# PGP-UK Genomics Report for ukDAC808

#### 1 Summary

This is the genome report was produced using collaborative research tools, including SNPedia and GetEvidence. This section shows an overview of all the small variants which were found in the genome for this individual, when compared with a reference genome. These variants are summarised in Table 1 and the pie-charts in Figures 2, 3 and 4.

This report was generated automatically and is not clinically approved. It is provided for <u>personal and research purposes</u> only.

This document contains hyperlinks, shown in grey, that will take you to external websites where you can find more detailed explanations. Some of the technical terms are also explained in more detail in the Ensembl Glossary. We would welcome your feedback about this report, for example, if you would like more information about anything or if any of the links have become inactive. You can contact us on: pgp-uk@ucl.ac.uk.

This summary shows an overview of all the variants which were found in the genome for this individual. The "variants remaining after filtering" refers to any differences in the DNA identified when compared to the reference genome. Of these, the majority will have already been found in some other sequenced individual and put on a database (existing variants) while others have not yet been annotated (novel variants).

"Overlapped genes" refers to the number of times where a variant was found in a region of the genome containing a gene. The diagram in Figure 1 is a simplification of the usual gene structure. "Exon" refers to the part of the gene which goes on to form a protein, and variants in this part of the gene are more likely to cause changes in the shape of the protein. Upstream, downstream, intronic and intergenic variants are more likely to alter the regulation of that gene but will not change the protein itself.

A transcript for a protein-coding gene can include the exons, introns and other gene features that are transcribed and important for gene function but might not be translated into the final protein. Not all transcripts are for protein-coding genes, with many containing non-coding RNAs that can be overlapping other genes, in introns or in intergenic regions.

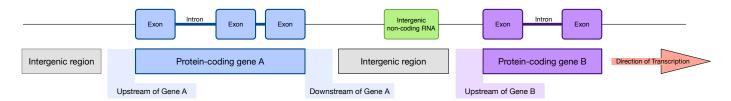


Figure 1: Diagram of gene structure indicating locations of potential variants

Feature	Count
Lines of input read	4952831
Variants filtered out	0
Novel / existing variants	$476133\ (9.6)\ /\ 4465386\ (90.4)$
Overlapped genes	56608
Overlapped transcripts	67375
Overlapped regulatory features	166703

Table 1: Variant calling summary

There are several different types of genomic variants. The most common change is when one single building block of the DNA (called a nucleotide) is changed, called a single nucleotide variants (SNV). Other variant types include insertions, where the DNA in the individual is longer than the reference sequence due to the insertion of one or more nucleotides; and deletions, where a few nucleotides are missing compared to the reference sequence.

Some of these changes will have no effect on the protein, while some changes may alter the protein function to varying degrees. The PolyPhen analysis software attempts to quantify the effect each mutation will have on the protein function. This ranges from "benign" where no change to the protein function is expected, to "probably damaging" where it is predicted that the mutation will affect protein function. It is nevertheless important to note that what is "damaging" for the protein is not necessarily damaging for the individual.

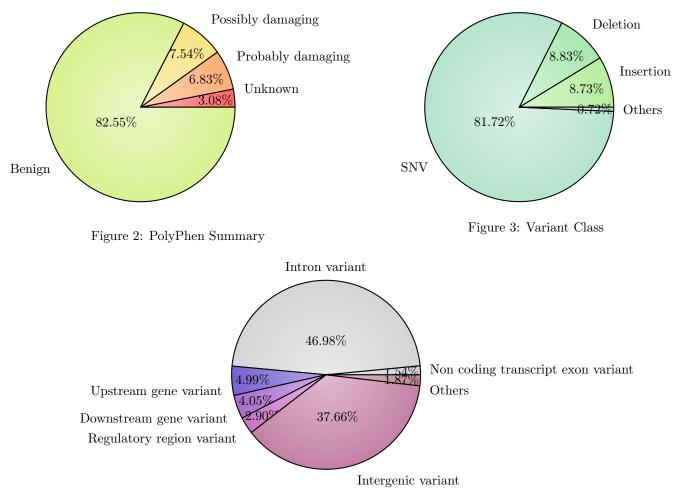


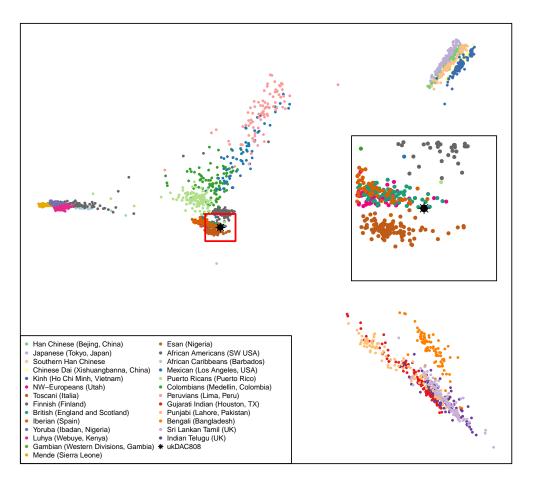
Figure 4: Consequence type

## 2 Ancestry

This plot shows the distribution of the genomes of different populations. Data from several studies which used whole genome sequencing was used to see the relationships between the genomes of the populations. It shows how closely related certain populations are genetically: Groups which cluster closely are more genetically similar than groups which are further apart. The black star symbol shows where this PGP-UK participant sits in relation to other populations, indicating their ancestry and their most closely related populations according to genetic sequence.

Based on the populations defined in the 1000 genomes project (1kGP), the ancestry composition for this individual is inferred to be 100.0 percent European [British in England and Scotland].

Please note that this analysis is limited by the populations available in the 1kGP data. If there are European subpopulations reported, and the ancestry of the participant does not correspond to any of the 1kGP populations, the closest 1kGP sampled subpopulation will be shown (even though it might be different from the participant's actual ancestry).



### Ancestry ukDAC808

Figure 5: Ancestry Principal Component Analysis

## 3 Traits (based on SNPedia information)

Existing research has associated many variants with phenotypic traits, some of which can be perceived as beneficial while others appear to have a harmful effect. Some traits are complex and can be affected by several variants. It is likely that some of these would confer a higher risk while others a lower risk of trait manifestation. These can not be combined linearly to produce an actual risk of disease.

It is important to note that in most cases genomic data is probabilistic, not deterministic- i.e. having a genetic predisposition for a disease is not a diagnosis; rather, it shows an increased likelihood of developing that disease. Also, one person can have both potentially beneficial and harmful variants in the same gene, or associated with the same disease.

Some variants can also affect certain populations more, or will only affect a particular gender. For example, a variant for higher risk of endometriosis in the sequence of a male will not directly affect that person, but can be passed on to descendants.

While many traits are the result of a unique variant, many are the combination of several variants throughout the genome. In SNPedia, these are called genosets. These can integrate some of the information already present in the single variant tables, or be the combination of variants that have no phenotypic effect on their own, but contribute to a trait when together.

The variants in the following tables are sorted by magnitude. This is an subjective measure defined in SNPedia to highlight the perceived importance of the genotype described. At the moment this scale goes from 0 to 10. You can read more about it by visiting their explanatory webpage.

As our knowledge grows, the interpretation of the effect of certain variants might change. Clicking on the links in the genome report tables will take you to websites containing more information about each variant.

Mag.	Identifier	Genotype	Summary	ExAC	GetEvidence	ClinVar
2.1	rs2511989	(A;G)	0.63x decreased age-related macular degeneratio		Link	
2.1	rs6505162	(A;A)	0.43x decreased risk for esophageal cancer	Link		
2	rs11045585	(A;A)	24% chance (lower than average) of docetaxel-in		Link	
2	rs11635424	(A;A)	<0.70x risk for restless legs		Link	
2	rs12593813	(A;A)	<0.71x risk for restless legs		Link	
2	rs12979860	(C;C)	$\sim 80\%$ of such hepatitis C patients respond to tr		Link	Link
2	rs1799884	(G;G)	Mothers have typical Birth-Weight babies. Sligh			
2	rs1864163	(A;G)	Associated with higher HDL cholesterol		Link	
2	rs2073963	(T;T)	Reduced risk of baldness			
2	rs2241423	(A;G)	0.79 decreased risk for obesity			
2	rs2542052	(C;C)	Better odds of living to 100			
2	rs3738579	(C;T)	0.5x decreased risk for cervical cancer: HNSCC:			
2	rs3764261	(G;T)	Associated with higher HDL cholesterol		Link	Link
2	rs4149268	(G;G)	Associated with higher HDL cholesterol		Link	
2	rs6495446	(T;T)	0.64x reduced risk for chronic kidney disease			
2	rs6855911	(A;G)	0.62x decreased risk for gout		Link	
2	rs763110	(C;T)	$\sim 0.80$ x reduced cancer risk			Link
2	rs7776725	(T;T)	Stronger bones		Link	
2	rs801114	(T;T)	0.78x decreased Basal Cell Carcinoma risk.		Link	
2	rs9272346	(A;G)	0.3x risk type-1 diabetes		Link	
2	rs9642880	(G;G)	Slightly lower risk of Bladder Cancer.		Link	
1.8	rs1128535	(A;G)	0.77x risk for Crohn's disease			
1.8	rs3814113	(C;T)	0.8x decreased risk for ovarian cancer		Link	
1.8	rs6897932	(C;T)	0.91x decreased risk for multiple sclerosis	Link	Link	Link
1.6	rs1061170	(T;T)	Lower risk for AMD: generally longer live than	Link	Link	Link
1.5	rs1026732	(A;G)	0.70x risk for restless legs		Link	
1.5	rs1063192	(C;C)	0.71x reduced risk of myocardial infarction			

#### 3.1 Possibly Beneficial Traits

Mag.	Identifier	Genotype	Summary	ExAC	GetEvidence	ClinVar
1.5	rs11136000	(C;T)	0.84x decreased risk for Alzheimer's disease		Link	
1.5	rs11212617	(A;C)	Somewhat increased likelihood of treatment succ			Link
1.5	rs16991615	(A;G)	Slight increase (11 months) in avg age at menop	Link	Link	
1.5	rs309375	(G;G)	Smaller mosquito bites			
1.5	rs3784709	(C;T)	0.71x risk of developing restless legs syndrome		Link	
1.5	rs4149274	(C;C)	Associated with higher HDL (good) cholesterol.			
1.5	rs4489954	(G;T)	0.69x risk risk of developing restless legs syn		Link	
1.5	rs4939883	(C;T)	Associated with higher HDL cholesterol		Link	
1.5	rs9939609	(T;T)	Lower risk of obesity and Type-2 diabetes		Link	
1.4	rs1165205	(A;T)	0.85x decreased gout risk		Link	
1.4	rs2294008	(C;C)	Lower risk of gastric and bladder cancer	Link	Link	
1.4	rs6700125	(C;C)	0.7x decreased risk for ALS			
1.2	rs11246226	(A;C)	Decreased risk of schizophrenia in limited stud		Link	
1.1	rs2293347	(G;G)	Among NSCLC patients: better Gefitinib response	Link		Link
1.1	rs4988235	(T;T)	Can digest milk			Link
1	rs182549	(T;T)	Can digest milk.			Link
1	rs2952768	(C;C)	Less drug dependence: decreased effectiveness o			Link
1	rs7850258	(A;G)	Typical odds of developing primary hypothyroidi			
1	rs800292	(C;T)	1% decreased risk of macular degeneration	Link	Link	Link
0.1	rs891512	(G;G)	Lower blood pressure than those with an A allel	Link		
0	rs10427255	(T;T)	Lowest odds of photic sneeze reflex			
0	rs1047781	(A;A)	ABH blood group "Secretor" status if Japanese	Link	Link	Link
0	rs1126809	(A;G)	Slight increase in skin cancer risk	Link	Link	Link
0	rs16990018	(A;A)	PrP Codon 171 Asn - Non-pathogenic variant	Link		Link
0	rs17244841	(A;A)	More responsive to statin treatment		Link	Link
0	rs1800562	(G;G)	Not a C282Y hemochromatosis carrier.	Link	Link	Link
0	rs28933385	(G;G)	Prion protein Codon 200 (E) - Non pathogenic va			Link
0	rs312481	(C;C)	Better response to certain calcium channel bloc			
0	rs5065	(A;A)	1.12x risk on diuretic; if hypertensive: better	Link	Link	Link
0	rs6259	(G;G)	Best inverse correlation between tea-drinking:	Link	Link	
0	rs74315403	(G;G)	PrP codon 178 (D) - non pathogenic variant			Link
0	rs7495174	(A;A)	Blue/gray eyes more likely		Link	
0	rs7997012	(A;A)	~18% more likely to respond to citalopram		Link	Link
0	rs9951307	(A;G)	0.10 decreased risk for brain edema after a str			

### 3.2 Possibly Harmful Traits

Mag.	Identifier	Genotype	Summary	ExAC	GetEvidence	ClinVar
3	rs1021737	(T;T)	Significantly higher plasma total homocysteine	Link	Link	Link
3	rs10897346	(C;C)	If depressed: 2.6x more likely to not respond t			
3	rs13266634	(C;C)	Increased risk for type-2 diabetes	Link	Link	Link
3	rs2237717	(T;T)	Reduced abilities related to neurocognition and			
3	rs2306402	(C;C)	1.18x increased risk for late-onset Alzheimer's			
3	rs2981582	(C;T)	1.3x higher risk of ER+ breast cancer		Link	
3	rs7754840	(C;C)	1.3x increased risk for type-2 diabetes		Link	
2.5	rs1799971	(A;G)	Stronger cravings for alcohol. if alcoholic: na	Link	Link	Link
2.5	rs187238	(G;G)	Hypertension increases risk 3.75x for sudden ca			
2.5	rs2943634	(C;C)	Slightly higher risk of ischemic stroke		Link	
2.5	rs339331	(T;T)	Prostate cancer risk			
2.5	rs5888	(C;T)	3x higher risk for age-related macular degenera	Link		
2.5	rs664143	(C;T)	Higher risk for number of cancers			
2.5	rs7574865	(G;T)	1.3x risk of rheumatoid arthritis; 1.55x risk o		Link	Link
2.5	rs8034191	(C;T)	1.27x lung cancer risk		Link	
2.4	rs2274223	(G;G)	1.9x increased risk for stomach and esophageal	Link	Link	Link
2.4	rs7966230	(G;G)	Slightly lower levels of plasma VWF			
2.3	rs1859962	(G;G)	1.28x increased risk for prostate cancer		Link	
2.2	rs2004640	(G;T)	1.4x increased risk for SLE		Link	Link
2.2	rs2231137	(G;G)	$\sim$ 1.5-3x increased risk for ischemic stroke	Link	Link	Link
2.2	rs944289	(T;T)	1.69x increased thyroid cancer risk		Link	
2.1	rs1050152	(T;T)	2.1x increased risk of Crohn's disease	Link	Link	Link
2.1	rs10811661	(T;T)	1.2x increased risk for type-2 diabetes		Link	
2.1	rs17070145	(C;C)	Reduced memory abilities			Link
2.1	rs3746444	(C;C)	$\sim 1.2x$ increased risk for cancer	Link		
2.1	rs5186	(A;C)	$\sim$ 1.4x increased risk of hypertension	Link	Link	Link
2.1	rs646776	(A;A)	1.2x risk of coronary artery disease		Link	
2.1	rs795484	(A;G)	Increased morphine dose requirement and postope			
2	rs10086908	(C;T)	1.7x increased risk for prostate cancer			
2	rs1024611	(C;T)	Increased risk of exercise induced ischemia			Link
2	rs10248420	(A;A)	7x less likely to respond to certain antidepres		Link	
2	rs1042838	(G;T)	1.28x risk for endometrial ovarian cancer; over	Link	Link	
2	rs10513789	(G;T)	Increased risk of Parkinson's disease			
2	rs1051730	(C;T)	1.3x increased risk of lung cancer	Link	Link	Link
2	rs10984447	(A;A)	>1.17x increased risk for multiple sclerosis		Link	
2	rs1160312	(A;G)	1.6x increased risk of Male Pattern Baldness.		Link	
2	rs11983225	(T;T)	7x less likely to respond to certain antidepres		Link	
2	rs12037606	(A;A)	1.52x risk of developing Crohn's disease			
2	rs1219648	(A;G)	1.20x risk for breast cancer		Link	
2	rs12567232	(A;G)	Increased risk for Crohn's Disease		Link	
2	rs12696304	(G;G)	Prone to aging faster: at least in European pop			
2	rs1360780	(C;T)	1.3x increased risk for depression		Link	
2	rs16942	(A;G)	Very slightly increased breast cancer risk	Link	Link	Link
2	rs16944	(A;A)	Increased risk for osteoarthritis		Link	
2	rs17228212	(C;T)	1.26x increased risk for heart disease		Link	
2	rs1734791	(A;A)	1.4x increased risk for lupus			
2	rs17576	(G;G)	Higher risk for lung cancer: and COPD in smoker	Link	Link	
2	rs17696736	(A;G)	1.34x risk of type-1 diabetes		Link	
2	rs1800896	(A;G)	1.6x increased prostate cancer risk			
2	rs2056116	(G;G)	1.41x risk for breast cancer			
2	rs2143340	(C;T)	Increased risk of dyslexia and poor reading per			
2	rs2156921	(A;G)	1.29x increased risk for depression			
2	rs2201841	(C;T)	1.5x increased risk for Crohn's disease; 2x inc		Link	

Mag.	Identifier	Genotype	Summary	ExAC	GetEvidence	ClinVar
2	rs2230201	(G;G)	>1.4x risk of lupus	Link		
2	rs2235015	(G;G)	Somewhat less likely to respond to certain anti	Link	Link	
2	rs2235040	(G;G)	7x less likely to respond to certain antidepres	Link	Link	
2	rs2235067	(G;G)	7x less likely to respond to certain antidepres			
2	rs2305795	(A;G)	1.28x higher risk of narcolepsy compared to (G;			Link
2	rs2383206	(A;G)	1.4x increased risk for heart disease			
2	rs2383207	(A;G)	Increased risk for heart disease			
2	rs241448	(C;T)	1.51x increased risk for Alzheimer's	Link		Link
2	rs2420946	(C;T)	1.20x risk for breast cancer			
2	rs25487	(A;G)	2x higher risk for skin cancer; possibly other	Link	Link	Link
2	rs2736100	(T;T)	Higher risk of Interstitial lung disease: and t		Link	
2	rs3025039	(T;T)	2.6x increased risk for ARMD in a Taiwanese pop			
2	rs3738919	(A;C)	1.94x risk of developing rheumatoid arthritis			
2	rs4027132	(A;A)	1.51x increased risk of developing bipolar diso			
2	rs4148739	(A;A)	7x less likely to respond to certain antidepres		Link	
2	rs4402960	(G;T)	1.2x increased risk for type-2 diabetes: ~1x ri		Link	Link
2	rs4444903	(A;G)	3.5x risk of hep-cancer in cirrhosis patients;			Link
2	rs4633	(C;T)	Higher risk for endometrial cancer	Link	Link	Link
2	rs4792311	(A;G)	Increased risk of prostate cancer	Link	Link	Link
2	rs5174	(A;G)	1.3x increased risk for heart disease	Link	Link	Link
2	rs520354	(A;A)	Increased risk in men for biliary conditions			
2	rs5759167	(T;T)	Higher prostate cancer risk		Link	
2	rs587776825	(-;C)	Associated with MODY3; maturity onset of diabet	Link		Link
2	rs629242	(C;T)	Somewhat higher risk for prostate cancer			
2	rs6441286	(G;T)	1.54x chance of developing primary biliary cirr		Link	
2	rs6457617	(C;T)	2.3x risk of rheumatoid arthritis		Link	
2	rs6498169	(A;A)	>1.14x risk of multiple sclerosis		Link	
2	rs663048	(G;T)	Significantly increased risk of developing lung	Link	Link	
2	rs6807362	(C;C)	Increased autism risk	Link	Link	
2	rs6896702	(T;T)	Increased risk of developing Parkinson's Diseas			
2	rs6908425	(C;C)	1.95x increased risk of developing Crohn's dise		Link	
2	rs6997709	(G;T)	1.2x higher risk for hypertension			
2	rs699	(C;C)	Increased risk of hypertension	Link	Link	Link
2	rs7216389	(T;T)	1.5x increased risk for Childhood Asthma.		Link	
2	rs7794745	(A;T)	Slightly increased risk for autism		Link	Link
2	rs7807268	(C;G)	1.3x risk for Crohn's disease		Link	
2	rs854560	(A;T)	Higher risk for heart disease: diabetic retinop	Link	Link	Link
2	rs965513	(A;G)	1.77x increased thyroid cancer risk		Link	
2	rs9954153	(G;G)	$\sim 5x$ higher risk for Fuchs' dystrophy: a corneal			
2.0	rs4911414	(G;T)	2-4x higher risk of sun sensitivity if part of		Link	
1.9	rs7923837	(A;G)	1.6x risk for T2D			
1.8	rs2278206	(T;T)	1.16x increased risk for asthma	Link	Link	
1.8	rs37973	(A;G)	Among asthmatics: 1.5x more likely to show less			Link
1.6	rs11523871	(A;C)	1.6x increased breast cancer risk for women ove	Link	Link	
1.6	rs2981745	(C;T)	1.6x increased risk for breast cancer in female			
1.6	rs3764880	(A;A)	1.2 - 1.8x increased tuberculosis risk	Link	Link	
1.6	rs3775948	(C;G)	Slightly higher risk for gout			
1.6	rs4959039	(G;G)	1.6x higher risk for multiple sclerosis			
1.5	rs10859871	(A;C)	Slight ( $~1.2x$ ) increase in endometriosis risk			
1.5	rs11171739	(C;T)	1.34x risk of developing Type-1 diabetes		Link	
1.5	rs1169300	(A;G)	~1.5x increased lung cancer risk			
1.5	rs12210050	(C;T)	Slightly higher risk for basal cell carcinoma		Link	
1.5	rs12431733	(C;T)	Slightly increased risk of developing Parkinson		Link	
1.5	rs13149290	(C;C)	Slightly increased risk of developing prostate			
1.5	rs1801020	(C;T)	1.31x increased risk of heart disease	Link		Link

Mag.	Identifier	Genotype	Summary	ExAC	GetEvidence	ClinVa
1.5	rs1801274	(T;T)	Complex; generally greater risk for cancer prog	Link	Link	Link
1.5	rs1867277	(A;G)	1.5x increased risk for thyroid cancer			
1.5	rs2007153	(G;G)	Increased risk of schizophrenia in limited stud			
1.5	rs2177369	(C;C)	1.5x increased risk for Alzheimer's disease			
1.5	rs2240340	(A;G)	Slightly increased $(1.5x)$ risk for RA	Link		
1.5	rs2241880	(C;T)	1.4x increased risk for Crohn's disease in Cauc	Link	Link	Link
1.5	rs2280714	(A;G)	1.4x increased risk of SLE			
1.5	rs2464196	(C;T)	~1.5x increased lung cancer risk	Link	Link	Link
1.5	rs2697962	(A;G)	Slightly increased risk of developing Parkinson			
1.5	rs2881766	(G;T)	Slightly increased risk for pregnancy-induced h			
1.5	rs3087243	(A;G)	Increased risk for auto-immune diseases		Link	
1.5	rs3212227	(A;A)	1.43x increased risk of developing psoriasis an			
1.5	rs358806	(A;C)	0.86x increased risk of developing Type-2 diabe		Link	
1.5	rs3814570	(C;T)	1.3x increased risk for Crohn's disease with il			
1.5	rs3825776	(A;G)	1.3x increased risk for ALS		Link	
1.5	rs393152	(A;A)	Increased risk of both PD and AD	Link	Link	
1.5	rs4464148	(C;T)	1.10x increased risk for colorectal cancer	171117	1711117	
1.5	rs46404148	(C,T) (C;T)	Increased risk of schizophrenia in limited stud			
$1.5 \\ 1.5$	rs4845618		1.7x increased melanoma risk			
1.5 1.5	rs4845018 rs486907	(G;T)	1.7x increased melanoma risk 1.5x increased prostate cancer risk	Link	Link	Link
$1.5 \\ 1.5$	rs480907 rs5219	(A;G)	_	Link	Link	Link
		(C;T)	1.3x increased risk for type-2 diabetes	LIIIK	LIIIK	LIIIK
1.5	rs5746059	(A;G)	Slightly higher fat mass		т. 1	
1.5	rs642961	(A;G)	1.68x increased risk of cleft lip		Link	
1.5	rs6601764	(C;T)	1.16x increased risk of developing Crohn's dise		Link	
1.5	rs699473	(C;C)	~1.5x increased brain tumor risk			
1.5	rs7774434	(C;T)	Slightly increased risk of developing primary b			
1.5	rs872071	(A;G)	~1.5x increased risk for chronic lymphocytic le		Link	
1.5	rs9652490	(A;G)	Slightly increased risk of developing Parkinson		Link	
1.5	rs995030	(G;G)	Non-protective against testicular cancer		Link	
1.4	rs10865331	(A;A)	1.4x higher risk for ankylosing spondylitis			
1.4	rs1126497	(T;T)	1.4x increased risk for breast cancer	Link	Link	Link
1.4	rs1545843	(A;A)	1.4x increased risk for depression (for those u			
1.4	rs2046210	(C;T)	1.4x increased breast cancer risk		Link	Link
1.4	rs3131296	(G;G)	1.4x increased risk for schizophrenia		Link	
1.4	rs3184504	(C;T)	Slightly increased risk for celiac disease	Link	Link	
1.4	rs4977756	(G;G)	1.93x higher risk for glioma development		Link	
1.4	rs6010620	(G;G)	1.4x higher risk for glioma development; but th		Link	
1.34	rs17465637	(C;C)	1.34x higher risk for myocardial infarction	Link	Link	
1.3	rs10947262	(C;C)	1.3x increased risk for osteoarthritis			
1.3	rs1260326	(C;T)	Slightly higher risk for gout	Link	Link	Link
1.3	rs1434536	(A;G)	1.29x increased breast cancer risk			
1.3	rs1746048	(C;C)	1.03 increased risk for coronary heart disease		Link	
1.3	rs2024513	(A;G)	1.3x higher risk for schizophrenia (among Han C			
1.3	rs34330	(C;T)	1.3x higher risk for endometrial cancer (in Chi			
1.2	rs11037909	(C,T) (T;T)	1.47x type II diabetes risk	Link		
1.2	rs1800693	(A;G)	Slight (1.2x) increase in risk for multiple scl	Link	Link	Link
1.2	rs2076295	(G;T)	One copy of the risk allele (G): slightly incre	11111	171111	
1.2	rs3740878	(G,1) (A;A)	1.46x type II diabetes risk; common	Link		Link
1.2	rs449647	(A;A) (A;T)	Possibly lower levels of ApoE	AIIIK		LIIIK
1.2 1.2	rs4686484		Slightly increased risk for celiac disease			
		(A;A)			Linl-	
1.15	rs748404	(C;T)	Very slightly increased risk (1.15) for lung ca		Link	
1.1	rs1110912	(C;C)	1.3x high blood pressure risk			
1.1	rs11650494	(A;G)	Slightly higher prostate cancer risk		T 1 1	
1.1	rs13387042	(A;G)	1.12x increased risk for breast cancer	T	Link	T
1.1	rs2295190	(G;T)	Slightly increased risk for ovarian cancer in w	Link	Link	Link

Mag.	Identifier	Genotype	Summary	ExAC	GetEvidence	ClinVar
1.1	rs249954	(C;T)	Potentially increased risk of Breast Cancer			Link
1.1	rs34516635	(G;G)	Less longevity for Ashkenazi Jewish women.	Link		Link
1.1	rs4324715	(C;T)	1.5x increased testicular cancer risk for men			
1.1	rs6800901	(T;T)	1.3x multiple myeloma risk			
1.1	rs688034	(C;T)	1.1x risk higher risk for coronary artery disea		Link	
1.1	rs6897876	(C;T)	Slight increase in testicular cancer risk for m			
1.1	rs7412	(C;T)	More likely to gain weight if taking olanzapine	Link	Link	Link
1.1	rs925391	(C;C)	More likely to go bald; common			
1.09	rs12050604	(A;C)	Very slightly increased risk for lung cancer			
1.07	rs2291834	(C;C)	Very slightly higher risk for myocardial infarc			
1	rs10504861	(G;G)	Major allele: normal risk of migraine			
1	rs10761659	(A;G)	1.2x risk of Crohn's disease		Link	
1	rs1143674	(A;A)	1.3x increased autism risk	Link		
1	rs2546890	(A;A)	Higher risk of multiple sclerosis			
1	rs3194051	(A;A)	>1.1x risk of type-1 diabetes	Link	Link	Link
1	rs6932590	(T;T)	1.1x increased risk for schizophrenia		Link	
1	rs987525	(A;C)	2.5x increased risk for cleft lip		Link	
0.1	rs601338	(A;G)	Susceptible to Norovirus infections	Link	Link	Link
0	rs1042173	(T;T)	Among alcoholics: likely to be heavier drinkers			
0	rs1495965	(A;A)	1.2x higher risk for spondylitis			
0	rs3813929	(C;C)	Possible weight gain if taking olanzapine		Link	Link
0	rs4293393	(T;T)	$1.25 \mathrm{x}$ Increased Risk of CKD for T allele in			
0	rs6314	(C;C)	Higher risk for RA	Link	Link	
0	rs7787082	(G;G)	7x less likely to respond to certain antidepres		Link	

#### 3.3 Genosets (Multi-variant Phenotypes)

Magnitude	Identifier	Summary
3	gs241	Lighter green: brown or hazel eye color
2.5	gs281	Part of the $88\%$ of the population claimed not t
2.5	gs285	You will lose 2.5x as much weight on a low fat
2.4	gs297	Lower heart attack risk than average
2	gs101	Probably able to digest milk
2	gs239	Reduced conversion of beta-carotene to retinol
2	gs249	Parkinson's Disease Risk
2	gs269	APOE $E2/E3$
1.2	gs184	Able to taste bitterness.
1	gs182	CYP2D6*39
0	gs158	CYP1A2 normal metabolizer

## 4 Raw Data

The raw data used to create this report has been assigned the identifier ERS1176623 in the European Nucleotide Archive (ENA) hosted at the European Bioinformatics Institute (EBI).

These data will not be accessible unless the report is approved. This will happen by default one month after the report is issued, or if the report is approved for immediate release within the one month period. Participants can also withdraw from the study at any time in which case the report and the data will not be released and will be deleted.

If the data has already been released, it can be accessed at: http://www.ebi.ac.uk/ena/data/view/ERS1176623

### 5 Report Metadata

Resource	Version	Website
Genome	GRCh38	Link
BWA	0.7.12	Link
SAMtools	1.3	Link
GATK	3.4-46	Link
PLINK	v1.90b3.35	Link
VEP	88	Link
SNPedia	30-Jul-2017	Link
ExAC	v0.3.1	Link
GetEvidence	16-Dec-2016	Link
ClinVar	16-Dec-2016	Link

Table 5: Analysis Pipeline Versions

Report generated on August 2, 2017.