove Theorem 3.2. To this end, we use the notions of a blocked path and valid adjustment set, which we restate for clarity.

**Definition B.1.** Consider a path \( \pi \) of causal graph \( G \). A set of nodes \( Z \) blocks \( \pi \), if \( \pi \) contains a triple of consecutive nodes connected in one of the following ways: \( N_i \rightarrow Z \rightarrow N_j \), \( N_i \leftarrow Z \rightarrow N_j \), with \( N_i, N_j \notin Z \), \( Z \in Z \), or \( N_i \rightarrow M \leftarrow N_j \) and neither \( M \) nor any descendent of \( M \) is in \( Z \).

Using this definition, we define the concept of a valid adjustment set.

**Definition 3.1** (valid adjustment set). Let \( G \) be a causal graph and let \( X, Y \) be disjoint (sets of) nodes in \( G \). A set of nodes \( S \) is a valid adjustment set for \( (X, Y) \), if (i) no element in \( S \) is a descendant in \( G \) of any node \( W \notin X \) which lies on a proper causal path from \( X \) to \( Y \), and (ii) \( S \) blocks all non-causal paths from \( X \) to \( Y \) in \( G \).

Definition 3.1 is a useful graphical criterion for the identifiability of counterfactual distributions. In fact, the following theorem holds:

**Theorem B.2** (Theorem 4 by Shpitser et al. [2010]). Let \( G \) be a causal graph, \( A, W \) be two (not necessarily disjoint) sets of nodes in \( G \), such that \( (A \cup W) \cap Y = \emptyset \). Define the set \( X = W \setminus A \). Suppose that a set of nodes \( S \) satisfies the adjustment criterion relative to \( (A \cup W, Y) \) in \( G \). Then, it holds \( Y_a^w, X \perp \! \! \! \perp A, X | S \) for any intervention \( A, X \leftarrow a', x' \).

### B.3 \( d \)-separation and conditional independence

In this section, we discuss a well-known criterion for conditional independence, which we will then use to prove Theorem 3.2. We use the notion of a blocked path, as in Definition B.3 and the concept of \( d \)-separation as follows.

**Definition B.3** (\( d \)-Separation). Consider a causal graph \( G \). Two sets of nodes \( X \) and \( Y \) of \( G \) are said to be \( d \)-separated by a third set \( S \) if every path from any node of \( X \) to any node of \( Y \) is blocked by \( S \).

We use the notation \( X \perp_{G} Y | S \) to indicate that \( X \) and \( Y \) are \( d \)-separated by \( S \) in \( G \). We use Definition B.3 as a graphical criterion for conditional independence [Pearl, 2000].

**Lemma B.4** (Markov Property). Consider a causal graph \( G \), and suppose that two sets of nodes \( X \) and \( Y \) of \( G \) are \( d \)-separated by \( S \). Then, \( X \) is independent of \( Y \) given \( S \) in any model induced by the graph \( G \).

The Markov Property is also referred to as \( d \)-separation property. We use the notation \( X \perp_{G} Y | S \) to indicate that \( X \) and \( Y \) are \( d \)-separated by \( S \) in \( G \).

### B.4 Proof of Theorem 3.2

Before discussing the proof of Theorem 3.2, we prove an additional auxiliary result. All the following theoretical results are novel and based on the previous sections.
Lemma B.5. Let $G$ be a causal graph, $A$, $X$ be two disjoint sets of nodes in $G$. Suppose that any r.v. $X \subseteq X$ is defined by a structural equation of the form $X = g(p(a), U_X)$, with $pa(X) \subseteq X \cup A$ and $U_X$ exogenous independent noise. Furthermore, suppose that $g$ is injective in $U_X$. Then, for any observational values $a, x$ with $\mathbb{P}_{X,A}(x, a) \neq 0$, there exist a value $u$ in the support of $U_X$ such that $\mathbb{P}_{U_X|A=a,X=x}(u) = \mathbb{I}_u$. Here, $\mathbb{I}_u$ is the Dirac $\delta$-distribution.

Proof. In this proof, we use the following additional notation. Consider a set of nodes $T \subseteq A \cup X$, and let $a$ be a point in the support of $A$. We denote with $ap$ the restriction of $a$ to $A \cap T$. Similarly, for a point $x$ in the support of $X$, we denote with $xp$ the restriction of $x$ to $X \cap T$. Furthermore, given two points $a, x$ as above, we define $a \cup x$ as the only point in the support of $A \cup X$ such that $(a \cup x) \mid_A = a$ and $(a \cup x) \mid_X = x$ respectively.

Furthermore, we denote with $X = g_X(pa(X), U_X)$ the structural equation for the joint random variable $X$. In this equation, $U_X$ corresponds to the joint exogenous noise. Note that the function $g_X(pa(X), U_X)$ is injective in $U_X$, by the injectivity assumption on the structural equation $g_X(pa(X), U_X)$ for each $X \subseteq X$. Define the function $h(p)$ over the support of $pa(X)$ as

$$h(a, x) := \begin{cases} u \text{ such that } g_X((a \cup x) \mid_{pa(X)}, u) = x & \text{if } \mathbb{P}_{pa(X), U_X}((a \cup x) \mid_{pa(X)}, u) \neq 0 \\ 0 & \text{otherwise} \end{cases}$$

Note that by the injectivity of $g_X$, the function $h$ as above is well-defined. Since $pa(X) \subseteq X \cup A$, then it holds $\mathbb{P}_{U_X|A=a,X=x}(u) = \mathbb{I}_{h(a, x)}$. The claim follows, by defining $u := h(a, x)$. \hfill \Box

We use Lemma B.5 to prove the following result.

Lemma B.6. Under the assumptions of Lemma B.5, fix a set $T \subseteq A \cup X$. Then, for any intervention $A \leftarrow a'$ and observational values $a, x$ with $\mathbb{P}_{X,A}(x, a) \neq 0$, there exist a point $t$ in the support of $T$ such that $\mathbb{P}_{T_{a'}\mid A=a,X=x}(t) = \mathbb{I}_t$.

Proof. In this proof, we use the notation introduced in Lemma B.5. Consider a set of nodes $T \subseteq A \cup X$, and let $a$ be a point in the support of $A$. We denote with $ar$ the restriction of $a$ to $A \cap T$. We prove the claim with a induction argument. To this end, we define $depth(T)$ as the length of the longest directed path $\pi$ in $G$ from any node in $A \cup U_T$ to any node in $T$ (see Figure 6).\footnote{Note that since $pa(X) \in X \cup A$ for all $X \in T$, then any such path $\pi$ only consists of nodes that are in $X \cup A \cup U_T$.}

(Base case). We first assume that with $depth(T) = 0$. We further assume w.l.o.g. that $T \cap A = \emptyset$, since the variables $T \cap A$ are fixed by the intervention $A \leftarrow a'$. Denote with $T = g_T(pa(T), U_T)$ the structural equation for the joint random variable $T$. Since $depth(T) = 0$, it holds $pa(T) \subseteq A$.\footnote{Here, $pa(T)$ can also be the empty set. In this case, $T = U_T$ is an exogenous random variable and the proof follows.} Then, the random variable $T_{a'}$ is defined by the formula $T_{a'} = g_T(a' \mid T, U_T)$. By Lemma B.5, there exist a value $u'$ in the support of $U_T$ such that $\mathbb{P}_{U_T|A=a,X=x}(u') = \mathbb{I}_{u'}$. It follows that $\mathbb{P}_{T_{a'}\mid A=a,X=x}(t) = \mathbb{I}_t$ with $t := g_T(a' \mid T, u')$.
(Inductive step). We assume by induction that the claim holds for all $T$ such that $\text{depth}(T) \leq k$, and we prove the claim for a set $T$ of depth $\text{depth}(T) = k + 1$. Since $\text{depth}(\text{pa}(T)) < \text{depth}(T) = k + 1$, by the inductive hypothesis, it holds

$$\mathbb{P}_{\text{pa}(T), t'} | A = a, X = x | (X) = 1,$$  \hfill (4)

for a point $t'$ in the support of $\text{pa}(T)$. Again, consider the structural equation $T = g_T(\text{pa}(T), U_T)$ for the joint random variable $T$. In this case, the random variable $T_a'$ is defined by the formula $T_a' = g_T(\text{pa}(T), U_T)$. By Lemma B.5, there exist a value $u'$ in the support of $U_T$ such that $\mathbb{P}_{U_T = a, X = x}(u) = 1$.

Combining this observation with Eq. (4), we get $\mathbb{P}_{T_a', t} | A = a, X = x(t) = 1_{t'}$. The claim follows by defining $t := g_T(t', u')$. 

We use the results above to prove a third auxiliary lemma.

**Lemma B.7.** Under the assumptions of Lemma B.5 and Lemma B.6, for any intervention $A \leftarrow a'$ and observational values $a, x$, there exist an intervention $X \leftarrow x'$ such that

$$\mathbb{P}_{Y_{a', x'}} | A = a, X = x(y) = \mathbb{P}_{Y_a} | A = a, X = x(y).$$

**Proof.** First note that by lemma B.6 there exists a point $x'$ such that $\mathbb{P}_{X_{a'} | A = a, X = x}(y) = 1$. We choose this point to define the intervention $X \leftarrow x'$ as in the statement of this lemma. Denote with $G_{a'}$ the causal graph after performing the intervention $A \leftarrow a'$. Furthermore, denote with $G_{a', x'}$ the causal graph after performing the interventions $A \leftarrow a'$ and $X \leftarrow x'$. We first prove that for any node $V$ of $G_{a'}$ and $G_{a', x'}$ it holds

$$\mathbb{P}_{V | A = a, X = x}(v) = \mathbb{P}_{V | A = a, X = x, X_{a'} = x'}(v).$$  \hfill (5)

In fact, it holds

$$\mathbb{P}_{V | A = a, X = x}(v) = \int \mathbb{P}_{V | A = a, X = x, X_{a'} = x'}(v) dP_{X_{a'} | A = a, X = x}(v')$$

(by marginalization)

$$= \int \mathbb{P}_{V | A = a, X = x, X_{a'} = x'}(v) dP_{x'}$$

(by lemma B.6 with $T = X$)

$$= \mathbb{P}_{V | A = a, X = x, X_{a'} = x'}(v).$$

(by marginalization)

Hence, Eq. (5) holds. It follows that

$$\mathbb{P}_{Y_{a', x'} | A = a, X = x}(y) = \mathbb{P}_{Y_{a'} | A = a, X = x, X_{a'} = x'}(y)$$

(Eq. (5) with $V = Y_{a', x'}^*$)

$$= \mathbb{P}_{Y_{a'} | A = a, X = x}(y)$$

(Axiom of Consistency [Pearl, 2000])

$$= \mathbb{P}_{Y_{a'} | A = a, X = x}(y).$$

(Eq. (5) with $V = Y_{a'}$)

Hence, the claim holds.

We now prove our main result.

**Proof of Theorem 3.2.** Following the notation of Theorem B.2, define the set $X = W \setminus A$. Note that, using this notation, the assumption $Y \perp \!
\perp A, W \mid S$ can be written as $Y \perp \!
\perp A, X \mid S$. Suppose that it holds

$$\mathbb{P}_{Y_{a', x'} | A = a, X = x}(y) = \int \mathbb{P}_{Y | A = a', X = x', S = s}(y) dP_S | A = a, X = x(s)$$  \hfill (6)

for any intervention $A, X \leftarrow a', x'$, and for any possible value $w$ attained by $W$. Assuming that Eq. (6) holds, we have that

$$\mathbb{P}_{Y_{a', x'} | A = a, X = x}(y) = \int \mathbb{P}_{Y | A = a', X = x', S = s}(y) dP_S | A = a, X = x(s)$$

(assuming Eq. (6))

$$= \int \mathbb{P}_{Y | A = a, X = x', S = s}(y) dP_S | A = a, X = x(s)$$

(Y $\perp \!
\perp A, X \mid S$

(assuming Eq. (6))  \hfill (7)
We first prove that if the HSCIC can be used to promote conditional independence, using a similar technique as Park and Muandet [2020]. The following theorem holds.

**Proof.** Theorem 3.4

Note that the claim follows from Eq. (8). The proof of Theorem 3.2 thus boils down to proving Eq. (6).

$\mathbb{P}_{Y^*_w \mid W=w}(y) = \int \mathbb{P}_{Y^*_w(A=\omega),X=x}(y)d\mathbb{P}_T|W=w(t)$  \hspace{1cm} (by conditioning)  

= $\int \mathbb{P}_{Y^*_w(A=\omega),X=x}(y)d\mathbb{P}_T|W=w(t)$  \hspace{1cm} (by Lemma B.7)  

= $\int \mathbb{P}_{Y^*_w(A=\omega),X=x}(y)d\mathbb{P}_T|W=w(t)$  \hspace{1cm} (by Eq. (7))  

= $\mathbb{P}_{Y^*_w(A=\omega),X=x}(y)$  \hspace{1cm} (by Lemma B.7)  

Note that the claim follows from Eq. (8). The proof of Theorem 3.2 thus boils down to proving Eq. (6).

$\mathbb{P}_{Y^*_w(A=\omega),X=x}(y)$  

= $\int \mathbb{P}_{Y^*_w(A=\omega),X=x,S}(y)d\mathbb{P}_S|A=a,X=x(s)$  \hspace{1cm} (by conditioning)  

= $\int \mathbb{P}_{Y^*_w(A=\omega),X=x,S}(y)d\mathbb{P}_S|A=a,X=x(s)$  \hspace{1cm} (by Theorem B.2)  

= $\int \mathbb{P}_{Y^*(A=\omega),X=x',S}(y)d\mathbb{P}_S|A=a,X=x(s)$  \hspace{1cm} (Axiom of Consistency [Pearl, 2000])

and Eq. (6) follows.  

C Proof of Theorem 3.4

We prove that the HSCIC can be used to promote conditional independence, using a similar technique as Park and Muandet [2020]. The following theorem holds.

**Theorem 3.4 (Theorem 5.4 by Park and Muandet [2020]).** If the kernel $k$ of $\mathcal{H}_X \otimes \mathcal{H}_{A \cup W}$ is characteristic\(^{13}\), $\text{HSCIC}(Y, A \cup W \mid S) = 0$ almost surely if and only if $Y \perp A \cup W \mid S$.

**Proof.** By definition, we can write $\text{HSCIC}(Y, A \cup W \mid S) = H_{Y, A \cup W \mid S}$, where $H_{Y, A \cup W \mid S}$ is a real-valued deterministic function. Hence, the HSCIC is a real-valued random variable, defined over the same domain $\Omega_S$ of the random variable $X$.

We first prove that if $\text{HSCIC}(Y, A \cup W \mid S) = 0$ almost surely, then it holds $Y \perp A \cup W \mid S$. To this end, consider an event $\Omega' \subseteq \Omega_X$ that occurs almost surely, and such that it holds $(H_{Y, A \cup W \mid X}(\omega)) = 0$ for all $\omega \in \Omega'$. Fix a sample $\omega \in \Omega'$, and consider the corresponding value $s_\omega = S(\omega)$, in the support of $S$. It holds

$\int k(y \otimes [a, w], \cdot) d\mathbb{P}_{Y,A \cup W \mid S=s_\omega} = \mu_{Y, A \cup W \mid S=s_\omega}$  \hspace{1cm} (by definition)  

$= \mu_{Y \mid S=s_\omega} \otimes \mu_{A \cup W \mid S=s_\omega}$  \hspace{1cm} (since $\omega \in \Omega'$)  

$= \int k_Y(y, \cdot) d\mathbb{P}_{Y \mid S=s_\omega} \otimes \int k_{A \cup W}([a, w], \cdot) d\mathbb{P}_{A \cup W \mid S=s_\omega}$  \hspace{1cm} (by definition)  

$= \int k_Y(y, \cdot) \otimes k_{A \cup W}([a, w], \cdot) d\mathbb{P}_{Y \mid S=s_\omega} \mathbb{P}_{A \cup W \cup W \mid S=s_\omega}$,  \hspace{1cm} (by Fubini’s Theorem)

\(^{13}\)The tensor product kernel $k$ is characteristic if $\mathbb{P}_{Y, A \cup W} \rightarrow \mathbb{E}_{Y, [a, w]}[k(\cdot, y \otimes [a, w])]$ is injective.
with \( k_Y \) and \( k_{A \cup W} \) the kernels of \( H_Y \) and \( H_{A \cup W} \) respectively. Since the kernel \( k \) of the tensor product space \( H_Y \otimes H_{A \cup W} \) is characteristic, then the kernels \( k_Y \) and \( k_{A \cup W} \) are also characteristic. Hence, it holds \( \mathbb{P}_{Y, A \mid S = s_w} = \mathbb{P}_{Y \mid S = s_w} \mathbb{P}_{A \mid S = s_w} \) for all \( \omega \in \Omega' \). Since the event \( \Omega' \) occurs almost surely, then \( \mathbb{P}_{Y, A \mid S = s_w} = \mathbb{P}_{Y \mid S = s_w} \mathbb{P}_{A \mid S = s_w} \) almost surely, that is \( Y \perp A \cup W \mid S \).

Assume now that \( Y \perp A \cup W \mid S \). By definition there exists an event \( \Omega'' \subseteq \Omega_S \) such that \( \mathbb{P}_{Y, A \cup W \mid S = s_w} = \mathbb{P}_{Y \mid S = s_w} \mathbb{P}_{A \cup W \mid S = s_w} \) for all samples \( \omega \in \Omega'' \), with \( s_w = S(\omega) \). It holds

\[
\begin{align*}
\mu_{Y, A \cup W \mid S = s_w} & = \int k(y \otimes [a, w], \cdot) d\mathbb{P}_{Y, A \cup W \mid S = s_w} \\
& = \int k(y \otimes [a, w], \cdot) d\mathbb{P}_{Y \mid S = s_w} \mathbb{P}_{A \cup W \mid S = s_w} \\
& = \int k_Y(y, \cdot) k_{A \cup W}([a, w], \cdot) d\mathbb{P}_{Y \mid S = s_w} \mathbb{P}_{A \cup W \mid S = s_w} \\
& = \int k_Y(y, \cdot) d\mathbb{P}_{Y \mid S = s_w} \otimes \int k_{A \cup W}([a, w], \cdot) d\mathbb{P}_{A \cup W \mid S = s_w} \\
& = \mu_{Y \mid S = s_w} \otimes \mu_{A \cup W \mid S = s_w}.
\end{align*}
\]

The claim follows. \( \square \)

## D Conditional kernel mean embeddings and the HSCIC

The notion of conditional kernel mean embeddings has already been studied in the literature. We show that, under stronger assumptions, our definition is equivalent to the definition by Park and Muandet [2020]. In this section, without loss of generality we will assume that \( W = \emptyset \) and we will refer to the conditioning set as \( Z \).

### D.1 Conditional kernel mean embeddings and conditional independence

We show that, under stronger assumptions, the HSCIC can be defined using the Bochner conditional expected value. The Bochner conditional expected value is defined as follows.

**Definition D.1.** Fix two random variables \( Y, Z \) taking value in a Banach space \( \mathcal{H} \), and denote with \( (\Omega, \mathcal{F}, \mathbb{P}) \) their joint probability space. Then, the Bochner conditional expectation of \( Y \) given \( Z \) is any \( \mathcal{H} \)-valued random variable \( X \) such that

\[
\int_E Y d\mathbb{P} = \int_E X d\mathbb{P}
\]

for all \( E \in \sigma(Z) \subseteq \mathcal{F} \), with \( \sigma(Z) \) the \( \sigma \)-algebra generated by \( Z \). We denote with \( \mathbb{E} [Y \mid Z] \) the Bochner expected value. Any random variable \( X \) as above is a version of \( \mathbb{E} [Y \mid Z] \).

The existence and almost sure uniqueness of the conditional expectation are shown in Din-culeanu [2000]. Given a RKHS \( \mathcal{H} \) with kernel \( k \) over the support of \( Y \), Park and Muandet [2020] define the corresponding conditional kernel mean embedding as

\[
\mu_{Y \mid Z} := \mathbb{E} [k(\cdot, y) \mid Z].
\]

Note that, according to this definition, \( \mu_{Y \mid Z} \) is an \( \mathcal{H} \)-valued random variable, not a single point of \( \mathcal{H} \). Park and Muandet [2020] use this notion to define the HSCIC as follows.

**Definition D.2** (The HSCIC according to Park and Muandet [2020]). Consider (sets of) random variables \( Y, A, Z \), and consider two RKHS \( \mathcal{H}_Y, \mathcal{H}_A \) over the support of \( Y \) and \( A \)
respectively. The HSCIC between $Y$ and $A$ given $Z$ is defined as the real-valued random variable

$$\omega \mapsto \|\mu_{Y,A|Z}(\omega) - \mu_{Y|Z}(\omega) \otimes \mu_{A|Z}(\omega)\|,$$

for all samples $\omega$ in the domain $\Omega_Z$ of $Z$. Here, $\|\cdot\|$ the metric induced by the inner product of the tensor product space $H_Y \otimes H_Z$.

We show that, under more restrictive assumptions, Definition D.2 can be used to promote conditional independence. To this end, we use the notion of a regular version.

**Definition D.3** (Regular Version, following Definition 2.4 by Çinlar [2011]). Consider two random variables $Y$, $Z$, and consider the induced measurable spaces $(\Omega_Y, \mathcal{F}_Y)$ and $(\Omega_Z, \mathcal{F}_Z)$. A regular version $Q$ for $P_{Y,Z}$ is a mapping $Q: \Omega_Z \times \mathcal{F}_Y \rightarrow [0, +\infty]: (\omega, y) \mapsto Q_\omega(y)$ such that: (i) the map $\omega \mapsto Q_\omega(x)$ is $\mathcal{F}_A$-measurable for all $y$; (ii) the map $y \mapsto Q_\omega(y)$ is a measure on $(\Omega_Y, \mathcal{F}_Y)$ for all $\omega$; (iii) the function $Q_\omega(y)$ is a version for $\mathbb{E}[\mathbb{1}_{Y=y}|Z]$.

The following theorem shows that the random variable as in Definition D.2 can be used to promote conditional independence.

**Theorem D.4** (Theorem 5.4 by Park and Muandet [2020]). With the notation introduced above, suppose that the kernel $k$ of the tensor product space $H_X \otimes H_A$ is characteristic. Furthermore, suppose that $P_{Y,A|X}$ admits a regular version. Then, $\|\mu_{Y,A|Z}(\omega) - \mu_{Y|Z}(\omega) \otimes \mu_{A|Z}(\omega)\| = 0$ almost surely if and only if $Y \perp \!\!\!\perp A | Z$.

Note that the assumption of the existence of a regular version is essential in Theorem D.4. In this work, HSCIC is not used for conditional independence testing but as a conditional independence measure.

### D.2 Equivalence with our approach

The following theorem shows that under the existence of a regular version, conditional kernel mean embeddings can be defined using the Bochner conditional expected value. To this end, we use the following theorem.

**Theorem D.5** (Following Proposition 2.5 by Çinlar [2011]). Following the notation introduced in Definition D.3, suppose that $P_{Y|Z}(\cdot | Z)$ admits a regular version $Q_\omega(y)$. Consider a kernel $k$ over the support of $Y$. Then, the mapping

$$\omega \mapsto \int k(\cdot, y) dQ_\omega(y)$$

is a version of $\mathbb{E}[k(\cdot, y) | Z]$.

As a consequence of Theorem D.5, we prove the following result.

**Lemma D.6.** Fix two random variables $Y$, $Z$. Suppose that $P_{Y|Z}$ admits a regular version. Denote with $\Omega_Z$ the domain of $Z$. Then, there exists a subset $\Omega \subseteq \Omega_Z$ that occurs almost surely, such that $\mu_{Y|Z}(\omega) = \mu_{Y|Z=Z(\omega)}$ for all $\omega \in \Omega$. Here, $\mu_{Y|Z=Z(\omega)}$ is the embedding of conditional measures as in Section 2.

**Proof.** Let $Q_\omega(y)$ be a regular version of $P_{Y|Z}$. Without loss of generality we may assume that it holds $P_{Y|Z}(y | \{Z = Z(\omega)\}) = Q_\omega(y)$. By Theorem D.5 there exists an event $\Omega \subseteq \Omega_Z$ that occurs almost surely such that

$$\mu_{Y|Z}(\omega) = \mathbb{E}[k(\cdot, \cdot) | Z](\omega) = \int k(y, \cdot) dQ_\omega(y),$$

(10)
for all $\omega \in \Omega$. Then, for all $\omega \in \Omega$ it holds
\[
\mu_{Y|Z}(\omega) = \int k(x, \cdot) dQ_{\omega}(x) = \int k(x, \cdot) dP_{X|A}(x \mid \{A = A(\omega)\}) = \mu_{X|A = A(\omega)},
\]
(it follows from Eq. (10))
\[
\|\mu_{X,A|Z}(\omega) - \mu_{X|Z}(\omega) \otimes \mu_{A|Z}(\omega)\| = (H_{Y,A|Z} \circ Z)(\omega).
\]
(by definition as in Section 2)
as claimed. \hfill \Box

As a consequence of Lemma D.6, we can prove that the definition of the HSCIC by Park and Muandet [2020] is equivalent to ours. The following corollary holds.

**Corollary D.7.** Consider (sets of) random variables $Y$, $A$, $Z$, and consider two RKHS $H_Y$, $H_A$ over the support of $Y$ and $A$ respectively. Suppose that $P_{Y,A|Z}(\cdot \mid Z)$ admits a regular version. Then, there exists a set $\Omega \subseteq \Omega_A$ that occurs almost surely, such that
\[
\|\mu_{X,A|Z}(\omega) - \mu_{X|Z}(\omega) \otimes \mu_{A|Z}(\omega)\| = (H_{Y,A|Z} \circ Z)(\omega).
\]
Here, $H_{Y,A|Z}$ is a real-valued deterministic function, defined as
\[
H_{Y,A|Z}(z) := \|\mu_{Y,A|Z} = z - \mu_{Y|Z} = z \otimes \mu_{A|Z} = z\|,
\]
and $\|\cdot\|$ is the metric induced by the inner product of the tensor product space $H_X \otimes H_A$.

We remark that the assumption of the existence of a regular version is essential in Corollary D.7.

**E The cross-covariance operator**

In this section, we show that under additional assumptions, our definition of conditional KMEs is equivalent to the definition based on the cross-covariance operator, under more restrictive assumptions. The definition of KMEs based on the cross-covariance operator requires the use of the following well-known result.

**Lemma E.1.** Fix two RKHS $H_X$ and $H_Z$, and let $\{\varphi_i\}_{i=1}^\infty$ and $\{\psi_j\}_{j=1}^\infty$ be orthonormal bases of $H_X$ and $H_Z$ respectively. Denote with $\text{HS}(H_X, H_Z)$ the set of Hilbert-Schmidt operators between $H_X$ and $H_Z$. There is an isometric isomorphism between the tensor product space $H_X \otimes H_Z$ and $\text{HS}(H_X, H_Z)$, given by the map
\[
T: \sum_{i=1}^\infty \sum_{j=1}^\infty c_{i,j} \varphi_i \otimes \psi_j \mapsto \sum_{i=1}^\infty \sum_{j=1}^\infty c_{i,j} \langle \cdot, \varphi_i \rangle_{H_X} \psi_j.
\]

For proof of this result see i.e., Park and Muandet [2020]. This lemma allows us to define the cross-covariance operator between two random variables, using the operator $T$.

**Definition E.2** (Cross-Covariance Oprarator). Consider two random variables $X$, $Z$. Consider corresponding mean embeddings $\mu_{X,Z}$, $\mu_X$ and $\mu_Z$, as defined in Section 3. The cross-covariance operator is defined as $\Sigma_{X,Z} := T(\mu_{X,Z} - \mu_X \otimes \mu_Z)$. Here, $T$ is the isometric isomorphism as in Lemma E.1.

It is well-known that the cross-covariance operator can be decomposed into the covariance of the marginals and the correlation. That is, there exists a unique bounded operator $\Lambda_{Y,Z}$ such that
\[
\Sigma_{Y,Z} = \Sigma_{Y,Y}^{1/2} \circ \Lambda_{Y,Z} \circ \Sigma_{Z,Z}^{1/2}
\]
Using this notation, we define the normalized conditional cross-covariance operator. Given three random variables $Y$, $A$, $Z$ and corresponding kernel mean embeddings, this operator is defined as

$$\Lambda_{Y,A|Z} := \Lambda_{Y,A} - \Lambda_{Y,Z} \circ \Lambda_{Z,A}. \tag{11}$$

This operator was introduced by Fukumizu et al. [2007]. The normalized conditional cross-covariance can be used to promote statistical independence, as shown in the following theorem.

**Theorem E.3** (Theorem 3 by Fukumizu et al. [2007]). Following the notation introduced above, define the random variable $\tilde{A} := (A, Z)$. Let $P_Z$ be the distribution of the random variable $Z$, and denote with $L^2(P_Z)$ the space of the square integrable functions with probability $P_Z$. Suppose that the tensor product kernel $k_Y \otimes k_A \otimes k_Z$ is characteristic. Furthermore, suppose that $H_Z + \mathbb{R}$ is dense in $L^2(P_Z)$. Then, it holds

$$\Lambda_{Y,\tilde{A}|Z} = 0 \text{ if and only if } Y \perp \perp A \mid Z.$$

Here, $\Lambda_{Y,\tilde{A}|Z}$ is an operator defined as in Eq. (11).

By Theorem E.3, the operator $\Lambda_{Y,\tilde{A}|Z}$ can also be used to promote conditional independence. However, CIP is more straightforward since it requires less assumptions. In fact, Theorem E.3 requires to embed the variable $Z$ in an RKHS. In contrast, CIP only requires the embedding of the variables $Y$ and $A$.

## F Random fourier features

Random fourier features is an approach to scaling up kernel methods for shift-invariant kernels [Rahimi and Recht, 2007]. Recall that a shift-invariant kernel is a kernel of the form $k(z, z') = h_k(z - z')$, with $h_k$ a positive definite function.

Fourier features are defined via the following well-known theorem.

**Theorem F.1** (Bochner’s Theorem). For every shift-invariant kernel of the form $k(z, z') = h_k(z - z')$ with $h_k(0) = 1$, there exists a probability density function $P_k(\eta)$ such that

$$k(z, z') = \int e^{-2\pi i \eta^T (z - z')} dP_k.$$

Since both the kernel $k$ and the probability distribution $P_k$ are real-valued functions, the integrand in Theorem F.1 can be replaced by the function $\cos \eta^T (z - z')$, and we obtain the following formula

$$k(z, z') = \int \cos \eta^T (z - z') dP_k = \mathbb{E} \left[ \cos \eta^T (z - z') \right], \tag{12}$$

where the expected value is taken with respect to the distribution $P_k(\eta)$. This equation allows to approximate the kernel $k(z, z')$, via the empirical mean of points $\eta_1, \ldots, \eta_l$ sampled independently according to $P_k$. In fact, it is possible to prove exponentially fast convergence of an empirical estimate for $\mathbb{E} \left[ \cos \eta^T (z - z') \right]$, as shown in the following theorem.

**Theorem F.2** (Uniform Convergence of Fourier Features, Claim 1 by Rahimi and Recht [2007]). Following the notation introduced above, fix any compact subset $\Omega$ in the domain of $k$, and consider points $\eta_1, \ldots, \eta_l$ sampled independent according to the distribution $P_k$. Define the function

$$\hat{k}(z, z') := \frac{1}{l} \sum_{j=1}^{l} \cos \eta_j^T (z - z'),$$

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for all \((z, z') \in \Omega\). Then, it holds

\[
P \left( \sup_{z, z'} \left| \hat{k}(z, z') - k(z, z') \right| \geq \varepsilon \right) \leq 2^8 \sigma_k \frac{\text{diam}(\Omega)}{\varepsilon} \exp \left\{ -\frac{\varepsilon^2 l}{4(d+1)} \right\}.
\]

Here \(\sigma_k^2\) is the second moment of the Fourier transform of the kernel \(k\), and \(d\) is the dimension of the arrays \(z\) and \(z'\).

By Theorem F.2, the estimated kernel \(\hat{k}\) is a good approximation of the true kernel \(k\) on the set \(\Omega\).

Similarly, we can approximate the Kernel matrix using Random Fourier features. Following the notation introduced above, define the function

\[
\zeta_{k,l}(z) := \frac{1}{\sqrt{l}} \left[ \cos \eta_{1}^T z, \ldots, \cos \eta_{l}^T z \right]
\]

with \(\eta_{1}, \ldots, \eta_l\) sampled independent according to the distribution \(\mathbb{P}_k\).

We can approximate the Kernel matrix using the functions defined as in Eq. (13). Consider \(n\) samples \(z_1, \ldots, z_n\), and denote with \(Z\) the \(n \times l\) matrix whose \(i\)-th row is given by \(\zeta_{k,l}(z_i)\). Similarly, denote with \(Z^*\) the \(l \times n\) matrix whose \(i\)-th column is given by \(\zeta_{k,l}^*(z_i)\). Then, we can approximate the kernel matrix as \(\hat{K}_Z \approx ZZ^*\).

We can also use this approximation to compute the kernel ridge regression parameters as in Section 3 using the formula \(\hat{W}_Y(z) \approx (ZZ^* - n\lambda I)^{-1} \left[ k_Z(z, z_1), \ldots, k_Z(z, z_n) \right]^T\). Avron et al. [2017] argue that the approximate kernel ridge regression, as defined above, is an accurate estimate of the true distribution. Their argument is based on proving that the matrix \(ZZ^* - n\lambda I\) is a good approximation of \(\hat{K}_Z - n\varepsilon I\). The notion of good approximation is clarified by the following definition.

**Definition F.3.** Fix two Hermitian matrices \(A\) and \(B\) of the same size. We say that a matrix \(A\) is a \(\gamma\)-spectral approximation of another matrix \(B\), if it holds \((1 - \gamma)B \preceq A \preceq (1 + \gamma)B\). Here, the \(\preceq\) symbol means that \(A - (1 - \gamma)B\) is positive semi-definite, and that \((1 + \gamma)B - A\) is positive semi-definite.

Avron et al. [2017] prove that \(ZZ^* - n\lambda I\) is a \(\gamma\)-approximation of \(\hat{K}_Z - n\varepsilon I\), if the number of samples \(\eta_1, \ldots, \eta_l\) is sufficiently large.

**Theorem F.4** (Theorem 7 by Avron et al. [2017]). Fix a constant \(\gamma \leq 1/2\). Consider \(n\) samples \(z_1, \ldots, z_n\), and denote with \(\hat{K}_Z\) the corresponding kernel matrix. Suppose that it holds \(\|\hat{K}_Z\|_2 \geq n\lambda\) for a constant \(\lambda > 0\). Fix \(\eta_1, \ldots, \eta_l\) samples with

\[
l \geq \frac{8}{3\gamma^2 \lambda} \ln \frac{16 \text{ tr}_\lambda(\hat{K}_Z)}{\gamma}
\]

Then, the matrix \(ZZ^* - n\lambda I\) is a \(\gamma\)-approximation of \(\hat{K}_Z - n\varepsilon I\) with probability at least \(1 - \gamma\), for all \(\gamma \in (0, 1)\). Here, \(\text{tr}_\lambda(\hat{K}_Z)\) is defined as the trace of the matrix \(\hat{K}_Z (\hat{K}_Z + n\lambda I)^{-1}\).

We conclude this section by illustrating the use of random Fourier features to approximate a simple Gaussian kernel. Suppose that we are given a kernel of the form

\[
k(z, z') := \exp \left\{ -\frac{1}{2} \sigma^2 \|z - z'\|_2^2 \right\}.
\]

Then, \(k(z, z')\) can be estimated as in Theorem F.2, with \(\eta_1, \ldots, \eta_l \sim \mathcal{N}(0, \Sigma)\), with \(\Sigma := \sigma^{-1}I\), with \(I\) the identity matrix. The functions \(\zeta_{k,l}(z)\) can be defined accordingly.
G Additional experiments and settings

This section contains detailed information on the experiments and additional results.

G.1 Dataset for model performance with the use of the HSCIC

The data-generating mechanism corresponding to the results in Fig. 2 is the following:

\[ Z \sim \mathcal{N}(0, 1) \quad A = Z^2 + \varepsilon_A \]

\[ L = \exp\left\{-\frac{1}{2}A^2\right\} \sin(2A) + 2Z\frac{1}{5}\varepsilon_L \]

\[ Y = \frac{1}{2} \exp\{-LZ\} \cdot \sin(2LZ) + 5A + \frac{1}{5}\varepsilon_Y, \]

where \( \varepsilon_A \sim \mathcal{N}(0, 1) \) and \( \varepsilon_Y, \varepsilon_L \text{i.i.d.} \sim \mathcal{N}(0, 0.1). \)

In the first experiment, Fig. 2 shows the results of feed-forward neural networks consisting of 8 hidden layers with 20 nodes each, connected with a rectified linear activation function (ReLU) and a linear final layer. Mini-batch size of 256 and the Adam optimizer with a learning rate of \( 10^{-3} \) for 1000 epochs were used.

G.2 Datasets and results for comparison with baselines

The comparison of our method CIP with the CF1 and CF2 is done on different simulated datasets. These will be referred to as Scenario 1 and Scenario 2. The data generating mechanism corresponding to the results in Fig. 2 (right) is the following:

\[ Z \sim \mathcal{N}(0, 1) \quad A = \exp\left\{\frac{1}{2}Z^2\right\} \cdot \sin(2Z) + \varepsilon_A \]

\[ L = (A + 0.1Z) \cdot \varepsilon_L \]

\[ Y = A + L + 0.1 \cdot \sin(Z) \]

where \( \varepsilon_A, \varepsilon_L \text{i.i.d.} \sim \mathcal{N}(0, 1) \) and \( \varepsilon_Y \text{i.i.d.} \sim \mathcal{N}(0, 0.1). \) This is referred to as Scenario 1. The data generating mechanism for Scenario 2 is the following:

\[ Z \sim \mathcal{N}(0, 1) \quad A = \exp\left\{-\frac{1}{2}A^2\right\} \cdot \varepsilon_L + 2Z \]

\[ L = \exp\left\{-\frac{1}{2}A^2\right\} \cdot \varepsilon_L + 2Z \]

\[ Y = \frac{1}{2} \sin(ZL) \cdot \exp\{-ZL\} + \frac{1}{5}\varepsilon_Y, \]

where \( \varepsilon_A, \varepsilon_L \text{i.i.d.} \sim \mathcal{N}(0, 1) \) and \( \varepsilon_Y \text{i.i.d.} \sim \mathcal{N}(0, 0.1). \) Fig. 2 (right) and Table 1 present the average and standard deviation resulting from 9 random seeds runs. For CIP, the same hyperparameters as in the previous setting are used. The MLPs implemented in CF1 and CF2 used for the prediction of \( \hat{Y} \) and the one used for the prediction of the \( L \) residuals in CF2 are all designed with similar architecture and training method. The MLP models consist of 8 hidden layers with 20 nodes each, connected with a rectified linear activation function (ReLU) and a linear final layer. During training, mini-batch size of 64 and the Adam optimizer with a learning rate of \( 10^{-3} \) for 200 epochs were used.
Table 1: Performance of the HSCIC against baselines CF1 and CF2 on two synthetic datasets. Notably, in both scenarios it is possible to select γ values for which CIP outperforms CF2 in MSE and VCF simultaneously.

| Scenario 1 | Scenario 2 |
|------------|------------|
| MSE $\times 10^0$ | HSCIC $\times 10^3$ | VCF $\times 10^2$ | MSE $\times 10^1$ | HSCIC $\times 10^2$ | VCF $\times 10^2$ |
| γ = 0.001 | 12 ± 9 | 45.38 ± 0.41 | 54.93 ± 7.50 | 0.0006 | 35.64 ± 0.32 | 5.60 ± 0.03 |
| γ = 0.01 | 16 ± 12 | 45.35 ± 0.41 | 54.57 ± 7.18 | 0.0019 | 35.44 ± 0.33 | 5.50 ± 0.03 |
| γ = 0.1 | 32 ± 20 | 45.11 ± 0.43 | 54.16 ± 7.58 | 0.11 ± 0.006 | 33.47 ± 0.36 | 4.46 ± 0.04 |
| γ = 0.2 | 81 ± 14 | 44.78 ± 0.47 | 53.59 ± 7.90 | 0.42 ± 0.02 | 31.38 ± 0.38 | 3.52 ± 0.04 |
| γ = 0.3 | 192 ± 33 | 43.92 ± 0.52 | 52.92 ± 7.54 | 0.82 ± 0.04 | 29.75 ± 0.34 | 2.50 ± 0.04 |
| γ = 0.4 | 384 ± 58 | 43.88 ± 0.57 | 52.06 ± 7.25 | 1.21 ± 0.05 | 28.63 ± 0.33 | 1.79 ± 0.03 |
| γ = 0.5 | 685 ± 133 | 42.26 ± 0.65 | 51.64 ± 7.40 | 1.56 ± 0.08 | 27.81 ± 0.26 | 1.1 ± 0.01 |
| γ = 0.6 | 1117 ± 165 | 42.47 ± 0.73 | 50.96 ± 7.36 | 1.84 ± 0.11 | 26.87 ± 0.22 | 0.79 ± 0.01 |
| γ = 0.7 | 1655 ± 223 | 42.11 ± 0.80 | 50.31 ± 7.44 | 2.11 ± 0.14 | 26.08 ± 0.20 | 0.49 ± 0.01 |
| γ = 0.8 | 2225 ± 296 | 41.87 ± 0.84 | 49.76 ± 7.25 | 2.37 ± 0.15 | 25.27 ± 0.18 | 0.31 ± 0.01 |
| γ = 0.9 | 2832 ± 372 | 41.52 ± 0.92 | 49.17 ± 7.41 | 2.58 ± 0.17 | 24.64 ± 0.16 | 0.21 ± 0.01 |
| γ = 1.0 | 3472 ± 422 | 38.37 ± 0.97 | 48.71 ± 7.55 | 2.77 ± 0.19 | 24.21 ± 0.15 | 0.14 ± 0.01 |
| CF1       | 10321 ± 72 | 41.37 ± 0.58 | 50 ± 0.00 | 4.59 ± 0.4478 | 25.01 ± 0.25 | 0 ± 0.00 |
| CF2       | 2728 ± 272 | 41.37 ± 0.92 | 59.50 ± 10.35 | 3.97 ± 0.3479 | 27.03 ± 0.35 | 2.62 ± 0.81 |

G.3 Datasets and results for multi-dimensional variables experiments

The data-generating mechanisms for the multi-dimensional settings of Fig. 3 are now shown. Given dimA = $D_1 \geq 2$, the datasets were generated from:

$$Z \sim \mathcal{N}(0, 1) \quad A_i = Z^2 + \varepsilon_A^i \quad \text{for } i \in \{1, D_1\}$$

$$L = \exp\left\{-\frac{1}{2} A_1\right\} + \sum_{i=1}^{D_1} A_i \cdot \sin(Z) + 0.1 \cdot \varepsilon_L$$

$$Y = \exp\left\{-\frac{1}{2} A_2\right\} \cdot \sum_{i=1}^{D_1} A_i + LZ + 0.1 \cdot \varepsilon_Y,$$

where $\varepsilon_L, \varepsilon_Y \sim \mathcal{N}(0, 0.1)$ and $\varepsilon_A^1, \ldots, \varepsilon_A^{D_1} \sim \mathcal{N}(0, 1)$. In this experiment, the mini-batch size chosen is 512 and the same hyperparameters are used as in the previous settings. The neural network architecture is trained for 800 epochs. Fig. 7 present the results corresponding to 10 random seeds with different values of the trade-off parameter γ corresponding to different values of dimA among {15, 100}. In all of the box plots, it is evident that there exists a trade-off between the accuracy and counterfactual invariance of the predictor. As the value of γ increases, there is a consistent trend of augmenting counterfactual invariance (as evidenced by the decrease in the VCF metric). Similarly to the previous boxplots visualizations, the boxes represent the interquartile range (IQR), the horizontal line is the median, and whiskers show the minimum and maximum values, excluding the outliers (determined as a function of the inter-quartile range). Outliers are represented in the plot as dots.
Figure 7: MSE, HSCIC, VCF for increasing dimension of $A$ on synthetic data from Section G.3 with dimA = 20 (left) and dimA = 100 (right). All other variables are one-dimensional.

G.4 Image dataset

The simulation procedure for the results shown in Section 4.2 is the following.

\[
\begin{align*}
\text{shape} & \sim \mathcal{P}(\text{shape}) \\
\text{y-pos} & \sim \mathcal{P}(\text{y-pos}) \\
\text{color} & \sim \mathcal{P}(\text{color}) \\
\text{orientation} & \sim \mathcal{P}(\text{orientation}) \\
\text{x-pos} & = \text{round}(x), \quad \text{where } x \sim \mathcal{N}(\text{shape} + \text{y-pos}, 1) \\
\text{scale} & = \text{round}\left(\frac{x \cdot \text{pos}}{24} + \frac{y \cdot \text{pos}}{24}\right) \cdot \text{shape} + \epsilon_S \\
Y & = e^{\text{shape}} \cdot \text{x-pos} + \text{scale}^2 \cdot \sin(\text{y-pos}) + \epsilon_Y,
\end{align*}
\]

where $\epsilon_S \sim \mathcal{N}(0, 1)$ and $\epsilon_Y \sim \mathcal{N}(0, 0.01)$. The data has been generated via a matching procedure on the original dSprites dataset.

In Table 2, the hyperparameters of the layers of the convolutional neural network are presented. Each of the convolutional groups also has a ReLU activation function and a dropout layer. Two MLP architectures have been used. The former takes as input the observed tabular features. It is composed by two hidden layers of 16 and 8 nodes respectively, connected with ReLU activation functions and dropout layers. The latter takes as input the concatenated outcomes of the CNN and the other MLP. It consists of three hidden layers of 8, 8 and 16 nodes, respectively.

G.5 Fairness with continuous protected attributes

The pre-processing of the UCI Adult dataset was based upon the work of Chiappa and Pacchiano [2021]. Referring to the causal graph in Fig. 8, a variational autoencoder [Kingma and Welling, 2014] was trained for each of the unobserved variables $H_m$, $H_d$ and $H_r$. The prior distribution of these latent variables is assumed to be standard Gaussian. The posterior distributions

Figure 8: Assumed causal graph for the Adult dataset, as in Chiappa and Pacchiano [2021]. The variables $H_m$, $H_d$, $H_r$ are unobserved, and jointly trained with the predictor $Y$. 

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Table 2: Architecture of the convolutional neural network used for the image dataset, as described in Section G.4.

| layer        | # filters | kernel size | stride size | padding size |
|--------------|-----------|-------------|-------------|--------------|
| convolution  | 16        | 5           | 2           | 2            |
| max pooling  | 1         | 3           | 2           | 0            |
| convolution  | 64        | 5           | 1           | 2            |
| max pooling  | 1         | 1           | 2           | 0            |
| convolution  | 64        | 5           | 1           | 2            |
| max pooling  | 1         | 2           | 1           | 0            |
| convolution  | 16        | 5           | 1           | 3            |
| max pooling  | 1         | 2           | 2           | 0            |

\( P(H_m | V), P(H_r | V), P(H_d | V) \) are modeled as 10-dimensional Gaussian distributions, whose means and variances are the outputs of the encoder.

The encoder architecture consists of a hidden layer of 20 hidden nodes with hyperbolic tangent activation functions, followed by a linear layer. The decoders have two linear layers with a hyperbolic tangent activation function. The training loss of the variational autoencoder consists of a reconstruction term (Mean-Squared Error for continuous variables and Cross-Entropy Loss for binary ones) and the Kullback–Leibler divergence between the posterior and the prior distribution of the latent variables. For training, we used the Adam optimizer with learning rate of \( 10^{-2} \), 100 epochs, mini-batch size 128.

The predictor \( \hat{Y} \) is the output of a feed-forward neural network consisting of a hidden layer with a hyperbolic tangent activation function and a linear final layer. In the training we used the Adam optimizer with learning rate \( 10^{-3} \), mini-batch size 128, and trained for 100 epochs. The choice of the network architecture is based on the work of Chiappa and Pacchiano [2021].

The estimation of counterfactual outcomes is based on a Monte Carlo approach. Given a data point, 500 values of the unobserved variables are sampled from the estimated posterior distribution. Given an interventional value for \( A \), a counterfactual outcome is estimated for each of the sampled unobserved values. The final counterfactual outcome is estimated as the average of these counterfactual predictions. In this experimental setting, we have \( k = 100 \) and \( d = 1000 \).

In the causal graph presented in Fig. 8, \( A \) includes the variables age and gender, \( C \) includes nationality and race, \( M \) marital status, \( D \) level of education, \( R \) the set of the working class, occupation, and hours per week and \( Y \) the income class. Compared to Chiappa and Pacchiano [2021], we include the race variable in the dataset as part of the baseline features \( C \). The loss function is the same as Eq. (1) but Binary Cross-Entropy loss (\( L_{BCE} \)) is used instead of Mean-Squared Error loss:

\[
L_{CIP}(\hat{Y}) = L_{BCE} (\hat{Y}) + \gamma \cdot HSCIC (\hat{Y}, \{\text{age, gender, marital status, education, work}\}|S)
\]

(14)

where the set \( S = \{\text{Race, Nationality}\} \) blocks all the non-causal paths from \( W \cup A \) to \( Y \). In this example we have \( W = \{C \cup M \cup D \cup R\} \). The results in Fig. 5 (right) refer to one run with conditioning set \( S = \{\text{Race, Nationality}\} \). The results correspond to 4 random seeds.

G.6 Illustrating the choice of \( \gamma \)

In Section 3.4, we propose to choose \( \gamma \) to obtain a maximal level of CI within a given tolerance on predictive performance. Here, we illustrate results from running the proposed procedure that dynamically selects \( \gamma \), adjusted to different predefined accuracy thresholds in a classification setting. Specifically, the algorithm chooses the largest \( \gamma \) value that yields an accuracy equal to
Table 3: Results of MSE and VCF (all times $10^2$ for readability) on synthetic data of CIP with trade-off parameters depending on the chosen accuracy threshold.

| Accuracy | VCF $\times 10^2$ | HSCIC $\times 10^2$ |
|----------|------------------|------------------|
| 90%      | 3.14 ± 0.92      | 4.51 ± 0.72      |
| 70%      | 3.01 ± 0.80      | 4.44 ± 0.65      |
| 1%       | 2.91 ± 0.92      | 4.39 ± 0.42      |

or better than the threshold. As described the algorithm operates on $\gamma$ values on a logarithmic scale, thereby ensuring a fine-grained search over a wide range of potential trade-off points. Table 3 shows the found trade-offs for tolerated accuracies of 90%, 70%, and 1% in the same setting as Section G.1.

G.7 Computational complexity and runtimes

In a dataset with $n = 1000$ data points from the setting discussed in Fig. 2, the average training time for one epoch without the regularization term is 0.003s and 1.112s with the regularization term. In these results, Adam optimizer with batch-size of 512 was used. By using smaller batch sizes, e.g. $n = 128$, the extra computational cost can further decrease. From a theoretical perspective, the estimation of the HSCIC requires kernel ridge regression (see Eq. (2) in our submission). In the high-dimensional image example, with a mini batch-size of 512, the average running time for an epoch with the regularization term is 64.03s and 34.01s without. Kernel ridge regression generally has a runtime that scales as $O(n^3)$ and memory requirements scaling like $O(n^2)$, where $n$ is the size of the dataset. However, these bounds can be significantly improved by using, i.e., random Fourier Features (see, i.e., Avron et al. 2017, Rahimi and Recht 2007) as detailed in Section F. In short, by using random Fourier features, the resulting approximate kernel ridge regression estimator can be computed in a runtime of $O(ns^2)$ with $O(ns)$ memory. Here, $s$ is a parameter determining the accuracy of the approximation. In practice, $s$ can be set to be much smaller than the problem size, resulting in a dramatic speed-up. Other methods for efficient kernel computation include the popular Nyström approximation (Drineas and Mahoney, 2005, Hsieh et al., 2014), and Memory-Efficient Kernel Approximation (MEKA) (Si et al., 2017). In this work, runtimes were still reasonable for all experimental settings, which is why we did not have to resort to these faster approximations.

H Comparison with additional baselines

In this section, we compare CIP with additional baselines. These include Veitch et al. [2021] and different heuristic methods.

H.1 Baseline experiments [Veitch et al., 2021]

We provide an experimental comparison against the method by Veitch et al. [2021]. To this end, we consider the following data-generating mechanism for the causal structure (see Fig. 1(b)):

$$Z \sim \mathcal{N}(0, 1) \quad A = \sin (0.1Z) + \varepsilon_A$$

$$X = \exp \left\{ -\frac{1}{2}A \right\} \sin (A) + \frac{1}{10} \varepsilon_X$$

$$Y = \frac{1}{10} \exp \{-X\} \cdot \sin (2XZ) + AA + \frac{1}{10} \varepsilon_Y,$$
Table 4: Results of the MSE, VCF of CIP and the baseline [Veitch et al., 2021] applied to the causal and anti-causal structure in Fig. 1(b-c). Although the graphical assumptions are not satisfied, CIP shows an overall decrease of VCF in both of the graphical structures, performing on par with the baseline Veitch et al. [2021] in terms of accuracy and counterfactual invariance.

|       | CIP          | Veitch et al. [2021]          |
|-------|--------------|------------------------------|
|       | MSE $\times 10^2$ | VCF $\times 10^2$ | MSE $\times 10^2$ | VCF $\times 10^2$ |
| $\gamma = 0.5$ | $4.58 \pm 0.31$ | $0.19 \pm 0.02$ | $4.50 \pm 0.40$ | $0.19 \pm 0.02$ |
| $\gamma = 1.0$ | $5.60 \pm 0.36$ | $0.18 \pm 0.01$ | $5.45 \pm 0.41$ | $0.18 \pm 0.02$ |

where $\varepsilon_X, \varepsilon_A \overset{i.i.d}{\sim} \mathcal{N}(0, 1)$ and $\varepsilon_Y \overset{i.i.d}{\sim} \mathcal{N}(0, 0.1)$. The data-generating mechanism of the anti-causal structure is the following (see Fig. 1(c)):

$$Z \sim \mathcal{N}(0, 1) \quad A = \frac{1}{5} \sin(Z) + \varepsilon_A$$

$$Y = \frac{1}{10} \sin(Z) + \varepsilon_Y$$

$$X = A + Y + \frac{1}{10} \varepsilon_X$$

where $\varepsilon_Y, \varepsilon_A \overset{i.i.d}{\sim} \mathcal{N}(0, 0.1)$ and $\varepsilon_X \overset{i.i.d}{\sim} \mathcal{N}(0, 1)$. We compare our method (CIP) against the method by Veitch et al. [2021] using different values for the trade-off parameter $\gamma$. In Fig. 1(b-c) the causal and anti-causal graphical settings proposed by Veitch et al. [2021] are presented. In both of these settings there is an unobserved confounder $Z$ between $A$ and $Y$. The graphical assumptions outlined in Theorem 3.2 of the CIP are not met in the graphical structures under examination, as the confounding path is not effectively blocked by an observed variable ($Z$ is unobserved). In light of this, it is assumed in our implementation that there is no unobserved confounder. In the graphical structure Fig. 1(b), CIP enforces $\text{HSIC}(\hat{Y}, A \cup X)$ to become small, gradually enforcing $\hat{Y} \perp \perp A \cup X$. HSIC is the Hilbert-Schmidt Independence Criterion, which is commonly used to promote independence (see, i.e., Fukumizu et al. [2007], Gretton et al. [2005]). Veitch et al. [2021] enforces as independence criterion $\text{HSIC}(\hat{Y}, A)$, which is implied by the independence enforced in CIP. In the anti-causal graphical setting presented in Fig. 1(c), the objective term used in CIP is $\text{HSCIC}(\hat{Y}, A | X)$, while in the method of Veitch et al. [2021] is $\text{HSCIC}(\hat{Y}, A | Y)$. In Table 4, the results of accuracy and VCF are presented.

In the experiments, the predictor $\hat{Y}$ is a feed-forward neural network consisting of 8 hidden layers with 20 nodes each, connected with a rectified linear activation function (ReLU) and a linear final layer. Mini-batch size of 256 and the Adam optimizer with a learning rate of $10^{-4}$ for 500 epochs were used.

H.2 Comparison baselines heuristic methods

We provide an experimental comparison of the proposed method (CIP) with some heuristic methods, specifically data-augmentation-based methods. We consider the same data-generating procedure and causal structure as presented in Section G.1. The heuristic methods considered
Table 5: Results of MSE and VCF (all times $10^2$ for readability) on synthetic data of CIP with trade-off parameters $\gamma = 0.5$ and $\gamma = 1$ with the heuristic methods data augmentation and causal-based data augmentation and naive prediction.

| Method                                      | VCF  $\times 10^2$ | MSE  $\times 10^3$ |
|---------------------------------------------|--------------------|---------------------|
| data augmentation                           | 3.12 ± 0.16        | 0.03 ± 0.01         |
| causal-based data augmentation              | 3.04 ± 0.16        | 0.13 ± 0.12         |
| CIP ($\gamma = 0.5$)                        | 1.05 ± 0.13        | 1.64 ± 0.22         |
| CIP ($\gamma = 1.0$)                        | 0.35 ± 0.19        | 2.50 ± 0.72         |
| naive prediction (ignore $A$)               | 9.01 ± 0.02        | 3.01 ± 0.91         |

are data augmentation and causal-based data augmentation. In the former, data augmentation is performed by generating $N = 50$ samples for every data-point by sampling new values of $A$ as $a_1, \ldots, a_N \sim \mathcal{F}_A$ and leaving $Z, L, Y$ unchanged. Differently, in the latter causal-based data augmentation method, we also take into account the causal structure given by the known DAG. Indeed, when manipulating the variable $A$, its descendants (in this example $L$) will also change. In this experiment, a predictor for $L$ as $\hat{L} = f_\theta(A, Z)$ is trained on 80% of the original dataset. In the data augmentation mechanism, for every data-point $\{a, x, z, y\}$, $N = 50$ samples are generated by sampling new values of $A$ as $a_1, \ldots, a_N \sim \mathcal{F}_A$, estimating the values of $L$ as $x_1 = f_\theta(a_1, z), \ldots, x_N = f_\theta(a_N, z)$, while leaving the values of $Z$ and $Y$ unchanged. Heuristic methods such as data-augmentation methods do not theoretically guarantee to provide counterfactually invariant predictors. The results of an empirical comparison are shown in Table 5 with the average and standard deviations after 5 random seeds. It can be shown that these theoretical insights are supported by experimental results, as the VCF metric measure counterfactual invariance is lower in both of the two settings of the CIP ($\gamma = \frac{1}{2}$ and $\gamma = 1$).

A dataset of $n = 3000$ is used, along with $k = 500$ and $d = 500$. The architecture for predicting $L$ and $Y$ are feed-forward neural networks consisting of 8 hidden layers with 20 nodes each, connected with a rectified linear activation function (ReLU) and linear final layer. Mini-batch size of 256 and the Adam optimizer with a learning rate of $10^{-3}$ for 100 epochs were used.