Transfer Learning Experiment

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1 The Idea

- Target Dataset is a small and labeled dataset from a certain domain
- Source Dataset(s) can be one or more large and labeled datasets from related domains
- If we would like to build a model for the Target Dataset using it alone as Training dataset, then the model will likely perform undesirably as the dataset is quite small!
- The idea is to make maximum use of the Source Datasets to augment the Target Dataset and classify data from its domain (this is Instance Transfer Learning)

2 The Data

- Two Source datasets TS3-Sapphire.arff and TS6-Sapphire.arff (This is for strain *Plasmodium vivax*)
- The TL algorithm supports having multiple source datasets as it assigns weights to instances as well as source datasets
- I have done **two experiments** with two different Target Datasets:
 - Data from the Venus Channel for assay TS6 (This is for strain Plasmodium falciparum)
 - this dataset has 1435 instances and the number of Active instances is 27
 - details of this experiment are in **experiment6.pdf**
 - Data from the Sapphire Channel for assay TS7 (This is for strain *Leishmania major*)
 - this dataset has 1326 instances and the number of Active instances is 12
 - details of this experiment are in **experiment7.pdf**
- Iteratively, I split each of these two datasets to create Target and Test datasets
- I started with a very small Target dataset of all Inactives and doubled the size at each iteration
- As the size increased, I started to include Active instances (with the same proportion for *Plasmodium falciparum* data)
- Remember. the Target dataset is used as Training dataset to build TL, NB, J48, SMO and KNN models

Exp. No.	Target Dataset (Training)	Test Dataset
1	Size=3 (3 Inactive + 0 Active)	Size=1432 (1405 Inactive + 27 Active)
2	Size=6 (6 Inactive + 0 Active)	Size=1429 (1402 Inactive + 27 Active)
3	Size=12 (12 Inactive + 0 Active)	Size=1423 (1396 Inactive + 27 Active)
4	Size=24 (24 Inactive + 0 Active)	Size=1411 (1384 Inactive + 27 Active)
5	Size=49 (48 Inactive + 1 Active)	Size=1386 (1360 Inactive + 26 Active)
6	Size=98 (96 Inactive + 2 Active)	Size=1337 (1312 Inactive + 25 Active)
7	Size=196 (192 Inactive + 4 Active)	Size=1239 (1216 Inactive + 23 Active)
8	Size=392 (384 Inactive + 8 Active)	Size=1043 (1024 Inactive + 19 Active)
9	Size=784 (768 Inactive + 16 Active)	Size=651 (640 Inactive + 11 Active)

Table 1: Details of Datasets for Strain Plasmodium falciparum

Exp. No.	Target Dataset (Training)	Test Dataset
1	Size=3 (3 Inactive + 0 Active)	Size=1323 (1311 Inactive + 12 Active)
2	Size=6 (6 Inactive + 0 Active)	Size=1320 (1308 Inactive + 12 Active)
3	Size=12 (12 Inactive + 0 Active)	Size=1314 (1302 Inactive + 12 Active)
4	Size=25 (24 Inactive + 1 Active)	Size=1301 (1290 Inactive + 11 Active)
5	Size=50 (48 Inactive + 2 Active)	Size=1276 (1266 Inactive + 10 Active)
6	Size=100 (96 Inactive + 4 Active)	Size=1226 (1218 Inactive + 8 Active)
7	Size=200 (192 Inactive + 8 Active)	Size=1239 (1122 Inactive + 4 Active)

Table 2: Details of Datasets for Strain $Leishmania\ major$

• Details of the various datasets are shown in the following tables:

2.1 Algorithm Names in Results Tables:

I have abbreviated column names in the tables to save display space:

TL = Transfer Learning,

NB = Naive Bayes,

J48 = The J48 Decision Trees,

SMO = Support Vector Machine,

IBk = Instance Based (k-Nearest Neighbour)

2.2 Column Names in Results Tables:

I have abbreviated column names in the tables to save display space. In order from left to right, they're as follows:

corr = Percentage of Correct guesses,

inco = Percentage of Incorrect guesses,

auc = Area Under the Curve (for class Active),

k = Kappa statistic,

mae = Mean Abs Error,

rmse = Root Mean Squared Error,

rae = Relative Abs Error,

rrse = Root Relative Squared Error,

prec = Precision (for class Active),

rec = Recall (for class Active),

fM = F-Measure (for class Active),

eR = Error Rate