

**A Vertebrate species**

Species Information

Taxon	GenBank ID
oryctolagus cuniculus	GCA_000003625.1

Total 2220 10/page < > Go to 1

Filter evidence status  Filter mapped to hg38

RS ID	Chr	Position	Ref	Alt	Evidence Status	Map(hg38)
rs3000161745	1	649376	C	T	NA	NA
rs3000180735	1	3038710	C	T	NA	NA
rs3000180735	1	3038710	C	T	NA	NA
rs3000180735	1	3038710	C	T	NA	NA
rs3000180735	1	3038710	C	T	NA	NA
rs3000180735	1	3038710	C	T	NA	NA
rs3000180735	1	3038710	C	T	NA	NA

Data Source Total Amount

European Variation Archive 981

Mapped Amount Evolutionary Distance

131 0.3769

**B Economic species**

Total 4490 10/page < > Go to 1

Trait  P-value <

Variant	Chr	Position	Ref	Alt	Trait	P-Value	Gene ID	Source	So
6_28019382_C/G	6	28019382	C	G	stem thickness	9.02e-8	Os06g0675200	GWAS Atlas	OSI
6_28356308_C/T	6	28356308	C	T	total lesion length	0.00000184	Os06g0680700	GWAS Atlas	OSI
6_28100960_A/G	6	28100960	A	G	rice bacterial blight disease resistance	0.00000215	Os06g0676700	GWAS Atlas	OSI
6_28019382_C/G	6	28019382	C	G	breaking strength of the basal internode	0.0000208	Os06g0675200	GWAS Atlas	OSI
6_28019382_C/G	6	28019382	C	G	stem diameter	0.000114	Os06g0675200	GWAS Atlas	OSI
6_28019382_C/G	6	28019382	C	G	stem diameter	0.000253	Os06g0675200	GWAS Atlas	OSI
6_28019382_C/G	6	28019382	C	G	stem diameter	0.000279	Os06g0675200	GWAS Atlas	OSI
6_28019382_C/G	6	28019382	C	G	stem diameter	0.000285	Os06g0675200	GWAS Atlas	OSI
6_27969431_C/G	6	27969431	C	G	percentage of head rice recovery	0.000822	Os06g0674000	GWAS Atlas	OSI

**C Primate species**

Gorilla ✓	Pan troglodytes ✓	Pongo abelii	Pongo pygmaeus	Bonobo

Total 57775 10/page < > Go to 1

Allele Frequency >

Variant (Mapped to hg38)	Gorilla	Pan troglodytes	Pongo abelii	Pongo pygmaeus
10_100048761_C/T	1	1	1	1
10_100065197_G/A	1	0.98	1	1
10_100081557_A/G	1	1	1	1
10_100185531_A/G	1	1	1	1
10_100329211_G/A	1	1	1	1
10_100354576_G/C	1	1	NA	NA
10_100487721_A/G	1	1	1	1
10_100488894_C/T	1	1	1	1
10_100506058_A/G	1	1	NA	NA
10_100509448_A/T	1	1	1	1

**Supplemental Figure S11.** Browse non-human species. (A) For vertebrate species, users can filter out records with "evidence status" and "hg38" (those mapped to the reference genome hg38 by MSA). (B) For economic species, filtering is available based on trait and p-value. (C) For primate species, users can refine results by allele frequency.