
Measuring statistical dependencies via maximum norm and characteristic functions

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Abstract

1 In this paper we focus on the problem of statistical dependence estimation. We propose
2 statistical dependence measure based on the maximum-norm of the absolute
3 value of difference between joint and product-marginal characteristic functions,
4 and its iterative estimation algorithm. The proposed measure is differentiable,
5 can be efficiently applied to high-dimensional data, and integrated into modern
6 machine learning pipelines. We also conduct experiments both with simulated and
7 real data, which reveal that the proposed measure can exploit statistical dependence
8 in non-linear data sets more efficiently, comparing to the previous work in this line
9 of research, and that it can improve real-data classification accuracy, when applied
10 for different tasks.

11 1 Introduction

12 The measurement of statistical dependence plays important role in various empirical learning methods
13 (e.g. hypothesis testing [1], feature selection and extraction [2, 3], information bottleneck methods
14 [4], causal inference [5], self-supervised learning [6], representation learning [7], among others).
15 Historically, earliest statistical dependence estimation ideas (e.g. conditional probability) share
16 nearly-common origin with the beginning of formal statistical reasoning itself. During last two
17 centuries ideas of correlation and (relative) entropy (including various generalizations) were proposed
18 and became very popular in numerous applications and theoretical developments. In recent years,
19 various other approaches (e.g. [1], [9, 10], [11]) induced or essentially contributed to several lines of
20 research of different popularity, formulating and investigating statistical dependence measures.

21 With the increasing growth of machine and deep learning, new statistical dependence estimation
22 methods, that are robust, applicable to noisy, high-dimensional, structured data, and which can be
23 efficiently integrated with modern machine learning and deep learning methods are helpful for the
24 development both of the theory and application.

25 In this article we focus on quantitative estimation of statistical dependencies, using characteristic
26 functions. We begin with the short review of some important previous approaches (Section 2),
27 devoting special attention to ones based on characteristic functions (Section 2.1). Afterwards, in
28 (Section 3), we formulate the proposed measure, its empirical estimator, and conduct preliminary
29 theoretical analysis, which are the main theoretical contribution of our paper. Section 4 is devoted
30 to the empirical investigation of the proposed measure. Therein we conduct experiments both with
31 simulated and real-world data sets, in order to empirically investigate its properties and applicability
32 in empirical inference scenarios. Finalizing Section 5 discusses and concludes this article.

2 Previous Work

During recent years, various approaches have been used in order to construct statistical dependence estimation methods. For example, information theory (mutual information [8], and generalisations), reproducing kernel Hilbert spaces (Hilbert-Schmidt independence criterion [1]), characteristic functions (distance correlation [9, 10]), and other (e.g. [11] copula-based kernel dependence measures, integral-probability-metric-reliant Sobolev independence criterion [12]). Further we will focus on characteristic-function-based methods.

2.1 Characteristic-function-based methods

Characteristic function (CF) of d_X -dimensional random vector X defined in some probability space $(\Omega_X, \mathcal{F}_X, \mathbb{P}_X)$ is defined as:

$$\phi_X(\alpha) := \mathbb{E}_X e^{i\alpha^T X}, \quad (1)$$

where $i = \sqrt{-1}$, $\alpha \in \mathbb{R}^{d_X}$. Having n i.i.d. realisations of X , corresponding empirical characteristic function (ECF) is defined as:

$$\widehat{\phi}_X(\alpha) := \frac{1}{n} \sum_{j=1}^n e^{i\langle \alpha, x_j \rangle}. \quad (2)$$

Having pair of two random vectors (X, Y) defined in another probability space $(\Omega_{X,Y}, \mathcal{F}_{X,Y}, \mathbb{P}_{X,Y})$ joint CF is defined as:

$$\phi_{X,Y}(\alpha, \beta) := \mathbb{E}_{X,Y} e^{i(\alpha^T X + \beta^T Y)}, \quad (3)$$

where $\alpha \in \mathbb{R}^{d_X}$ and $\beta \in \mathbb{R}^{d_Y}$. Similarly, having n i.i.d. realisations of (X, Y) , joint ECF is defined as:

$$\widehat{\phi}_{X,Y}(\alpha, \beta) := \frac{1}{n} \sum_{j=1}^n e^{i(\langle \alpha, x_j \rangle + \langle \beta, y_j \rangle)}. \quad (4)$$

Uniqueness theorem states that two random variables X and Y have the same distribution if and only if their CF's are identical [?]. Therefore, CF's can be regarded as an alternative description of distribution. Roughly speaking, CF can be regarded as Fourier transform of probability density function (PDF).

If for all $x \in \mathbb{R}^{d_X}$ and $y \in \mathbb{R}^{d_Y}$ cumulative distribution function (CDF) $F_{X,Y}(x, y)$ of (X, Y) factorises as,

$$F_{X,Y}(x, y) = F_X(x)F_Y(y), \quad (5)$$

where $F_X(x)$ and $F_Y(y)$ are marginal CDF's, X and Y are called independent (the same holds for probability density function, PDF). However, this criterion is impractical due to need of evaluation of potentially high-dimensional CDF or PDF, and often alternative independence criteria are more useful. Let us define

$$\Delta_{X,Y}(\alpha, \beta) := \phi_{X,Y}(\alpha, \beta) - \phi_X(\alpha)\phi_Y(\beta), \quad (6)$$

an its empirical counterpart:

$$\widehat{\Delta}_{X^n, Y^n}(\alpha, \beta) := \widehat{\phi}_{X^n, Y^n}(\alpha, \beta) - \widehat{\phi}_{X^n}(\alpha)\widehat{\phi}_{Y^n}(\beta), \quad (7)$$

where $(X^n, Y^n) := (x_j, y_j)_{j=1}^n$.

In terms of CF's, statistical independence of X and Y is equivalent to $\forall \alpha \in \mathbb{R}^{d_X}, \forall \beta \in \mathbb{R}^{d_Y}, \Delta_{X,Y}(\alpha, \beta) = 0$ [18].

Previously, $\Delta_{X,Y}(\alpha, \beta)$ was used as the basis (first in [9] for one-dimensional case, and afterwards extended and developed by [10] for bivariate multidimensional random vectors) for construction of statistical independence tests and measures. Distance covariance and distance correlation, proposed by [10] relies on weighted L^2 -norm analysis of (6). They select weighting function in such a way, that dependence measure can be expressed in terms of correlation of data-dependent distances. Recent result of [13] generalises [10] to multivariable case. [14] proposed computationally efficient algorithm for estimation of distance correlation measure, reducing computational complexity from $O(n^2)$ to $O(n \cdot \log n)$, where n is sample size.

71 **Motivation and Connection To Previous Work** Taking $\Delta_{X,Y}(\alpha, \beta) = 0$ (6) as the criterion of
 72 statistical independence we view the work [10] from the perspective of weighted L^p spaces, measuring
 73 statistical dependence by the corresponding L^p -norms of $|\Delta_{X,Y}(\alpha, \beta)|$ (6) (i.e. $\|\Delta_{X,Y}(\alpha, \beta)\|_{L^p} =$
 74 $\int |\Delta_{X,Y}(\alpha, \beta)|^p d\alpha d\beta)^{\frac{1}{p}}$).

75 Taking into account that [10] in high dimensions is affected with the curse of dimensionality [15], we
 76 focus on the limit case $p \rightarrow \infty$ (L^∞ space), which is associated to the supremum norm. This norm
 77 has several potential advantages.

78 We hypothesise, that its locality could be exploited to detect statistical independence more efficiently,
 79 comparing to case $p = 2$. In addition, numerically calculation of L^∞ norm would not require to
 80 directly calculate norm integral, since norm of L^p converges to supremum norm when $p \rightarrow \infty$. Also,
 81 from practical point of view maximization is convenient, because it is efficiently implemented in
 82 modern deep learning frameworks (e.g. Pytorch [16]). In addition, in our opinion it is worth to note,
 83 that applications of characteristic functions in machine learning are quite scarce, despite that they
 84 provide quite convenient theoretical proxy to access distributions.

85 3 Proposed Measure

86 The above considerations serves as the basis for constructing of a novel dependence measure, which
 87 we further refer to as Kac independence measure (KacIM). Having two random vectors X and Y ,
 88 KacIM is defined as

$$\kappa(X, Y) := \max_{\alpha \in \mathbb{R}^{d_X}, \beta \in \mathbb{R}^{d_Y}} |\Delta_{X,Y}(\alpha, \beta)|. \quad (8)$$

89 3.1 Basic Properties

90 **Theorem 1.** *KacIM (8) has the following properties:*

- 91 1. $\kappa(X, Y) = \kappa(Y, X)$,
- 92 2. $0 \leq \kappa(X, Y) \leq 1$,
- 93 3. $\kappa(X, Y) = 0$ iff $X \perp Y$.

94 *Proof.* Property 1. is obvious from definition (8) (commutativity of addition and multiplication), and
 95 property 2. directly follows from Cauchy inequality and that absolute value of CF is bounded by 1:

$$\begin{aligned} |\phi_{X,Y}(\alpha, \beta) - \phi_X(\alpha)\phi_Y(\beta)|^2 &= \mathbb{E}_{X,Y} |(e^{i\alpha^T X} - \phi_X(\alpha))(e^{i\beta^T Y} - \phi_Y(\beta))|^2 \leq \\ &\mathbb{E}_{X,Y} |(e^{i\alpha^T X} - \phi_X(\alpha))|^2 |(e^{i\beta^T Y} - \phi_Y(\beta))|^2 = (1 - |\phi_X(\alpha)|^2)(1 - |\phi_Y(\beta)|^2). \end{aligned}$$

96 Property 3. directly follows from properties of CF's (see e.g. [17], Corollary 14.1)¹. \square

97 Although (8) is not scale invariant in general, scale invariance can be achieved by assuming standardi-
 98 zation of X and Y .

99 3.2 Estimation

100 Having i.i.d. observations $(X^n, Y^n) := (x_j, y_j)$, $j = 1, 2, \dots, n$, an empirical estimator of (8) is
 101 defined via corresponding ECF's (4) and (2):

$$\hat{\kappa}(X^n, Y^n) := \max_{\alpha, \beta} |\hat{\Delta}_{X^n, Y^n}(\alpha, \beta)| = \max_{\alpha, \beta} |\hat{\phi}_{X^n, Y^n}(\alpha, \beta) - \hat{\phi}_{X^n}(\alpha)\hat{\phi}_{Y^n}(\beta)|. \quad (9)$$

102 By *Levy continuity theorem* [18] ECF pointwise converges to CF. Therefore empirical estimator (9)
 103 almost surely converges into KacIM (8). In practice, KacIM (8) can be estimated iteratively by
 104 Algorithm 1².

¹This property also is known as Kac's theorem [18]. Although it is quite simple mathematical fact, this provides the basis of the proposed measure's name.

²Pytorch [16] implementation can be accessed from https://github.com/povidanius/kac_independence_measure

Algorithm 1 KacIM gradient estimation.

Require: Number of iterations N , batch size n_b , gradient-based optimiser $\text{GradOpt}([parameters], \cdot)$, initial $\alpha \in \mathbb{R}^{d_X}, \beta \in \mathbb{R}^{d_Y}$.
for iteration=1 to N **do**
 Sample data batch $(X^n, Y^n) := (x_i, y_i)_{i=1}^{n_b}$.
 Normalize (X^n, Y^n) to zero mean and unit variance (scale invariance).
 Calculate $\hat{\Delta}_{\alpha, \beta}((X^n, Y^n))$.
 Perform one maximization iteration of $\hat{\Delta}_{\alpha, \beta}(X^n, Y^n)$: $\hat{\Delta}_{\alpha, \beta}(X^n, Y^n)$ via $\alpha, \beta \rightarrow \text{GradOpt}([\alpha, \beta], \hat{\Delta}_{\alpha, \beta}(X^n, Y^n))$.
end for

Algorithm 1 requires to initialise α and β , and gradient-based (local) optimiser. In our implementation we use uniform initialisation of parameters, and decoupled weight decay regularization optimizer [19]. We also empirically observed that normalisation of parameters α and β on to unit sphere increases estimation stability. After the estimation of KacIM via Algorithm 1, the evaluation the estimator has computation complexity $O(n)$, where n is sample size.

3.3 Interpretation and connection to the related approaches

Interpretation as maximum covariance. Since (8) can be reformulated as

$$\kappa(X, Y) = \max_{\alpha, \beta} |\text{cov}(e^{i\alpha^T X}, e^{i\beta^T Y})|, \quad (10)$$

by Euler's formula, it corresponds to the maximum pseudocovariance between complex exponents $e^{i\alpha^T X} = \cos(\alpha^T X) + i \cdot \sin(\alpha^T X)$ and $e^{i\beta^T Y} = \cos(\beta^T Y) + i \cdot \sin(\beta^T Y)$. Since $\text{var}(e^{i\alpha^T X}) = \phi(2\alpha) - \phi(\alpha)^2$, one can also define the normalised version of KacIM (refine or remove this):

$$\kappa_{\text{norm}}(X, Y) = \max_{\alpha, \beta} |\text{corr}(e^{i\alpha^T X}, e^{i\beta^T Y})| = \max_{\alpha, \beta} \frac{|\text{cov}(e^{i\alpha^T X}, e^{i\beta^T Y})|}{\sqrt{|\phi_X(2\alpha) - \phi_X^2(\alpha)| |\phi_Y(2\beta) - \phi_Y^2(\beta)|}}. \quad (11)$$

Interpretation in Gaussian case. In special case when both X and Y are zero mean Gaussian random vectors we have $\phi_X(\alpha) = e^{-\frac{1}{2}\alpha^T \Sigma_X \alpha}$, $\phi_Y(\beta) = e^{-\frac{1}{2}\beta^T \Sigma_Y \beta}$, $\phi_{X,Y}(\alpha, \beta) = e^{-\frac{1}{2}(\alpha^T \Sigma_X \alpha + \beta^T \Sigma_Y \beta + 2\alpha^T \Sigma_{X,Y} \beta)}$. Therefore

$$\kappa(X, Y) = \max_{\alpha, \beta} |e^{-\frac{1}{2}(\alpha^T \Sigma_X \alpha + \beta^T \Sigma_Y \beta)} (e^{-\alpha^T \Sigma_{X,Y} \beta} - 1)|. \quad (12)$$

Assuming constant $\alpha^T \Sigma_X \alpha$ and $\beta^T \Sigma_Y \beta$, the maximization corresponds to the maximization of $\alpha^T \Sigma_{X,Y} \beta$, which coincides with canonical correlation analysis [21]. Here Σ_X , Σ_Y , and $\Sigma_{X,Y}$ are covariance matrices of X , Y , and cross-covariance matrix between X and Y , respectively.

Mutual information For the neural estimation of mutual information its variational (Donsker-Varadhan) representation $I(X, Y) = \max_{\theta} \mathbb{E}_{X,Y} f(x, y|\theta) - \log \mathbb{E}_X \mathbb{E}_Y e^{f(x,y|\theta)}$ [20] is often used, since it allows to avoid density estimation (here $f(x, y|\theta)$ is neural network with parameters θ). The estimation is also iterative process, similar to Algorithm 1. In this case, optimisation is conducted over the space of neural network parameters, which often is substantially larger than the number of parameters needed to estimate KacIM (i.e. $d_x + d_y$ parameters).

4 Experiments

Further we will conduct empirical investigation of KacIM in order to investigate its behaviour in simulations and practical applications.

4.1 Generated data

Non-linear statistical dependence detection. We begin with simulated multivariate data with additive and multiplicative noise.

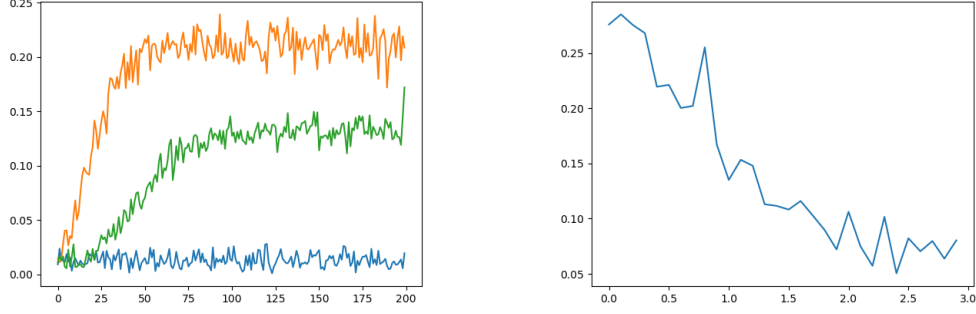


Figure 1: Left figure: KacIM evaluation for independent data (blue), additive (orange) and multiplicative (green) noise scenarios (x axis - iteration, and y - corresponding value of KacIM). Right figure: noise level (x axis) vs final iteration KacIM value (y axis). KacIM values for larger noise levels saturates as in tail of graph

Figure 1 reflects KacIM values during iterative adaptation (200 iterations). In the case of independent data, both x_i and y_i ($d_x = 512$, $d_y = 4$) are sampled from gaussian distribution, independently. In the case of dependent data, an additive noise and multiplicative noise, the dependent variable is generated according to $y_i = \sin(Px_i) + \cos(Px_i) + \lambda\epsilon_i$ ($\lambda = 1.00$) and $y_i = (\sin(Px_i) + \cos(Px_i))\epsilon_i$, respectively, where P is $d_x \times d_y$ random projection matrix, $\epsilon_i \sim N(0, 1)$ and $\epsilon_i \perp x_i$.

When data is independent, both in additive and multiplicative cases, due to independence, estimator (9) is resistant to maximisation, and oscillates near zero. On the other hand, when the data is not independent, the condition (6) is violated and maximization of estimator (9) is possible.

Noise variance effect In this simulation we use the same additive noise setting as in previous paragraph, but evaluate all noise levels $\lambda \in [0.1, 3.0]$, with step 0.1. Figure 1 empirically shows that value of KacIM negatively correlates with noise level, and therefore the proposed measure is able not only to detect whether independence is present, but also to quantitatively evaluate it.

Comparison with distance correlation We also evaluated distance correlation [10] on the same generated samples of data, comparing it with KacIM. From Figure. 2 we see that as data dimensionality grows, for independent data, the values of measure not only is significantly larger than zero, but it also grows like values of measure of dependent data. This empirically demonstrates that distance correlation is affected by the curse of dimensionality. On the other side, KacIM even for larger dimensions oscillates near zero for independent data, and significantly deviates from zero for dependent data case, as indicated in right component of Figure. 2.

4.2 Feature extraction

Previous work in the field of supervised feature extraction, which rely on dependency-based cost functions, include [2, 3, 22] (HSIC),....().

Let us denote by $T := (x_i, y_i)_{i=1}^N$ a supervised-learning dataset of N pairs of d_x -dimensional inputs x_i , and d_y -dimensional one-hot-encoded outputs y_i .

In feature extraction experiments we will use a set of classification data sets from OpenML [23], which cover different domains. We use KacIM in order to conduct supervised linear feature extraction by seeking

$$W^* = \arg \max_W \hat{\kappa}_N(Wx, y) - \lambda \text{Tr}\{(W^T W - I)^T (W^T W - I)\}, \quad (13)$$

where the regularisation term, controlled by multiplier $\lambda \geq 0$, enforces semi-orthogonality of projection matrix W^* , and $\text{Tr}\{.\}$ denotes matrix trace operator.

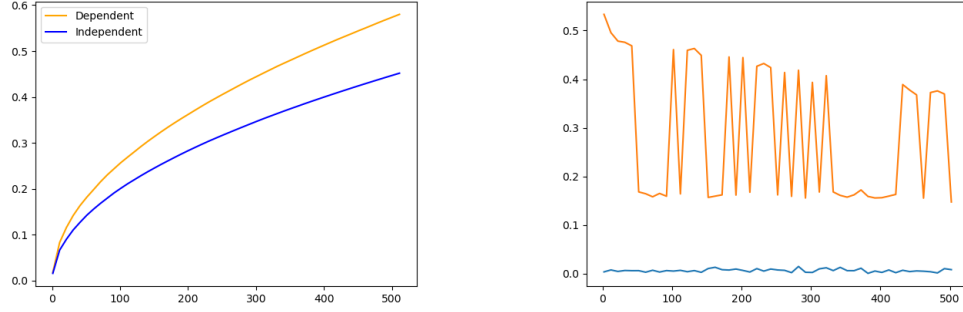


Figure 2: The dimension of data is on the x axis, and on y axis is evaluation of distance correlation (left) and KacIM (right). Blue graph corresponds of independent data of dimension, indicated by x axis, and orange one corresponds to dependent data.

Dataset	$N/d_x/n_c$	Raw	KacIMFE	NCA
isolet	(7797,617,26)	0.9261	0.9437	0.9477
madelon	(2600,500,2)	0.6015	0.5484	0.5685
prnn-viruses	(61,18,4)	0.6452	0.9265	0.9355
ionosphere	(351,34,2)	0.8807	0.9278	0.9375
micro-mass	(360,1300,10)	0.8778	0.9282	0.8944
clean1	(476,168,2)	0.7689	0.9888	0.9790
robot-failures-lp2	(47,90,5)	0.4583	0.6067	0.5833
waveform-5000	(5000,40,3)	0.8692	0.8017	0.8516
spambase	(4601,57,2)	0.6906	0.8285	0.8705
gina-agnostic	(3468,970,2)	0.8512	0.7894	0.8080
scene	(2407,299,2)	0.8895	0.9707	0.9336
tokyo1	(959,44,2)	0.7250	0.8995	0.9062
one-hundred-plants-shape	(1600,64,100)	0.1013	0.4913	0.4688

Table 1: Classification accuracies. N denotes full data set size, d_x - input dimensionality, and n_c - number of classes. In this table feature dimension is equal to a half of original input dimension. Best accuracies that are also statistically significant (Wilcoxon’s signed rank test [25], 25 runs, p -value threshold 0.01) are indicated in bold text.

In all the experiments (13) the cost function is optimised iteratively (250 iterations), learning rate was set to 0.007, simultaneously optimising parameters of KacIM (α and β) and projection matrix W . After the optimisation, the feature extraction is conducted by $f(x) = W^*x$, where x is original input vector, and f are corresponding feature vector.

In each experiment, we randomly split all the datasets in training and testing sets of equal size, and report accuracy, measured on the testing set as the performance measure. We set λ to 1.0 to quickly ensure orthogonal projection matrices, and further proceed to dependence maximization stage. In order to quantitatively evaluate features, we use logistic regression classifier accuracy, measured on the testing set. The logistic regression classifier is trained using the data from the training set.

We compare our approach with the two baselines: raw features (RAW column in Table 1) and neighborhood component analysis [24] (NCA column in Table 1).

The classification accuracies, reported in Table 1 demonstrate that KacIM-based feature extraction procedure (KacIMFE column) indeed allows to increase classification accuracy when applied to real data sets from different domains, including high-dimensional and ill-defined ones (e.g. *micro-mass* dataset). In contrast to our feature extraction approach, NCA explicitly optimises for classification accuracy, rather than more abstract dependency of features $f(x)$ with the dependent variable y .

Mode	Average accuracy (%)
Without regularisation	93.01
With regularisation	93.34

Table 2: Melanoma classification accuracy comparison of regularised and not regularised model. Bold text indicates that model with regulariser was more accurate (Wilcoxon’s signed rank test [25], 30 runs, p -value threshold 0.04))

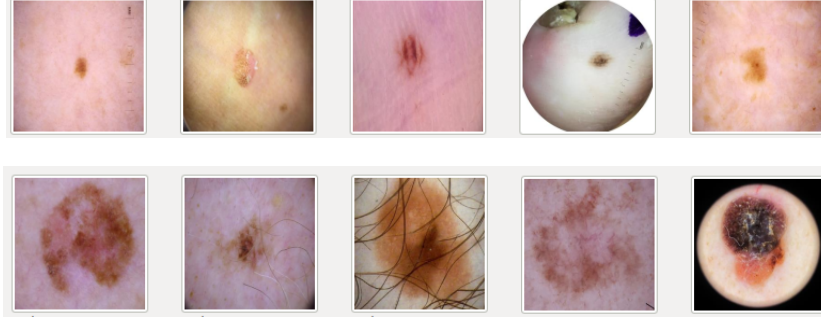


Figure 3: Top figure - benign moles, bottom figure - malignant tumors.

178 4.3 Ensemble Redundancy Regularisation

179 We investigate lung x-ray classification task. It is a binary classification data set, consisting of 5856
180 images, which should be classified as healthy or pneumonia (e.g. Figure 3). For classification we
181 train Resnet18 with 3 classification heads, and majority voting output.

182 As a baseline we minimize average cross-entropy loss, on all three heads. We compare it with the
183 same model, but which loss includes additive redundancy regularization term, $\kappa(z_1, z_2) + \kappa(z_1, z_3) +$
184 $\kappa(z_2, z_3)$, where z_i are internal output of i -th classification head. We expect that this regularization
185 term will make classification heads rely on independent features, thereby redundancy of (bias-
186 variance).

187 We will follow the same protocol as previously, in each experiment splitting the data into random
188 training and testing set (in 80%/20% proportion), training both models on the first one, and testing on
189 the second one. We compare average accuracies of both models, and evaluate statistical significance
190 via Wilcoxon signed rank test.

191 4.4 Regularisation

192 In regularisation experiments we investigate skin lesion classification task. It is a binary classification
193 data set, consisting of 10605 images, which should be classified as benign or malignant (e.g. Figure 3).

194 We use ResNet18 backbone model (pretrained on ImageNet) with added classification head. Fur-
195 ther we train this model with batches of 128 elements. We denote our classification network as
196 $f(\phi(x|\theta_0)|\theta_1)$, where θ_0 are parameters of ResNet18, θ_1 are classification head parameters, and x is
197 224×224 input image. For optimisation we use decoupled weight decay regularization optimizer [19]
198 with learning rate set to 0.0002, and weight decay parameter set to 0.00001 (3 epochs). The internal
199 learning rate of estimator (seeefalogo) was set to 0.07 and weight decay parameters to 0.01.

200 We will investigate additive regularizer, which maximises dependency of bottleneck the feature $\phi(x|\theta_0)$
201 and target variable y (one-hot encoding):

$$Cost(\theta_0, \theta_1, W) := (1 - \rho)CE(f(\phi(x|\theta_0)|\theta_1), y) - \rho\kappa(\phi(x|\theta_0), y), \quad (14)$$

202 where $CE(., .)$ is cross-entropy loss, $\rho \geq 0$ is regularisation parameter (in our experiments $\rho = 0.2$).
203 During backward pass, this regularizer is designed to directly transfer information from y to the
204 output ResNet18 $\phi(.\mid\theta_0)$, and we hypothesise that this could provide possibility to learn more
205 discriminative features.

Mode	Average accuracy (%)
Without regularisation	93.01
With regularisation	93.34

Table 3: Melanoma classification accuracy comparison of regularised and not regularised model. Bold text indicates that model with regulariser was more accurate (Wilcoxon’s signed rank test [25], 30 runs, p -value threshold 0.04))

In each experiment we train classifier 30 times with randomly splitted training and testing data (9000 images for training, and 1605 for testing). The average accuracies reported in Table 3, that application (14) slightly (but with statistical significance) increased classification accuracy.

5 Conclusion

In this article we propose statistical dependence measure, KacIM, which corresponds to the L^∞ norm of the absolute value of difference between joint characteristic function and the product of marginal ones. The proposed measure, in theory can detect non-linear statistical dependence between a pairs of random variables of possibly different dimension, extended to various directions (e.g. kernels, multiple variables), applied to several machine learning tasks (e.g. feature extraction, regularisation, among others). On the other side, it raises a corresponding set of unanswered questions, both theoretical and empirical.

For example, although it converges, the variance of the estimator sometimes is high and it is still remains unclear how to control it, also the interpretability when it approaches its maximal value remains insufficiently clear. However empirical experiments with simulated data reveals, that increasing independence between two random variables is reflected in a decreasing trend on the estimated values of the proposed dependence measure(e.g. Figure 1).

Therefore, parameter initialization, meta-parameter (e.g. stopping criteria, batch size) selection are needed in order to evaluate it efficiently.

Beside demonstrated applications in Section 4, the proposed measure is differentiable and thereby can be integrated with various modern deep-learning methods, applied to high-dimensional and structured data. We see exploration and comparative analysis of KacIM in causality, information bottleneck theory, self-supervised learning, and other modern problems, where dependence measures define a criterion of optimisation, as future work.

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298 **A Appendix**

299 Optionally include extra information (complete proofs, additional experiments and plots) in the
300 appendix. This section will often be part of the supplemental material.