

Bayesian Lasso logistic regression

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Basic information

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Github Link: https://github.com/weilu6/STAT447_Bayesian

Theme: Comparison between Bayesian classification models

Compare the simple Logistic regression and Lasso logistic regression with spike-and-slab priors.

Motivation: The classic Lasso regression performs feature selections inherently by shrinking coefficients to zero, but it is difficult to evaluate the significance of these coefficients. Instead, Bayesian Lasso provides an alternative to measure the uncertainty of parameters by credible intervals [1]. There are many types of priors on parameters such as the Laplace priors, the adaptive priors, the spike-and-slab priors, etc [2]. This project will focus on spike-and-slab priors and use the logistic link function to perform classification on the Pumpkin Seeds Dataset. Previous study shows that the frequentist's logistic regression achieves 87.92% accuracy on the same dataset[3]. This project aims to develop a calibrated hierarchical Lasso model to accurately classify the pumpkin seeds with less features. In addition, the posterior distributions of selected/unselected parameters under different situations (e.g. mis-specification, small sample size) will be investigated compared to the baseline model.

Potential approaches:

First, we split the dataset into training data and testing data. Exploratory data analysis will be performed on the training data, followed by data preprocessing. A simple Bayesian Logistic model will work as an baseline for comparison. The Lasso logistic model will be built following the recipe and references. After the initial evaluation, potential studies might be: (1) to further decrease the prior sensitivity. (2) to investigate the performance of models and the selected features when the data size becomes smaller.

Candidate datasets:

Main dataset: <https://www.kaggle.com/datasets/muratkokludataset/pumpkin-seeds-dataset/data>

```
suppressPackageStartupMessages(require(readxl))
pumpkin_df <- read_xlsx("Data/Pumpkin_Seeds_Dataset.xlsx")
head(pumpkin_df)
```

A tibble: 6 x 13

	Area	Perimeter	Major_Axis_Length	Minor_Axis_Length	Convex_Area	Equiv_Diameter
	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
1	56276	888.	326.	220.	56831	268.
2	76631	1068.	417.	234.	77280	312.
3	71623	1083.	436.	211.	72663	302.
4	66458	992.	382.	223.	67118	291.
5	66107	998.	384.	220.	67117	290.
6	73191	1041.	406.	231.	73969	305.

i 7 more variables: Eccentricity <dbl>, Solidity <dbl>, Extent <dbl>,
Roundness <dbl>, Aspect_Ration <dbl>, Compactness <dbl>, Class <chr>

Backup dataset: <https://www.kaggle.com/datasets/erdemtaha/cancer-data/data>

```
cancer_df <- read_csv("Data/Cancer_Data.csv")
head(cancer_df)
```

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean
1	842302	M	17.99	10.38	122.80	1001.0
2	842517	M	20.57	17.77	132.90	1326.0
3	84300903	M	19.69	21.25	130.00	1203.0
4	84348301	M	11.42	20.38	77.58	386.1
5	84358402	M	20.29	14.34	135.10	1297.0
6	843786	M	12.45	15.70	82.57	477.1

	smoothness_mean	compactness_mean	concavity_mean	concave.points_mean
1	0.11840	0.27760	0.3001	0.14710
2	0.08474	0.07864	0.0869	0.07017
3	0.10960	0.15990	0.1974	0.12790
4	0.14250	0.28390	0.2414	0.10520
5	0.10030	0.13280	0.1980	0.10430
6	0.12780	0.17000	0.1578	0.08089

	symmetry_mean	fractal_dimension_mean	radius_se	texture_se	perimeter_se
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1	0.2419		0.07871	1.0950	0.9053	8.589
2	0.1812		0.05667	0.5435	0.7339	3.398
3	0.2069		0.05999	0.7456	0.7869	4.585
4	0.2597		0.09744	0.4956	1.1560	3.445
5	0.1809		0.05883	0.7572	0.7813	5.438
6	0.2087		0.07613	0.3345	0.8902	2.217
	area_se	smoothness_se	compactness_se	concavity_se	concave.points_se	
1	153.40	0.006399	0.04904	0.05373		0.01587
2	74.08	0.005225	0.01308	0.01860		0.01340
3	94.03	0.006150	0.04006	0.03832		0.02058
4	27.23	0.009110	0.07458	0.05661		0.01867
5	94.44	0.011490	0.02461	0.05688		0.01885
6	27.19	0.007510	0.03345	0.03672		0.01137
	symmetry_se	fractal_dimension_se	radius_worst	texture_worst	perimeter_worst	
1	0.03003		0.006193	25.38	17.33	184.60
2	0.01389		0.003532	24.99	23.41	158.80
3	0.02250		0.004571	23.57	25.53	152.50
4	0.05963		0.009208	14.91	26.50	98.87
5	0.01756		0.005115	22.54	16.67	152.20
6	0.02165		0.005082	15.47	23.75	103.40
	area_worst	smoothness_worst	compactness_worst	concavity_worst		
1	2019.0	0.1622		0.6656	0.7119	
2	1956.0	0.1238		0.1866	0.2416	
3	1709.0	0.1444		0.4245	0.4504	
4	567.7	0.2098		0.8663	0.6869	
5	1575.0	0.1374		0.2050	0.4000	
6	741.6	0.1791		0.5249	0.5355	
	concave.points_worst	symmetry_worst	fractal_dimension_worst	X		
1		0.2654	0.4601		0.11890	NA
2		0.1860	0.2750		0.08902	NA
3		0.2430	0.3613		0.08758	NA
4		0.2575	0.6638		0.17300	NA
5		0.1625	0.2364		0.07678	NA
6		0.1741	0.3985		0.12440	NA

References:

- [1]: Park, T., & Casella, G. (2008). The Bayesian Lasso. *Journal of the American Statistical Association*, 103(482), 681–686. <https://doi.org/10.1198/016214508000000337>
- [2]: Chen, S. M., Bauer, D. J., Belzak, W. M., & Brandt, H. (2021). Advantages of Spike and Slab Priors for Detecting Differential Item Functioning Relative to Other Bayesian Regularizing

Priors and Frequentist Lasso. *Structural Equation Modeling: A Multidisciplinary Journal*, 29(1), 122–139. <https://doi.org/10.1080/10705511.2021.1948335>

[3]: KOKLU, M., SARIGIL, S., & OZBEK, O. (2021). The use of machine learning methods in classification of pumpkin seeds (*Cucurbita pepo* L.). *Genetic Resources and Crop Evolution*, 68(7), 2713-2726. Doi: <https://doi.org/10.1007/s10722-021-01226-0>