Quick Permutation Test: feature filtering of n-gram data

Piotr Sobczyk^{1*}, Michał Burdukiewicz², Chris Lauber³, Paweł Mackiewicz²
*Piotr.Sobczyk@pwr.edu.pl

¹Wrocław University of Technology, Institute of Mathematics and Computer Science, Poland ²University of Wrocław, Department of Genomics, Poland

³Dresden University of Technology, Institute of Medical Informatics and Biometry, Poland

Introduction

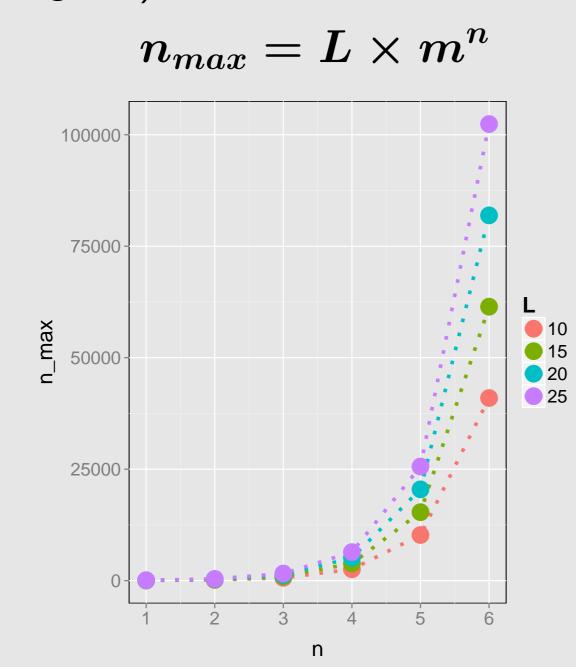
N-grams (k-tuples) are vectors of n characters derived from input sequence(s). They may form continuous sub-sequences or be discontinuous. Another important n-gram parameter is its position. Instead of just counting n-grams, one may want to count how many n-grams occur at a given position in multiple (e.g. related) sequences.

	P1	P2	P3	P4	P5	P6
S1	3	1	2	1	2	2
S2	1	2	3	3	4	1
S3	4	1	3	1	1	1
	Sar	nple	sea	lllen	ces	

P1_1	P2_1	P3_1	P4_1	P5_1	P6_1	P1_2	P2_2	P3_2	P4_2	P5_2
0	1	0	1	0	0	0	0	1	0	1
1	0	0	0	0	1	0	1	0	0	0
0	1	0	1	1	1	0	0	0	0	0
A fraction of possible unigrams with position information.										

Curse of dimensionality

Number of possible positioned n-grams (not taking into account distances between elemnts of n-gram):



Permutation test

During permutation tests class labels are randomly exchanged during computation of significance statistic. p-values are defined as:

p-value
$$=rac{N_{T_P>T_R}}{N}$$

where $N_{T_P>T_R}$ is number of times when T_P (permuted test statistic) was more extreme than T_R (test statistic for non-permuted data).

Permutation tests are model and statistic independent, but computationally expensive (especially precise estimation of low p-values, because the number of permutations is inversely proportional to the interval between p-values).

QuiPT algorithm

If probability that target equals 1 is $oldsymbol{p}$ and probability that feature equals 1is q and feature and target are independent then each of them has the following probabilities

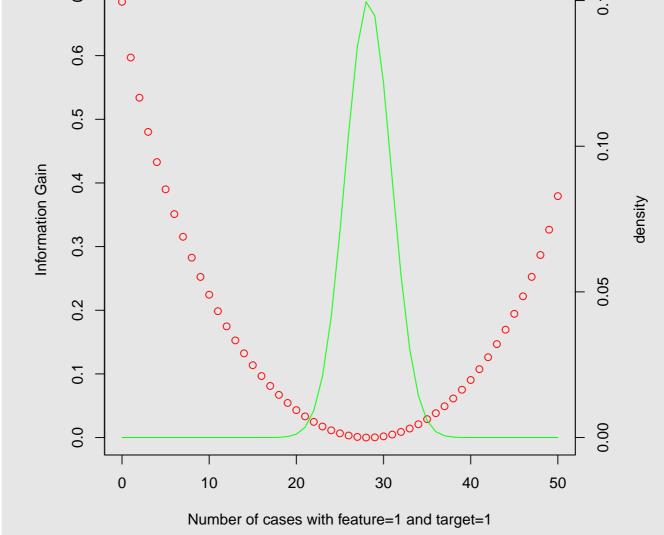
$$P(Target, Feature) = (1, 1)) = p \cdot q$$
 $P(Target, Feature) = (1, 0)) = p \cdot (1 - q)$
 $P(Target, Feature) = (0, 1)) = (1 - p) \cdot q$
 $P(Target, Feature) = (0, 0)) = (1 - p) \cdot (1 - q)$

$$F(n_{1,1},n_{1,0},n_{0,1},n_{0,0}) = inom{n}{n_{1,1}} (p \cdot q)^{n_{1,1}} n - n_{1,1} \ inom{p \cdot (1-q)^{n_{1,0}}}{n_{1,0}} \ inom{n - n_{1,1} - n_{1,0}}{n_{0,1}} \ inom{n - n_{1,1} - n_{1,0} - n_{0,1}}{n_{0,0}} \ inom{n - n_{1,1} - n_{1,0} - n_{0,1}}{n_{0,0}} \ inom{n - n_{1,1} - n_{1,0} - n_{0,1}}{n_{0,0}} \ inom{n - n_{0,1}}{n_{0,0}} \ inom{n - n_{0,0}}{n_{0,0}} \ \ inom{n - n_{0,0}}{n_{0,0}} \ \ inom{n - n_{0,$$

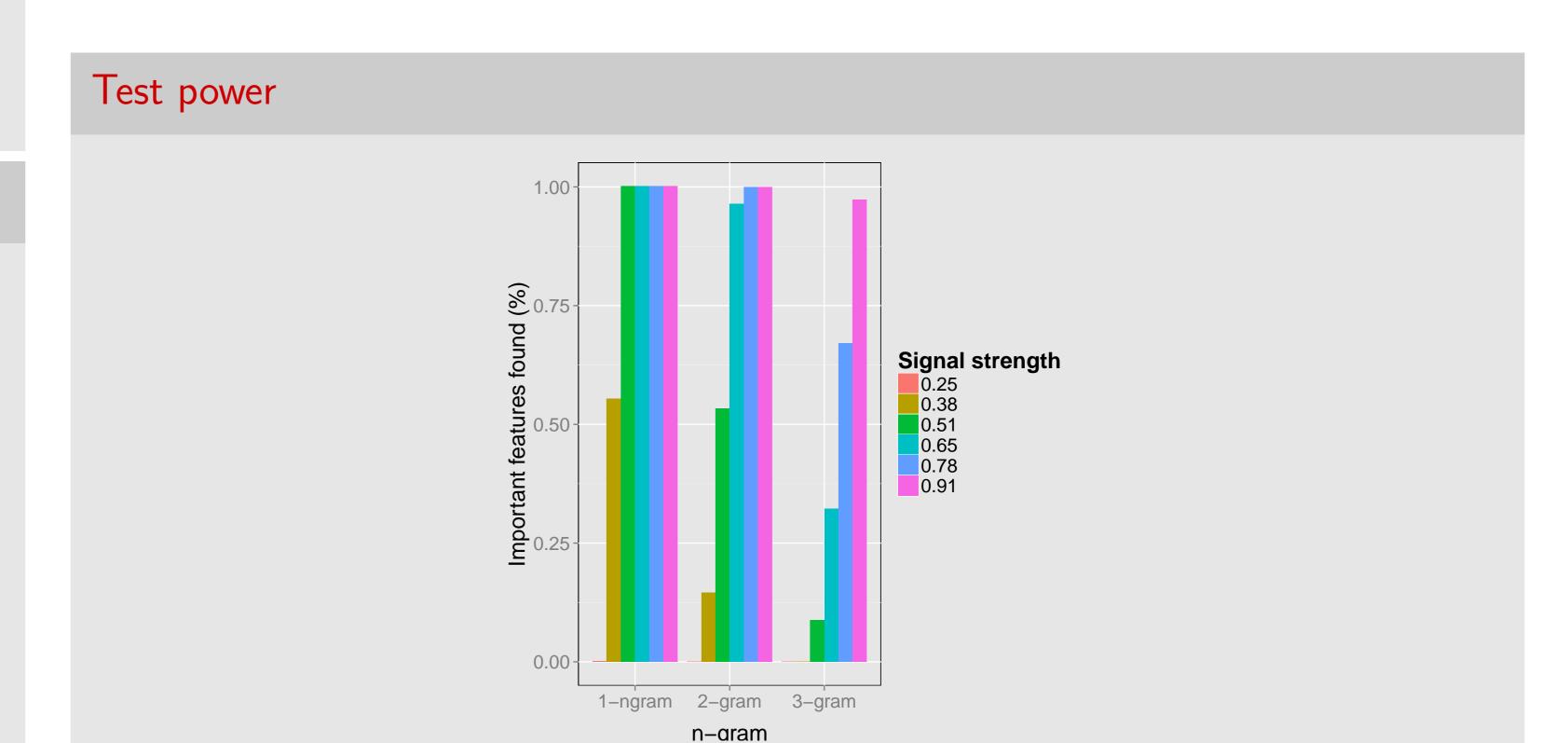
In addition to this: $n_{1,\cdot}=n_{1,1}+n_{1,0}$ and $n_{\cdot,1}=n_{1,1}+n_{0,1}$ are known

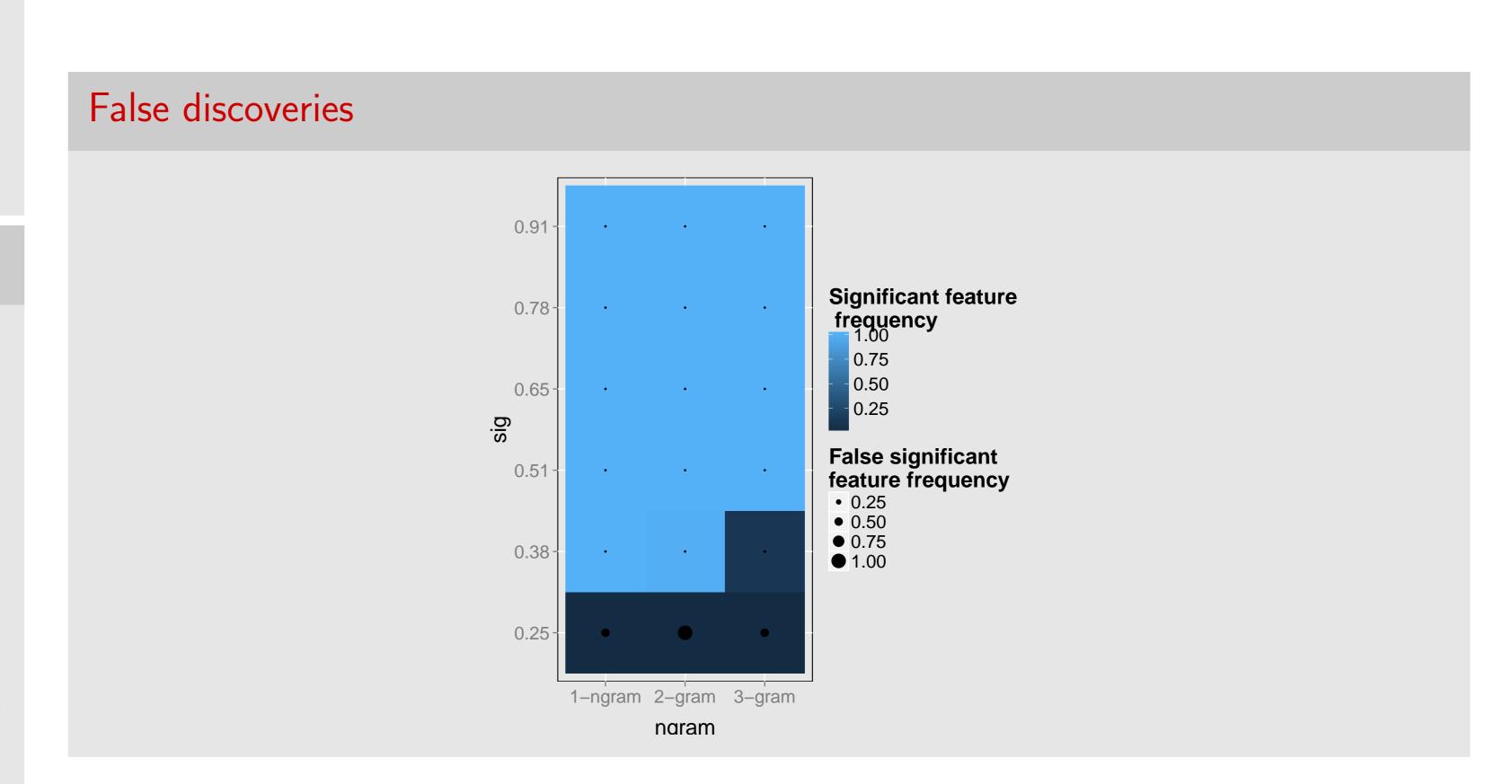
and fixed.





٦	arget	Feature	Freq
1 0		0	40
2 1		0	10
3 0		1	25
4 1		1	40





Summary

Quick permutation test is a powerful and quick equivalent of permutation test in binary feature-binary target testing scenario.

Avaibility

biogram R package: http://cran.r-project.org/web/packages/biogram/

Bibliography