



Global Alliance
for Genomics & Health

Beacon Network

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2 years ago



Global Alliance for Genomics and Health
Beacon of Beacons

Beacon: all Chromosome: 15 Position: 41087869 Alt: A

Beacon	Result
Wellcome Trust Sanger Institute	No
EMBL-EBI	No
UniProt	No
NCBI ClinVar	No
NCBI	No
Leiden Open Variation	No
AMPLab	Yes

Search [all beacons](#) for allele

GRCh37 ▾

13 : 32936732 G > C

Search

Response [All](#) [None](#)

- ☒ Found 14
- ☒ Not Found 37
- ☐ Not Applicable 14



Organization [All](#) [None](#)

- ☒ AMPLab, UC Berkel...
- ☒ BGI
- ☒ BioReference Labor...
- ☒ Brazilian Initiative o...
- ☒ BRCA Exchange
- ☒ Broad Institute
- ☒ Centre for Genomic...
- ☒ Centro Nacional de...
- ☒ Curoverse
- ☒ EMBL European Bio...
- ☒ Global Alliance for ...
- ☒ Google
- ☒ Institute for System...
- ☒ Instituto Nacional d...



AMPLab - 1000 Genomes Project

Hosted by AMPLab, UC Berkeley

Not Found



Biobase - HGMD

Hosted by University of California, Santa Cruz

Found



BioReference

Hosted by BioReference Laboratories

Not Found



BRCA Exchange

Hosted by BRCA Exchange

[Hide Metadata](#)

Found

Clinical significance citations [PMID:23108138](#)

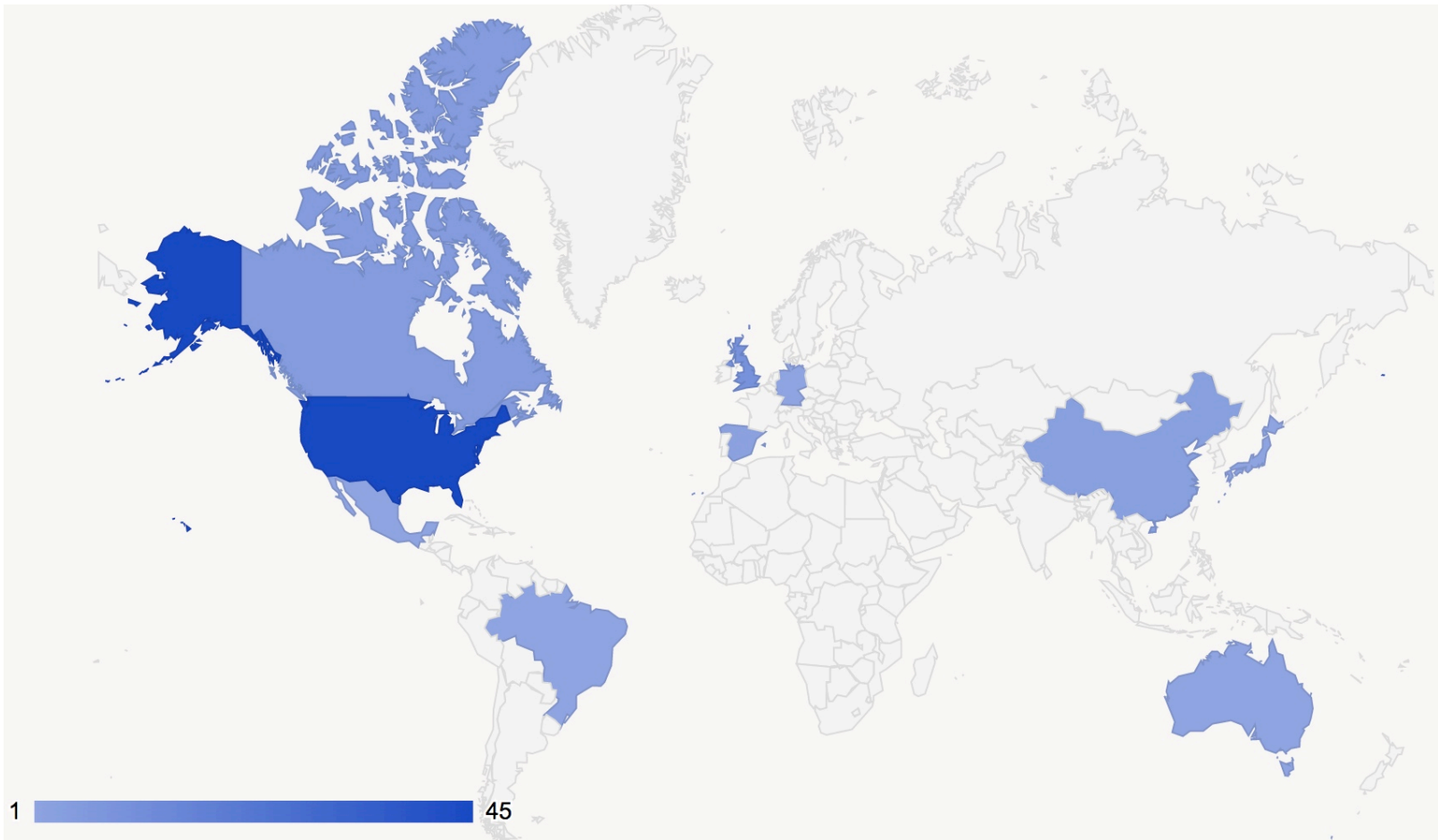
HGVS cDNA c.7878G>C

Condition ID type OMIM

Date last evaluated 8/10/15

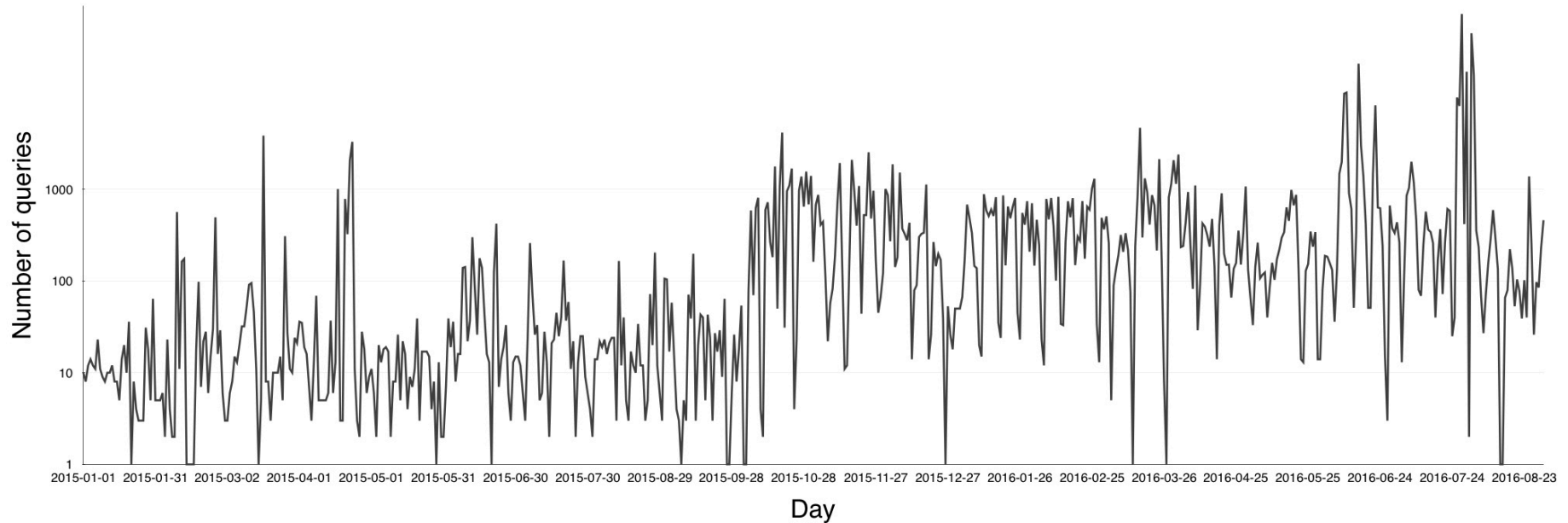
Adoption

- **25+** of the world's top genomic organizations, **60+** beacons.



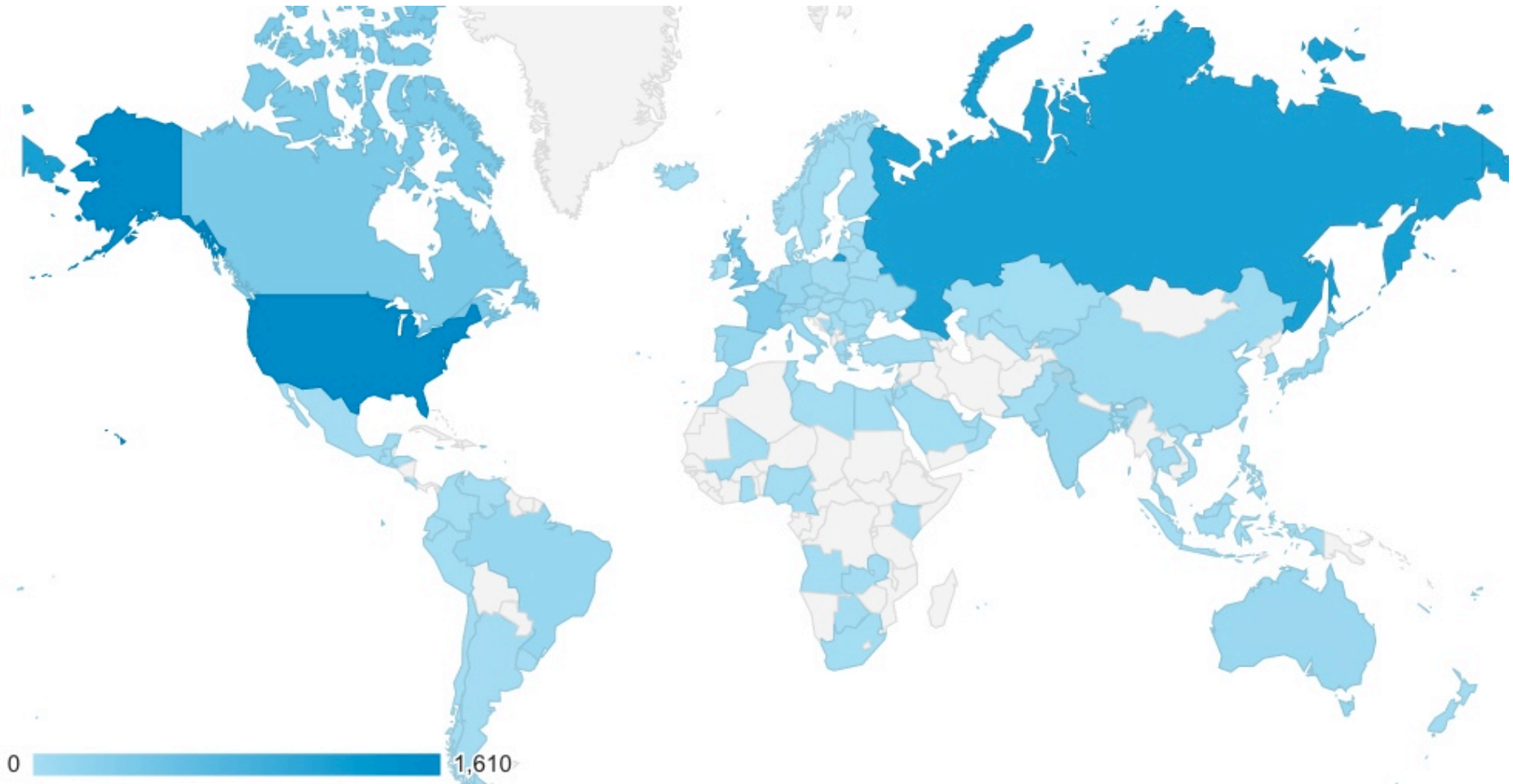
Usage

- **400K+** queries received → **2M+** queries dispatched.



Usage

- **6K+** users in **100+** countries.

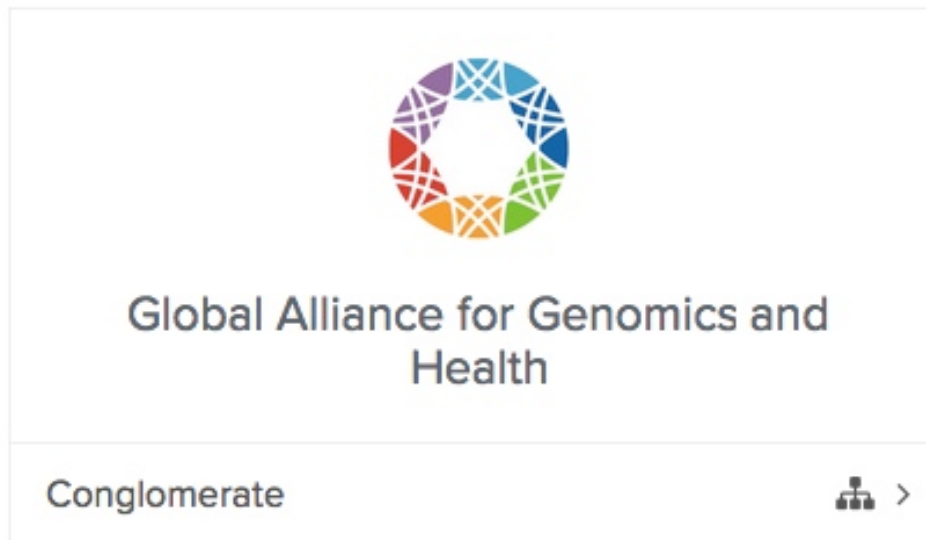


Usage

- Other metrics:
 - Size/growth.
 - Requests.
 - User visits.
 - Assemblies, chromosomes, variants.
 - Harmfulness.
 - Rarity.
 - Genes.
 - Disorders, clinical abnormalities.


Features

- Federation of queries across beacons.
- Registry of public beacons.
- Unified beacon API.
- Input/output conversion.
- Distribution of queries through the network.
- Aggregation.



Features

- Metadata support.

 BRCA Exchange Hosted by BRCA Exchange		Hide Metadata Found
Clinical significance citations	PMID:23108138	
HGVS cDNA	c.7878G>C	
Condition ID type	OMIM	
Date last evaluated	8/10/15	
HGVS protein	p.(Trp2626Cys)	
Assertion method citation	http://enigmaconsortium.org/documents/ENIGMA_Rules_2015-03-26.pdf	
ClinVarAccession	SCV000244475	
alt	C	
Condition category	Disease	
Comment on clinical significance	IARC class based on posterior probability from multifactorial likelihood analysis, thresholds for class as per Plon et al. 2008 (PMID: 18951446). Class 5 based on posterior probability = 1	
Reference sequence	NM_000059.3	
Gene symbol	BRCA2	
URL	C">http://hci-exlovd.hci.utah.edu/variants.php?select_db=BRCA2&action=search_all&search_Variant%2FDNA=c.7878G>C	
chrom	13	
Abbrev AA change	W2626C	
ref	G	
Assertion method	ENIGMA BRCA1/2 Classification Criteria (2015)	
BIC Nomenclature	8106G>C	
Clinical significance	Pathogenic	


Features

- External system integration (e.g. secured beacons).

Search [1000 Genomes Project - Yoruban Trio](#) for allele

GRCh37 ▾	1: 695744 G > T	Search
----------	-----------------	--------

Response	All None	
<input checked="" type="checkbox"/> Found	1	
<input type="checkbox"/> Not Found	0	

	1000 Genomes Project - Yoruban Trio Hosted by DNASTack	Found More info...
---	--	--

Features

- External system integration (e.g. secured beacons).

Search [1000 Genomes Project - Yoruban Trio](#) for allele

GRCh37 ▾	1 : 695744 G > T	Search
----------	------------------	--------

Response	All	None
<input checked="" type="checkbox"/> Found	1	
<input type="checkbox"/> Not Found	0	



1000 Genomes Project - Yoruban Trio

Hosted by DNASTack

Found

[More info...](#)



☐ Remember me

[Forgot password?](#)

[Register for an account](#)

Log in

Features

- External system integration (e.g. secured beacons).

Search **1000 Genomes Project - Yoruban Trio** for allele

GRCh37 ▾

1 : 695744 G > T

Search

Response

All None

☒ Found 1

☐ Not Found 0



1000 Genomes Project - Yoruban Trio

Hosted by DNASTack

Found

More info...



username or email

password

☐ Remember me

[Forgot password?](#)

[Register for an account](#)

Log in



Yoruban Trio

Hosted by Yoruban Trio

Found

Datasets 1

YRI.trio.2010_03.genotypes.vcf

Found

AA

.

DP

115

Tools

- REST API
 - <https://beacon-network.org/api/>

Endpoint	Description	Example
/beacons	Lists beacons.	Go to example
/beacons/{beacon}	Shows a beacon.	Go to example
/organizations	Lists organizations.	Go to example
/organizations/{organization}	Shows an organization.	Go to example
/responses	Queries beacons.	Go to example
/responses/{beacon}	Queries a beacon.	Go to example
/chromosomes	Lists supported chromosome.	Go to example
/alleles	Lists supported alleles.	Go to example
/references	Lists supported reference genomes.	Go to example

<https://beacon-network.org/#/developers/api/beacon-network>

- Open-source client library (Java)
 - <https://github.com/mcupak/beacon-network-client>
 - <https://mcupak.github.io/beacon-network-client/>

```
BeaconNetworkClient bnc = new BeaconNetworkClientImpl();  
BeaconResponseDto response = bnc.getResponse(ChromosomeDto.CHR13, 32936732L,  
                                             AlleleDto.C, ReferenceDto.HG19, "amplab");
```

```
public interface BeaconNetworkClient
```

Beacon Network API. All methods throw either operation specific exceptions or `InternalException`.

Method Summary

All Methods

Instance Methods

Abstract Methods

Modifier and Type

Method and Description

`com.dnastack.bob.service.dto.BeaconDto`

`getBeacon`(`java.lang.String beaconId`)
Shows information on the beacon specified by its id.

`java.util.List<com.dnastack.bob.service.dto.BeaconDto>`

`getBeacons`()
Lists available beacons.

Tools

- Open-source CLI
 - <https://github.com/mcupak/beacon-network-cli>

```
$ java -jar cli.jar beacon -help
```

Beacon Network CLI Usage:

```
[beacon | organization | response] : Use one of these commands. To get help on  
each command, Type -h with the  
corresponding command.  
-u (--url) VAL : Beacon Network URL (default =  
https://beacon-network.org/api/)
```

Tools

- Embeddable widget
 - <https://beacon-network.org/#/developers/embed>

```
<iframe src="https://beacon-network.org:443/#/widget?rs=GRCh37&chrom=13&pos=32936732&ref=G&allele=C"
style="width: 500px; height: 700px; overflow: auto;"
marginwidth="0" marginheight="0" frameborder="0" vspace="0" hspace="0">
</iframe>
```

Beacon Network

Response

AllNone

☒ Found11

☐ Not Found22

☐ Not Applicable16

Organization

AllNone

☒ AMPLab, UC Berkeley


☒ BGI

☒ BioReference Laboratories

☒ Brazilian Initiative on Precision Medicine


☒ BRCA Exchange

Show All

 Biobase - HGMD

Hosted by University of California, Santa Cruz

Found

 BRCA Exchange

Hosted by BRCA Exchange

Show MetadataFound

Future

- Support for the latest version of the Beacon API.
- User accounts.
- Secured beacon support.
- Programmatic beacon registration, beacon management.
- Workflow for contacting beacon operators.
- Continued support for tooling.
- **Beacon Network specification.**

Questions?

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 info@beacon-network.org

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 miro@dnastack.com