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Exact Partial Information Decompositions for Gaussian Systems Based on Dependency Constraints

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Abstract: The Partial Information Decomposition, introduced by Williams P. L. et al. (2010), provides a theoretical framework to characterize and quantify the structure of multivariate information sharing. A new method (I_{dep}) has recently been proposed by James R. G. et al. (2017) for computing a two-predictor partial information decomposition over discrete spaces. A lattice of maximum entropy probability models is constructed based on marginal dependency constraints, and the unique information that a particular predictor has about the target is defined as the minimum increase in joint predictor-target mutual information when that particular predictor-target marginal dependency is constrained. Here, we apply the I_{dep} approach to Gaussian systems, for which the marginally constrained maximum entropy models are Gaussian graphical models. Closed form solutions for the I_{dep} PID are derived for both univariate and multivariate Gaussian systems. Numerical and graphical illustrations are provided, together with practical and theoretical comparisons of the I_{dep} PID with the minimum mutual information partial information decomposition (I_{mmi}), which was discussed by Barrett A. B. (2015). The results obtained using I_{dep} appear to be more intuitive than those given with other methods, such as I_{mmi} , in which the redundant and unique information components are constrained to depend only on the predictor-target marginal distributions. In particular, it is proved that the I_{mmi} method generally produces larger estimates of redundancy and synergy than does the *I*_{dep} method. In discussion of the practical examples, the PIDs are complemented by the use of tests of deviance for the comparison of Gaussian graphical models.

Keywords: partial information decomposition; mutual information; unique information; dependency constraints; Gaussian graphical models; maximum entropy

1. Introduction

The Partial Information Decomposition (PID) [1] provides a theoretical framework to characterize and quantify the structure of multivariate information sharing. That is, given a *target* variable *Y*, and a number of *predictor* variables X_i the PID attempts to describe the mutual information between the target and predictors $I({X_i}; Y)$ in terms of that which is unique to each predictor, as well as that which is shared (redundant) or synergistic between subsets of predictors. However, while the PID framework provides a theoretical structure for this sharing, practical applications require measures to quantify the different terms. Although a number of different candidate measures have been proposed, this remains an open area of research [2–8].

In James et al. [2] recently proposed a measure based on dependency constraints, denoted I_{dep} , which quantifies the unique information conveyed by a single predictor. In the case of two predictors, this is sufficient to obtain all four terms of the full PID; for higher order systems some terms remain indeterminate. For larger systems, there are a number of noted concerns with the PID

approach. For three predictors, it has been shown that the proposed axioms and lattice cannot result in a non-negative decomposition. A specific counter example has been demonstrated [9], and an alternative view based on an intuitive interpretation of the relationship between PID and secret sharing schemes also demonstrated the same issue [10]. Despite this, systems with two predictors can still be of theoretical and practical interest, so we focus here on that specific case [2,5,11].

The I_{dep} measure was derived and presented for discrete systems [2]. However, there are many applications in which continuous variables might be subjected to the same analysis, and the PID approach has been considered for Gaussian systems [6,12]. I_{dep} is derived from considering dependency constraints imposed within a lattice of maximum entropy probability models. Here, we apply the same logic to derive I_{dep} in the case of continuous Gaussian variables. In this case, the maximum entropy probability models are Gaussian graphical models [13–15], also termed covariance selection models [16]. We provide closed form expressions for the two predictor I_{dep} PID, for both univariate and multivariate continuous Gaussian predictors and target. Code implementing these measures is provided as the Supplementary Materials.

First, we provide a brief review of the PID (Section 1.1) and the discrete I_{dep} measure (Section 1.2). In Section 2, we derive I_{dep} for univariate Gaussian variables, and in Section 3 extend to multivariate Gaussian variables.

1.1. The Partial Information Decomposition

The partial information decomposition was introduced in [1] as a method to decompose mutual information in a multivariate system in terms of redundancies and synergies within and between subsets of predictors. Formally, the PID is developed as the Mobiüs inversion of a shared information measure over the lattice of antichains of predictor variables. We refer the reader to [1] for the full details.

In this manuscript, we focus on the case of two predictors, X_0 , X_1 , and a target Y. In this case, the mutual information $I(X_0, X_1; Y)$ is decomposed into four terms:

- red, the information about Y that is shared, common or redundant between X_0 and X_1 ,
- unq0, the information about Y that is available only from *X*₀,
- unq1, the information about Y that is available only from *X*₁,
- syn, the information about Y that is only available when X_0 and X_1 are observed together.

These terms satisfy the following intuitive relationships:

$$I(X_0, X_1; Y) = \operatorname{red} + \operatorname{unq0} + \operatorname{unq1} + \operatorname{syn}$$
(1)

$$I(X_0; Y) = \operatorname{red} + \operatorname{unq} 0 \tag{2}$$

$$I(X_1;Y) = \text{red} + \text{unq1} \tag{3}$$

Given the existence of these three constraints in terms of classical mutual information values, there is only one degree of freedom left to specify the bivariate PID. With any of the four terms quantified, the remaining three can be easily calculated. The initial formulation of [1] was based on quantifying redundancy, and deriving the other quantities, but others have focussed on quantifying unique information or synergy directly.

1.1.1. The Partial Information Decomposition for Gaussian Variables

The original definition of the PID and most of the subsequent work referenced above focussed on discrete variables. However, there are many applications where continuous-valued Gaussian variables are interesting subjects for information theoretic analysis. For example, simplified model systems [17,18] or empirical data analysis [19,20]. In [12], all discrete PID measures available at the time were considered and their principles applied to multivariate Gaussian systems, where one univariate component of the Gaussian is denoted the target. It was shown [12] that for a univariate target, if red, unq0 and unq1 depend only on the predictor-target marginal (X_0 , Y), (X_1 , Y) distributions, then there is a unique non-negative PID for which the redundancy is given by the minimum mutual information (MMI). Several proposed discrete PID measures fall into this class [1,3–5,21], so for Gaussian systems these approaches are all equivalent and equal to the MMI PID. The full bivariate MMI PID is defined as follows:

$$\operatorname{red} = \min\{I(X_0; Y), I(X_1; Y)\}$$
(4)

$$unq0 = \begin{cases} 0, & \text{if } I(X_0; Y) < I(X_1; Y) \\ I(X_0; Y) - I(X_1; Y), & \text{otherwise} \end{cases}$$
(5)

unq1 =
$$\begin{cases} 0, & \text{if } I(X_1; Y) < I(X_0; Y) \\ I(X_1; Y) - I(X_0; Y), & \text{otherwise} \end{cases}$$
(6)

$$syn = \begin{cases} I(X_0, X_1; Y) - I(X_1; Y), & \text{if } I(X_0; Y) < I(X_1; Y) \\ I(X_0, X_1; Y) - I(X_0; Y), & \text{otherwise} \end{cases}$$
(7)

The MMI PID takes the redundancy component to be the minimum of the two mutual informations between the target and the predictors. Hence, one of the unique information components will always be zero. The MMI PID has been used also with Gaussian systems involving multivariate time series [12,22]; for an alternative approach, see [23].

Another recently proposed measure, I_{ccs} , exploits the additivity of local or pointwise entropy to calculate the *common change in surprisal* provided by multiple predictors. By considering the signs of the local predictor-target information values, and the sign of the set theoretic intersection provided by local co-information, it is possible to sum up only pointwise terms that unambiguously correspond to redundant or overlapping information. I_{ccs} is calculated on the maximum entropy distribution subject to pairwise marginal constraints (i.e., including the (X_0, X_1) distribution). It therefore does not satisfy the Barrett conditions, and is not equivalent to the MMI PID. It also does not provide a non-negative PID, even in the two predictor case, since it is possible for one predictor to provide a unique negative contribution at the pointwise level (since pointwise mutual information is not non-negative). While there is no closed form expression for I_{ccs} for Gaussians, it has been implemented using Monte Carlo methods [6]. I_{dep} [2] is also not invariant to changes in the predictor-predictor marginal distribution, and therefore does not reduce to MMI in the Gaussian case either.

1.2. Unique Information via Dependency Constraints

In [2] a method is proposed to quantify the unique information conveyed by a predictor variable. They start from a lattice of maximum entropy models subject to marginal constraints, where the lattice structure comes from the hierarchy of marginal constraints. This lattice is illustrated in Figure 1.

For example, U_1 represents the maximum entropy distribution, having probability density function (p.d.f.) $g(x_0, x_1, y)$, under the constraints that the univariate marginals match exactly the univariate marginals of the original distribution, which has p.d.f. $f(x_0, x_1, y)$. That is: $g(x_0) = f(x_0)$, $g(x_1) = f(x_1)$, g(y) = f(y). U_2 represents the maximum entropy distribution subject to the constraints $g(x_0, x_1) = f(x_0, x_1)$, g(y) = f(y). U_5 represents the maximum entropy distribution subject to the constraints $g(x_0, x_1) = f(x_0, x_1)$, $g(x_0, y) = f(x_0, y)$, and so on. The lattice structure arises from the higher order constraints enforcing corresponding lower order constraints, so that for example imposing a bivariate marginal constraint such as $g(x_0, y) = f(x_0, y)$ means also that the lower order constraints $g(x_0) = f(x_0)$ and g(y) = f(y) also hold. Note that in [2] there is an additional model U_9 for the full distribution including third order interactions. We focus here on Gaussian systems which are fully determined by their first and second order moments, and so do not feature any triple-wise interactions. Therefore, model U_9 does not appear in our lattice for Gaussian systems.

For the sake of brevity in the sequel, rather than speaking of imposing a constraint of the form $g(x_0, y) = f(x_0, y)$, for example, we will speak of 'adding the constraint X_0Y' .

The colored edges correspond to adding a pairwise marginal constraint. Blue edges represent the constraint X_0X_1 , i.e., preserving the pairwise dependency between X_0 and X_1 . Green and red labelled edges correspond to the addition of the X_0Y and the X_1Y dependencies respectively. For each model $U_1 \dots U_8$ we calculate the mutual information between predictors and target under that model, $I_{U_i}(X_0, X_1; Y)$. The unique information in X_0 is then obtained as the minimum change in I_{U_i} along all the green edges due to the addition of the X_0Y constraint to the model below. Similarly, the unique information in X_1 can be obtained as the minimum change in I_{U_i} along all the red edges due to the addition of the X_1Y constraint to the model below. Therefore, for example, the edge value d is equal to $I_{U_5} - I_{U_2}$ and the edge value f is equal to $I_{U_6} - I_{U_2}$.

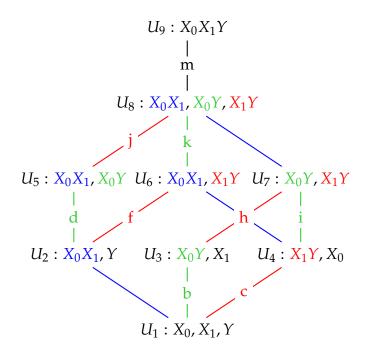


Figure 1. A dependency lattice of models (based on [2]). Edges coloured green (b, d, i, k) correspond to adding the constraint X_0Y to the model immediately below. Edges coloured red (c, f, h, j) correspond to adding the constraint X_1Y to the model immediately below.

If the edge labels in Figure 1 represent the change in mutual information along that edge, then the I_{dep} PID is given by:

$$unq0 = \min\{b, d, i, k\},\tag{8}$$

$$\operatorname{red} = I(X_0; Y) - \operatorname{unq0},\tag{9}$$

$$unq1 = I(X_1; Y) - red,$$
 (10)

$$syn = I(X_0, X_1; Y) - I(X_1; Y) - unq0$$
(11)

or:

$$unq1 = \min\{c, f, h, j\},\tag{12}$$

$$\operatorname{red} = I(X_0; Y) - \operatorname{unq1},\tag{13}$$

$$unq0 = I(X_0; Y) - red, \tag{14}$$

$$syn = I(X_0, X_1; Y) - I(X_0; Y) - unq1$$
(15)

(10)

It is shown in [2] that this approach is consistent; the same PID results from either of the two forms above. They also show that the resulting PID satisfies the core axioms of symmetry, self-redundancy, monotonicity, local positivity and the identity axiom [2].

2. An I_{dep} PID for Univariate Gaussian Predictors and Target

Since we will find in Section 2.2 that the required maximum entropy distributions that are described in Section 1.2 are Gaussian graphical models we begin with a brief discussion of such models.

2.1. Gaussian Graphical Models

The independence graph for a probability distribution on three univariate random variables, X_0, X_1, Y has three vertices and three possible edges, as described in Table 1. Let $\mathbf{Z} = \begin{bmatrix} X_0 & X_1 & Y \end{bmatrix}^T$.

Graphical models represent the conditional independences present in a probability distribution, as described in Table 1.

Table 1. Graphical models and independences for the probability distribution of **Z**. The vertices for random variables, X_0 , X_1 , Y are denoted by 0, 1, 2, respectively. Edges are denoted by pairs of vertices, such as (1, 2). In the column of independences, for example, $1 \perp 2 \mid 0$ indicates that X_1 and Y are conditionally independent given X_0 (based on [13], p. 61).

Model	Independences	Edge Set	Diagram	Description
<i>G</i> ₁	1.1.2 0, 0.1.2 1	{}	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	Mutual independence
	1.1.2 0			
G ₂	2.11.0 1, 2.11.1 0	$\{(0,1)\}$	$ \begin{array}{c} \bullet \\ 0 \\ 1 \end{array} $	Independent subsets
G_3	10 2, 12 0	$\{(0,2)\}$		Independent subsets
G_4	01 2, 02 1	$\{(1,2)\}$	$ \begin{array}{c} 0 \\ 0 \\ 1 \\ 2 \end{array} $	Independent subsets
G_5	1.1.1.2 0	$\{(0,1),(0,2)\}$	$ \begin{array}{c} 1 \\ \bullet \\ 0 \\ 1 \end{array} $	One independence
G_6	0⊥⊥2 1	$\{(0,1),(1,2)\}$		One independence
G ₇	0.1112	$\{(0,2),(1,2)\}$	$ \begin{array}{c} 1 \\ \bullet \\ 0 \\ 1 \end{array} $	One independence
G ₈	None	$\{(0,1),(0,2),(1,2)\}$		Complete interdependence

Suppose that **Z** has a multivariate Gaussian distribution with mean vector μ_Z , positive definite covariance matrix Σ_Z and p.d.f. $f(x_0, x_1, y)$. There is no loss of generality in assuming that each component of **Z** has mean zero and variance equal to 1 [12]. If we let the covariance (correlation) between X_0 and X_1 be p, between X_0 and Y be q and between X_1 and Y be r, then the covariance (correlation) matrix for **Z** is

$$\Sigma_Z = \begin{bmatrix} 1 & p & q \\ p & 1 & r \\ q & r & 1 \end{bmatrix}$$
(16)

and we require that |p|, |q|, |r| are each less than 1, and to ensure positive definiteness we require also that $|\Sigma_Z| > 0$.

Conditional independences are specified by setting certain off-diagonal entries to zero in the inverse covariance matrix, or concentration matrix, $K = \Sigma^{-1}$ ([13], p. 164). Given our assumptions about the covariance matrix of *Z*, this concentration matrix is

$$K = \frac{1}{|\Sigma_Z|} \begin{bmatrix} 1 - r^2 & qr - p & pr - q \\ qr - p & 1 - q^2 & pq - r \\ pr - q & pq - r & 1 - p^2 \end{bmatrix},$$
(17)

where $|\Sigma_Z| = 1 - p^2 - q^2 - r^2 + 2pqr$.

We now illustrate using these Gaussian graphical models how conditional independence constraints also impose constraints on marginal distributions of the type required, and we use the Gaussian graphical models G_8 and G_6 to do so.

Since **Z** is multivariate Gaussian and has a zero mean vector, the distribution of **Z** is specified via its covariance matrix Σ_Z . Hence, fitting any of the Gaussian graphical models $G_1 \dots G_8$ involves estimating the relevant covariance matrix by taking the conditional independence constraints into account. Let $\hat{\Sigma}_i$ and \hat{K}_i be the covariance and concentration matrices of the fitted model G_i , (i = 1...8).

We begin with the saturated model G_8 which has a fully connected graph and no constraints of conditional independence. Therefore, there is no need to set any entries of the concentration matrix K to zero, and so $\hat{\Sigma}_8 = \Sigma_Z$. That is: model G_8 is equal to the given model for **Z**.

Now consider model G_6 . In this model there is no edge between X_0 and Y and so X_0 and Y are conditionally independent given X_1 . This conditional independence is enforced by ensuring that the [1, 3] and [3, 1] entries in \hat{K}_6 are zero. The other elements in \hat{K}_6 remain to be determined. Therefore \hat{K}_6 has the form

$$\hat{K}_{6} = \begin{bmatrix} \hat{k}_{00} & \hat{k}_{01} & 0\\ \hat{k}_{01} & \hat{k}_{11} & \hat{k}_{12}\\ 0 & \hat{k}_{12} & \hat{k}_{22} \end{bmatrix} .$$
(18)

Given the form of \hat{K}_6 , $\hat{\Sigma}_6$ has the form

$$\hat{\Sigma}_{6} = \begin{bmatrix} 1 & p & \hat{\sigma}_{02} \\ p & 1 & r \\ \hat{\sigma}_{02} & r & 1 \end{bmatrix},$$
(19)

where $\hat{\sigma}_{02}$ is to be determined. Notice that only the [1, 3] and [3, 1] entries in $\hat{\Sigma}_6$ have been changed from the given covariance matrix Σ_Z , since the [1, 3] and [3, 1] entries of \hat{K}_6 have been set to zero. An exact solution is possible. The inverse of $\hat{\Sigma}_6$ is

$$\hat{K}_{6} = \hat{\Sigma}_{6}^{-1} = \frac{1}{|\hat{\Sigma}_{6}|} \begin{bmatrix} 1 - r^{2} & \hat{\sigma}_{02}r - p & pr - \hat{\sigma}_{02} \\ \hat{\sigma}_{02}r - p & 1 - \hat{\sigma}_{02}^{2} & p\hat{\sigma}_{02} - r \\ pr - \hat{\sigma}_{02} & p\hat{\sigma}_{02} - r & 1 - p^{2} \end{bmatrix}$$
(20)

Since the [1, 3] entry in \hat{K}_6 must be zero, we obtain the solution that $\hat{\sigma}_{02} = pr$, and so the estimated covariance matrix for model G_6 is

$$\hat{\Sigma}_{6} = \begin{bmatrix} 1 & p & pr \\ p & 1 & r \\ pr & r & 1 \end{bmatrix}.$$
(21)

The estimated covariance matrices for the other models can be obtained exactly using a similar argument.

Model G_6 contains the marginal distributions of X_0 , X_1 , Y, (X_0, X_1) and (X_0, Y) . It is important to note that these marginal distributions are exactly the same as in the given multivariate Gaussian

distribution for **Z**, which has covariance matrix Σ_Z . To see this we use a standard result on the marginal distribution of a sub-vector of a multivariate Gaussian distribution [24], p. 63.

The covariance matrix of the marginal distribution (X_0, X_1) is equal to the upper-left 2 by 2 sub-matrix of $\hat{\Sigma}_6$, which is also equal to the same sub-matrix in Σ_Z in (16). This means that this marginal distribution in model G_6 is equal to the corresponding marginal distribution in the distribution of Z. The covariance matrix of the marginal distribution (X_0, Y) is equal to the lower-right 2 by 2 sub-matrix of $\hat{\Sigma}_6$, which is also equal to the same sub-matrix in Σ_Z in (16), and so the (X_0, Y) marginal distribution in model G_6 matches the corresponding marginal distribution in the distribution of Z. Using similar arguments, such equality is also true for the other marginal distributions in model G_6 .

Looking at (17), we see that setting to [1, 3] of *K* entry to zero gives q = pr. Therefore, simply imposing this conditional independence constraint also gives the required estimated covariance matrix $\hat{\Sigma}_{6}$.

It is generally true ([13], p.176) that applying the conditional independence constraints is sufficient and it also leads to the marginal distributions in the fitted model being exactly the same as the corresponding marginal distributions in the given distribution of **Z**. For example, in (19) we see that the only elements in $\hat{\Sigma}_6$ that are altered are the [1, 3] and [3, 1] entries and these entries corresponds exactly to the zero [1, 3] and [3, 1] entries in \hat{K}_6 . That is: the location of zeroes in \hat{K}_6 determines which entries in $\hat{\Sigma}_6$ will be changed; the remaining entries of $\hat{\Sigma}_Z$ are unaltered and therefore this fixes the required marginal distributions. Therefore, in Section 2.2, we will determine the required maximum entropy solutions by simply applying the necessary conditional independence constraints together with the other required constraints.

We may express the combination of the constraints on marginal distributions and the constraints imposed by conditional independences as follows [16]. For model G_k , the (i, j)th entry of $\hat{\Sigma}_Z$ is given by

$$\hat{\Sigma}_{Z}[i,j] = \Sigma_{Z}[i,j], \text{ for } i = j \text{ and } (i,j) \in E_{k},$$

where E_k is the edge set for model G_k (see Table 1). For model G_k , the conditional independences are imposed by setting the (i, j)th entry of \hat{K} to zero whenever $(i, j) \notin E_k$.

Before moving on to derive the maximum entropy distributions, we consider the conditional independence constraints in model G_3 . In model G_3 we see from Table 1 that this model has no edge between X_0 and X_1 and none between X_1 and Y. Hence, X_0 and X_1 are conditionally independent given Y and also X_1 and Y are conditionally independent given X_0 . Hence, in K in (17) we set the [1, 2] and [2, 3] (and the [2, 1] and [3, 2]) entries to zero to enforce these conditional independences. That is: p = qr and r = pq. Taken together these equations give that p = 0 and r = 0, and so the estimated covariance matrix for model G_3 is

$$\hat{\Sigma}_{3} = \begin{bmatrix} 1 & 0 & q \\ 0 & 1 & 0 \\ q & 0 & 1 \end{bmatrix}.$$
(22)

We also note that model U_3 in Figure 1 also possesses the same conditional independences as G_3 . This is true for all of the maximum entropy models U_i , and so when finding the nature of these models in the next section we apply in each case the conditional independence constraints satisfied by the graphical model G_i .

2.2. Maximum Entropy Distributions

We are given the distribution of *Z* which is multivariate Gaussian with zero mean vector and covariance matrix Σ_Z in (16), and has p.d.f. $f(\mathbf{z}) \equiv f(x_0, x_1, y)$. For each of the models $U_1 \dots U_8$, we will determine the p.d.f. of the maximum entropy solution $g(\mathbf{z}) \equiv g(x_0, x_1, y)$ subject to the constraints

$$\int_{\mathbb{R}^3} \mathbf{z} g(\mathbf{z}) \, d\mathbf{z} = \mathbf{0}, \quad \int_{\mathbb{R}^3} g(\mathbf{z}) \, d\mathbf{z} = 1, \quad g(\mathbf{z}) > 0, \tag{23}$$

and the separate constraint for model U_i

$$\int_{\mathbb{R}^3} \mathbf{z} \mathbf{z}^T g(\mathbf{z}) \, d\mathbf{z} = \hat{\Sigma}_i, \tag{24}$$

as well as the conditional independence constraints given in Table 2.

Table 2. Conditional independence constraints satisfied by the Gaussian graphical models $G_1 \dots G_7$ that are applied when determining the maximum entropy models $U_1 \dots U_7$.

$U_1: p = qr, q = pr, r = pq$		
$U_2: q = pr, r = pq$	$U_3: p = qr, r = pq$	$U_4: p = qr, q = pr$
$U_5: r = pq$	$U_6: q = pr$	$U_7: p = qr$

We begin with model U_8 . As shown in the previous section, the estimated covariance matrix for model U_8 , $\hat{\Sigma}_8$, is equal to the covariance matrix of **Z**, Σ_Z . By a well-known result [25], the solution is that U_8 is multivariate Gaussian with mean vector zero and covariance matrix, Σ_Z . That is: U_8 is equal to the given distribution of **Z**.

For model U_5 , the conditional independence constraint is r = pq and so

$$\hat{\Sigma}_5 = \begin{bmatrix} 1 & p & q \\ p & 1 & pq \\ q & pq & 1 \end{bmatrix}.$$
(25)

Hence, using a similar argument to that for U_8 , the maximum entropy solution for model U_5 is multivariate Gaussian with zero mean vector and covariance matrix $\hat{\Sigma}_5$, and so is equal to the model G_5 .

In model U_3 , the conditional independence constraints are p = qr, r = pq and so p = 0 and r = 0. Therefore,

$$\hat{\Sigma}_3 = \begin{bmatrix} 1 & 0 & q \\ 0 & 1 & 0 \\ q & 0 & 1 \end{bmatrix}$$
(26)

and the maximum entropy solution for U_3 is multivariate Gaussian with zero mean vector and covariance matrix $\hat{\Sigma}_3$, and so is equal to G_3 . The derivations for the other maximum entropy models are similar, and we state the results in Proposition 1.

Proposition 1. The distributions of maximum entropy, $U_1 \dots U_8$, subject to the constraints (23)–(24) and the conditional independence constraints in Table 2, are trivariate Gaussian graphical models $G_1 \dots G_8$ having mean vector **0** and with the covariance matrices $\hat{\Sigma}_i$, $(i = 1, \dots, 8)$, given above in Table 3.

The estimated covariance matrices in Table 3 were inverted to give the corresponding concentration matrices, which are also given in Table 3. They indicate by the location of the zeroes that the conditional independences have been appropriately applied in the derivation of the results in Proposition 1.

It is important to check that the relevant bivariate and univariate marginal distributions are the same in all of the models in which a particular constraint has been added For example, the X_0X_1 constraint is present in models U_2 , U_5 , U_6 , U_8 . The marginal bivariate X_0X_1 distribution has zero mean vector and so is determined by the upper-left 2 by 2 sub-matrix of the estimated covariance matrices,

 $\hat{\Sigma}_i$ ([24], p. 63). Inspection of Table 3 shows that this sub-matrix is equal to $\begin{bmatrix} 1 & p \\ p & 1 \end{bmatrix}$ in all four models. Thus, the bivariate distribution of (X_0, X_1) is the same in all four models in which this dependency constraint is fitted. It is also the same as in the original distribution, which has covariance matrix Σ_Z .

in (16). Further examination of Table 3 shows equivalent results for the (X_0, Y) and (X_1, Y) bivariate

marginal distributions. The univariate term Y is present in all eight models. The univariate distribution of Y has mean zero and so is determined by the [3, 3] element of the estimated covariance matrices $\hat{\Sigma}_i$ ([24], p. 63). Looking at the $\hat{\Sigma}_i$ column, we see that the variance of Y is equal to 1 in all eight models, and so the marginal distribution of Y is the same in all eight models. In particular, this is true in the original distribution, which has covariance matrix Σ_Z in (16).

Table 3. Covariance matrices, with corresponding concentration matrices, for the Gaussian graphical models which were derived as maximum entropy probability models in Proposition 1.

Model	$\hat{\Sigma}_i$	\hat{K}_i
$U_1: X_0, X_1, Y$	$\begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}$	$\begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}$
$U_2: X_0 X_1, Y$	$\begin{bmatrix} 1 & p & 0 \\ p & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}$ $\begin{bmatrix} 1 & 0 & q \\ 0 & 1 & 0 \\ q & 0 & 1 \\ \end{bmatrix}$	$\frac{1}{1-p^2} \begin{bmatrix} 1 & -p & 0\\ -p & 1 & 0\\ 0 & 0 & 1-p^2 \end{bmatrix}$
$U_3: X_0Y, X_1$	$ \begin{bmatrix} 1 & 0 & q \\ 0 & 1 & 0 \\ - & 0 & 1 \end{bmatrix} $	$\frac{1}{1-q^2} \begin{vmatrix} 1 & 0 & -q \\ 0 & 1-q^2 & 0 \\ 0 & 0 & 1 \end{vmatrix}$
$U_4: X_1Y, X_0$	$\begin{bmatrix} q & 0 & 1 \\ 1 & 0 & 0 \\ 0 & 1 & r \\ 0 & r & 1 \end{bmatrix}$	$\frac{1}{1-p^2} \begin{bmatrix} 1 & -p & 0 \\ -p & 1 & 0 \\ 0 & 0 & 1-p^2 \end{bmatrix}$ $\frac{1}{1-q^2} \begin{bmatrix} 1 & 0 & -q \\ 0 & 1-q^2 & 0 \\ -q & 0 & 1 \end{bmatrix}$ $\frac{1}{1-r^2} \begin{bmatrix} 1-r^2 & 0 & 0 \\ 0 & 1 & -r \\ 0 & -r & 1 \end{bmatrix}$
$U_5: X_0 X_1, X_0 Y$	$\begin{bmatrix} 1 & p & q \\ p & 1 & pq \\ q & pq & 1 \end{bmatrix}$	$\frac{1}{(1-p^2)(1-q^2)} \begin{bmatrix} 1-p^2q^2 & (q^2-1)p & (p^2-1)q \\ (q^2-1)p & 1-q^2 & 0 \\ (p^2-1)q & 0 & 1-p^2 \end{bmatrix}$
$U_6: X_0 X_1, X_1 Y$	$\begin{bmatrix} 1 & p & pr \\ p & 1 & r \\ pr & r & 1 \end{bmatrix}$	$\frac{1}{(1-p^2)(1-r^2)} \begin{bmatrix} 1-r^2 & (r^2-1)p & 0\\ (r^2-1)p & 1-p^2r^2 & (p^2-1)r\\ 0 & (p^2-1)r & 1-p^2 \end{bmatrix}$
$U_7: X_0Y, X_1Y$	$\begin{bmatrix} 1 & qr & q \\ qr & 1 & r \\ q & r & 1 \end{bmatrix}$	$\frac{1}{(1-q^2)(1-r^2)} \begin{bmatrix} 1-r^2 & 0 & (r^2-1)q \\ 0 & 1-q^2 & (q^2-1)r \\ (r^2-1)q & (q^2-1)r & 1-q^2r^2 \end{bmatrix}$
$U_8: X_0X_1, X_0Y, X_1Y$	$\begin{bmatrix} 1 & p & q \\ p & 1 & r \\ q & r & 1 \end{bmatrix}$	$\frac{1}{ \Sigma_Z } \begin{bmatrix} 1 - r^2 & qr - p & pr - q \\ qr - p & 1 - q^2 & pq - r \\ pr - q & pq - r & 1 - p^2 \end{bmatrix}$

2.3. Mutual Information

Some required results involving mutual information will now be stated. They will be used to find expressions for the total mutual information of each model and also in constructing the I_{dep} and I_{mmi} PIDs.

$$I(X_0, X_1; Y) = \frac{1}{2} \log \left(\frac{1 - p^2}{1 - p^2 - q^2 - r^2 + 2pqr} \right),$$
(27)

$$I(X_0; Y) = \frac{1}{2} \log\left(\frac{1}{1-q^2}\right),$$
(28)

$$I(X_1; Y) = \frac{1}{2} \log\left(\frac{1}{1 - r^2}\right),$$
(29)

$$I(X_0; Y|X_1) = \frac{1}{2} \log \left(\frac{(1-p^2)(1-r^2)}{1-p^2-q^2-r^2+2pqr} \right),$$
(30)

$$I(X_1; Y|X_0) = \frac{1}{2} \log \left(\frac{(1-p^2)(1-q^2)}{1-p^2-q^2-r^2+2pqr} \right).$$
(31)

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Application of (27) with the covariance matrices given in Table 3 gives the following expressions for the total mutual information $I(X_0, X_1; Y)$ for the maximum entropy models derived in Proposition 1.

2.4. The Idep PID for Univariate Gaussian Predictors and Target

The I_{dep} PID for Gaussian predictors and target will now be constructed; for details, see (8)–(11) in Section 1.2.

Using the results in Table 4 together with the dependency lattice in Figure 2, we may write down expressions for all the required edge values, and they are given in Table 5.

Table 4. Expressions for the predictors-target mutual information for the eight models in the dependency lattice of Figure 2, as described in Table 3.

$U_8: \frac{1}{2} \log \left(\frac{1-p^2}{1-p^2-q^2-r^2+2pqr} \right)$	$U_4: I(X_1; Y)$
$U_7: \frac{1}{2} \log \left(\frac{1-q^2r^2}{(1-q^2)(1-r^2)} \right)$	$U_3:I(X_0;Y)$
$U_6: I(X_1; Y) = \frac{1}{2} \log\left(\frac{1}{1-r^2}\right)$	$U_2: 0$
$U_5: I(X_0; Y) = \frac{1}{2} \log\left(\frac{1}{1-q^2}\right)$	$U_1:0$
$U_{8}: X_{0}X_{1}, X_{0}X_{1}, X_{0}X_{1}, X_{0}X_{1}, X_{0}X_{1}, X_{0}Y = U_{5}: X_{0}X_{1}, X_{0}Y = U_{6}: X_{0}X_{1}, X_{1}X_{1}$ $U_{5}: X_{0}X_{1}, X_{0}Y = U_{6}: X_{0}X_{1}, X_{1}X_{1}X_{1}X_{1}X_{1}X_{1}X_{1}X_{1}$	$X_{1}Y U_{7}: X_{0}Y, X_{1}Y$ $ $ $h \qquad i$ $ $ $X_{1} \qquad U_{4}: X_{1}Y, X_{0}$ c

Figure 2. A dependency lattice of models (based on [2]). Edges coloured green (b, d, i, k) correspond to adding the term X_0Y to the model immediately below. Edges coloured red (c, f, h, j) correspond to adding the term X_1Y to the model immediately below. The two relevant sub-lattices are shown here.

Table 5. Expression for the edge values in the dependency lattice in Figure 2 that are used to determine the unique informations.

$$b = I(X_0; Y) = \frac{1}{2} \log\left(\frac{1}{1-q^2}\right) \qquad c = I(X_1; Y) = \frac{1}{2} \log\left(\frac{1}{1-r^2}\right)$$
$$d = I(X_0; Y) \qquad f = I(X_1; Y)$$
$$i = \frac{1}{2} \log\left(\frac{1-q^2r^2}{(1-q^2)(1-r^2)}\right) - I(X_1; Y) \qquad h = \frac{1}{2} \log\left(\frac{1-q^2r^2}{(1-q^2)(1-r^2)}\right) - I(X_0; Y)$$
$$k = \frac{1}{2} \log\left(\frac{1-p^2}{(1-p^2-q^2-r^2+2pqr)}\right) - I(X_1; Y) \qquad j = \frac{1}{2} \log\left(\frac{1-p^2}{(1-p^2-q^2-r^2+2pqr)}\right) - I(X_0; Y)$$

By making use of the edge values given in Table 5 together with (8)–(11) from Section 1.2 the I_{dep} PID can be constructed. We now state some results for the I_{dep} PID for univariate Gaussian predictors, X_0 , X_1 , and target, Y, with proofs given in Appendix A.

Proposition 2. For two univariate Gaussian predictors, X_0 , X_1 , and one univariate Gaussian target, Y, the I_{dep} PID, defined in Table 5, and (8)–(11) in Section 1.2, has the following properties.

- (a) The I_{dep} PID possesses consistency as well as the core axioms of non-negativity, self-redundancy, monotonicity, symmetry and identity.
- (b) When unq0 is equal to b or d, the the redundancy component is zero.
- (c) When unq0 is equal to *i*, the redundancy and both unique informations are constant with respect to the correlation between the two predictors.
- (*d*) When the correlations between each predictor and the target are both non-zero, then unq0 is equal to either *i* or to *k*.
- (e) When unq0 is equal to k, the synergy component is zero.
- (f) The redundancy component in the I_{mmi} PID is greater than or equal to the redundancy component in the I_{dep} PID with equality if, and only if, at least one of the following conditions holds: (i) either predictor and the target are independent; (ii) either predictor is conditionally independent of the target given the other predictor.
- (g) The synergy component in the I_{mmi} PID is greater than or equal to the synergy component in the I_{dep} PID with equality if, and only if, at least one of the following conditions holds: (i) either predictor and the target are independent; (ii) either predictor is conditionally independent of the target given the other predictor.
- (h) The I_{dep} and I_{mmi} PIDs are identical when either X_0 and Y are conditionally independent given X_1 or X_1 and Y are conditionally independent given X_0 , and in particular they are identical for models $U_1 \dots U_6$. In model U_7 the synergy component of I_{dep} is zero.

The I_{mmi} PID is defined in (4)–(7) in Section 1.1. We now consider examples of the I_{dep} PID as well as comparisons between the I_{mmi} and I_{dep} PIDs in the following subsections.

2.5. Some Examples

Example 1. We consider the I_{dev} PID when $q = corr(X_0, Y) = 0, r \neq 0, p \neq 0$.

When q = 0, we see from Table 4 that b = d = i = 0 and k > 0, so unq0 = 0, and since $I(X_0; Y) = 0$ the redundancy component is also zero. The unique information, unq1, and the synergy component, syn, are equal to

$$I(X_1; Y) = \frac{1}{2} \log \frac{1}{1 - r^2}, \quad I(X_0; Y | X_1) = \frac{1}{2} \log \left(\frac{(1 - p^2)(1 - r^2)}{1 - p^2 - r^2} \right),$$

respectively. The I_{mmi} PID is exactly the same as the I_{dep} PID.

Example 2. We consider the I_{dep} PID when $r = corr(X_1, Y) = 0, q \neq 0, p \neq 0$.

When r = 0, we see from Table 5 that

$$b = d = i = \frac{1}{2}\log\frac{1}{1 - q^2}$$

and also that $k > \{b, d, i\}$ because

$$(1-p^2)(1-q^2) > 1-p^2-q^2$$
,

since $p \neq 0, q \neq 0$. It follows that $unq0 = \frac{1}{2} \log \frac{1}{1-q^2}$, and that the synergy component is equal to

$$\frac{1}{2} \log \left(\frac{(1-p^2)(1-q^2)}{1-p^2-q^2} \right).$$

Since $I(X_1; Y) = 0$, from (29), the redundancy component is zero, as is unq1. The I_{mmi} PID is exactly the same as the I_{dep} PID.

Example 3. We consider the I_{dep} PID when $p = corr(X_0, X_1) = 0, q \neq 0, r \neq 0$.

Under the stated conditions, it is easy to show that b < i and i < k and so the minimum edge value is attained at *i*. Using the results in Table 5 and (29)–(31), we may write down the I_{dep} PID as follows.

$$unq0 = \frac{1}{2} \log \left(\frac{1 - q^2 r^2}{1 - q^2} \right)$$

$$unq1 = \frac{1}{2} \log \left(\frac{1 - q^2 r^2}{1 - r^2} \right)$$

$$red = \frac{1}{2} \log \left(\frac{1}{1 - q^2} \right) - \frac{1}{2} \log \left(\frac{1 - q^2 r^2}{1 - q^2} \right) = \frac{1}{2} \log \left(\frac{1}{1 - q^2 r^2} \right)$$

$$syn = I(X_0; Y | X_1) - unq0 = \frac{1}{2} \log \left(\frac{(1 - q^2)(1 - r^2)}{(1 - q^2 - r^2)(1 - q^2 r^2)} \right)$$

For this situation, the I_{mmi} PID takes two different forms, depending on whether or not |q| < |r|. Neither form is the same as the I_{dep} PID.

Example 4. Compare the I_{mmi} and I_{dep} PIDs when p = -0.2, q = 0.7 and r = -0.7.

The PIDs are given in the following table.

PID	unq0	unq1	red	syn
I _{dep}	0.2877	0.2877	0.1981	0.4504
I _{mmi}	0	0	0.4587	0.7380

There is a stark contrast between the two PIDs in this system. Since |q| = |r|, the I_{mmi} PID has two zero unique informations, whereas I_{dep} has equal values for the uniques but they are quite large. The I_{mmi} PID gives much larger values for the redundancy and synergy components than does the I_{dep} PID. In order to explore the differences between these PIDs, 50 random samples were generated from a multivariate normal distribution having correlations p = -0.2, q = 0.7, r = -0.7. The sample estimates of p, q, r were $\hat{p} = -0.1125$, $\hat{q} = 0.6492$, $\hat{r} = -0.6915$ and the sample PIDs are

PID	unq0	unq1	red	syn
I _{dep}	0.2324	0.3068	0.1623	0.4921
I _{mmi}	0	0.0744	0.3948	0.7245

We now apply tests of deviance in order to test model U_i within the saturated model U_8 . The null hypothesis being tested is that model U_i is true (see Appendix E). The results of applying tests of deviance ([13], p. 185), in which each of models $U_1 \dots U_7$ is tested against the saturated model U_8 , produced approximate p values that were close to zero ($p < 10^{-11}$) for all but model U_7 , which had a p value of 10^{-6} . This suggests that none of the models $U_1 \dots U_7$ provides an adequate fit to the data and so model U_8 provides the best description. The results of testing U_6 and U_7 within model U_8

gave strong evidence to suggest that the interaction terms X_0Y and X_0Y are required to describe the data, and that each term makes a significant contribution in addition to the presence of the other term. Therefore, one would expect to find fairly sizeable unique components in a PID, and so the I_{dep} PID seems to provide a more sensible answer in this example. One would also expect synergy to be present, and both PIDs have a large, positive synergy component.

Example 5. Prediction of grip strength

Some data concerning the prediction of grip strength from physical measurements was collected from 84 male students at Glasgow University. Let Y be the grip strength, X_0 be the bicep circumference and X_1 the forearm circumference. The following correlations between each pair of variables were calculated: corr $(X_1, Y) = 0.7168$, corr $(X_0, Y) = 0.6383$, corr $(X_0, X_1) = 0.8484$, and PIDs applied with the following results.

PID	unq0	unq1	red	syn
I _{dep}	0.0048	0.1476	0.3726	0
I _{mmi}	0	0.1427	0.3775	0.0048

The I_{dep} and I_{mmi} PIDs are very similar, and the curious fact that unq0 in I_{dep} is equal to the synergy in I_{mmi} is no accident, It is easy to show this connection theoretically by examining the results in (30)–(31) and Table 5; that is, the sum of unq0 and syn in the I_{dep} PID or the sum of unq1 and syn in the I_{dep} PID is equal to the synergy value in the I_{mmi} PID. This happens because the I_{mmi} PID must have a zero unique component.

These PIDs indicate that there is almost no synergy among the three variables, which makes sense because the value of $I(X_0; Y|X_1)$ is close to zero, and this suggests that X_0 and Y are conditionally independent given X_1 . On the other hand, $I(X_1; Y|X_0)$ is 0.1427 which suggests that X_1 and Y are not conditionally independent given X_0 , and so both terms X_0X_1 and X_0Y are of relevance in explaining the data, which is the case in model U_6 . This model has $I(X_0; Y|X_1) = 0$ and therefore no synergy and also a zero unique value in relation to X_0 . The results of applying tests of deviance ([13], p. 185), in which each of models $U_1 \dots U_7$ is tested within the saturated model U_8 , show that the approximate p values are close to zero ($p < 10^{-13}$) for all models except U_5 and U_6 . The p value for the model U_5 is 3×10^{-5} , while the p value for the test of U_6 against U_8 is approximately 0.45. Thus, there is strong evidence to reject all the models except model U_6 and this suggests that model U_6 provides a good fit to data, and this alternative viewpoint provides support for the form of both PIDs.

2.6. Graphical Illustrations

We present some graphical illustrations of the I_{dep} PID and compare it to the I_{mmi} PID; see Sections 1.1.1 and 1.2 for definitions of these PIDs.

Since q = r, both the I_{mmi} unique informations are zero in Figure 3a. The redundancy component is constant, while the synergy component decreases towards zero. In Figure 3b, we observe change-point behaviour of I_{dep} when p = 0.25. For p < 0.25 the unique components of I_{dep} are equal, constant and positive. The redundancy component is also constant and positive with a lower value than the corresponding component in the I_{mmi} PID. The synergy component decreases towards zero and reaches this value when p = 0.25. The I_{dep} synergy is lower that the corresponding I_{mmi} synergy for all values of p.

At p = 0.25, the synergy "switches off" in the I_{dep} PID, and stays "off" for larger values of p, and then the unique and redundancy components are free to change. In the range 0.25 , the redundancy increases and takes up all the mutual information when <math>p = 1, while the unique informations decrease towards zero. The I_{dep} and I_{mmi} profiles show different features in this case. The "regime switching" in the I_{dep} PID is interesting. As mentioned in Proposition 2, the minimum edge value occurs with unq0 = i or k. When unq0 = k the synergy must be equal to zero, whereas when

unq0 = *i* the synergy is positive and the values of the unique informations and the redundancy are constant. Regions of zero synergy in the I_{dep} PID are explored in Figure 5.

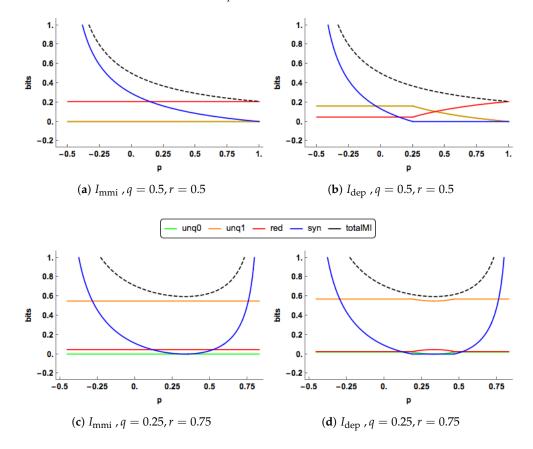


Figure 3. The I_{mmi} & I_{dep} PID components are plotted for a range of values of the correlation (*p*) between the two predictors. Two combinations of the correlations (*q*, *r*) between each predictor and the target are displayed. The total mutual information $I(X_0, X_1; Y)$ is also shown as a dashed black curve.

In Figure 4a,b, there are clear differences in the PID profiles between the two methods. The I_{dep} synergy component switches off at p = 0.5 and is zero thereafter. For p < 0.5, both the I_{dep} uniques are much larger than those of I_{mmi} , which are zero, and I_{mmi} has a larger redundancy component. For p > 0.5, the redundancy component in I_{dep} increases to take up all of the mutual information, while the unique information components decrease towards zero. In contrast to this, in the I_{mmi} PID the redundancy and unique components remain at their constant values while the synergy continues to decrease towards zero.

The PIDs are plotted for increasing values of q = r in Figure 4c,d when p = 0.25. The I_{mmi} and I_{dep} profiles are quite different. As q increases, the I_{mmi} uniques remain at zero, while the I_{dep} uniques rise gradually. Both the I_{mmi} redundancy and synergy profiles rise more quickly than their I_{dep} counterparts, probably because both their uniques are zero. In the I_{dep} PID, the synergy switches on at p = 0.5 and it is noticeable than all the I_{dep} components can change simultaneously as q increases.

One of the characteristics noticed in Figures 3 and 4 is the 'switching behaviour' of the I_{dep} PID in that there are kinks in the plots of the PIDs against the correlation between the predictors, *p*: the synergy component abruptly becomes equal to zero at certain values of *p*, and there are other values of *p* at which the synergy moves from being zero to being positive.

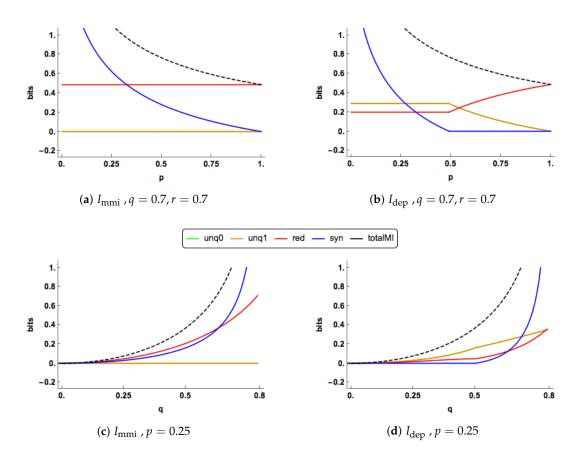


Figure 4. In (**a**,**b**), the $I_{mmi} \& I_{dep}$ PIDs are plotted for a range of values of the correlation (*p*) between the two predictors. One combination of the correlations (*q*, *r*) between each predictor and the target are displayed. In (**c**,**d**), the $I_{mmi} \& I_{dep}$ PID are plotted for a range of allowable values of *q*, where *q* is equal to *r*, for *p* = 0.25. The total mutual information $I(X_0, X_1; Y)$ is also shown as a dashed black curve.

In Proposition 2, it is explained for the I_{dep} PID that when both predictor-target correlations are non-zero the minimum edge value occurs at edge value *i* or *k*. When the synergy moves from zero to a positive value, this means that the minimum edge value has changed from being *k* to being equal to *i*, and vice-versa. For a given value of *p*, one can explore the regions in (*q*, *r*) space at which such transitions take place. In Figure 5, this region of zero synergy is shown, given four different values of *p*. The boundary of each of the regions is where the synergy component changes from positive synergy to zero synergy, or vice-versa.

The plots in Figure 5 show that synergy is non-zero (positive) whenever q and r are of opposite sign. When the predictor-predictor correlation, p, is 0.05 there is also positive synergy for large regions, defined by qr - p > 0, when q and r have the same sign. As p increases the regions of zero synergy change shape, initially increasing in area and then declining as p becomes quite large (p = 0.75). As p is increased further the zero-synergy bands narrow and so zero synergy will only be found when q and r are close to being equal.

When *p* is negative, the corresponding plots are identical to those with positive *p* but rotated counter clockwise by $\pi/2$ about the point q = 0, r = 0. Hence, synergy is present when *q* and *r* have the same sign. When *q* and *r* have opposite signs, there is also positive synergy for regions defined by qr - p < 0.

The case of p = 0 is of interest and there are no non-zero admissible values of q and r (where the covariance matrix is positive definite) where the synergy is equal to zero. Hence the system will have synergy in this case unless q = 0 or r = 0. This can be seen from the I_{dep} synergy expression in Example 3.



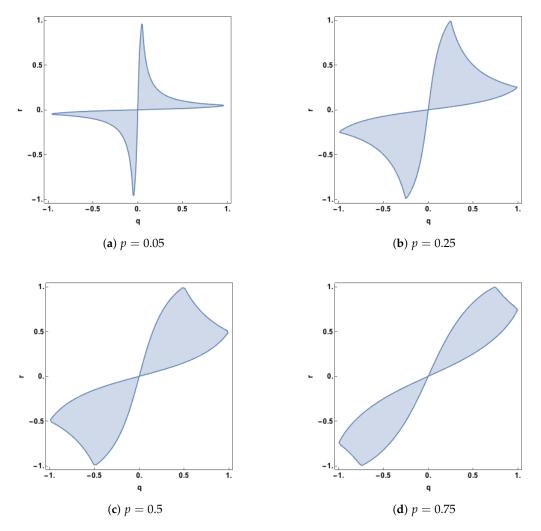


Figure 5. Regions in (q, r) space in which the synergy component in the I_{dep} PID is equal to zero, plotted for four different values of *p*. Also, the determinant of Σ_Z is positive.

3. Multivariate Continuous Predictors and Target

We now extend the results developed in Section 2 and consider the case where the three continuous variables X_0 , X_1 , Y become random vectors \mathbf{X}_0 , \mathbf{X}_1 , \mathbf{Y} , of dimensions n_0 , n_1 , n_2 , respectively, with mean vectors equal to a zero vector of lengths n_0 , n_1 , n_2 , respectively, and covariance matrices equal to an identity matrix of the respective sizes $n_0 \times n_0$, $n_1 \times n_1$, $n_2 \times n_2$. The fact that there is no loss of generality in making these assumptions will be explained in Section 3.4. We stack these random vectors into the random vector \mathbf{Z} , where \mathbf{Z} has dimension $n_0 + n_1 + n_2$, and assume that \mathbf{Z} has a multivariate Gaussian distribution with p.d.f. $f(\mathbf{x}_0, \mathbf{x}_1, \mathbf{y})$, mean vector $\mathbf{0}$ and covariance matrix given by

$$\Sigma_{Z} = \begin{bmatrix} I_{n_{0}} & P & Q \\ P^{T} & I_{n_{1}} & R \\ Q^{T} & R^{T} & I_{n_{2}} \end{bmatrix},$$
(32)

where the matrices *P*, *Q*, *R* are of size $n_0 \times n_1$, $n_0 \times n_2$, $n_1 \times n_2$, respectively, and are the cross-covariance (correlation) matrices between the three pairings of the three vectors **X**₀, **X**₁, **Y**, and so

$$\mathbb{E}(\mathbf{X}_0\mathbf{X}_1^T) = P, \quad \mathbb{E}(\mathbf{X}_0\mathbf{Y}^T) = Q, \quad \mathbb{E}(\mathbf{X}_1\mathbf{Y}^T) = R, \tag{33}$$

defined on \mathbb{R}^m , where $m = n_0 + n_1 + n_2$.

3.1. Properties of the Matrices P, Q, R, and the Inverse Matrix of Σ_Z

We require some matrix results, which will be proved in Appendix B.

Lemma 1. Suppose that a symmetric matrix M is partitioned as

$$M = \begin{bmatrix} M_{11} & M_{12} & M_{13} \\ M_{12}^T & M_{22} & M_{23} \\ M_{13}^T & M_{23}^T & M_{33} \end{bmatrix},$$

where the diagonal blocks M_{11} , M_{22} , M_{33} are symmetric and square. Then if M is positive definite these diagonal blocks are also positive definite, and so nonsingular.

Lemma 2. When the covariance matrix Σ_Z in (32) is positive definite then the following matrices are also positive definite, and hence nonsingular:

$$I_{n_1} - P^T P$$
, $I_{n_0} - PP^T$, $I_{n_2} - R^T R$, $I_{n_1} - RR^T$, $I_{n_2} - Q^T Q$, $I_{n_0} - QQ^T$.

Also, the determinant of each of these matrices is positive and bounded above by unity, and it is equal to unity if, and only if, the matrix involved is the zero matrix. Furthermore,

$$\begin{vmatrix} I_{n_0} & P \\ P^T & I_{n_1} \end{vmatrix} = |I_{n_1} - P^T P|.$$

With these results in place, we now present the inverse of Σ_Z , which is equal to the concentration matrix *K*. It was determined by solving simultaneous equations for block matrices and we omit the details. It is

$$K = \Sigma_Z^{-1} = \begin{bmatrix} A & U & V \\ U^T & B & W \\ V^T & W^T & C \end{bmatrix},$$
(34)

where

$$U = (I_{n_0} - QQ^T)^{-1}(QR^T - P)B$$
(35)

 $V = A(PR - Q)(I_{n_2} - R^T R)^{-1}$ (36)

$$W = (I_{n_1} - P^T P)^{-1} (P^T Q - R)C$$
(37)

$$A = \left[I_{n_0} - PP^T - (PR - Q)(I_{n_2} - R^T R)^{-1}(PR - Q)^T\right]^{-1}$$
(38)

$$B = \left[I_{n_1} - RR^T - (QR^T - P)^T (I_{n_0} - QQ^T)^{-1} (QR^T - P)\right]^{-1}$$
(39)

$$C = \left[I_{n_2} - Q^T Q - (P^T Q - R)^T (I_{n_1} - P^T P)^{-1} (P^T Q - R)\right]^{-1}$$
(40)

The various inverses used in (35)–(40) are valid for the following reasons. The matrix Σ_Z is positive definite, and so its inverse is also positive definite. By Lemma 1, *A*, *B*, *C* are positive definite and so invertible, which means in turn that their inverses are invertible. From Lemma 2, we have that the matrices $I_{n_1} - P^T P$, $I_{n_0} - QQ^T$, $I_{n_2} - R^T R$ are invertible. Therefore the sub-matrices in the inverse of Σ_Z in (35)–(40) are well-defined.

3.2. Block Gaussian Graphical Models

As in Section 2.1, we will consider graphical models to express the conditional independences in the probability distribution for Z, although each graph will still have three vertices, with each vertex representing one of the random vectors, X_0 , X_1 , Y. Each graph can be thought of as a block independence graph. This means that only dependences between pairs of vectors will be represented, while there will be no dependences among the variables within each of the three random vectors, since they are mutually independent. The models which express conditional dependences have the same format as in Table 1 in Section 2.1 and we use the same notation again here, the only difference being to express X_0 , X_1 , Y in a bold font. We term these models 'block graphical models' since we are treating each random vector as the block containing a number of mutually independent random variables. Here is an illustration of such a model:

The model in Figure 6 is the block version of model G_6 from Table 1 and denoted as X_0X_1, X_1Y . A product term, such as X_0X_1 , encapsulates correlations between each random variable in X_0 and each random variable in X_1 . For example, in Figure 6 there are 12 correlations between the elements of X_0 and X_1 , and 6 correlations between the elements of X_1 and Y. Using the block notation provides some simplicity, for otherwise one would be required to write expressions such as

$$\mathbf{X}_{1}\mathbf{Y} = X_{11}Y_{1} + X_{11}Y_{2} + X_{12}Y_{1} + X_{12}Y_{2} + X_{13}Y_{1} + X_{13}Y_{2}$$

for the set of constraints within each block interaction term. The block graphical models in the multivariate version of the dependency lattice are given in Figure 7. We now define the conditional independence constraints for the block versions of model $G_1 \dots G_8$ in Table 1 and determine some of their estimated covariance matrices. The block version of model G_8 has no conditional independences. Hence, no block zeroes are imposed on the concentration matrix K, and the estimated covariance matrix for this model is $\hat{\Sigma}_8 = \Sigma_Z$, which means that model G_8 is equal to the given distribution of **Z**.

Consider the block version of model G_7 , in Table 1. In G_7 , X_0 and X_1 are conditionally independent given **Y** and so we apply the constraint U = 0 in the concentration matrix K in (34). From (35), this block constraint is

$$(I_{n_0} - QQ^T)^{-1}(QR^T - P)B = 0.$$
(41)

Given the results stated at the end of Section 3.1, we can pre-multiply by $I_{n_0} - QQ^T$ and post-multiply by B^{-1} in (41) to obtain the required block constraint as $QR^T - P = 0$, that is $P = QR^T$. Hence, the estimated covariance matrix for block model G_7 is

$$\hat{\Sigma}_{7} = \begin{bmatrix} I_{n_{0}} & QR^{T} & Q \\ RQ^{T} & I_{n_{1}} & R \\ Q^{T} & R^{T} & I_{n_{2}} \end{bmatrix}.$$
(42)

Applying a similar argument to the expressions for *V* and *W* in (36) and (37), it can be shown that for the conditions for conditional independence between X_0 and Y given X_1 in model G_6 the block constraint is Q = PR, while for the conditional independence between X_1 and Y given X_0 in model G_5 the block constraint is $R = P^T Q$. Hence, the estimated covariance matrices for block models G_5 and G_6 are

$$\hat{\Sigma}_{5} = \begin{bmatrix} I_{n_{0}} & P & Q \\ P^{T} & I_{n_{1}} & P^{T}Q \\ Q^{T} & Q^{P} & I_{n_{2}} \end{bmatrix}, \text{ and } \hat{\Sigma}_{6} = \begin{bmatrix} I_{n_{0}} & P & PR \\ P^{T} & I_{n_{1}} & R \\ R^{T}P^{T} & R^{T} & I_{n_{2}} \end{bmatrix}.$$
(43)

In block model G_2 both of the conditional independences defining block models G_5 and G_6 are present. Therefore, the conditional independence constraints are $R = P^T Q$ and Q = PR. Combining them gives $R = P^T PR$, which may be written as $(I_{n_1} - P^T P)R = 0$. By Lemma 2, we may pre-multiply by the inverse of $I_{n_1} - P^T P$ to obtain R = 0, which in turn implies that Q = 0. Hence the conditional independence constraints for block model G_2 are Q = 0, R = 0 and so the estimated covariance matrix is

$$\hat{\Sigma}_{2} = \begin{bmatrix} I_{n_{0}} & P & 0\\ P^{T} & I_{n_{1}} & 0\\ 0 & 0 & I_{n_{2}} \end{bmatrix}.$$
(44)

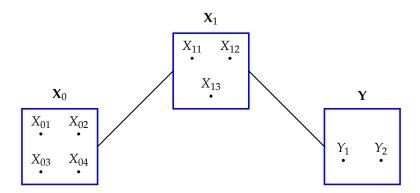


Figure 6. An illustration of a block graphical model for the random vectors X_0 , X_1 and Y. X_0 contains four random variables, while X_1 has three and Y has two. This model expresses the conditional independence of X_0 and Y given X_1 . In this model, the bivariate marginals X_0X_1 and X_1Y , as well as lower-order marginals, are fixed.

The other estimated covariance matrices can be derived in a similar fashion. As in Section 2.1, it is the case that applying the conditional independence constraints also ensures that the required marginal distributions in each of the block graphical models are equal to the corresponding marginal distributions in the given distribution of **Z**. We also note, in particular, that model M_2 in Figure 7 has the same conditional independences as those present in block graphical model G_2 , and this is true for each of the maximum entropy models M_i , and so when finding the form of these models in the next section we apply in each case the conditional independence constraints satisfied by the block graphical model G_i .

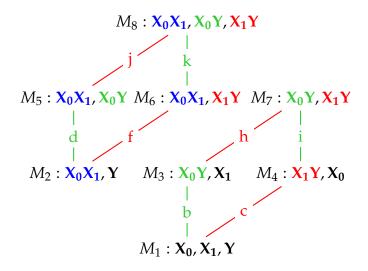


Figure 7. A dependency lattice of block graphical models. Edges coloured green (b, d, i, k) correspond to adding the set of constraints within X_0Y to the model immediately below. Edges coloured red (c, f, h, j) correspond to adding the set of constraints within X_1Y to the model immediately below. The two relevant sub-lattices are shown here.

3.3. Maximum Entropy Distributions

We are given the distribution of *Z* which is multivariate Gaussian with zero mean vector and covariance matrix Σ_Z in (32), and has p.d.f. $f(\mathbf{z}) \equiv f(x_0, x_1, y)$. For each of the models $M_1 \dots M_8$, we will determine the p.d.f. of the maximum entropy solution $g(\mathbf{z}) \equiv g(x_0, x_1, y)$ subject to the constraints

$$\int_{\mathbb{R}^3} \mathbf{z} g(\mathbf{z}) \, d\mathbf{z} = \mathbf{0}, \quad \int_{\mathbb{R}^3} g(\mathbf{z}) \, d\mathbf{z} = 1, \quad g(\mathbf{z}) > 0, \tag{45}$$

and the separate constraint for model M_i

$$\int_{\mathbb{R}^3} \mathbf{z} \mathbf{z}^T g(\mathbf{z}) \, d\mathbf{z} = \hat{\Sigma}_i,\tag{46}$$

as well as the conditional independence constraints given in Table 6.

Table 6. Conditional independence constraints satisfied by the block Gaussian graphical models $G_1 \dots G_8$ that are applied when determining the maximum entropy models $M_1 \dots M_8$.

$M_1: P = QR^T, Q = PR, R = P^TQ$		
$M_2: Q = PR, R = P^T Q$	$M_3: P = QR^T, R = P^T Q$	$M_4: P = QR^T, Q = PR$
$M_5: R = P^T Q$	$M_6: Q = PR$	$M_7: P = QR^T$

For model M_8 , the estimated covariance matrix $\hat{\Sigma}_8 = \Sigma_Z$ and so the maximum entropy distribution M_8 is equal to block graphical model G_8 , which is equal to the given distribution of **Z**. Similarly, the maximum entropy model M_7 is equal to the block graphical model G_7 , which is multivariate Gaussian with zero mean vector and covariance matrix $\hat{\Sigma}_7$, defined in (42), and so on. Hence we can state our results in Proposition 3.

Proposition 3. The distributions of maximum entropy, $M_1 \dots M_8$, subject to the constraints (45)–(46) and the conditional independence constraints in Table 6 are block Gaussian graphical models $G_1 \dots G_8$ having mean vector **0** and with the covariance matrices $\hat{\Sigma}_i$, $(i = 1, \dots 8)$, given below in Table 7.

Model	$\hat{\Sigma}_i$	Model	$\hat{\Sigma}_i$
$M_1: \mathbf{X}_0, \mathbf{X}_1, \mathbf{Y}$	$\begin{bmatrix} I_{n_0} & 0 & 0 \\ 0 & I_{n_1} & 0 \\ 0 & 0 & I_{n_2} \end{bmatrix}$	$M_5: \mathbf{X}_0 \mathbf{X}_1, \mathbf{X}_0 \mathbf{Y}$	$\begin{bmatrix} I_{n_0} & P & Q \\ P^T & I_{n_1} & P^T Q \\ Q^T & Q^T P & I_{n_2} \end{bmatrix}$
$M_2: \mathbf{X}_0 \mathbf{X}_1, \mathbf{Y}$	$\begin{bmatrix} I_{n_0} & P & 0 \\ P^T & I_{n_1} & 0 \\ 0 & 0 & I_{n_2} \end{bmatrix}$	$M_6: \mathbf{X}_0 \mathbf{X}_1, \mathbf{X}_1 \mathbf{Y}$	$\begin{bmatrix} I_{n_0} & P & PR \\ P^T & I_{n_1} & R \\ R^T P^T & R^T & I_{n_2} \end{bmatrix}$
$M_3: \mathbf{X}_0 \mathbf{Y}, \mathbf{X}_1$	$\begin{bmatrix} I_{n_0} & 0 & Q \\ 0 & I_{n_1} & 0 \\ Q^T & 0 & I_{n_2} \end{bmatrix}$	$M_7: \mathbf{X}_0 \mathbf{Y}, \mathbf{X}_1 \mathbf{Y}$	$\begin{bmatrix} I_{n_0} & QR^T & Q \\ RQ^T & I_{n_1} & R \\ Q^T & R^T & I_{n_2} \end{bmatrix}$
$M_4: \mathbf{X}_1 \mathbf{Y}, \mathbf{X}_0$	$\begin{bmatrix} I_{n_0} & 0 & 0 \\ 0 & I_{n_1} & R \\ 0 & R^T & I_{n_2} \end{bmatrix}$	$M_8: \mathbf{X}_0\mathbf{X}_1, \mathbf{X}_0\mathbf{Y}, \mathbf{X}_1\mathbf{Y}$	$\begin{bmatrix} I_{n_0} & P & Q \\ P^T & I_{n_1} & R \\ Q^T & R^T & I_{n_2} \end{bmatrix}$

Table 7. Covariance matrices for the Gaussian block graphical models in Proposition 3.

We can now check by inspecting the $\hat{\Sigma}_i$ entries in Table 7 that particular marginal distributions involving two blocks, such as $\mathbf{X}_1 \mathbf{Y}$, are the same in all of the models and also equal to the marginal distribution in the given distribution. For example, the block interaction term $\mathbf{X}_1 \mathbf{Y}$ is present in models M_4, M_6, M_7, M_8 . The distribution of $[\mathbf{X}_1 \ \mathbf{Y}]^T$ is multivariate normal with mean vector equal to a zero vector and covariance matrix given by the bottom right 2 by 2 block matrix in $\hat{\Sigma}_Z$ ([24], p. 63). For each of these four models, we can see by inspection that this covariance matrix is $\begin{bmatrix} I & R \\ R^T & I \end{bmatrix}$, and so the $(\mathbf{X}_1, \mathbf{Y})$ marginal distribution is the same in all of these four models in which this particular block interaction term has been fitted. Since M_8 is equal to the given distribution of \mathbf{Z} , it follows that this

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marginal distribution is the same as in the given distribution. Similar checks can be made regarding the other marginal distributions involving two blocks to find that a similar conclusion applies also to them. We can also check that the single-block terms, such as X_1 have the same distribution. This term has been fitted in all eight models. Since the mean vector of X_1 is a zero vector, its distribution is determined by its covariance matrix. This is given by the central [2, 2] sub-matrix in $\hat{\Sigma}_Z$. Inspection of the fitted $\hat{\Sigma}_Z$ covariance matrices in Table 7 reveals that the relevant matrix I_{n_1} is the same in all eight models. Hence this block marginal distribution is fixed in the eight maximum entropy distributions.

3.4. Mutual Information

It was claimed in Section 3 that there is no loss of generality in assuming that the mean vectors of X_0, X_1, Y are a zero vector and that their covariance matrices are an identity matrix, of the required sizes. We will now demonstrate this, by calculating the mutual information $I(X_0, X_1; Y)$, using the general form of covariance matrix (which is partitioned conformably to Σ_Z in (32)):

$$\Sigma = \begin{bmatrix} \Sigma_{00} & \Sigma_{01} & \Sigma_{02} \\ \Sigma_{01}^T & \Sigma_{11} & \Sigma_{12} \\ \Sigma_{02}^T & \Sigma_{12}^T & \Sigma_{22} \end{bmatrix} = \begin{bmatrix} \Sigma_{00}^{\frac{1}{2}} & 0 & 0 \\ 0 & \Sigma_{11}^{\frac{1}{2}} & 0 \\ 0 & 0 & \Sigma_{22}^{\frac{1}{2}} \end{bmatrix}^T \begin{bmatrix} I_{n_0} & P & Q \\ P^T & I_{n_1} & R \\ Q^T & R^T & I_{n_2} \end{bmatrix} \begin{bmatrix} \Sigma_{00}^{\frac{1}{2}} & 0 & 0 \\ 0 & \Sigma_{11}^{\frac{1}{2}} & 0 \\ 0 & 0 & \Sigma_{22}^{\frac{1}{2}} \end{bmatrix}, \quad (47)$$

where

$$P = \Sigma_{00}^{-\frac{1}{2}} \Sigma_{01} \Sigma_{11}^{-\frac{1}{2}}, \quad Q = \Sigma_{00}^{-\frac{1}{2}} \Sigma_{02} \Sigma_{22}^{-\frac{1}{2}}, \quad R = \Sigma_{11}^{-\frac{1}{2}} \Sigma_{12} \Sigma_{22}^{-\frac{1}{2}}.$$
 (48)

Since Σ_{ii} (i = 0, 1, 2) is positive definite (by Lemma 1) it has a positive definite square root $\Sigma_{ii}^{\frac{1}{2}}$ ([26], pp. 405–406). Therefore, using standard properties of determinants,

$$|\Sigma| = |\Sigma_{00}||\Sigma_{11}||\Sigma_{22}| \begin{vmatrix} I_{n_0} & P & Q \\ P^T & I_{n_1} & R \\ Q^T & R^T & I_{n_2} \end{vmatrix}.$$
 (49)

From [24], we can state the following marginal distributions.

$$X_0 \sim N(0, \Sigma_{00}), \quad X_1 \sim N(0, \Sigma_{11}), \quad Y \sim N(0, \Sigma_{22}),$$

and

$$[\mathbf{X}_0 \ \mathbf{X}_1]^T, \quad [\mathbf{X}_0 \ \mathbf{Y}]^T, \quad [\mathbf{X}_1 \ \mathbf{Y}]^T$$
(50)

are multivariate normal with covariance matrices

$$\begin{bmatrix} \Sigma_{00} & \Sigma_{01} \\ \Sigma_{01}^T & \Sigma_{11} \end{bmatrix}, \begin{bmatrix} \Sigma_{00} & \Sigma_{02} \\ \Sigma_{02}^T & \Sigma_{22} \end{bmatrix}, \begin{bmatrix} \Sigma_{11} & \Sigma_{12} \\ \Sigma_{12}^T & \Sigma_{22} \end{bmatrix},$$
(51)

respectively. The formula for the entropy of a multivariate Gaussian distribution is required. For a *k*-dimensional random variable **W** following a Gaussian distribution with mean vector μ and covariance matrix Σ , the entropy in the distribution of *W* is [25]

$$H(\mathbf{W}) = \frac{k}{2} + \frac{k}{2}\log(2\pi e) + \frac{1}{2}\log|\Sigma|,$$
(52)

Using the formula for entropy in (52), and using a similar argument to that which produced (49), we may write

$$H(\mathbf{Y}) = \frac{n_2}{2} + \frac{n_2}{2}\log(2\pi e) + \frac{1}{2}\log|\Sigma_{22}|,$$
(53)

$$H(\mathbf{X}_{0}, \mathbf{X}_{1}) = \frac{(n_{0} + n_{1})}{2} + \frac{(n_{0} + n_{1})}{2}\log(2\pi e) + \frac{1}{2}\log(|\Sigma_{00}||\Sigma_{11}|) + \frac{1}{2}\log\begin{vmatrix} I_{n_{0}} & P\\ P^{T} & I_{n_{1}} \end{vmatrix}$$
(54)

$$H(\mathbf{X}_{0}, \mathbf{X}_{1}, \mathbf{Y}) = \frac{m}{2} + \frac{m}{2}\log(2\pi e) + \frac{1}{2}\log(|\Sigma_{00}||\Sigma_{11}||\Sigma_{22}|) + \frac{1}{2}\log\begin{vmatrix} I_{n_{0}} & P & Q \\ P^{T} & I_{n_{1}} & R \\ Q^{T} & R^{T} & I_{n_{2}} \end{vmatrix}$$
(55)

Therefore, applying a version of (27) with vector arguments, and using (32), the total mutual information is given by

$$I(\mathbf{X}_{0}, \mathbf{X}_{1}; \mathbf{Y}) = \frac{1}{2} \log \left| I_{n_{1}} - P^{T} P \right| - \frac{1}{2} \log \left| \begin{matrix} I_{n_{0}} & P & Q \\ P^{T} & I_{n_{1}} & R \\ Q^{T} & R^{T} & I_{n_{2}} \end{matrix} \right| = \frac{1}{2} \log \frac{\left| I_{n_{1}} - P^{T} P \right|}{|\Sigma_{Z}|},$$
(56)

using the fact that (Lemma 2)

$$\begin{vmatrix} I_{n_0} & P \\ P^T & I_{n_1} \end{vmatrix} = \begin{vmatrix} I_{n_1} - P^T P \end{vmatrix}.$$

Therefore, we have demonstrated that the mutual information between the predictors and the target does not depend on the either the mean vectors or the covariance matrices of the individual random vectors X_0 , X_1 and Y. Using (52), the distributional results (50)–(51) and similar arguments to that leading to (49), we state formulae for the other required mutual informations.

$$I(\mathbf{X}_{0}; \mathbf{Y}) = \frac{1}{2} \log \frac{1}{\left| I_{n_{2}} - Q^{T} Q \right|},$$
(57)

$$I(\mathbf{X}_{1};\mathbf{Y}) = \frac{1}{2}\log\frac{1}{\left|I_{n_{2}} - R^{T}R\right|},$$
(58)

$$I(\mathbf{X}_{0}; \mathbf{Y} | \mathbf{X}_{1}) = \frac{1}{2} \log \frac{\left| I_{n_{1}} - P^{T} P \right| \left| I_{n_{2}} - R^{T} R \right|}{|\Sigma_{Z}|},$$
(59)

$$I(\mathbf{X}_1; \mathbf{Y} | \mathbf{X}_0) = \frac{1}{2} \log \frac{\left| I_{n_1} - P^T P \right| \left| I_{n_2} - Q^T Q \right|}{|\Sigma_Z|}$$
(60)

Expressions for the total mutual information, $I(\mathbf{X}_0, \mathbf{X}_1; \mathbf{Y})$, for models $M_1 \dots M_8$ are provided in Table 8.

Table 8. Expressions for the predictors-target mutual information for the eight models in the dependency lattice in Figure 7, that are stated in Table 7.

$$M_{8}: \frac{1}{2} \log \frac{\left|I_{n_{1}} - P^{T}P\right|}{\left|\Sigma_{Z}\right|} \qquad M_{4}: I(\mathbf{X}_{1}; \mathbf{Y})$$
$$M_{7}: \frac{1}{2} \log \frac{\left|I_{n_{1}} - RQ^{T}QR^{T}\right|}{\left|I_{n_{2}} - Q^{T}Q\right|\left|I_{n_{2}} - R^{T}R\right|} \qquad M_{3}: I(\mathbf{X}_{0}; \mathbf{Y})$$
$$M_{6}: I(\mathbf{X}_{1}; \mathbf{Y}) = \frac{1}{2} \log \frac{1}{\left|I_{n_{2}} - R^{T}R\right|} \qquad M_{2}: 0$$
$$M_{5}: I(\mathbf{X}_{0}; \mathbf{Y}) = \frac{1}{2} \log \frac{1}{\left|I_{n_{2}} - Q^{T}Q\right|} \qquad M_{1}: 0$$

3.5. The I_{dev} PID for Multivariate Gaussian Predictors and Targets

Using the expressions for the total mutual information between predictors and the target in Table 8, formulae for the edge values that are used in the construction of the I_{dep} PID, are given in Table 9. They are computed by subtracting the mutual informations of the relevant models in Figure 7; for example, *k* is computing by subtracting the mutual information in model M_6 from that in model M_8 .

Table 9. Expression for the edge values in the dependency lattice in Figure 7 that are used to determine the unique informations.

$$b = d = I(\mathbf{X}_{0}; \mathbf{Y}) = \frac{1}{2} \log \frac{1}{|I_{n_{2}} - Q^{T}Q|} \qquad c = f = I(\mathbf{X}_{1}; \mathbf{Y}) = \frac{1}{2} \log \frac{1}{|I_{n_{2}} - R^{T}R|}$$
$$i = \frac{1}{2} \log \frac{|I_{n_{1}} - RQ^{T}QR^{T}|}{|I_{n_{2}} - Q^{T}Q||I_{n_{2}} - R^{T}R|} - I(\mathbf{X}_{1}; \mathbf{Y}) \qquad h = \frac{1}{2} \log \frac{|I_{n_{1}} - RQ^{T}QR^{T}|}{|I_{n_{2}} - Q^{T}Q||I_{n_{2}} - R^{T}R|} - I(\mathbf{X}_{0}; \mathbf{Y})$$
$$k = \frac{1}{2} \log \frac{|I_{n_{1}} - P^{T}P|}{|\Sigma_{Z}|} - I(\mathbf{X}_{1}; \mathbf{Y}) \qquad j = \frac{1}{2} \log \frac{|I_{n_{1}} - P^{T}P|}{|\Sigma_{Z}|} - I(\mathbf{X}_{0}; \mathbf{Y})$$

Given the edge values in Table 9, we can form the I_{dep} PID for multivariate Gaussian predictors and targets.

$$\operatorname{unq0} = \min\{b, d, i, k\}, \qquad \operatorname{red} = I(\mathbf{X}_0; \mathbf{Y}) - \operatorname{unq0}, \qquad (61)$$

$$unq1 = I(\mathbf{X}_1; \mathbf{Y}) - red, \qquad syn = I(\mathbf{X}_0; \mathbf{Y} | \mathbf{X}_1) - unq0.$$
(62)

We now state some results for this PID. Proofs are given in Appendix C.

Proposition 4. For two multivariate Gaussian predictors, X_0, X_1 , and one multivariate Gaussian target, Y, the PID defined in Table 9 and (61)–(62) has the following properties.

- (a) This I_{dep} PID possesses consistency as well as the core axioms of non-negativity, self-redundancy, monotonicity, symmetry and identity.
- (b) When unq0 is equal to b or d, the the redundancy component is zero.
- (c) When unq0 is equal to *i*, the redundancy and both unique informations are constant with respect to the correlation matrix P between the two predictors, X_0, X_1 .
- (d) When neither predictor and the target are independent, then unq0 is equal to either i or to k.
- (e) When unq0 is equal to k, the synergy component is zero.
- (f) The redundancy component in the I_{mmi} PID is greater than or equal to the redundancy component in the I_{dep} PID with equality if, and only, if at least one of the following conditions holds: (i) either predictor and the target are independent; (ii) either predictor is conditionally independent of the target given the other predictor.
- (g) The synergy component in the I_{mmi} PID is greater than or equal to the synergy component in the I_{dep} PID with equality if, and only, if at least one of the following conditions holds: (i) either predictor and the target are independent; (ii) either predictor is conditionally independent of the target given the other predictor.
- (h) The I_{dep} and I_{mmi} PIDs are identical when either X_0 and Y are conditionally independent given X_1 or X_1 and Y are conditionally independent given X_0 , and in particular they are identical for models $M_1 \dots M_6$. In model M_7 the synergy component of I_{dep} is zero.

3.6. Examples and Illustrations

The multivariate version of the I_{dep} PID was implemented using the edge coefficients in Table 9 together with the PID rules in (61)–(62). The matrices, *P*, *Q*, *R*, were given an equi-correlation structure in which all the entries were equal within each matrix:

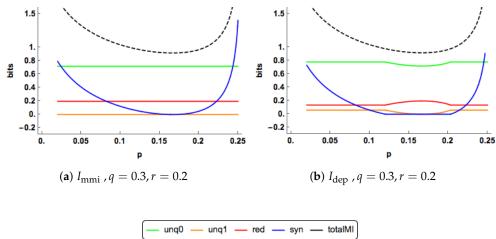
$$P = p \mathbf{1}_{n_0} \mathbf{1}_{n_1}^T, \quad Q = q \mathbf{1}_{n_0} \mathbf{1}_{n_2}^T, \quad R = r \mathbf{1}_{n_1} \mathbf{1}_{n_2}^T, \tag{63}$$

where p, q, r denote here the constant correlations with each matrix and $\mathbf{1}_n$ denotes an n-dimensional vector whose entries are each equal to unity.

Taking p = 0.1, q = 0.2, q = 0.3, $n_0 = 4$, $n_1 = 3$, $n_0 = 2$, respectively, the covariance (correlation) matrix Σ_Z was computed and plots produced of the PIDs as displayed below. The covariance matrix is positive definite only for limited ranges of p, q, r. The I_{mmi} PID was computed using the formulae in Section 2.5, but replacing X_0 , X_1 , Y by their vector counterparts X_0 , X_1 , Y, respectively.

Figure 8 shows some plots of the multivariate I_{mmi} and I_{dep} PIDs as a function of p, for particular values of q and r. These plots display similar characteristics to those shown in Figure 3, Section 2.6. Some further plots are displayed in Figure 9. This time the PIDs are shown for increasing values of q(=r), for two values of p. Again, these plots have similar characteristics to those considered in Figure 4, Section 2.6.

(n_0,n_1,n_2)	(p,q,r)	PID	unq0	unq1	red	syn
(3,4,3)	(-0.15, 0.15, 0.15)	<i>I</i> dep	0.1227	0.1865	0.0406	2.4772
		I _{mmi}	0	0.0638	0.1632	2.6000
(4, 4, 2)	(-0.2, -0.2, 0.3)	<i>I</i> dep	0.0893	0.7293	0.1889	0.0087
		I _{mmi}	0	0.6401	0.2782	0.0980
(4, 2, 4)	(-0.1, 0.15, -0.2)	<i>I</i> dep	0.2336	0.1899	0.0883	0.0345
		I _{mmi}	0.0437	0	0.2782	0.2234



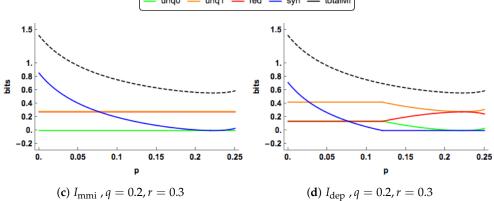


Figure 8. The I_{mmi} and I_{dep} PID components are plotted for a range of values of the correlation (*p*) between the two predictors. Two combinations of the correlations (*q*, *r*) between each predictor and the target are displayed. The total mutual information $I(X_0, X_1; Y)$ is also shown as a dashed black curve.

Values of n_0 , n_1 , n_2 , p, q, r were chosen, ensuring that the covariance matrix was positive definite, and the equi-correlation structure defined in (49) was used. The PID results are presented in the table above.

We see rather different compositions of PID components in the three examples as well as some differences between the two methods. For the first system, both methods have a very large value for the synergy component, with I_{dep} having larger values for the unique informations than I_{mmi} but lower redundancy. The two methods produce fairly similar PIDs for the second system, although there are some differences of about 0.09 bit in all of the components. The third system has strong differences between the I_{dep} and I_{mmi} PIDs. I_{dep} has large values for the two unique components along with small values for redundancy and synergy, whereas I_{mmi} has large values for redundancy and synergy and very small values for the uniques.

In these examples, the dimensions of the predictors and target have an impact on the resulting PIDs as well as the correlations.

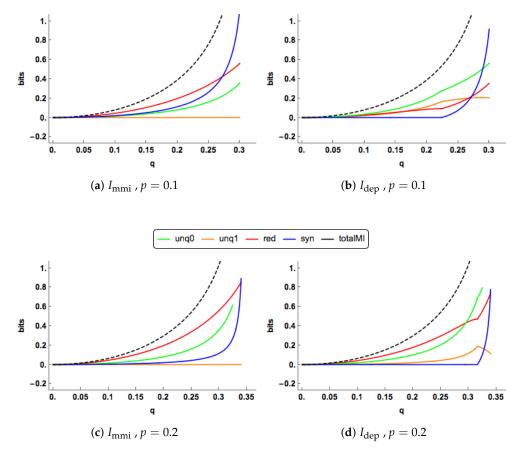


Figure 9. The I_{mmi} and I_{dep} PID components are plotted for a range of values of the correlation (*q*) between the predictor \mathbf{X}_0 and the target \mathbf{Y} . Two combinations of the correlations (*q*, *r*) between each predictor and the target are displayed. The total mutual information $I(X_0, X_1; Y)$ is also shown as a dashed black curve.

Example 6. Prediction of calcium contents

The multivariate I_{dep} PID and the I_{mmi} PID were applied using data ([27], p. 145) on 73 women involving one set of predictors X_0 (Age, Weight, Height), another set of two predictors X_1 (Diameter of os calcis, Diameter of radius and ulna), and target **Y** (Calcium content of heel and forearm). The following results were obtained.

PID	unq0	unq1	red	syn
<i>I</i> dep	0.4077	0.0800	0.0232	0.1408
I _{mmi}	0.3277	0	0.1032	0.2209

Both PIDs indicate the presence of synergy and a large component of unique information due to the variables in X_0 . The I_{dep} PID shows more unique information but less redundancy and synergy than I_{mmi} . To explore matters further, deviance tests ([13], p. 185) were performed which compared models $M_1 \dots M_7$ against the saturated model M_8 . In all seven tests the *p*-values are very small indeed $(p < 2 \times 10^{-4})$, indicating that there is very strong evidence to reject models $M_1 \dots M_7$ and that model M_8 provides the best explanation of the data. The test of M_6 against M_8 provided very strong evidence in favour including the term **XY** in model M_8 , and so there is little surprise that both PIDs have a large value for unq0. The test of model M_5 against model M_8 gave very strong evidence that **X**₁**Y** should also be included in model M_8 . On this occasion, this has not led to a large value for unq1, so perhaps these two terms combine to produce synergy and redundancy.

The PIDs were also computed with the same X_0 and Y but taking X_1 to be another set of four predictors (Surface area, Strength of forearm, Strength of leg, Area of os calcis). The following results were obtained.

PID	unq0	unq1	red	syn
<i>I</i> dep	0.3708	0.0186	0.0601	0
	0.3522	0	0.0787	0.0186

In this case, the I_{mmi} and I_{dep} PIDs are very similar, with the main component being due to unique information due to the variables in \mathbf{X}_0 . The I_{dep} PID indicates zero synergy and almost zero unique information due to the variables in \mathbf{X}_1 Again, deviance tests were performed. In six of the seven tests the approximate *p*-values are very small indeed (less than 4×10^{-5}). The exception is model M_5 for which the deviance test has an approximate *p*-value of 0.98, indicating that model M_5 provides the simplest explanation for the data and an extremely good fit to the data. In model M_5 , it is expected that there will be zero synergy as well as a zero unique component due to the variables in \mathbf{X}_1 , and this matches quite well the information produced in the PIDs.

When working with real or simulated data it is important to use the correct correlation matrix. In order to use the results given in Table 9 and (61)–(62) it is essential that the input covariance matrix has the structure of Σ_Z , as given in (32). The computational approach used here is described in Appendix D.

4. Discussion

We have applied the I_{dep} method to obtain bivariate partial information decompositions for Gaussian systems with both univariate and multivariate predictors and targets. We give closed form solutions for all PID terms in both these cases, to allow easy computation of a PID from a covariance or correlation matrix. The main properties enjoyed by I_{dep} for Gaussian systems are the same as those defined in [2]. The characteristics of the I_{dep} PIDs for Gaussian system have been illustrated by graphical exploration as well as numerical examples.

Given that the I_{dep} method employs a lattice of probability models, Gaussian graphical models, it seems natural when attempting to understand the form of a particular PID to consider formal statistical tests in order to determine which of the models in the lattice best fits the data. Therefore, deviance tests have been used for this purpose. They provide a useful complementary approach, as demonstrated in the examples considered.

There are now three approaches to the PID for Gaussian systems, I_{nmi} [12], I_{ccs} [6] and I_{dep} [2] as developed here. While they may agree in some cases, these methods are in general all distinct. For I_{dep} and I_{ccs} the redundancy and unique information values are not invariant to the predictor-predictor marginal distribution (here p or P), and so they are not equivalent to I_{mmi} . Here, we proved that

redundancy and synergy measured with I_{mmi} are never less than those measured with I_{dep} , and are equal in specific circumstances regarding marginal and conditional independence (see Propositions 2 and 4 (f, g)). We note that if the full system or data matches any one of the models $U_1 \dots U_6$ (or $M_1 \dots M_6$) then a conditional independence condition is met. This forces one unique component and the synergy component to be equal to zero, and I_{mmi} and I_{dep} are identical. Therefore, in practice, if any of these models provide an acceptable fit to the data, then the I_{dep} and I_{mmi} PIDs are likely to be quantitatively very similar. If models U_7, U_8 (or M_7, M_8) provide a better fit to the data, then all four components can be non-zero. By considering the perspective of multivariate linear regression based on the conditional distribution of **Y** given **X**₀ and **X**₁, as in [21] for the univariate case, synergy is expected to be present in model M_8 , and I_{dep} and I_{mmi} can diverge. This is also the case with model M_7 , although here the synergy component of I_{dep} is zero. A more thorough comparison of the behaviour of these different measures across families of Gaussian systems could help to illustrate their different interpretations and perhaps shed light on the different approaches to the PID in the discrete case.

As noted, while the I_{mmi} PID has the property that the redundancy component does not depend on the correlation between the predictors this is not true in general for the I_{dep} PIDs. When there is positive synergy in the I_{dep} PIDs it is the case that redundancy and unique information are invariant to predictor-predictor dependence (*p*), but when synergy is zero this does not hold (see e.g., Figure 3). However, as shown, the I_{dep} redundancy and synergy terms are always less than or equal to the corresponding I_{mmi} terms. From considering the arguments related to the best fitting models, it seems that in some cases the I_{mmi} approach may overstate redundancy. Further, for I_{mmi} it is by definition not possible for two predictors to both carry unique information. Considering the properties of Gaussian systems and simple noisy additive linear systems this seems unintuitive: if the predictors are independent or anti-correlated but with fixed correlation with the target it seems more natural that, across samples, they each provide a positive unique information contribution to an estimate of the target (see e.g., Example 4). Similarly, it also seems intuitive that in a Gaussian setting the amount of information shared between two predictors with fixed target correlation should increase as the correlation between the predictors increases (Figure 3d). Further, one would imagine, in general that it should be possible for two variables to carry the same amount of information, but for that information to be different.

Both of these considerations suggest that the dependence on the predictor-predictor marginals in both I_{dep} and I_{ccs} seems to be more natural for Gaussian systems. The invariance to the predictorpredictor marginals was a foundational assumption in the derivation of the method presented in [5], and was based on a decision theoretic operationalization of unique information. However, a game theoretic extension of this approach in [6] suggests that this invariance is not a natural requirement for a measure of shared information. In addition, for Gaussian systems there are existing classical variance-based approaches to the problem, such as commonality analysis [28,29], based on semi-partial correlation, or path analysis [30], which could provide another perspective on the problem. Systematically comparing these methods is an interesting area for future work.

The I_{dep} PIDs presented provide a non-negative decomposition of a joint predictor-target mutual information for Gaussian systems. This could have broad applications, from an exploratory statistical tool, to analysis of complex systems and networks. Gaussian models or approximations have been used for computing information-theoretic statistics from experimental data in neuroscience and neuroimaging [19,20]. For example, in neuroimaging there are often statistical effects of a stimulus observed in multiple recorded responses (for example different brain regions, or different temporal offsets from stimulation). Methods such as the PID can provide a practical tool to relate two such modulations and so give insight into whether they are likely to reflect the same or different brain processes. Similarly, if multiple stimulus features or aspects are presented in an experiment, the PID can be applied to quantify how much of the neural response is commonly predicted from both stimulus features, uniquely available from each or synergistically available only from the combination.

The I_{dep} method of [2] is a very general one and it could be applied to systems other than discrete systems [2] or the Gaussian PIDs developed here. For example, the I_{dep} method could be used with other types of graphical model, such as mixed discrete-continuous systems [13,15] based on the CG model, and also in multivariate autoregressive modelling of time series data [12,31,32] using graphical models [33–35]. We look forward to engaging in further exploration of the potential of the I_{dep} method.

Supplementary Materials: The following are available online at http://www.mdpi.com/1099-4300/20/4/240/s1.

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Appendix A. Proof of Proposition 2

Examination of the edge values in Table 5 shows that the PID derived here satisfies Equations (B2)–(B5) of [2], on taking m = 0. Therefore, the properties of consistency, non-negativity, self-redundancy, monotonicity and identity follow for this new PID using the arguments given in [2]. Therefore, we consider only parts (b)–(h).

(b) This is true because both *b* and *d* are equal to $I(X_0; Y)$.

(c) When unq0 = i, the unique informations and the redundancy components are

$$\frac{1}{2}\log\frac{1-q^2r^2}{1-q^2}, \quad \frac{1}{2}\log\frac{1-q^2r^2}{1-r^2}, \quad \frac{1}{2}\log\frac{1}{1-q^2r^2}$$

respectively, and all these terms are independent of *p*.

(d) We are given that $q \neq 0, r \neq 0$. Now,

$$b - i = \frac{1}{2} \log \frac{1}{1 - q^2 r^2} > 0$$

when $q \neq 0, r \neq 0$, and since |q| < 1, |r| < 1. Hence the minimum of the edge values is not *b* or *d*, which leaves only *i* and *k* as possibilities.

(e) From (30) and the expression for k in Table 5, we see that the synergy component

$$I(X_0; Y|X_1) - unq0$$

is equal to zero.

(f, g) We will use the definitions of the I_{dep} and I_{mmi} PIDs in (8)–(11) and (4)–(7) and also the bivariate and conditional mutual informations in (28)–(31). We denote the I_{mmi} redundancy and synergy components by R_m and S_m , respectively, using R_d and S_d for the corresponding I_{dep} components. Denote unq0 and unq1 in I_{dep} as U_{0d} , U_{1d} , respectively. We note that

$$I(X_0; Y) = 0 \iff q = 0$$
, and $I(X_0; Y|X_1) = 0 \iff q = pr$. (A1)

First, suppose that $I(X_0; Y) < I(X_1; Y)$. If $I(X_0; Y) = 0$, then q = 0 and 0 = b = i < k in I_{dep} . Hence,

$$R_m = R_d = 0$$
, and $S_m = S_d = I(X_0; Y | X_1)$.

If $I(X_0; Y) \neq 0$, then $R_m = I(X_0; Y)$ and $S_m = I(X_0; Y|X_1)$. In I_{dep} , the redundancy and synergy components are

$$R_d = I(X_0; Y) - U_{0d}$$
, and $S_d = I(X_0; Y|X_1) - U_{0d}$.

It follows that $R_m \ge R_d$ and $S_m \ge S_d$ with equality iff $U_{0d} = 0$. From (d), it follows for I_{dep} that $U_{0d} = i$ or $U_{0d} = k$. Since $I(X_0; Y) \ne 0$, i > 0 so $U_{0d} = 0$ iff k = 0, which from Table 5 and (A1) is true iff $I(X_0; Y|X_1) = 0$, in which case $S_m = S_d = 0$ and $R_m = R_d = I(X_0; Y)$. Hence result.

The proof when $I(X_0; Y) > I(X_1; Y)$ is similar and is omitted, although it is worth noting that

$$I(X_1; Y) = 0 \iff r = 0$$
, and $I(X_1; Y|X_0) = 0 \iff r = pq$. (A2)

When $I(X_0; Y) = I(X_1; Y)$, then

$$R_m = I(X_0; Y) = I(X_1; Y)$$
, and $R_d = I(X_0; Y) - U_{0d} = I(X_1; Y) - U_{1d}$

$$S_m = I(X_0; Y|X_1) = I(X_1; Y|X_0)$$
, and $S_d = I(X_0; Y|X_1) - U_{0d} = I(X_0; Y|X_1) - U_{1d}$.

Therefore $R_m \ge R_d$ and $S_m \ge S_d$ with equality iff $U_{0d} = U_{1d} = 0$. From the argument above, this happens iff

$$I(X_0; Y) = I(X_1; Y) = 0$$
, or $I(X_0; Y|X_1) = I(X_1; Y|X_0) = 0$.

In this case, the I_{dep} and I_{mmi} redundancy and synergy components are equal if, and only if, each of X_0 and X_1 is independent of Y, and each of X_0 and X_1 is conditionally independent of Y given the other predictor.

(h) When $I(X_0; Y|X_1) = 0$, the unq0 and syn components are zero in both the I_{dep} and I_{mmi} PIDs. From (A1), p = qr, and from Table 5 we see that in the I_{dep} PID, i = k < b, and so $U_{1d} = I(X_1; Y|X_0)$ and $R_d = I(X_0; Y)$. Since $(X_0; Y|X_1) = 0$, it follows from

$$I(X_0, X_1; Y) = I(X_0; Y) + I(X_0; Y|X_1) = I(X_1; Y) + I(X_1; Y|X_1)$$

that $I(X_0; Y) \leq I(X_1; Y)$ and so in the I_{mmi} PID, red = $I(X_0; Y)$ and $unq1 = I(X_1; Y|X_0)$. It follows that the I_{dep} and I_{mmi} PIDs are identical.

The proof when $I(X_1; Y; X_0) = 0$ is very similar and it is omitted. Model U_6 has $I(X_0; Y|X_1) = 0$, model U_5 has $I(X_1; Y|X_0) = 0$, and models $U_1 \dots U_4$ have at least one of these conditions. Hence result. In model U_7 , p = qr. If $q \neq 0$, $r \neq 0$, it follows from (16) and Table 5 that

$$|\Sigma_Z| = (1 - q^2)(1 - r^2)$$
, and $I(X_0; Y|X_1) = k = i < b$ (if $q \neq 0$).

Therefore, in the I_{dep} PID unq0 =k and so syn =0. If q = 0 then from Table 5, $b = i = k = I(X_0; Y|X_1) = 0$, and so syn =0. If r = 0, then $b = i = k = I(X_0; Y|X_1)$, and so syn = 0. Hence result.

Appendix B. Proof of Matrix Lemmas

We begin by stating some some useful results from matrix algebra ([26], p. 472), ([36], p. 475). Suppose that a symmetric matrix *M* is partitioned as

$$M = \begin{bmatrix} A & B \\ B^T & C \end{bmatrix},$$

where *A* and *C* are symmetric and square. Then

- (i) The matrix *M* is positive definite if and only if *A* and $C B^T A^{-1} B$ are positive definite.
- (ii) The matrix *M* is positive definite if and only if *C* and $A BC^{-1}B^{T}$ are positive definite.
- (iii) $|M| = |A||D B^T A^{-1}B|.$

Proof of Lemma 1

If we write *M* as

$$M = \begin{bmatrix} A & M_{13} \\ M_{13}^T & M_{33} \end{bmatrix}, \text{ with } A = \begin{bmatrix} M_{11} & M_{12} \\ M_{12}^T & M_{22} \end{bmatrix}$$

then if *M* is positive definite it follows from (i) that *A* and M_{33} are both positive definite. Applying result (i) to the matrix *A* then shows that M_{11} and M_{22} are also positive definite. A positive definite matrix is nonsingular. Hence result.

Proof of Lemma 2

Given that Σ_Z is positive definite, we note that

$$\begin{bmatrix} I_{n_0} & P \\ P^T & I_{n_1} \end{bmatrix}, \begin{bmatrix} I_{n_0} & Q \\ Q^T & I_{n_2} \end{bmatrix}, \begin{bmatrix} I_{n_1} & R \\ R^T & I_{n_2} \end{bmatrix}$$

are principal sub-matrices of Σ_Z and so they are positive definite ([26], p. 397). From (i, ii), it follows that the matrices

$$I_{n_1} - P^T P$$
, $I_{n_0} - PP^T$, $I_{n_2} - R^T R$, $I_{n_1} - RR^T$, $I_{n_2} - Q^T Q$, $I_{n_0} - QQ^T$.

are positive definite.

Suppose that $I_n - X^T X$ is positive definite, where X is a $p \times n$ matrix. Then the matrix $X^T X$ is positive semi-definite and so has non-negative eigenvalues, $\lambda_1, \lambda_2, ..., \lambda_n$. The eigenvalues of $I_n - X^T X$ are $\{1 - \lambda_i : i = 1, 2, ..., n\}$. Since $I_n - X^T X$ is positive definite we know that $1 - \lambda_i > 0$ for i = 1, 2, ..., n. It follows that $0 < 1 - \lambda_i \le 1$ for i = 1, 2, ..., n. Since the determinant of a square matrix is the product of its eigenvalues we have that

$$|I_n - X^T X| = \prod_{i=1}^n (1 - \lambda_i),$$

and so $0 < |I_n - X^T X| \le 1$. It also follows that $|I_n - X^T X| = 1$ if, and only if, all the eigenvalues of $X^T X$ are equal to zero, which means that $X^T X$ is the zero matrix. Taking $X = P, P^T, Q, Q^T, R, R^T$ in turn gives the required result.

Application of (iii) gives the result that

$$\begin{vmatrix} I_{n_0} & P \\ P^T & I_{n_1} \end{vmatrix} = |I_{n_1} - P^T P|.$$

Appendix C. Proof of Proposition 4

Examination of the edge values in Table 9 shows that the multivariate I_{dep} PID derived here satisfies Equations B2–B5 of [2], on taking m = 0. Therefore, the properties of consistency, non-negativity, self-redundancy, monotonicity and identity follow for this new PID using the arguments given in [2]. Hence, we focus attention only on parts (b–h).

(b) This is true since both *b* and *d* are equal to $I(\mathbf{X}_0; Y)$.

(c) When unq0 = i, then the expressions for the unique informations and the redundancy, given in Table 9 and (61)–(62) do not depend on the matrix *P*

(d) From (48) and the assumption that neither Σ_{02} nor Σ_{12} is equal to a zero matrix, it follows that QR^T is not equal to a zero matrix.

From Table 9,

$$b-i = \frac{1}{2} \log \frac{1}{|I_{n_1} - RQ^T QR^T|}.$$

From Table 5, we have that the covariance matrix under model M_7 is

$$\Sigma_7 = \begin{bmatrix} I_{n_0} & QR^T & Q \\ RQ^T & I_{n_1} & R \\ Q^T & R^T & I_{n_2} \end{bmatrix}.$$

Applying a similar argument to that in the proof of Lemma 2, it follows that $|I_{n_1} - RQ^TQR^T|$ is positive and bounded above by unity. Also it is equal to unity only when QR^T is equal to the zero matrix having n_0 rows and n_1 columns. Since this is not the case, it follows that

$$0 < |I_{n_1} - RQ^T Q R^T| < 1$$
(A3)

and so b > i. Therefore the minimum does not occur at b or d, leaving only i and k as the remaining possibilities.

(e) From (59) and the entry for k in Table 9, we see that the synergy component is equal to zero when unq0 = k.

(f, g) The proofs are very similar to those for (f, g) in Proposition 2 and so they are omitted. The following results are useful. From (57)–(60), we can state the following results.

 $I(\mathbf{X}_0; \mathbf{Y}) = 0$ iff \mathbf{X}_0 and \mathbf{Y} are independent, iff the matrix *P* is a zero matrix.

 $I(\mathbf{X}_1; \mathbf{Y}) = 0$ iff \mathbf{X}_1 and \mathbf{Y} are independent, iff the matrix *R* is a zero matrix.

 $I(\mathbf{X}_0; \mathbf{Y} | \mathbf{X}_1) = 0$ iff \mathbf{X}_0 and \mathbf{Y} are conditionally independent given \mathbf{X}_1 , iff Q = PR, from (36).

 $I(\mathbf{X}_1; \mathbf{Y} | \mathbf{X}_0) = 0$ iff \mathbf{X}_1 and \mathbf{Y} are conditionally independent given \mathbf{X}_0 , iff $R = P^T Q$, from (37).

(h) When $I(\mathbf{X}_0; \mathbf{Y} | \mathbf{X}_1) = 0$, the unq0 and syn components components are zero in both the I_{dep} and I_{mmi} PIDs, and so in the I_{dep} PID, unq1 = $I(\mathbf{X}_1; \mathbf{Y} | \mathbf{X}_0)$ and red = $I(\mathbf{X}_0; \mathbf{Y})$. Since $I(\mathbf{X}_0; \mathbf{Y} | \mathbf{X}_1) = 0$, it follows that $I(\mathbf{X}_0; \mathbf{Y}) \leq I(\mathbf{X}_1; \mathbf{Y})$ and so in the I_{mmi} PID, red = $I(\mathbf{X}_0; \mathbf{Y})$ and unq1 = $I(\mathbf{X}_1; \mathbf{Y} | \mathbf{X}_0)$. It follows that the I_{dep} and I_{mmi} PIDs are identical.

The proof when $I(\mathbf{X}_1; \mathbf{Y}; \mathbf{X}_0) = 0$ is very similar and it is omitted. Model M_6 has $I(\mathbf{X}_0; \mathbf{Y} | \mathbf{X}_1) = 0$, model M_5 has $I(\mathbf{X}_1; \mathbf{Y} | \mathbf{X}_0) = 0$, and models $M_1 \dots M_4$ have at least one of these conditions. Hence result.

In model M_7 , $P = QR^T$, from Table 6. Also from (48)

$$QR^{T} = \Sigma_{00}^{-\frac{1}{2}} \Sigma_{02} \Sigma_{22}^{-1} \Sigma_{12}^{T} \Sigma_{11}^{-\frac{1}{2}}.$$

Provided that neither Σ_{02} nor Σ_{12} is equal to a zero matrix, it follows from (32), Table 8 and (A3) that

$$|\Sigma_Z| = |I_{n_2} - Q^T Q| |I_{n_2} - R^T R|$$
, and $I(X_0; Y|X_1) = k = i < b$.

Therefore, in the I_{dep} PID, unq0 =k and so syn =0. If Σ_{02} is equal to a zero matrix, then from Table 9, $b = i = k = I(X_0; Y|X_1) = 0$, and so syn = 0. Similarly, if Σ_{12} is equal to a zero matrix then $b = i = k = I(X_0; Y|X_1)$, and so syn =0. Hence result.

Appendix D. Computation of the Multivariate I_{dep} PID

Given a multivariate data having two different sets of predictors, X_0 , X_1 and a target, Y, the special formulae presented in Table 9 and (61)–(62) can be used to compute the I_{dep} PID. In order to ensure that the input data have the required format one can use the following procedure.

Suppose that the general covariance matrix is

$$\Sigma = \begin{bmatrix} \Sigma_{00} & \Sigma_{01} & \Sigma_{02} \\ \Sigma_{01}^T & \Sigma_{11} & \Sigma_{12} \\ \Sigma_{02}^T & \Sigma_{12}^T & \Sigma_{22} \end{bmatrix}.$$
 (A4)

Then the required matrices, P, Q, R, can be obtained from this covariance matrix using the following formulae based on (48). The transposes are used here since the extracted square root matrix used here is not symmetric.

$$P = \left[\Sigma_{00}^{-\frac{1}{2}}\right]^T \Sigma_{01} \Sigma_{11}^{-\frac{1}{2}}, \quad Q = \left[\Sigma_{00}^{-\frac{1}{2}}\right]^T \Sigma_{02} \Sigma_{22}^{-\frac{1}{2}}, \quad R = \left[\Sigma_{11}^{-\frac{1}{2}}\right]^T \Sigma_{12} \Sigma_{22}^{-\frac{1}{2}}.$$
 (A5)

Therefore, the procedure involves: (a) extracting the block diagonal matrices, Σ_{00} , Σ_{11} , Σ_{22} , in (A4), (b) finding each square root as an upper triangular matrix by Cholesky decomposition, (c) inverting the upper triangular matrix using a 'backsolve' method and (d) applying the formulae in (A5). Code written in R was used here and in the other examples.

Appendix E. Deviance Tests

We give some details of the deviance tests that have been performed in Sections 2.6 and 3.6. The following notes are based on ([14], p. 40). See also [13]. Suppose that a random vector \mathbf{Z} , of dimension q, follows a Gaussian graphical model \mathcal{M} having mean vector $\boldsymbol{\mu}$ and covariance matrix Σ . Suppose that a sample of N observations is available and that the sample covariance matrix (with divisor N) is S. Let the estimated covariance matrix for model \mathcal{M} be $\hat{\Sigma}$. Then the maximised log likelihood under model \mathcal{M} is

$$\hat{l}_m = -Nq\ln(2\pi)/2 - N\ln|\hat{\Sigma}|/2 - Nq/2.$$

Under the full (or saturated) model \mathcal{M}_f , $\hat{\Sigma} = S$ and so the maximised likelihood under \mathcal{M}_f is

$$\hat{l}_f = -Nq\ln(2\pi)/2 - N\ln|S|/2 - Nq/2.$$

The deviance of a model is defined to be

$$G^2 = 2(\hat{l}_f - \hat{l}_m) = N \ln \frac{|\hat{\Sigma}|}{|S|}$$

and G^2 can be used as a test statistic when testing model \mathcal{M} within the saturated model \mathcal{M}_f . The null distribution of G^2 has an asymptotic chi-squared distribution with degrees of freedom given by the difference in the number of edges between model \mathcal{M} and the saturated model, \mathcal{M}_f . This test is an example of a generalised likelihood ratio test which can be used to compare nested statistical models. A p value can be calculated as $p = \Pr(G^2 \ge G^2_{obs} | \mathcal{M} \text{ is true})$, where G^2_{obs} is the observed value of the test statistic G^2 . When p < 0.01 we may say that there is strong evidence against model \mathcal{M} , the implication being that this model does not provide an acceptable fit to the data. On the other if p > 0.1 we may say that there is little evidence against model \mathcal{M} , with the implication being that this model fit to the data. We may say that there is moderate evidence if 0.01 , and weak evidence if <math>0.05 in the borderline case.

It should be noted that this test is approximate and its performance improves the larger the sample is. There are exact tests in some cases and there are correction factors that can improve the approximation [13,14]. These were not used in this study because the results are so clear cut.

When model M_0 is a special case of (or nested within) model M_1 and it is required to test model M_0 within model M_1 then the deviance test statistic is

$$D = N \ln \frac{|\hat{\Sigma}_0|}{|\hat{\Sigma}_1|}$$

and the null distribution of D also has an asymptotic chi-squared distribution with degrees of freedom equal to the difference in the number of edges between M_0 and M_1 .

There is an interesting connection between the test statistic *D* and some of the edge values in Figure 1. For each of the edge values in the set $\{b, c, d, f, j, k\}$, the edge value is equal to the corresponding value of the test statistic *D* divided by $2N \ln 2$. This is not the case for edge values *h* and *i*.

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