

Computational Intelligence Modelling of Pharmaceutical Tableting Processes Using Bio-inspired Optimization Algorithms

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Abstract

In pharmaceutical development, it is very useful to exploit the knowledge of the causal relationship between product quality and critical material attributes (CMA) in developing new formulations and products, and optimizing manufacturing processes. With the big data captured in the pharmaceutical industry, computational intelligence (CI) models could potentially be used to identify critical quality attributes (CQA), CMA and critical process parameters (CPP). The objective of this study was to develop computational intelligence models for pharmaceutical tableting processes, for which bio-inspired feature selection algorithms were developed and implemented for optimisation while artificial neural network (ANN) was employed to pre-

dict the tablet characteristics such as porosity and tensile strength. Various pharmaceutical excipients (MCC PH 101, MCC PH 102, MCC DG, Mannitol Pearlitol 200SD, Lactose, and binary mixtures) were considered. Granules were also produced with dry granulation using roll compaction. The feed powders and granules were then compressed at various compression pressures to produce tablets with different porosities, and the corresponding tensile strengths were measured. For the CI modelling, the efficiency of seven bio-inspired optimization algorithms were explored: grey wolf optimization (GWO), bat optimization (BAT), cuckoo search (CS), flower pollination algorithm (FPA), genetic algorithm (GA), particle swarm optimization (PSO), and social spider optimization (SSO). Two-thirds of the experimental dataset was randomly chosen as the training set, and the remaining was used to validate the model prediction. The model efficiency was evaluated in terms of the average reduction (representing the fraction of selected input variables) and the mean square error (MSE). It was found that the CI models can well predict the tablet characteristics (i.e. porosity and tensile strength). It was also shown that the GWO algorithm was the most accurate in predicting porosity. While the most accurate prediction for the tensile strength was achieved using the SSO algorithm. In terms of the average reduction, the

GA algorithm resulted in the highest reduction of inputs (i.e. 60%) for predicting both the porosity and the tensile strength.

Keywords: Tableting, Computational Intelligence; Critical Process Parameters; Predictive modelling; Die Compaction; Bio-inspired Optimization; Artificial Neural Network.

1. Introduction

Over 70% pharmaceutical products are in the tablet form, which is manufactured with die compaction of formulated blends. Most pharmaceutical blends need to go through granulation processes to improve their processabilities. Two types of granulation processes are generally used: wet granulation and dry granulation. In recent years, dry granulation has become one of the primary production processes for pharmaceutical tablets ([1], [2], [3]) due to its distinct advantages for heat and moisture sensitive materials. Roll compaction is a widely used dry granulation technique that consists of two main steps: 1) roll compaction of raw material to obtain ribbons and 2) milling the ribbons into granules. The granules are then used to produce tablets. It is well recognized that the variation in composition and the quality of tablets are determined by material properties and process conditions during

die filling, compaction and ejection [4].

One of the grand challenges in pharmaceutical development is to identify the causal relationship between material properties, intermediate properties, process variables and final product properties[5]. For example, it is well known that comparing tablets produced by direct compression, the tablets made with granules produced using roll compaction generally have lower tensile strengths, i.e. there is a loss in tableability [6], [7]. Several attempts were then made to explain this behavior. For example, Malkowska and Khan [8] proposed the concept of work hardening that was defined as the increase of resistance to permanent deformation of a material with the amount of deformation and argued that work hardening led to the observed reduction in tableability of pharmaceutical excipients. Sun and Himmelpach [9] showed that tableability of MCC powders reduced with increasing granule size, and suggested that granule size enlargement was primarily responsible for the reduction in tableability, because larger granules tend to pack less efficiently due to smaller binding/contact areas, resulting in a reduced tensile strength of the tablets. This illustrates that many factors (i.e. features) play a role in determining the quality of final products. However, the contribution of each feature varies from process to process, and from product to product. It is

of practical and scientific importance to identify the dominant ones, i.e. to select the dominant feature in formulation design and product development.

Feature selection is very important in many engineering practices, due to the abundance of noisy, irrelevant, or misleading features. In the development of predictive models, the redundant and irrelevant features could degrade the model performance during the learning process [10]. Feature selection is particularly useful when the number of features is large and not all of them are required for describing the data and for further exploring the data features in experiments [11]. This process leads to reduction in the dimensionality of feature space for a successful prediction task. Feature selection also helps understanding data, reducing computation requirements, and reducing the effect of the curse of dimensionality [12]. The selected features will improve the prediction model performance and provide a faster and more cost-effective prediction while maintaining the predictability (i.e. accuracy) [13]. Identification and selection of the relevant features is, however, a complex problem. Feature selection is considered as a multiobjective task that minimizes both the selected features and the prediction error. These two objectives are normally contradicted and the optimal solution needs to be sought in the presence of a tradeoff between them [14]. Hence a robust

optimisation technique becomes essential in feature selection.

For optimisation, various heuristic techniques mimicing the social behaviour of biological and physical systems of insects, birds, animals, fish in nature were proposed [15]. For instance, genetic algorithm (GA) was the first evolutionary algorithm introduced in the literature that mimics the natural evolution process of a population of initial individuals [16]. Particle swarm optimization (PSO) was one of the well-known swarm algorithms based on the movement and the social behaviour of birds within a flock [17]. Artificial bee colony (ABC) was a numerical optimization algorithm based on the foraging behavior of honeybees [18]. A virtual bee algorithm (VBA) was developed to optimise the numerical function in 2-D using a swarm of virtual bees, which move randomly in the search space and interact to find food sources [19] [20]. Artificial fish swarm (AFS) algorithm was introduced to mimic the stimulant reaction by controlling the tail and fin [21].

However, the application of these optimisation techniques in feature selection is still very limited for complex pharmaceutical development and manufacturing, where a feature is a measurable property contributing to the process performance and product quality, and the number of features involved is generally very large, even though the contributions of some features are very

small. Therefore, the purpose of this study is to develop predictive CI models for the tablet manufacturing process with pharmaceutical powders and roll-compacted granules, for which various bio-inspired optimization algorithms are employed for feature selection and an artificial neural network is applied to predict tablet properties, such as porosity and tensile strength. The efficiency of these bio-inspired optimization algorithms for feature selection will be evaluated for maximising feature reduction (minimizing the number of selected features, or identifying the most important features) while obtaining comparable or even better prediction.

2. Bio-inspired optimization algorithms

In this study, seven bio-inspired optimisation algorithms are considered: 1) Grey Wolf Optimization (GWO), 2) Bat Algorithm (BAT), 3) Cuckoo Search (CS), 4) Flower Pollination Algorithm (FPA), 5) Social Spider Optimization (SSO), 6) Genetic Algorithm (GA), and 7) Particle Swarm Optimization (PSO). For completeness, a brief introduction of these algorithms is presented in this section, together with a brief description of the artificial neural network.

2.1. Grey Wolf Optimization (GWO)

Grey wolf optimization (GWO) is a new evolutionary computation technique that is successfully applied in many optimization tasks but still suffers from the lack of fast and global convergence. Grey wolves manifest a social behaviour - living in groups or packs (of 5 to 12 on average) with a very rigid hierarchy made of four different classes [22]:

1. The leaders (denoted as α wolves), consisting of one male and one female wolves. They are responsible for making necessary decisions on all pack activities, such as hunting, resting, and travelling. All other wolves in the pack obey their decisions.
2. *Beta* (β) wolves. They help the α wolves in decision-making and other group actions, and are the best candidates to be elected as the next α wolf if a α wolf dies or becomes too old.
3. *Delta* (δ) wolves. They are obeying the α and β wolves. There are various types of *delta* wolves with different duties, such as *hunters*, *sentinels*, *Scouts*, and *caretakers*.
4. *Omega* (ω) wolves. They are considered as the lowest ranked wolves and have to obey the different dominant wolves.

Each grey wolf decides its position and movement to better areas. GWO is incorporated with a risk probability that mimics the events of wolves crashing with their foes. Moreover, each grey wolf has a particular sensing (coverage) range, indicated as the visual distance. In each iteration t , the new positions X of the wolves are determined using equations (1) and (2) [53]. The best fitted solution is assigned to α , the second-best solution to β , and the third-best to γ . All other solutions are assigned to ω .

$$\vec{X}(t+1) = \vec{X}_p(t) + \vec{A} \cdot \vec{D}, \quad (1)$$

$$\vec{D} = |\vec{C} \cdot \vec{X}_p(t) - \vec{X}(t)|, \quad (2)$$

where \vec{D} is the fitness value of each hunt agent, \vec{A} , and \vec{C} demonstrate coefficient vectors are calculated as equations (3) and (4), \vec{X}_p depicts a prey position, and \vec{X} represents a wolf position.

$$\vec{A} = 2\vec{A} \cdot \vec{R}_1 - \vec{a}, \quad (3)$$

$$\vec{C} = 2\vec{R}_2, \quad (4)$$

where \vec{a} is the updating parameter, R_1 and R_2 are the random vectors with a value in $[0, 1]$.

The α , β , and δ wolves are defined, imposing the other agents to upgrade their positions accordingly. Thus, the wolves' positions are updated as

$$\begin{aligned}\vec{D}_\alpha &= |\vec{C}_1 \cdot \vec{X}_\alpha - \vec{X}|, \\ \vec{D}_\beta &= |\vec{C}_2 \cdot \vec{X}_\beta - \vec{X}|, \\ \vec{D}_\delta &= |\vec{C}_3 \cdot \vec{X}_\delta - \vec{X}|\end{aligned}\tag{5}$$

$$\begin{aligned}\vec{X}_1 &= |\vec{X}_\alpha - \vec{A}_1 \cdot \vec{D}_\alpha|, \\ \vec{X}_2 &= |\vec{X}_\beta - \vec{A}_2 \cdot \vec{D}_\beta|, \\ \vec{X}_3 &= |\vec{X}_\delta - \vec{A}_3 \cdot \vec{D}_\delta|\end{aligned}\tag{6}$$

$$\vec{X}(t+1) = \frac{\vec{X}_1 + \vec{X}_2 + \vec{X}_3}{3},\tag{7}$$

where t is the iteration number. α , β and δ are assigned with random numbers in $[0, 1]$.

2.2. Bat Algorithm (BAT)

BAT algorithm is a meta-heuristic technique for solving complex optimization problems. Bats use echolocation behavior for seeking the prey and detect/avoid the obstacles. The bats use time delay in the reflection of the emitted loud sound pulse from the surrounding object for navigation in dark [24]. And the magnitude of the emitted sound fluctuates from a high value when looking at the prey and a low value when flying near their prey [25]. The bats' positions can be determined from [26], [27]:

$$F_i = F_{min} + (F_{max} - F_{min})\beta, \quad (8)$$

$$V_i^t = V_i^{t-1} + (X_i^t - X^*)F_i, \quad (9)$$

$$X_i^t = X_i^{t-1} + V_i^t, \quad (10)$$

where β indicates the random vector defined in the uniform distribution, X^* denotes the best position obtained so far, F_{min} outlines the minimum frequency, F_{max} shows the maximum frequency, and V_i depicts the speed vector. Moreover, a local exploration is performed by employing the random movement as follows

$$X_{new} = X_{old} + \epsilon A^t, \quad (11)$$

where ϵ is a random number in $[-1, 1]$, and A^t represents a bat loudness at the t time and is given as

$$A_i^{t+1} = \alpha A^t, \quad (12)$$

where α is a constant that can be experimentally determined, R_i controls the BAT local search and is determined by:

$$R_i^{t+1} = R_i^0 [1 - \exp(-\gamma t)], \quad (13)$$

where R_i^0 denotes the first pulse emission which has a value larger than 0.

2.3. Cuckoo Search (CS)

Cuckoo search (CS) is a heuristic search technique inspired by the cuckoos reproduction strategy [28]. Cuckoos replace their eggs in the nests of other host birds that may be of different kinds. Once the host bird notices that the eggs are not its own, it will either crush the egg or leave the nest to another. Various studies showed that the flight behavior of many animals and insects

had the typical characteristics of Lèvy flights. Lèvy flights provide a random walk while their random steps are drawn from a Lèvy distribution for large steps [28]. In the CS optimisation, a new solution X_i is based on Lèvy flight and given as

$$X_i^{t+1} = X_i^t + \vartheta \oplus Lèvy(\beta), \quad (14)$$

where ϑ represents the step size associated with the scale problem that is set to 1 in most of the cases, \oplus indicates entry wise multiplications.

Previous studies on the flight behaviour of different animals and insects [29]. Lèvy flights for large moves could be defined as

$$Lèvy \sim u = t^{-\lambda}, (1 < \lambda \leq 3). \quad (15)$$

Hence, the successive jumps of the cuckoo form a random walk, and the Lèvy walk should generate some new solutions close to the best solution, which will advance the local search process [28]. The CS local search can be formulated as

$$X_i^{new} = X_i^{old} + 2 * r * (X_i^{old} - X_{best}), \quad (16)$$

This can be applied to obtain new cuckoo solutions based on Equation

(14) [28]. In Equation (16), r indicates the random number picked from the Lèvy distribution, X_{best} denotes the current best solution, X_i^{old} represents an old solution, and X_i^{new} demonstrates a newly generated solution.

The strength of the CS lies in the procedure to discard the wrong solution, so a fraction pa of solutions are ignored, and the updated solutions are obtained by

$$X_i^{new} = X_i^{old} + rand_1 * (rand_2 > pa) * (X_a - X_b), \quad (17)$$

where X_i^{new} is the new nest (solution), X_i^{old} is the old nest to be neglected, $rand_1$ and $rand_2$ represent two random numbers drawn from a uniform distribution, pa denote the probability of finding the nest, X_a and X_b are two randomly selected of the current nests [28].

2.4. Flower Pollination Algorithm (FPA)

Flower pollination algorithm (FPA) is inspired from the flower pollination process of plants that dictates the ultimate reproduction. FPA is typically related to the pollen transfer by pollinators [30]. Pollination is normally carried out in two modes: *cross pollination* (global search) and *self pollination* (local search) , which are described in detail as follows [31]:

1. *Cross pollination* is referred to as the transfer of the pollen of flowers of different plants separated by a long distance by pollinators that can fly a long distance (i.e. also known as global pollination) [31]. In the cross pollination, it is the pollinators that assure the pollination and proliferation of the optimal solution g_* . The initial rule may be formulated as

$$X_i^{t+1} = X_i^t + L(X_i^t - g_*), \quad (18)$$

where X_i^t represents the vector of a i solution at t iteration, g_* demonstrates the present best solution, and L describes the pollination strength that randomly pulled from the Lèvy distribution.

2. *Self pollination* is defined as the implantation of one flower from the pollen of the identical flower or different flowers of the identical plant, which usually happens when there is no pollinator available. The local pollination and flower constancy are expressed as

$$X_i^{t+1} = X_i^t + \epsilon(X_j^t - X_k^t), \quad (19)$$

where X_j^t and X_k^t demonstrate two random solutions, and ϵ denotes the uniform distribution function.

Since local pollination can be substantial, denoted by the fraction (p), in the aggregate pollination actions (in this study, it is assumed that $p = 0.5$). A switching probability $p \in [0, 1]$ controls the local and global pollination.

2.5. *Social Spider Optimization (SSO)*

Social spider optimization (SSO) is a population-based algorithm and one of the swarm algorithms recently proposed by Cuevas et. al [33]. SSO algorithm is extracted from the social behaviour of the spider's colony in nature. A SSO algorithm consists of two main components: *social members* and *communal web*. Social members are divided into male and female spiders. The social behaviour and cooperative interaction depend on the spider gender. The number of female spiders accounts for at least 65% of the total colony members. Female spiders present an attraction or dislike to other spiders according to their vibrations that are circulated through the communal web. More details on the SSO implementation can be found in references [34], [35], [36]. The SSO algorithm consists of the following steps:

1. ***Population initialization.*** The most important property of the SSO is the tendentious female population. The number of females N_f

is randomly chosen within the range [65% - 90%], the number of male spiders N_m is then calculated by

$$N_f = \text{floor}[(0.9 - \text{rand} * 0.25).N], \quad (20)$$

$$N_m = N - N_f, \quad (21)$$

where $\text{floor}[x]$ function gives the largest integer less than or equal to x . Thence, the complete population S consists of N spiders, which is then splitted into two categories female (F) and male (M). The F group contains the female members ($F = f_1, f_2, f_3, \dots, f_{N_f}$), whereas M group contains the male members ($M = m_1, m_2, m_3, \dots, m_{N_m}$).

The female spider's position f_i and the male spider's position m_i are randomly selected between the defined initial parameters (lower limit p^{low} and upper limit p^{high}) as follows

$$f_{i,j}^t = p_j^{low} + \text{rand} * (p_j^{high} - p_j^{low}), \quad (22)$$

$$m_{i,j}^t = p_j^{low} + \text{rand} * (p_j^{high} - p_j^{low}), \quad (23)$$

where i , and j are the parameter and individual indexes respectively, $rand$ is a random number generator between [0 and 1].

2. ***Fitness evaluation.*** In the SSO algorithm, every spider (individual) has a weight w_i that defines the solution goodness, a fitness value of any spider i is calculated as

$$w_i = \frac{J(s_i) - s_{worst}}{s_{best} - s_{worst}}, \quad (24)$$

where $J(s_i)$ represents the obtained fitness of the spider s_i position, the values of s_{worst} and s_{best} describe the worst and the best fitness values (minimization problem) as defined by

$$s_{best} = \max_{i \in [1, 2, \dots, N]}(J(s_i)), \quad (25)$$

$$s_{worst} = \min_{i \in [1, 2, \dots, N]}(J(s_i)), \quad (26)$$

3. ***Vibration modelling in the communal web.*** Vibration modelling is utilized to transfer the data between the colony members. The encoded waves rely on the weight and distance of each spider [35].

4. ***Mating.*** It is a vital process to ensure the colony survival and allow the information exchange between all the members. Social spider mating is achieved between the dominant male and female members in the colony. However, the dominant male spider determines the location of one or several female members within a particularized range. Thereafter it mates with them to produce offspring [35].

2.6. Genetic Algorithm (GA)

Genetic algorithm (GA) mimics the natural evolution process of a population of initial solutions (individuals) [37]. Some individuals undertake crossover and mutation operations to produce better individuals (offsprings) that become the next generation of the population. To determine which individuals should participate in the crossover and mutation, a selection process takes place to select the fittest individuals according to a predetermined fitness function [37].

The crossover operation randomly chooses pairs of these selected individuals to breed. The mutation of some individuals keeps diversity among the population. The larger the population size, the higher the probability to reach a better solution. However, evaluating the fitness of all individuals in a large population is very computer-intensive and needs tremendous

computational resources.

2.7. Particle Swarm Optimization (PSO)

Particle swarm optimization (PSO) is one of the well-known swarm technique based on the movement and the social behaviour of birds within a flock. In PSO, each solution is considered as a particle with specific characteristics (position, fitness, and a speed vector), which defines the motion of the particle [17]. PSO is widely used for optimization and feature selection. PSO uses a number of particles (N) that constitute a swarm moving in the search space, looking for the optimal solution [38]. In the PSO algorithm, the position of a particle is determined as [17]

$$X_{i,d}^{t+1} = X_{i,d}^t + V_{i,d}^{t+1}, \quad (27)$$

where X is the particle position vector, V is the particle velocity vector, i is the index of a particle, t is the time or iteration counter, d is the dimension, $X_{i,d}^t$ is the position of particle i at iteration t in the d dimension, $X_{i,d}^{t+1}$ is the new position of particle i at iteration $t + 1$ in the d dimension, $V_{i,d}^{t+1}$ is the velocity of particle i at iteration $t + 1$ in the d dimension and is given as

$$V_{i,d}^{t+1} = w * V_{i,d}^t + c_1 * rand_1^t (pbest_{i,d}^t - X_{i,d}^t) + c_2 * rand_2^t (gbest_d^t - X_{i,d}^t), \quad (28)$$

where w is the inertia weight factor, $gbest_d^t$ is the most optimist position of the swarm at time t along dimension d , c_1 and c_2 are parameters representing loyalty and selfishness of particles. Normally, $c_1 = c_2 = 2$, while $rand_1^t$ and $rand_2^t$ are random numbers distributed uniformly over $[0, 1]$.

2.8. Artificial Neural Network (ANN)

Artificial neural networks (ANN) are regarded as generalizations of mathematical models of biological nervous systems. In ANN, the effects of the synapses are described by weighted connections that modulate the effect of the associated input signals, and a transfer function that represents the non-linear characteristic exhibited by neurons. The neuron impulse is then determined as the weighted sum of the input signals transformed by the transfer function.

The learning ability of ANN is performed by adapting the weights using the chosen learning algorithm. The behaviour of the neural network depends mainly on the interaction between the different neurons. The basic architecture consists of three types of neuron layers: input, hidden, and output [40].

Various ANN architectures can be created according to the characteristics and specification of the applications [41]. The feed-forward ANN is one of the commonly used architectures, in which the signal flow is from input to output units strictly in a feed-forward direction and the data is processed over multiple units without feedback connections.

3. The computational intelligence model

In this study, the artificial neural network (ANN) is used as a regression model to evaluate the final prediction performance, while seven bio-inspired optimization algorithms (GWO, BAT, CS, FPA, SSO, GA, and PSO) described above are incorporated in the ANN for feature selection, as illustrated in Figure 1. The proposed optimization algorithms for feature selection work in a wrapper-based manner. The central point of the wrapper methods is the use of ANN as regression to ensure the quality of selected features during the feature selection process.

Each optimization algorithm is run for 20 times to test the convergence capability. The performance of the optimization algorithms are evaluated using the following indicators:

1. *Average feature reduction (R^*)* defines the mean size of the reduced

features to the aggregate amount of features as follows

$$R^* = \left(1 - \frac{\text{size}(g_*^i)}{N_t}\right) \times 100, \quad (29)$$

where g_*^i is the best solution that obtained in the i -th application of the algorithm, N_t represents the total number of features in a given dataset. For example, the dataset used in this study has 3 features, *i.e.* $N_t = 3$, and if the best solution selects 2 features, then $R^* = (1 - \frac{2}{3}) * 100 = 33.3$

2. *Mean square error (MSE)* measures the average of squared errors between the actual output and predicted ones and is given as

$$MSE = \frac{\sum_{i=1}^n (P_i - O_i)^2}{n}, \quad (30)$$

where O_i and P_i are the observed and predicted values respectively, n is the total number of samples, and i denotes the i -th number of sample in dataset.

3. *Standard deviation (Std)* measures how much the sets differ from the mean value. Std represents the optimizer convergence to the same optimal and ensures repeatability of the results [42].

A generic representation of the fitness function assessing both regression performance and feature reduction is also introduced as

$$f_{\theta} = \alpha * E + (1 - \alpha) \frac{\sum_i \theta_i}{N}, \quad (31)$$

where f_{θ} is the fitness function of a vector θ with N elements of value 0 or 1, representing unselected or selected features, E is the prediction error, and α is a constant controlling the importance of regression performance to the number of features selected and balancing the trade-off between exploration and exploitation. Normally, at the beginning of optimization (α) has its maximum value to allow for maximum exploration, while, at the end of optimization, it has the minimum value for more exploitation of search space.

Each bio-inspired algorithm is initialized using n random agents (solutions) with each agent representing a given selected feature combination. Then each algorithm is iteratively applied for a number of iterations aiming to converge to a good solution. The individual solution is represented as a continuously valued vector with the same dimension as a number of attributes in the given dataset. The solution vector values are normalised so that their values are in the range $[0, 1]$. For the solution fitness function evaluation, these solutions are converted to their binary representations by

$$y_{ij} = \begin{cases} 0 & (x_{ij} < 0.5) \\ 1 & (x_{ij} \geq 0.5) \end{cases} \quad (32)$$

where x_{ij} is the continuous value of the solution number i in dimension j , and y_{ij} is the discrete representation of solution vector x .

4. Experimental

This study focuses on powder compaction, a typical manufacturing process used for a wide range of products, such as pharmaceutical tablets and catalyst pellets. A variety of widely used pharmaceutical excipients [43] are considered, including micro-crystalline cellulose of different grades: Avicel PH-101, Avicel PH-102 and DG (FMC, Biopolymer, USA); (Pearlitol 200 SD, Roquette, UK), and Lactose (Granulac 140, Meggle GmbH, Germany). Moreover, binary mixtures of MCC 102 and lactose with various mass fractions are also produced using a mixer (TURBULA T2F, Wab, UK). In total, three binary mixtures (see Table 1), named mixture 1, 2 and 3, based on their compositions are considered, for which the mass fractions of MCC 102 are 25%, 50% and 75%, respectively. The samples were mixed for 15 minutes

at a consistent speed of 34 min⁻¹.

Furthermore, dry granulation using roll compaction is performed to produce granules from feed powders and their mixtures. Ribbons are produced using a custom-made gravity fed roll compactor with a roll gap of 1.2 mm and roll speed of 1 rpm (for which, two smooth rolls of 200 mm in diameter and 46 mm in width are used), and are then milled using a cutting mill (SM 100, Retsch, Germany) at a speed of 1,500 rpm, for which a screen size of 4 mm is used. The produced granules are then sieved using a sieving tower into different granule size classes (0-90, 90-250, 250-500, 500-1000, 1000-1400, 1400-2360 μ m), for which the upper size limit is used to represent the granular size.

The feed powders, their mixture and granules are then compressed at various compression pressures to produce tablets of various porosities, for which the corresponding tensile strengths are measured. Cylindrical tablets are produced with an Instron universal testing machine equipped with a 30 kN load cell. A die of 11.28 mm of diameter is used. A powder mass of 300 mg is manually poured into the die and compressed to different maximum pressures at the room temperature (23°C, 45±2% RH) and a compaction speed of 5 mm/min. Diametrical compression tests on the produced tablets

are performed using a diametric strength tester (Erweka TBH30, Erweka GmbH, Germany), in which the the crushing load is determined. The tablet tensile strength (σ_t) is then calculated according to [44]

$$\sigma_t = \frac{2 * F}{\pi * D * H}, \quad (33)$$

where F is the maximum load required to break the tablet, D and H are the tablet diameter and height, respectively. All experiments are run in triplicate.

5. Results and discussion

5.1. Dataset generation

A large dataset (see supplement) is then created using the material, particle/granule size and compaction pressure as the inputs (i.e. 3 inputs), while the porosity and the tensile strength of the produced tablets as independent outputs (i.e. 2 outputs), as exemplified in Table 1. It is worth noting that the tensile strength and tablet porosity are generally inter-related (e.g. the lower the porosity, the higher the tensile strength) for tablets made of feed powders and their mixtures. However, this inter-relationship does not necessarily hold for roll-compacted granules, as discussed in the Introduction section. Hence

these two tablet properties are treated as independent outputs in this study in order to explore if the CI models can identify these behaviours from the data produced with both feed powders and granules.

The legend used to identify the different materials for the CI model is specified as follows:

- Material 1 = MCC PH 101 powder.
- Material 2 = MCC PH 102 powder.
- Material 3 = MCC DG powder.
- Material 4 = Mannitol Pearlitol 200 SD powder.
- Material 5 = MCC PH 101 Granules.
- Material 6 = Binary mixture: 75% MCC102 + 25% Lactose.
- Material 7 = Granules binary mixture 75% MCC102 + 25% Lactose.
- Material 8 = Binary mixture: 50% MCC102 + 50% Lactose.
- Material 9 = Granules binary mixture 50% MCC102 + 50% Lactose.
- Material 10 = Binary mixture: 25% MCC102 + 75% Lactose.
- Material 11 = Granules binary mixture 25% MCC102 + 75% Lactose.

5.2. Model construction

Two-thirds of the dataset are randomly chosen to train the models (*training set*) and the remaining part (*testing set*) is used to test the model prediction. The *training set* is used to evaluate the ANN throughout the optimization process with the bio-inspired algorithms and used in the prediction model. The *testing* data is kept hidden from the optimization and only used during the prediction process. The optimisation process is run 20 times with each algorithm in order to get an average performance for the prediction error and accurately assess statistical evaluation indicators. The bio-inspired optimization algorithms are used for feature selection, so that only the most significant features are fed into the ANN.

In the CI model, ANN is used for the regression purpose (prediction of a continuous output), and two approaches are created:

1. A model is used to evaluate all the possible solutions during the bio-inspired feature selection process (i.e. this ANN regression model is used to ensure the goodness of the selected features). At this stage, ANN is utilised for the bio-inspired optimization algorithm to reach the optimal solution (best-selected features).
2. A predictive model is established for the testing data using the selected

features. At this stage, the model is built using the inputs selected by the bio-inspired optimization algorithm as the train set.

5.3. Model prediction

The results presented in Figures 2 and 3 show the MSE values using each optimizer for 20 different runs, while Figure 6 shows the average feature reduction of the two outputs. It can be seen that the GWO algorithm was the most accurate in predicting porosity, while the SSO algorithm achieved the most accurate prediction for the tensile strength. In addition, the GA algorithm led to the highest reduction of features - 60% - with an average MSE of 7.2 for predicting porosity and 5.1 for predicting tensile strength.

Figures 4 and 5 show the standard deviation values obtained by all the bio-inspired algorithms. The minimum value of the *std* measure is obtained by GWO for the porosity as shown in Figure 4 and obtained by SSO for the tensile strength as shown in Figure 5. Furthermore, it is clear that the GWO and SSO algorithms have in general standard deviation (*std*) less than the ones obtained from the GA and PSO algorithms, which indicates the capability of GWO and SSO to converge to optimal or near-optimal solutions. In Figures 7 and 8, the majority (9 from 14) of the seven bio-inspired algorithms identify the compaction pressure as the most importance

input for both porosity and tensile strength as shown in Figures 7 and 8.

Overall, it can be concluded that the GWO algorithm obtains the best compromise between MSE, the standard deviation, and feature reduction for predicting both the porosity and the tensile strength. Each optimizer is run for 20 different runs to ensure convergence capability.

The leader selection for a given swarm has a very great impact on the explorative/exploitative ability of each optimizer. The GWO optimizer keeps track of the best three solutions found, but the SSO optimizer has track of the N best solutions found. Therefore, the GWO and SSO bio-inspired algorithms perform differently on the same dataset. GWO performs better in predicting porosity in some runs and SSO performs better in predicting porosity in other runs. But over the 20 runs, the average performance of the GWO algorithm was the most accurate in predicting porosity and the average performance of the SSO algorithm achieved the most accurate prediction for the tensile strength. The different performances of the various bio-inspired optimization algorithms are primarily due to the intrinsic nature in modelling, as detailed in the recent publications [45], [46], [47] [48], [49], [50], [51], [52], [53], [54].

5.4. Model Prediction

Figure 7 shows the comparison between the experimental results and predicted results from the SSO model. This model was chosen here for demonstration because it is the most accurate algorithm for prediction of tensile strength with an average MSE (1.375) and average reduction (20%). More specifically, Figure 7 presents the tensile strength of the tablets as a function of compaction pressure for pure powder (a), Granules (b) and mixtures (c). It is shown that there is an increase in tensile strength with the increase of compaction pressure for all the materials and granules sizes investigated. Moreover, the prediction for pure powders as MCC PH 101 (Figure 9-a) and mixtures (Figure 9-c) are generally more accurate than the granule tensile strength predictions (Figure 9-b) where more scattered results were observed. Similarly, Figures 10, 11, and 12 show the comparison between compaction experimental results and prediction using the GWO due to its highest accuracy (MSE of 4.832) for porosity prediction as discussed in the previous section (5.2). There is an almost exponential decrease of porosity with the increase of compaction pressure. Interestingly, the GWO gives almost identical values to the measured ones for all the materials with no distinction between pure powders (Figure 10), mixtures (Figure 11) or granules (Figure

12). This proves that the GWO model can predict the tablet porosity during powder compaction for a wide range of materials with high accuracy.

6. Conclusions

The robustness of CI models that integrate artificial neural network (ANN) with bio-inspired feature selection algorithms for predicting tablet manufacturing processes was evaluated. In particular, tablet properties such as porosity and tensile strength were predicted based upon powder characteristics. Seven bio-inspired optimization algorithms for feature selection were applied. The modelling efficiency was evaluated in terms of the average feature reduction and mean square error. It was found that the GWO algorithm was the most accurate in predicting porosity with equal accuracy for pure powders, mixtures, and granules, while the most accurate prediction of the tensile strength can be achieved using the SSO algorithm, in particular, the values for pure powders and mixtures were more accurately predicted than the granule tensile strength. Regarding the average feature reduction, GA obtained the highest reduction for predicting both the porosity and the tensile strength outputs, and could be more useful for identification of key features or the critical material attributes. Moreover, it was shown from the results

obtained with various optimisation algorithms that the most significant feature is the compaction pressure for both tensile strength and porosity, which is in broad agreement with the experimental observations reported in the literature.

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Table 1: Example of dataset generated for the model

Inputs			Outputs	
Material	Granule size (m)	Compaction pressure (MPa)	Porosity (%)	Tensile Strength (MPa)
1	59	412.500	7.199	15.786
1	59	406.100	7.333	15.354
1	59	384.800	8.547	15.266
1	59	355.600	8.551	15.100
..
5	500	268.679	8.095	11.081
5	500	159.511	12.582	7.803
5	500	78.139	23.237	3.133
..
7	500	140	18.336	3.728
7	500	140	18.515	3.810
7	800	20	42.772	0.203

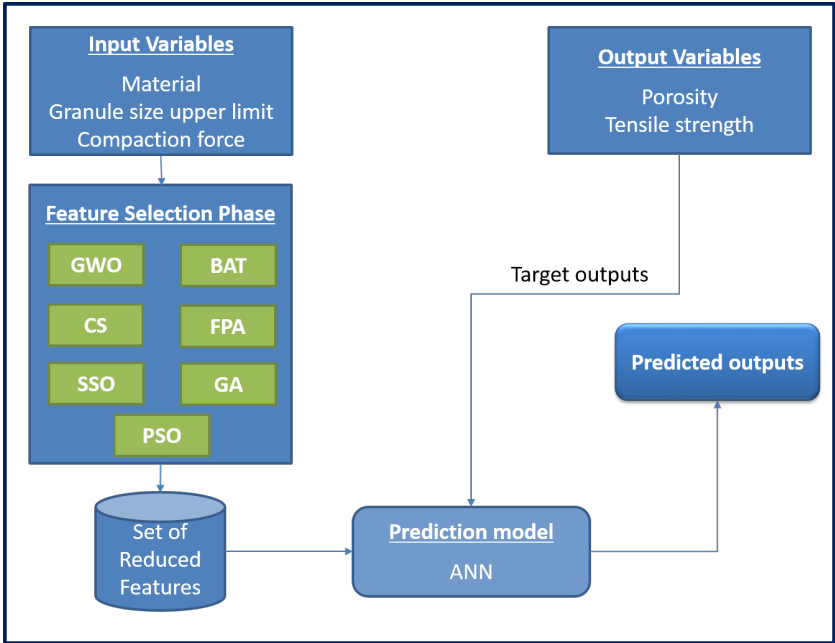


Figure 1: An overview of the proposed model

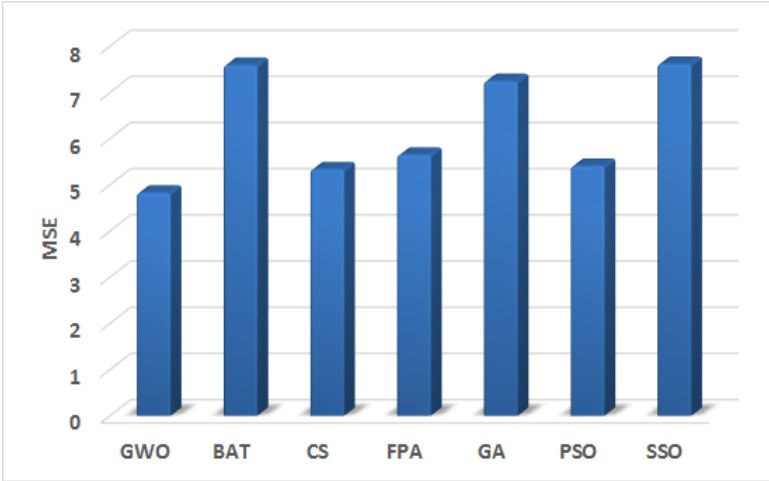


Figure 2: Average MSE for porosity

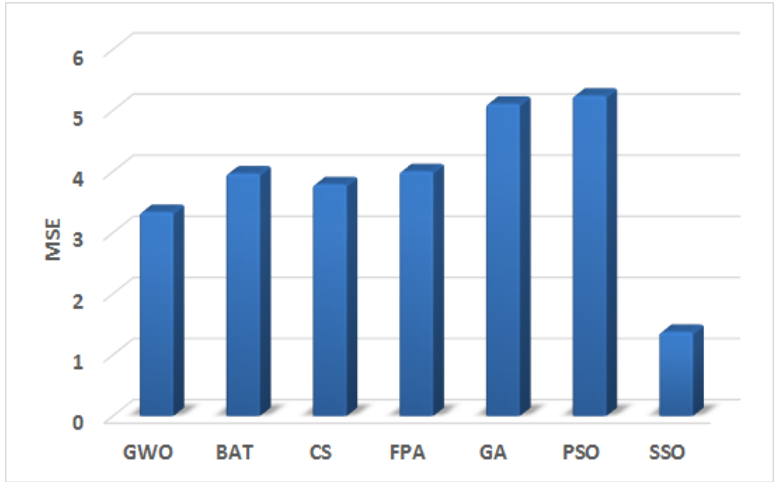


Figure 3: Average MSE for tensile strength

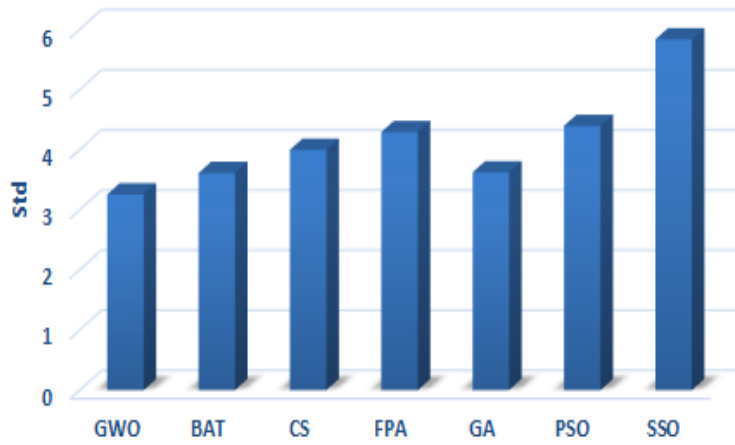


Figure 4: Average Std for porosity

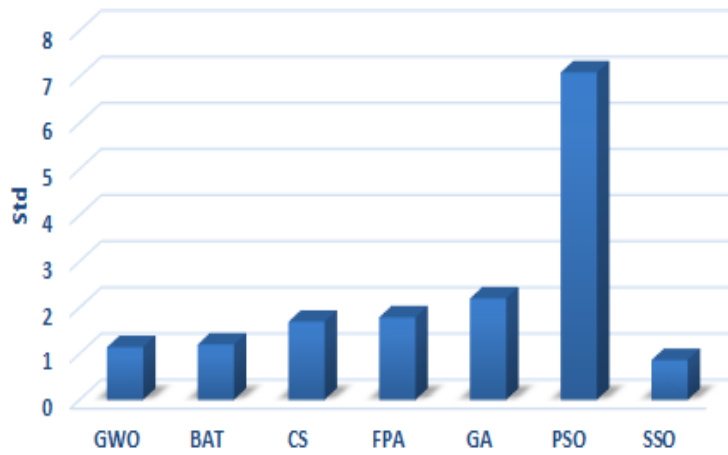


Figure 5: Average Std for tensile strength

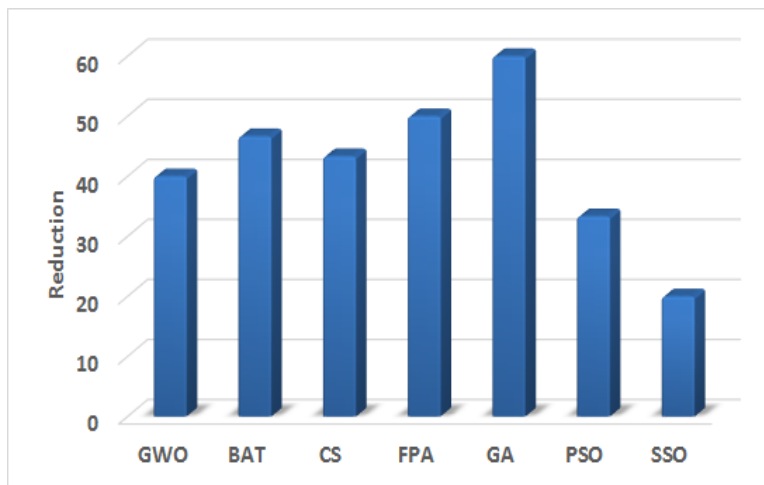
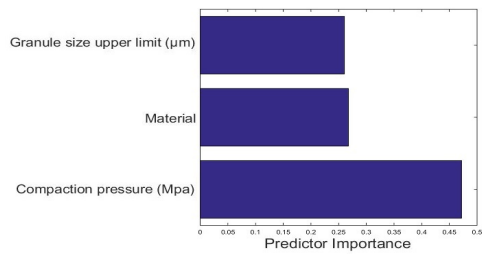
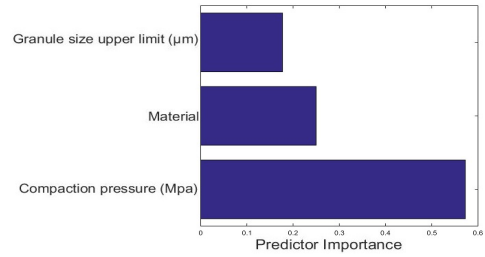


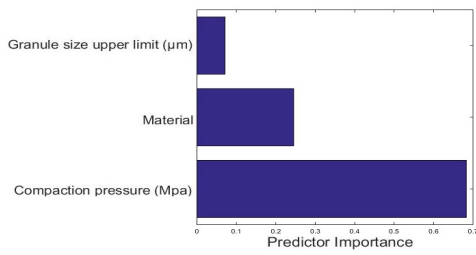
Figure 6: Average feature reduction for both tensile strength and porosity



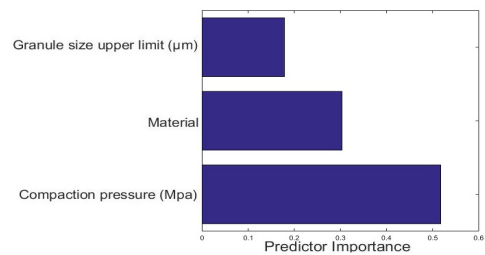
a) BAT predictor



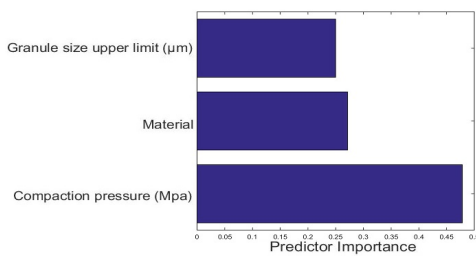
b) CS predictor



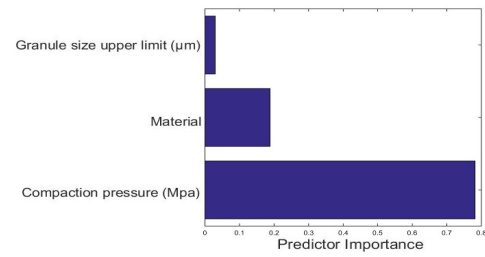
c) FPA predictor



d) GA predictor



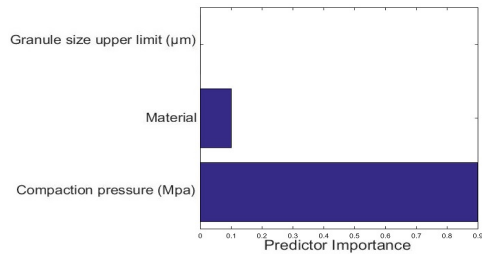
e) GWO predictor



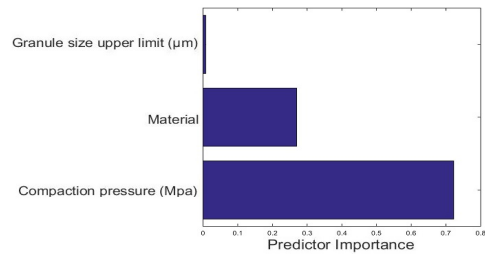
f) PSO predictor

g) SSO predictor

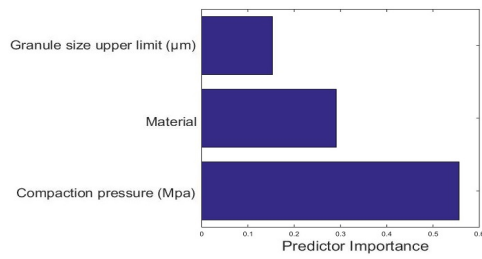
Figure 7: Feature importance for porosity



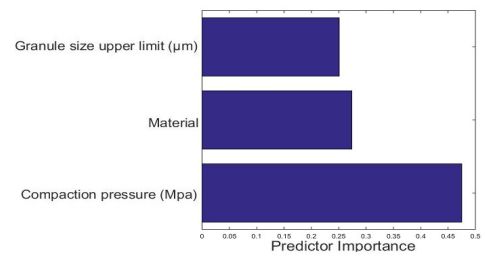
(a) BAT predictor



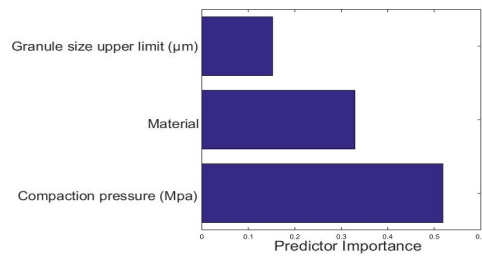
(b) CS predictor



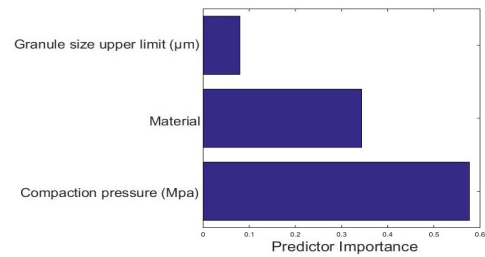
(c) FPA predictor



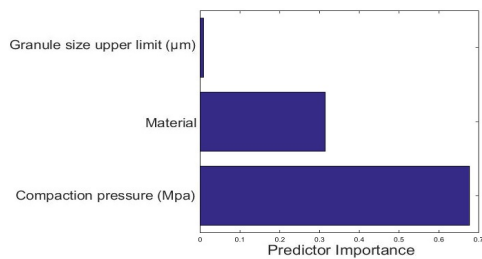
(d) GA predictor



(e) GWO predictor

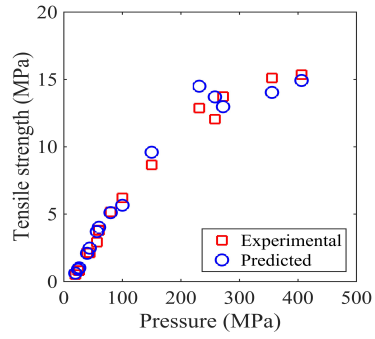


(f) PSO predictor

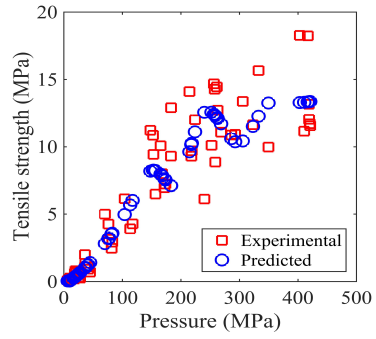


(g) SSO predictor

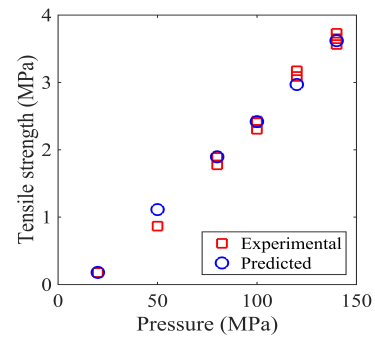
Figure 8: Feature importance for tensile strength



(a) Pure Mcc 101 powder



(b) MCC 101 Granules



(c) Mixtures of 75% Mcc and 25% lactose

Figure 9: Comparison of experimental and predicted data obtained with the SSO algorithm for different materials

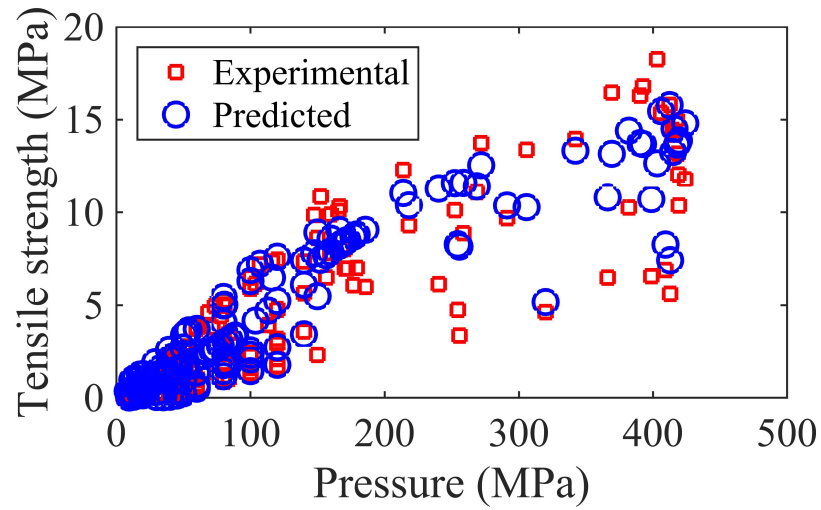


Figure 10: Comparison of experimental and predicted tensile strength obtained with the GWO algorithm for all materials considered

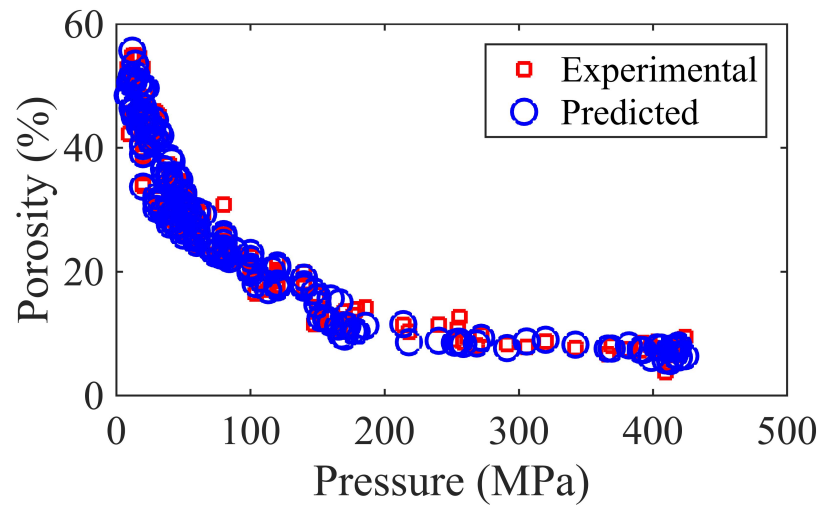


Figure 11: Comparison of experimental and predicted porosity obtained with the GWO algorithm for all materials considered

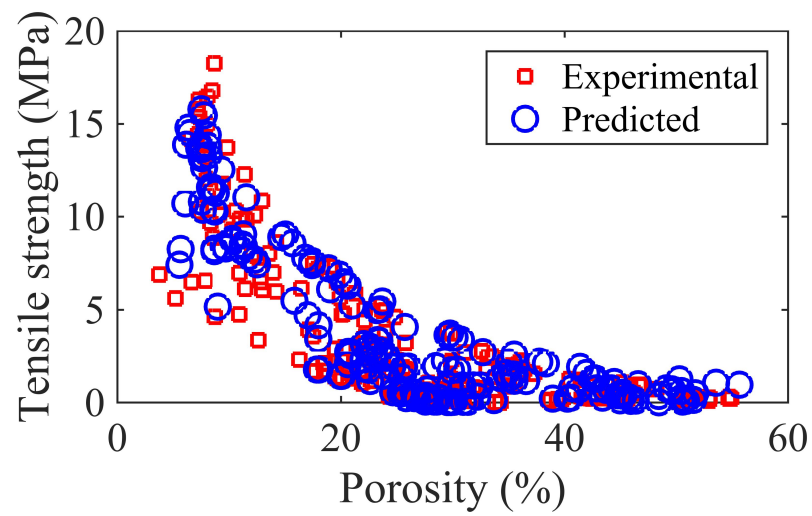


Figure 12: Comparison of experimental and predicted data on the variation of tensile strength with porosity obtained with the GWO algorithm for all materials considered