

BLAST and Finding Data

Sept 9, 2015

ANNOUNCEMENTS

BLAST

Steps in BLAST

BLAST

1. Build Lookup table

Preprocess: Build a *lookup table* of size $|\Sigma|^w$ for all w -length words in D

$$\Sigma = \{A, C, G, T\}$$

$$w = 2$$

→ $4^2 (=16)$ entries in lookup table

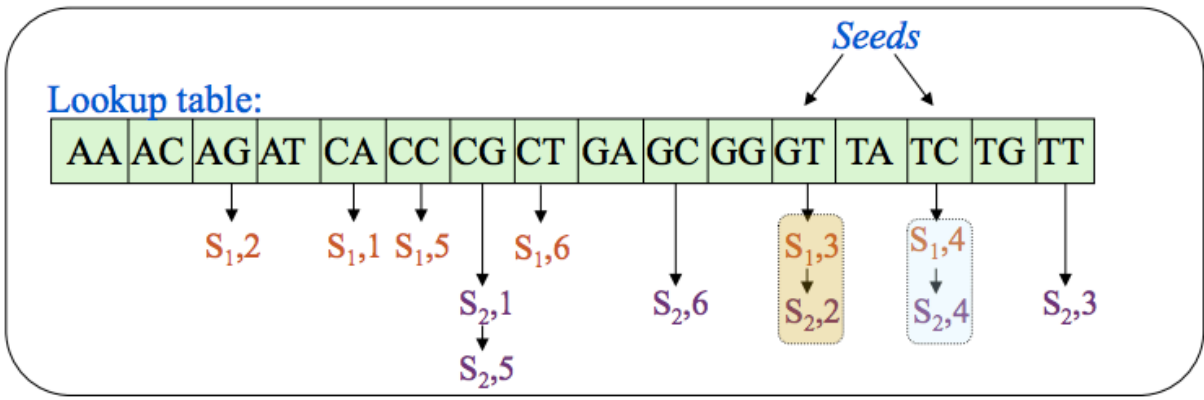
Lookup table:

AA	AC	AG	AT	CA	CC	CG	CT	GA	GC	GG	GT	TA	TC	TG	TT
----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----

BLAST

2. Filter low complexity and Identify Seeds

1 2 3 4 5 6 7
S₁: CAGTCTCT
S₂: CGTTCGC



BLAST

4. Rank and report

BLAST

Stats

$$E = Kmne^{-\lambda S}$$

BLAST

Stats

$$p = 1 - e^{-E}$$

BLAST

Is my p-value significant?

	H_0 true	H_0 false
Reject H_0	Type 1 error (false pos)	Correct!
Accept H_0	Correct!	Type 2 error (false neg)

BLAST

Is my p-value significant?

	H ₀ true	H ₀ false
Reject H ₀	Type 1 error (false pos)	Correct!
Accept H ₀	Correct!	Type 2 error (false neg)

BLAST null: There is no match between query and database entry

BLAST

Multiple testing correction

<http://www.r-fiddle.org/#/>

Finding Data

Read data

- <http://www.ebi.ac.uk/ena>
- <http://www.ncbi.nlm.nih.gov/sra>
- <http://metagenomics.anl.gov/?page=MetagenomeSelect>

Assembly (and other) Data

- <http://useast.ensembl.org/info/data/ftp/index.html>
- <http://www.ncbi.nlm.nih.gov/genome/>
- <http://datadryad.org/>
- <http://figshare.com/>

Finding Data

Human Stuff

- <http://www.ncbi.nlm.nih.gov/clinvar/>
- <http://www.ncbi.nlm.nih.gov/omim>
- <http://snpedia.com/index.php/SNPedia>

Journal	23andMe White Paper
Study Size	■■■
Replications	None
Contrary Studies	None
Applicable Ethnicities	European
Marker	rs2937573

A study of roughly 80,000 individuals with European ancestry who participated in 23andMe research surveys identified a genetic marker associated with sensitivity to the sound of other people chewing food. The marker rs2937573 is located near a [gene](#) (TENM2) that may play a role in the brain. Individuals with the GG [genotype](#) at rs2937573 had about 1.2 times higher odds of being sensitive to the sound of chewing, compared to individuals with the AG genotype. Individuals with the AA genotype had about 1.2 times lower odds of being sensitive.

Who	Genotype	Genetic Result
Kate MacManes, Lilly Mendel (Mom)	GG	Slightly higher odds of being sensitive to the sound of chewing.
Lauren MacManes, Owen MacManes, Patrick MacManes	AG	Typical odds of being sensitive to the sound of chewing.
Matthew MacManes, Greg Mendel (Dad)	AA	Slightly lower odds of being sensitive to the sound of chewing.

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rs2937573 [Homo sapiens]

1.

GCCCACTCAAAGTGGCAACTGCCCA[A/G]CACTGTGACTAAGTAAGATGGTGTA

Chromosome: 5:167044193

Gene: TENM2 (GeneView)

Functional Consequence: intron variant

Validated: by 1000G,by 2hit 2allele,by cluster,by frequency,by hapmap,by submitter

Global MAF: G=0.3990/1998

HGVs: NC_000005.10:g.167044193G>A, NC_000005.9:g.166471198G>A, XM_005265950.1:c.-189-29049G>A, XM_006714897.1:c.-189-29049G>A, XM_011534604.1:c.-189-29049G>A

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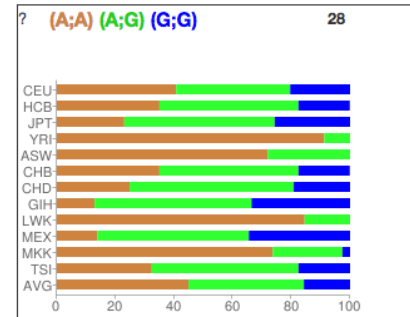
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The ALLEle FREquency Database

ALFRED is a resource of gene frequency data on human populations supported by the U. S. National Science Foundation.

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Polymorphism Information

Name	ALFRED UID	Locus Name	Locus Symbol
rs2937573	SI368946K	rs2937573 is intergenic between RPLP0P9 and ODD2	rs2937573

Fst	Avg Het	# Populations Typed
0.181	0.407	51

Synonyms: rs2937573 ;
 Frequency on Map: [GoogleMap](#) [Help](#)

Frequency Display Formats: [Graph](#) [Table](#)

Estimated Heterozygosity: [Graph](#)

Frequency Download: [Tab Delimited](#) [Arlequin](#) [Help](#)

External Resources: [dbSNP rs# Record](#) [PharmGKB Variant Information Record](#)

References: [See References](#)

Polymorphism Description: This is a A/G SNP

Alleles:

Allele Name	Allele Symbol	Description
A	A	5' - gtcaaaatggcaagtgccc A cactgtgactaagtaagatg - 3'
G	G	5' - gtcaaaatggcaagtgccc G cactgtgactaagtaagatg - 3'

References:
 - Kenneth K. Kidd et al. "Data unpublished".

