EvoImp: Multiple Imputation of Multi-label Classification data with a genetic algorithm

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Appendix S2 - Computational complexity of the proposed method

The computational complexity of EvoImp (O(Method)) is structured into three main processes:

- Initial population generation using Simple Imputation Methods $(O(SIM_i))$;
- Genetic operations: selection, crossover and mutation (O(GenOp));
- Classification to measure the performance of the individuals $(O(Class_i))$.

Therefore, the complexity of EvoImp can be described by Eq. 1:

$$O(Method) = \sum_{i=1}^{n} O(SIM_i) + O(GenOp) + \sum_{j=1}^{m} O(Class_j)$$
 (1)

where $i \in (1, ..., n)$ denotes the number of imputation methods adopted, and $j \in (1, ..., m)$ denotes the number of classification methods used for the multi-label learning tasks.

The $O(SIM_i)$ and O(GenOp) complexities for this problem studied had a low impact on the equation. This fact occurs because the attributes that influence these complexity have a low value (for example, the number of individuals, the number of generations, the number of missing values, and others).

On the other hand, the complexity of most multi-label classifiers depends on the size of the database (X) and especially on the number of labels (Y) [1]. In this case, $O(SIM_i)$ and O(GenOp) complexities are much smaller than $O(Class_j)$. This fact was confirmed during the computational experiments. Therefore, we can conclude that the complexity of the method is (Eq. 2):

$$O(Method) = MAX(O(Class_i))$$
 (2)

Among the classifiers chosen for this work, HOMER presented the worst performance, which can be justified by the balanced clustering process [2].

References

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