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## ABSTRACT

Optimisation problems often have multiple conflicting objectives that can be computationally and/or financially expensive. Monosurrogate Bayesian optimisation (BO) is a popular model-based approach for optimising such black-box functions. It combines objective values via scalarisation and builds a Gaussian process (GP) surrogate of the scalarised values. The location which maximises a cheap-to-query acquisition function is chosen as the next location to expensively evaluate. While BO is an effective strategy, the use of GPs is limiting. Their performance decreases as the problem input dimensionality increases, and their computational complexity scales cubically with the amount of data. To address these limitations, we extend previous work on BO by density-ratio estimation (BORE) to the multi-objective setting. BORE links the computation of the probability of improvement acquisition function to that of probabilistic classification. This enables the use of state-of-the-art classifiers in a BO-like framework. In this work we present MBORE: multi-objective Bayesian optimisation by densityratio estimation, and compare it to BO across a range of synthetic and real-world benchmarks. We find that MBORE performs as well as or better than BO on a wide variety of problems, and that it outperforms BO on high-dimensional and real-world problems.

### **CCS CONCEPTS**

• Computing methodologies  $\rightarrow$  Modeling methodologies; • Theory of computation  $\rightarrow$  Gaussian processes; • Applied computing  $\rightarrow$  Multi-criterion optimization and decision-making.

### **KEYWORDS**

Bayesian optimisation, Surrogate modelling, Scalarisation methods, Efficient multi-objective optimisation, Acquisition function

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### **1 INTRODUCTION**

Optimisation problems, particularly those in real-world settings, often have conflicting objectives that can be computationally and/or financially expensive to evaluate. For example, it is often desirable to maximise a robot's speed and minimise its energy consumption [50], or to maximise crop yield while minimising the environmental impacts the required agricultural development [40]. These tend to be *black-box* problems, i.e. they have no closed-form or derivative information available. In order to optimise these expensive blackbox functions, a common strategy is to create surrogate models for each of the objective functions and perform optimisation on these instead, choosing where to evaluate the expensive functions next based on locations that are predicted to yield high-quality solutions.

A plethora of multi-objective surrogate-based approaches have been proposed in recent years [2, 13, 57, 75, 78, 86]. These are usually inspired by, or directly use, the framework of single-objective Bayesian optimisation (BO). In BO, given an initial set of expensively evaluated solutions, a surrogate model, often a probabilistic model such as a Gaussian process (GP), is created for either each objective function (a multi-surrogate approach), or a scalarised representation of the multiple objectives (a mono-surrogate approach). In these approaches, an acquisition function, also known as an infill criterion, is optimised to suggest the next location to expensively evaluate. These attempt to balance the exploration of locations with high amounts of prediction uncertainty with the exploitation of locations that have good predicted values. They are also cheap-to-query and, therefore, transform the expensive optimisation problem into a sequence of cheaper ones followed by an expensive evaluation. Once a new candidate solution has been selected and evaluated, the surrogate models are retrained with the new solution and the process is repeated ad libitum.

Mono-surrogate approaches are known to be considerably faster than multi-surrogate approaches [12]. Using only one surrogate model offers large computational savings because the model of choice, GPs, scale cubically in the number of solutions the model is trained on [58], leading to large amounts of computation needed for larger numbers of solutions. Acquisition functions that have been designed for BO [e.g. 16, 42, 72, 80], can be used without alteration. Single-objective acquisition functions are usually cheap to evaluate, unlike those used in multi-surrogate approaches. These often need to carry out expensive multidimensional integrations, such as when calculating the expected hypervolume improvement criterion [19].

Despite the success of mono-surrogate approaches, they are not without limitations. The aforementioned cubic scaling of GPs leads to model training times increasing as the number of solutions grows. Similarly, GPs are known to suffer greatly from the curse of

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dimensionality. Specifically, the modelling ability of nonparametric regression depends exponentially on the problem dimensionality [28]. Approaches to address these fundamental problems with GPs usually involve either reducing the number solutions included in the GP models via *inducing point* methods [62, 66, 68, 79, 83], or making assumptions about the underlying structure of the function of interest to simplify the inference, such as assuming that it resides on a low-dimensional manifold [23, 43, 47, 60, 81, 82]. However, these necessarily lead to poorer modelling of target function [47].

An alternative approach to improving the modelling in BO is to use a different surrogate model, such as a random forest [35] or neural network [70]. However, these models do not usually come with principled ways to compute prediction uncertainty, limiting their applicability to Bayesian optimisation. Recently, Tiao et al. [78] proposed to carry out BO via density-ratio estimation (BORE), a generalisation of the classical tree-structured Parzen estimator [3]. They claimed that maximising the *expected* improvement over the best-seen function value in a probabilistic regression model, the most popular formulation of BO, is equivalent to maximising the class-posterior probability of classifier trained with a proper scoring rule [26]. Unfortunately, this claim is only partially correct [71]. Garnett [25] demonstrates that, in fact, maximising the class-posterior probability is equivalent to maximising the probability of improvement. Nevertheless, the result is important because it still facilitates the use of most state-of-the-art classifiers, such as neural networks or gradient boosted trees [10]. Naturally, this leads to a vast reduction in computational complexity as well as an increase in modelling capability of surrogate models available for use in BO. Therefore, this allows for higher-dimensional problems to be tackled by using more suitable models, e.g. neural networks, which are well-known for having universal function approximation guarantees [33].

In this work, we extend BORE to the multi-objective setting via the use of scalarising functions [12, 36], a mono-surrogate approach. Specifically, we compare MBORE, our proposed multiobjective version of BORE, using two different classification models: a fully-connected neural network and gradient boosted machines (XGBoost [10]) with the standard BO approach using a GP. We compare their performance when using three existing scalarisation approaches, augmented Tchebycheff [44], hypervolume improvement [57] and dominance ranking [57], as well as Pareto hypervolume contribution (PHC), a novel scalarisation method proposed in this work. Multi-objective BO and MBORE are evaluated on two synthetic benchmark suites, DTLZ [18] and WFG [34], across a range of problem dimensionalities and numbers of objectives, as well as a recently proposed real-world set of benchmark problems [76]. Additionally, we also investigate the performance of BO and MBORE on high-dimensional problems using the WFG benchmark, and empirically compare their computational costs.

Our main contributions can be summarised as follows:

- We present MBORE, a novel mono-surrogate multi-objective classification-based BO approach using scalarisation.
- (2) We also present PHC, a new scalariser that directly uses the hypervolume contribution of a solution to its Pareto shell.
- (3) We compare MBORE and BO, using two probabilistic classifiers and several popular scalarisation methods, across a range of synthetic and real-world test problems.

(4) Empirically, we show that MBORE is equal to or better than BO on a wide range of problems, particularly when using PHC, and that MBORE is considerably better than BO on the evaluated real-world and high-dimensional problems, all while using a fraction of the computational resources.

We begin in Section 2 by providing an overview of both singleand multi-objective BO, including a review of GPs, acquisition functions and scalarisation methods. Section 3 introduces BO by density-ratio estimation and, in Section 4, we extend it to the multiobjective setting. An extensive experimental evaluation of BO and MBORE is carried out in Section 5, along with a discussion of the results. We finish with concluding remarks in Section 6.

### 2 BAYESIAN OPTIMISATION FRAMEWORK

In this section we first describe single-objective Bayesian optimisation, including its two main components, the Gaussian process surrogate model and acquisition function. We then go on to show how BO can be extended to the multi-objective setting via scalarisation, giving examples of the scalarisation methods used in this work, along with the proposal of a new scalariser.

### 2.1 Bayesian Optimisation

Bayesian optimisation (BO) was first proposed for single-objective problems by Kushner [46], and later improved by Močkus et al. [55] and Jones et al. [42]. Interested readers should refer to [65] and [22] for recent and comprehensive surveys on the topic. BO is a global search strategy that sequentially samples design space at locations that are likely to contain the global optimum, while taking into account the prediction of a surrogate model and its associated uncertainty [42]. The single-objective optimisation problem can be defined as finding a minimum of an unknown objective function  $f : X \mapsto \mathbb{R}$ , defined on a compact domain  $X \subset \mathbb{R}^d$ :

$$\min_{\mathbf{x}\in\mathcal{X}} f(\mathbf{x}). \tag{1}$$

BO starts by using a space-filling algorithm, such as Latin hypercube sampling [53], to generate an initial set of solutions  $\{\mathbf{x}_i\}_{i=1}^t$ , and then expensively evaluates them with the objective function. These observations form the dataset  $\mathcal{D} = \{(\mathbf{x}_i, f_i \triangleq f(\mathbf{x}_i))\}_{i=1}^t$  that the initial surrogate model is trained with. Following model training, and at each subsequent iteration, the next location to expensively evaluate is chosen as the location  $\mathbf{x}'$  that maximises an acquisition function  $\alpha(\mathbf{x})$ , i.e.  $\mathbf{x}' = \operatorname{argmax}_{\mathbf{x} \in \mathcal{X}} \alpha(\mathbf{x})$ . The dataset is then augmented with  $\{\mathbf{x}', f(\mathbf{x}')\}$  and the process is repeated until budget exhaustion. The global minimum of f is then estimated to be the best-observed value thus far,  $f^* = \min\{f_i\}$ .

2.1.1 *Gaussian Processes.* Gaussian processes (GP) are a common choice of surrogate model due to their strengths in uncertainty quantification and function approximation [58]. They define a prior distribution over functions, such that any finite number of drawn function values are jointly Gaussian, with mean  $m(\mathbf{x})$  and covariance  $\kappa(\mathbf{x}, \mathbf{x}' | \theta)$ , with hyperparameters  $\theta$ . Without loss of generality, we use a zero-mean prior  $m(\mathbf{x}) = 0 \forall \mathbf{x} \in X$ ; see [17] for alternatives. Conditioning the GP prior distribution on data consisting of *t* sampled locations  $\mathcal{D} = \{(\mathbf{x}_i, f_i \triangleq f(\mathbf{x}_i))\}_{i=1}^t$  leads to a

posterior distribution that is also a GP:

$$p(f(\mathbf{x}) \mid \mathbf{x}, \mathcal{D}, \boldsymbol{\theta}) = \mathcal{N}(\mu(\mathbf{x}), \sigma^2(\mathbf{x}))$$
(2)

with mean and variance

$$\mu(\mathbf{x} \mid \mathcal{D}, \boldsymbol{\theta}) = \boldsymbol{\kappa}(\mathbf{x}, \mathbf{X}) \mathbf{K}^{-1} \mathbf{y}$$
(3)

$$\sigma^{2}(\mathbf{x} \mid \mathcal{D}, \boldsymbol{\theta}) = \kappa(\mathbf{x}, \mathbf{x}) - \boldsymbol{\kappa}(\mathbf{x}, \mathbf{X})^{\top} \mathbf{K}^{-1} \boldsymbol{\kappa}(\mathbf{X}, \mathbf{x}).$$
(4)

Here,  $\mathbf{X} \in \mathbb{R}^{t \times d}$  is a matrix of input locations in each row and  $\mathbf{y} = (f_1, f_2, \dots, f_t)^\top$  is the corresponding vector of expensive function evaluations. The kernel matrix  $\mathbf{K} \in \mathbb{R}^{t \times t}$  is  $\mathbf{K}_{ij} = \kappa(\mathbf{x}_i, \mathbf{x}_j \mid \boldsymbol{\theta})$  and  $\kappa(\mathbf{x}, \mathbf{X}) \in \mathbb{R}^t$  is given by  $[\kappa(\mathbf{x}, \mathbf{X})]_i = \kappa(\mathbf{x}, \mathbf{x}_i \mid \boldsymbol{\theta})$ . In this work we use a Matérn 5/2 kernel, as recommended for modelling realistic functions [69, 73]. The kernel's hyperparameters  $\boldsymbol{\theta}$  are learnt via maximising the log marginal likelihood [15, 58].

2.1.2 Acquisition Functions. Acquisition functions  $\alpha : X \mapsto \mathbb{R}$  measure the expected utility of expensively evaluating f at any location  $\mathbf{x}$ . Its maximiser is chosen as the next location to expensively evaluate. This strategy allows for an estimate of the global optimum to be generated cheaply via the surrogate model, rather than by repeatedly querying the expensive objective function. The probability of improvement (PI) [46] is one of the earliest infill criteria. It is the probability that the predicted value at a location  $\mathbf{x}$  is less than a threshold  $\tau$ , which, if the posterior distribution is Gaussian, can be expressed in closed form as

$$\alpha_{\mathrm{PI}}(\mathbf{x},\tau) = p(f(\mathbf{x}) < \tau \mid \mathcal{D}, \boldsymbol{\theta}) = \Phi(s).$$
(5)

Here,  $\tau$  is usually set to the best solution seen thus far,  $s = (\tau - \mu(\mathbf{x}))/\sigma(\mathbf{x})$  is the predicted improvement normalised by its uncertainty, and  $\Phi(\cdot)$  is the standard Gaussian cumulative density function. Its successor, the expected improvement (EI) [42, 55], is one of the most common acquisition functions and measures the expected positive predicted improvement over a threshold  $\tau$ . It is expressible in closed-form under the same assumptions [42]:

$$\alpha_{\rm EI}(\mathbf{x},\tau) = \sigma(\mathbf{x}) \left[ s\Phi(s) + \phi(s) \right],\tag{6}$$

where  $\phi(\cdot)$  is the standard Gaussian probability density function.

In practice, EI is often preferred because PI tends to be overlyexploitative [16, 42]. However, many other acquisition functions have been proposed, including optimistic strategies such as upper confidence bound [72], information-theoretic approaches [30, 31, 61, 63, 80], and  $\epsilon$ -greedy strategies [8, 16].

### 2.2 Multi-objective Bayesian Optimisation

Real-world optimisation problems often have multiple conflicting objectives, all of which must be minimised at the same time [14]. The multi-objective optimisation problem can be defined as simultaneously minimising  $M \ge 2$  unknown objective functions

$$\min_{\mathbf{x}\in\mathcal{X}}\mathbf{F}(\mathbf{x}) = (f^{1}(\mathbf{x}), \dots, f^{M}(\mathbf{x})), \tag{7}$$

where  $f^i$  is the *i*-th objective value and  $\mathbf{F} : \mathcal{X} \mapsto \mathbb{R}^M$ . Assuming that problem contains conflicting objectives, then there will not be a one unique solution. Instead, possibly infinitely many solutions may exist that trade off between the different objectives. The

trade-off between solutions is often characterised by a *dominance* relationship: **x** is said to dominate  $\mathbf{x}'$ , denoted  $\mathbf{x} < \mathbf{x}'$ , iff:

$$\forall i \in \{1, 2, \dots M\} \left( f^{i}(\mathbf{x}) \leq f^{i}(\mathbf{x}') \right) \land \exists i \left( f^{i}(\mathbf{x}) < f^{i}(\mathbf{x}') \right).$$
(8)

The set containing solutions that optimally trades off between the objectives is referred to as the Pareto set:

$$\mathcal{P} = \{ \mathbf{x} \mid \mathbf{x}' \not\prec \mathbf{x} \land \mathbf{x}, \mathbf{x}' \in \mathcal{X} \}, \tag{9}$$

where  $\mathbf{x}' \neq \mathbf{x}$  indicates that  $\mathbf{x}'$  does not dominate  $\mathbf{x}$ . Computation of  $\mathcal{P}$  is infeasible, due to its potentially infinite size. Thankfully, an approximation is often sufficient, leading to the goal of generating a good approximation  $\tilde{\mathcal{P}}$  to the Pareto set  $\mathcal{P}$ .

Knowles [44] presented the first mono-surrogate approach for multi-objective BO. They proposed to *scalarise* the objective function values  $(f^1, f^2, \ldots, f^M)$ , i.e. mapping them to a single value, via the use of a scalarising function. It uses the randomly-weighted normalised objective values in an augmented Tchebycheff function, drawing weightings from a set of predefined weights to scalarise the sets of objective values. These can then be used in lieu of the objective values, directly within the BO framework (Section 2.1). The location that maximises the expected improvement over the best-seen scalarisation is then chosen to be expensively evaluated.

2.2.1 Scalarisation functions. The augmented Tchebycheff scalarisation was the *de facto* choice for multi-objective BO. However, Rahat et al. [57] recently proposed several scalarisation that outperform it. Subsequentially, Chugh [12] conducted an exhaustive study of scalarisation methods for multi-objective BO, comparing those in [57] to several outside the BO literature. They show that the choosing the best scalarisation method is far from trivial and is, in fact, problem dependent. It is also noted that the ease with which a GP may model a landscape produced by a scalarisation is problem dependent, i.e. the same scalarisation may produce easier or harder landscapes for a GP to model, relative to other methods.

The main properties required for a scalarisation to be used in the BO framework is that it should preserve dominance relationships and allow for every Pareto optimal solution to be reached [36]. In doing so, the maximisation of a dominance-preserving scalarisation should lead to an improvement in the Pareto set [91]. Although many methods have been proposed, we focus on the popular augmented Tchebycheff method [44], two of the best-performing in [57], and a novel scalarisation proposed in this work.

Augmented Tchebycheff (AT). First presented for BO by Knowles [44], the augmented Tchebycheff method combines the Tchebycheff function with a weighted sum, scaled by a small positive value  $\rho$ :

$$g_{\rm AT}(\mathbf{x}, X) = \max_{i} \left[ w^{i} \bar{f}^{i} \right] + \rho \sum_{i=1}^{M} w^{i} \bar{f}^{i}, \tag{10}$$

where  $\bar{f}^i \in [0, 1]$  are the normalised versions of the observed objective values. At each BO iteration, the weights  $\mathbf{w} = (w^1, w^2, \dots, w^M)$ , where  $|\mathbf{w}| = 1$ , are sampled uniformly from a fixed set of evenly distributed weight vectors on the *M*-dimensional unit simplex.

*Hypervolume Improvement (HypI).* The hypervolume indicator  $H(X, \mathbf{r})$  measures the volume of space dominated by a set of solutions  $X = {\mathbf{x}_1, \mathbf{x}_2, ..., \mathbf{x}_t}$  relative to a reference vector  $\mathbf{r}$  [90]. It is a popular choice for comparing two sets of solutions in because

maximising it is equivalent to locating the optimal Pareto set [20]. In order to turn the hypervolume indicator into a scalariser, it is natural to consider the *contribution* of each set member, i.e. the amount of hypervolume gained by including a set member. However, only solutions that reside within the Pareto set of X will have a non-zero contribution, even if solutions dominated by the Pareto set dominate other solutions, they will all be assigned a value of zero. This would hinder the progress of BO because it would create a plateau that lacks spatial information about solution quality.

Rahat et al. [57] proposed a solution to the inherent problems of solely using the hypervolume contribution that they named the hypervolume improvement (HYPI). The members of X are first ranked according to Pareto shells in which they reside. Let the first Pareto shell  $\mathcal{P}^1$  be the Pareto set of X, i.e.  $\mathcal{P}^1 = \text{NONDOM}(X)$ , where NONDOM(X) returns the non-dominated members of X. Subsequent Pareto shells  $\mathcal{P}^{\lambda}$ , with ( $\lambda > 1$ ), can then be defined as

$$\mathcal{P}^{\lambda} = \operatorname{NONDOM}\left(X \setminus \bigcup_{i=1}^{\lambda-1} \mathcal{P}^{i}\right).$$
 (11)

Once the members of *X* have been ranked, the HYPI of a solution **x** is defined to be the hypervolume of the union of **x** and the first Pareto shell  $\mathcal{P}^{\dagger}$  that contains no solutions that dominate it:

$$g_{\mathrm{HYPI}}(\mathbf{x}, X) = H\left(\{\mathbf{x}\} \cup \mathcal{P}^{\dagger}, \mathbf{r}\right).$$
(12)

Therefore, each solution has a non-zero, dominance-preserving scalarisation, with values that provide a gradient towards non-dominated space.

Dominance Ranking (DomRank). An alternative way to compare multi-objective solutions is to count the number of solutions that dominate them, with the idea being that we should prefer solutions that are dominated less [21]. Rahat et al. [57] use this idea to form the DomRank scalarisation:

$$g_{\text{DOMRANK}}(\mathbf{x}, X) = 1 - \frac{|\text{DOM}(\mathbf{x}, X)|}{|X| - 1},$$
(13)

where  $DOM(\mathbf{x}, X)$  returns the set of solutions in *X* that dominate  $\mathbf{x}$ . The authors found that DomRank was performed similarly to other scalarisations, even though, in the degenerate setting, it is possible for all members of *X* to be non-dominated with respect to one another, resulting in equal scalarisations for all *X*. However, this should be expected to happen more frequently as the number of objectives *M* increases because the likelihood of one solution dominating another is inversely proportional to *M* [48].

*Pareto Hypervolume Contribution (PHC).* Inspired by the success of HYPI, we present an alternative scalariser that can directly use the hypervolume contribution of each solution: PHC. Let  $v(\mathbf{x}, \mathcal{P}^{\lambda})$  be a function that calculates the hypervolume contribution of a solution  $\mathbf{x}$  to the shell  $\mathcal{P}^{\lambda}$  it resides in:

$$v(\mathbf{x}, \mathcal{P}^{\lambda}) = H(\mathcal{P}^{\lambda}, \mathbf{r}) - H(\mathcal{P}^{\lambda} \setminus {\mathbf{x}}, \mathbf{r}).$$
(14)

The PHC of a solution  $\mathbf{x}$  can be then be calculated by taking its hypervolume contribution and adding the largest contribution from each subsequent shell:

$$g_{\text{PHC}}(\mathbf{x}, X) = v(\mathbf{x}, \mathcal{P}^{\lambda}) + \sum_{i=\lambda+1}^{N} \max\{v(\mathbf{x}', \mathcal{P}^{i}) \mid \mathbf{x}' \in \mathcal{P}^{i}\}, \quad (15)$$

where *N* is the total number of Pareto shells. If **x** dominates **x'**, then **x'** will be in a subsequent shell to **x** and, by definition,  $g_{PHC}(\mathbf{x}, X) > g_{PHC}(\mathbf{x}', X)$ . Therefore, it preserves dominance relationships between members of *X* and ensures monotonicity between shells.

# 3 BORE: BAYESIAN OPTIMISATION BY DENSITY-RATIO ESTIMATION

BO is a popular and successful strategy for both single- and multiobjective optimisation of expensive black-box problems. However, GPs, the surrogate models of choice in BO, present some notable limitations. Specifically, its  $O(n^3)$  computational complexity in the number *n* of solutions the GP is trained with, as well as additional assumptions which, often, do not apply to the functions that they are trying to model. GPs almost exclusively use stationary kernels, i.e. they only depend on the distance between solutions. This means that they are incapable of modelling different length-scales of data in different regions of X.

Motivated by these shortcomings, Bergstra et al. [3] presented a reformulation of BO by estimating the probability of improvement (5) acquisition function by density-ratio estimation. We note that the authors claimed that the expected improvement was being estimated, but this has since been proven to be incorrect [25, 71]. We start, following the exposition of [25], by considering two densities that depend on a threshold  $\tau = \Phi^{-1}(\gamma)$  that is the  $\gamma$ -th quantile of the observed objective values f, where  $0 < \gamma \leq 1$  and  $\gamma = \Phi(\tau) = p(f \leq \tau)$ :

$$b(\mathbf{x}) = p(\mathbf{x} \mid f < \tau) \tag{16}$$

$$\ell(\mathbf{x}) = p(\mathbf{x} \mid f \ge \tau). \tag{17}$$

Here, *b* is the probability density of observed values being less than the threshold  $\tau$ , and  $\ell$  of observations not being less than  $\tau$ . Using Bayes rule,  $\ell$  and *b* can be expressed as being proportional to PI:

$$b(\mathbf{x}) \propto p(f < \tau)p(\mathbf{x}) = \alpha_{\rm PI}(\mathbf{x}, \tau)p(\mathbf{x}) \tag{18}$$

$$\ell(\mathbf{x}) \propto p(f \ge \tau) p(\mathbf{x}) = (1 - \alpha_{\rm PI}(\mathbf{x}, \tau)) p(\mathbf{x}), \tag{19}$$

where  $p(\mathbf{x})$  is a prior density over X. Bergstra et al. [3] use the ratio of these densities as an acquisition function that monotonically increases with  $\alpha_{PI}(\mathbf{x}, \tau)$ :

$$\alpha_{\rm TPE}(\mathbf{x},\tau) = \frac{b(\mathbf{x})}{\ell(\mathbf{x})} \propto \frac{\alpha_{\rm PI}(\mathbf{x},\tau)}{1 - \alpha_{\rm PI}(\mathbf{x},\tau)},\tag{20}$$

where the prior density  $p(\mathbf{x})$  cancels under the assumption that it has support over all of X. Therefore, maximising  $\alpha_{\text{TPE}}(\mathbf{x}, \tau)$  is equivalent to maximising  $\alpha_{\text{PI}}(\mathbf{x}, \tau)$ .

This transforms the problem training a GP model and maximising PI (5), to one of estimating two densities and maximising their ratio (20). Using tree-based kernel density estimators [67] to estimate  $\ell$  and b, results in the tree-structured Parzen estimator (TPE) [3], and reduces the computational cost to O(n).

Tiao et al. [78] provide a full discussion on the weaknesses of TPE. Here, we highlight the most important. Density estimation in higher dimensions is notoriously difficult [74], and even the estimating the kernel's bandwidth and choosing the correct kernel, is far from trivial [77]. We also note that even with correct density estimation, the optimisation of (20) is often numerically unstable [85].

Instead of trying to first estimate the densities and then their ratio, one can directly estimate their ratio by exploiting results in

class-probability estimation [4, 11, 54, 74]. It can be shown [78] that the  $\gamma$ -relative density-ratio of (16) and (17),

$$r_{\gamma}(\mathbf{x}) = \frac{\ell(\mathbf{x})}{\gamma \ell(\mathbf{x}) + (1 - \gamma)b(\mathbf{x})},$$
(21)

is proportional to the probability of improvement (5):

$$\alpha_{\rm PI}(\mathbf{x}, \Phi^{-1}(\gamma)) \propto r_{\gamma}(\mathbf{x}). \tag{22}$$

We note that the TPE acquisition function (20) is a special case of (21), i.e.  $\alpha_{\text{TPE}}(\mathbf{x}, \tau) \equiv r_0(\mathbf{x})$ .

Next, we introduce a binary class label z such that

$$z = \begin{cases} 1 & \text{if } f < \tau \\ 0 & \text{if } f \ge \tau. \end{cases}$$
(23)

If the class-posterior probability of  $f < \tau$  given an observation **x** is defined to be  $\pi(\mathbf{x}) = p(z = 1 | \mathbf{x})$ , and noting that  $\ell(\mathbf{x}) = p(\mathbf{x} | z = 1)$  and  $b(\mathbf{x}) = p(\mathbf{x} | z = 0)$ , it can also be shown [78] that

$$r_{\gamma}(\mathbf{x}) \equiv \gamma^{-1} \pi(\mathbf{x}). \tag{24}$$

A probabilistic classifier, e.g. a neural network, can be used to estimate  $\pi(\mathbf{x})$  by learning a function  $\pi_{\boldsymbol{\xi}} : \mathcal{X} \mapsto [0, 1]$  with parameters  $\boldsymbol{\xi}$ . Provided that the classifier is trained with a proper scoring rule [26], such as the binary cross entropy (log loss), the relative density-ratio is approximated by

$$\pi_{\boldsymbol{\xi}}(\mathbf{x}) \approx \gamma r_{\boldsymbol{\gamma}}(\mathbf{x}). \tag{25}$$

This result links PI (5) to density-ratio estimation and onto classprobablity estimation, leading to the BO by density-ratio (BORE) framework. It transforms the problem from training a GP and maximising an acquisition function, into one of training a probabilistic classifier and maximising  $\pi_{\xi}(\mathbf{x})$ . To carry out BORE, a proportion  $\gamma$ is first chosen, and the best  $\gamma$ -th proportion of solutions are labelled as class 1, with the remaining labelled as class 0. A classifier  $\pi_{\xi}$  is then trained on the two sets of solutions, and  $\mathbf{x}' = \operatorname{argmax}_{\mathbf{x}} \pi_{\xi}(\mathbf{x})$ is chosen as the next location to be expensively evaluated.

The two main hyperparameters of BORE are the proportion  $\gamma$ of solutions to include in class 1 and the choice of  $\pi_{\xi}$ . Increasing  $\gamma$ encourages exploration because it will result in a worse threshold  $\tau$  from which to (effectively) calculate PI with, i.e. the likelihood of a solution exceeding a worse threshold will be higher. In the original BORE formulation Tiao et al. [78],  $\gamma$  is fixed throughout optimisation. Although it is fixed, exploitation increases as more solutions are evaluated because the class threshold will be more biased towards the better solutions collected during optimisation. The choice of  $\pi_{\mathcal{E}}$  controls the types of functions that can be modelled during optimisation. Multi-layer perceptrons (MLPs) are a natural choice due to their function approximation guarantees [33], their ability to scale to arbitrary problem dimensions, and for being endto-end differentiable. This latter point means that quasi-newton based methods, such as L-BFGS-B [9], can be used to optimise  $\pi_{\mathcal{E}}(\mathbf{x})$ . Ensemble-based methods are an attractive alternative to MLPs because they also scale well. They are not, however, end-toend differentiable meaning that non-gradient-based optimisation methods must be used such as, e.g. CMA-ES [29]. In this work we use MLPs and an ensemble method, gradient-boosted trees (XG-Boost), because they can be both trained using a proper scoring rule, thereby ensuring a good approximation to (25).

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Algorithm 1 Multi-objective BO by Density-Ratio Estimation

**Inputs**: Number of initial samples *S*, total budget *T*, scalarising function  $g(\cdot)$ , probabilistic classifier  $\pi_{\xi}(\cdot)$ , proportion  $\gamma$ . **Steps**:

1:  $X \leftarrow \text{SpaceFillingSampling}(X, S)$ ▶ Generate initial samples 2:  $\mathbf{f}_i \leftarrow \mathbf{F}(\mathbf{x}_i)$  for  $i \in \{1, \ldots, S\}$ ▶ Expensively evaluate samples 3: for  $t = S + 1 \rightarrow T$  do 4:  $g_i \leftarrow g(\mathbf{f}_i)$  for  $i \in \{1, \ldots, t\}$ ▶ Scalarise objective values  $\tau \leftarrow \Phi^{-1}(\gamma)$ 5: ▷ Calculate  $\gamma$ -th quantile of  $\{g_i\}_{i=1}^t$  $z_i \leftarrow \mathbb{I}[g_i < \tau] \text{ for } i \in \{1, \ldots, t\}$  Assign class labels 6:  $\boldsymbol{\xi^{\star}} \leftarrow \text{TrainClassifier}(\pi_{\boldsymbol{\xi}}, \{(\mathbf{x}_i, z_i)\}_{i=1}^t)$ 7: 8:  $\mathbf{x}_{t+1} \leftarrow \operatorname{argmax}_{\mathbf{x} \in \mathcal{X}} \pi_{\boldsymbol{\xi}^{\star}}(\mathbf{x}) \triangleright \operatorname{Maximise class-posterior probability}$ 9:  $\mathbf{f}_{t+1} \leftarrow \mathbf{F}(\mathbf{x}_{t+1})$ ▶ Expensively evaluate new solution

### 4 MBORE: MULTI-OBJECTIVE BORE

In this section, we introduce Multi-objective BO by Density-Ratio Estimation (MBORE). It extends BORE to the multi-objective setting by scalarising the previously-evaluated solutions and separating them into two classes via thresholding the scalarised values, thereby enabling a probabilistic classifier to be trained. In doing so, to the best of our knowledge, we present the first classificationbased multi-objective method using scalarisation for BO. Before discussing MBORE further, we first give an overview of related work in the field of classification-based multi-objective optimisation.

Classification-based multi-objective optimisation methods using surrogate models often focus on predicting the quality of an evolved population of solutions. Two main schemes are proposed, either predicting which Pareto shell a candidate solution would belong to, or whether the current population (set of previouslyevaluated solutions) dominates it. Rank-based support vector machines (SVM) [41, 49], for example, have been proposed [51, 64] to predict the shell a candidate solution belongs to. However, dominance prediction is a much more common strategy. Early works include using a one-class SVM to models whether solutions were dominated [52], and using a naive Bayes classifier to predict whether one solution dominates another [27]. More recently, the direction of research has moved towards learning to predict whether candidate solutions are dominated by an existing set of solutions, for example, by using k-nearest neighbours [89] or MLP classifiers [56, 87].

The classification-based approach we present in this paper, i.e. separating solutions into two classes based on their scalarisation, is conceptually similar to previous work on, e.g., classification based on dominance. However, there are some important differences and benefits from using scalarisation. Most importantly is that using scalarisation and the subsequent thresholding into classes, allows for PI (5) to be calculated via the BORE framework. This provides a much-needed theoretical motivation as to why a classification-based approach is suitable for multi-objective optimisation. The size of the proportion  $\gamma$  gives control over how many solutions are placed into each class. Contrastingly, if a set of solutions were completely non-dominated with respect to one another, using a domination-based approach would result in an empty class.

Motivated by the success of BORE in the single-objective setting, we present MBORE. Algorithm 1 outlines its general structure. It starts (line 1), identically to BO, with a space-filling design such as Latin hypercube sampling [53]. These samples  $X = {\mathbf{x}_i}_{i=1}^S$  are

then expensively evaluated with the objective functions, i.e.  $\mathbf{f}_i = \mathbf{F}(\mathbf{x}_i)$ . At each subsequent iteration of the algorithm, the objective values are scalarised (line 4) and the  $\gamma$ -th quantile  $\tau$  of the scalarised objective values is calculated (line 5). The solutions are then split into two classes (line 6), with solutions that have a scalarisation of less  $\tau$  labelled as class 1, and the remaining labelled as class 0. Next, a probabilistic classifier is trained (line 7), and the location that maximises its prediction is chosen as the next location to evaluate (lines 8 and 9). This process is then repeated until budget depletion.

Given the clear link to PI, and the strengths of MLPs and XGBoost in function approximation, one might expect MBORE to perform similarly to the traditional mono-surrogate approach (BO) on lowdimensional problems (e.g.  $d \le 10$ ) and to improve upon BO on problems that have a larger dimensionality. In the following section we investigate this by comparing BO to MBORE on a wide variety of low- and high-dimensional synthetic and real-world problems.

### 5 EXPERIMENTAL EVALUATION

The performance of MBORE is investigated using two probabilistic classifiers, XGBoost [10] (XGB) and a multi-layer perception (MLP). We compare it to the standard mono-surrogate BO approach [44], often referred to as ParEGO [44], using a GP (2) surrogate model and the EI (6) acquisition function. The methods are compared using two popular synthetic benchmarks, DTLZ [18] and WFG [34], along with a set of real-world problems [76]. Experiments are carried out using the typical number of problem dimensions, e.g.  $d \leq 10$ , with varying numbers of objectives *M*. A high-dimensional, e.g.  $d \in \{20, 50, 100\}$ , version of the WFG benchmark is also evaluated to assess optimisation performance in more challenging scenarios. Experiments are repeated for the scalarisation methods discussed in Section 2.2.1: augmented Tchebycheff (AT) [44], hypervolume improvement (HYPI) [57], dominance ranking (DomRank) [57], and our novel scalariser, Pareto hypervolume contribution (PHC) (15).

The two versions of MBORE (XGB and MLP) were created and trained using the same configurations as BORE [78, Appendix J]. A zero-mean GP with an ARD Matérn 5/2 kernel was used for BO. At each iteration, before new locations were selected, the hyperparameters of the GP were optimised by maximising the log likelihood [58] using L-BFGS-B [9] with a multi-restart strategy [84], and choosing the best set of hyperparameters from 10 restarts for the model. The weight vectors for the AT scalarisation were calculated via the Riesz' *s-Energy* method [7]; see the supplement for more details.

Input variables were normalised to reside in  $[0, 1]^d$  before they were used to train both MBORE and BO. Objective values similarly normalised on a per-objective basis. The models were initially trained on S = 2d observations generated by maximin Latin hypercube sampling [53] and then optimisation was carried out for a further 300 function evaluations. Each optimisation run was repeated 21 times from a different set of Latin hypercube samples. Initial sets of locations were common across all methods to enable statistical comparison. The next location to evaluate, i.e.  $\mathbf{x}' = \operatorname{argmax}_{\mathbf{x}} \pi_{\mathbf{\xi}}(\mathbf{x})$ , was selected for the MLP as in [78], and for XGB using bi-pop CMA-ES [29] with 10 restarts. EI (6) in BO was optimised with a multi-restart strategy of first sampling B = 1024d randomly-chosen locations and then optimising the best 10 of these with L-BFGS-B. The budget for optimising  $\pi_{\mathbf{\xi}}$  for both the MLP and XGB classifiers George De Ath, Tinkle Chugh, and Alma A. M. Rahat

Name	IDs	Problem configurations ( <i>d</i> , <i>M</i> )		
DTLZ	1-7	(2,2), (5,2), (5,3), (5,5), (10,2), (10,3), (10,5), (10,10)		
WFG	1-9	(6,2), (6,3), (8,2), (8,3), (10,2), (10,3), (10,5)		

Table 1: Benchmarks, problem IDs and configurations.

		XGB	MLP	GP
ІДУН	DTLZ	36	4	29
	WFG	37	0	36
DR	DTLZ	21	11	37
	WFG	36	0	45
PHC	DTLZ	36	12	22
	WFG	39	0	36
AT	DTLZ	13	8	43
	WFG	1	0	63

Table 2: Performance summary of MBORE (MLP and XGB) and BO (GP) for a given scalariser on the benchmarks. Table values correspond to the number of times each model was the best or statistically equivalent to the best model.

was also set to *B* to ensure fair comparison. Like Tiao et al. [78], we fix  $\gamma = 1/3$  for all experiments.

Performance is reported in terms of the hypervolume (HV) [91] of the estimated Pareto set; IGD+ [37] is also reported in the supplement. The BO optimisation pipeline was constructed using GPy-Torch [24] and BoTorch [1]. MBORE uses pagmo [5] for carrying out fast dominated sorting and hypervolume calculation, as well as pymoo [6] for the IGD+ calculation. Code for all methods, as well as the initial starting solutions, reference vectors and the estimated Pareto sets for HV and IGD+ is available online<sup>1</sup>.

### 5.1 Synthetic Benchmarks

The popular DTLZ [18] and WFG [34] benchmark problem suites were selected to compare MBORE and BO. The benchmarks were chosen because both the problem dimensionality *d* and the number of objectives *M* are configurable, allowing for a large range of problems to be generated. Table 1 outlines the benchmark problems used, including the specific problem numbers (IDs) and combinations of (*d*, *M*) used from each suite, resulting in 56 and 63 distinct test problems from DTLZ and WFG respectively. Following standard practice [6, 12], the WFG position and scale parameters (*k*, *l*) were set to (4, *d* – 4) for *M* = 2, and (2*M* – 1, *d* – (2*M* – 1)) for *M* > 2.

We start by comparing the performance of MBORE using the XGB and MLP classification models to mono-surrogate BO with a GP surrogate model. Table 2 shows the number of times a model (XGB, MLP, and GP) is best on each benchmark for a given scalarisation method. Specifically, a model is counted as being the best if it has the largest median hypervolume over the 21 optimisation runs, or it is statistically indistinguishable from the best method, according to a one-sided, paired Wilcoxon signed-rank test [45] with Holm-Bonerroni correction [32]. Models that are the best the

<sup>&</sup>lt;sup>1</sup>http://www.github.com/georgedeath/mbore



Figure 1: Hypervolume (HV) performance summary for the DTLZ and WFG benchmarks. Bar heights correspond to the proportion of times that a model and scalariser combination is best or statistically equivalent to the best method.

highest number of times on each benchmark for a given scalariser (table rows) are highlighted in grey. Interestingly, either XGB or GP is the best for each scalarisation method, with results relatively close for HYPI, DomRank (DR) and PHC. This highlights the efficacy of MBORE for general-purpose use in multi-objective BO, and suggests that it should be preferred over BO when using the HYPI and PHC scalarisation methods. Note that MBORE is outperforming BO even though it is approximating PI, an acquisition strategy that is generally regarded as being inferior to EI [16, 25]. We suspect that this is because the increased modelling capacity of XGB is outweighing the marginally worse performance of PI.

Given that Table 2 only compares performance between models for a given scalarisation, we now evaluate which combination of scalarisation and model performs the best on both the DTLZ and WFG benchmarks. Figure 1 summarises the performance for all combinations of model and scalariser. Bar heights correspond to the proportion of times each model and scalarisation combination was the best on each test problem. As can be seen from the figure, MBORE with XGB and our novel PHC scalariser (15) has the best overall performance for both benchmarks. Surprisingly, MBORE with the MLP classification method performs worse than the best performing method on all the 63 WFG test problems. However, this reflects the results presented in [78] that also show that XGB tends outperform MLP. We note that relative rankings of methods remain the same when using IGD+; see the supplement for details.

Next, we investigate the performance of the models with respect to a problem's dimensionality d and number of objectives M. Since PHC was the best-performing method, we limit this discussion to its results. Results for all scalarisers are available in the supplement. Figure 2 summarises the performance of each combination of model and scalariser for the two benchmarks in terms of d and M. The DTLZ results show that BO's performance deteriorates as dincreases, a trend mirrored for the other scalarisations. Conversely,





Figure 2: Hypervolume performance summary for the PHC scalariser on the DTLZ (*left*) and WFG (*right*) benchmarks given the problem's dimensionality (*upper*) and number of objectives (*lower*).



Figure 3: Hypervolume (HV) performance summary for the real-world benchmark problems. Bar heights correspond to the proportion of times that a model and scalariser combination is best or statistically equivalent to the best method.

there is no clear pattern regarding M for DLTZ. In the WFG benchmark the reverse is true, changing d does not result in a consistent trend, but the performance of BO decreases and XGB increases as M increases across the majority of scalarisation methods. We conjecture that this is related to the increased complexity of the WFG problems [34], compared to DTLZ. Therefore, increasing Mleads to an increase in the scalarisation landscape's complexity, resulting in features that GPs model poorly, such as discontinuities.

#### 5.2 Real-world Benchmark

Hand-crafted benchmarks, such as DTLZ and WFG, are known to have properties that are unlikely to appear in real-world applications [38, 88]. Therefore, to investigate the performance of MBORE in more realistic settings, we turn to the real-world benchmark of Tanabe and Ishibuchi [76]. It has 11 continuous-valued test problems taken from real-world problems, such as car side impact design [39] and water resource planning [59]; see [76] for detailed descriptions. The problem *RE3-4-3* was not included due to numerical instabilities; we focus on the remaining 10 problems.

Figure 3 summarises the performance of each combination of model and scalariser on the real-world problem benchmark. As



HV Performance Summary: WFG (High-dimensional)

Figure 4: Hypervolume (HV) performance summary for the high-dimensional WFG benchmark. Bar heights correspond to the proportion of times that a model and scalariser combination is best or statistically equivalent to the best method.

can be seen from the figure, MBORE with XGB and using the PHC scalarisation is by far the best method for optimising the real-world problems based on hypervolume. Interestingly, for DomRank and AT, using BO (GP) was more effective than MBORE, with HYPI and PHC roughly equal between the two methods; see the supplement for full details. These results again show that using MBORE is suitable for multi-objective optimisation and outperforms BO when using more effective scalarisation methods.

#### 5.3 High-dimensional Benchmark

BO often struggles on high-dimensional problems due to the surrogate models of choice, GPs, decreasing in modelling accuracy as the problem dimensionality increases [28]. Consequentially, comparisons of multi-objective methods often focus on lower numbers of both objectives M and d. In contrast to this, and in order to evaluate MBORE in more realistic settings where both d and M are comparatively large, we investigate optimisation performance on the WFG benchmark [34] with a large dimensionality  $d \in \{20, 50, 100\}$ . Due to the computational costs of training GPs in high dimensions, we limit the number of objectives to one configuration (M = 10); see the following section for a comparison of the computational costs.

Figure 4 summarises the performance on these problems. The efficacy of MBORE is repeated for the high-dimesional versions of the WFG problems, with the combination of MBORE, XGB and the PHC scalariser being the best on roughly two thirds of the problems. This is somewhat expected because, in the high-dimensional  $(d \ge 20)$  setting due to the aforementioned difficulties GPs can have with larger problem dimensions. Intriguingly, as shown in the supplement, the comparative performance for MLP-based MBORE increases with *d* for all four scalarisations.

### 5.4 Computational Timing

We also investigate the computational performance of the three methods, MBORE with XGB and MLP, and BO with a GP surrogate model. The cost of performing one iteration of each method was recorded for all scalarisers on the WFG benchmark with  $d \in \{10, 20, 50, 100\}$ . To ensure a fair comparison, each optimisation run was carried out on one core of an Intel Xeon E5-2640 v4 CPU. Figure 5 shows the timing results. The median computation time across all scalarisations for each iteration is shown, with shaded regions corresponding to the interquartile range. As theory necessarily



Figure 5: Computational time taken per iteration on the WFG benchmark for  $d \in \{10, 20, 50, 100\}$ . The median computation time over all scalarisations is shown as the solid lines, with the corresponding interquartile ranges shaded.

dictates [58], the GP's computation time increases with the number of solutions. Conversely, the computation time of XGB and MLP are roughly constant for all iterations for a given dimensionality.

### 6 CONCLUSIONS

In this work, we presented MBORE: a novel multi-objective algorithm for expensive optimisation problems by scalarisation. It replaces the traditional mono-surrogate BO pipeline by training a probabilistic classifier on previously-evaluated solutions that were thresholded into two classes based on their objective value's scalarised form. The classifier's predictions can then be shown to approximate the probability of improving over the given threshold. In addition to MBORE, we also introduce PHC, a dominancepreserving scalarisation method that uses a modified form of each solution's hypervolume contribution to its Pareto shell.

As demonstrated throughout, MBORE provides a strong alternative to mono-surrogate BO using GPs. This is particularly true for more difficult problems, such as the WFG and real-world benchmarks, as well as for problems with higher dimensionalities. Additionally, the computational costs of MBORE remain approximately constant as the number of solutions included in the model increases. We note that, we are not able to recommend MBORE over BO (or vice versa) for an arbitrary scalarisation, as shown by BO being consistently better with the AT scalarisation and MBORE with PHC. However, PHC consistently outperformed all other scalarisation methods. Therefore, we recommend the use of MBORE with XGB using the PHC scalarisation method as the new *de facto* choice for mono-surrogate-based multi-objective optimisation.

Future work includes extending MBORE to non-continuous spaces by using random forests. These are able to naturally adapt to discrete and categorical data without the need for, e.g. one-hot encoding. Additionally, we also seek to extend MBORE to the multisurrogate setting by replacing its multiple GP models with one or more probabilistic classifiers, thereby substantially reducing computational complexity.

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