

# Resource Summary Report

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## Japanese Single Nucleotide Polymorphisms

RRID:SCR\_013076

Type: Tool

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### Proper Citation

Japanese Single Nucleotide Polymorphisms (RRID:SCR\_013076)

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### Resource Information

**URL:** <http://snp.ims.u-tokyo.ac.jp/>

**Proper Citation:** Japanese Single Nucleotide Polymorphisms (RRID:SCR\_013076)

**Description:** JSNP is a database of Japanese Single Nucleotide Polymorphisms. It includes BLAST capability, keyword search, mapping information, and other tools that allow users to gather information on SNP's. SNPs are the most common form of DNA sequence variation. They are useful polymorphic markers to investigate genes susceptible to diseases or those related to drug responsiveness. Furthermore, a small subset of SNPs directly influences to the quality and/or quantity of the gene product, and increase a risk to certain diseases and to severe side effect by drugs. Through a discovery of a large number of SNPs, we would like to contribute to identification of disease-related genes and also to establish a diagnostic method to avoid drug side-effect.

**Synonyms:** JSNP

**Resource Type:** data or information resource, database

**Keywords:** FASEB list

**Resource Name:** Japanese Single Nucleotide Polymorphisms

**Resource ID:** SCR\_013076

**Alternate IDs:** nif-0000-03063

**Record Creation Time:** 20220129T080314+0000

**Record Last Update:** 20260905T133204+0000

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### Ratings and Alerts

No rating or validation information has been found for Japanese Single Nucleotide Polymorphisms.

No alerts have been found for Japanese Single Nucleotide Polymorphisms.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 42 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [RRID](#) .

Roberts R, et al. (2020) A Journey through Genetic Architecture and Predisposition of Coronary Artery Disease. *Current genomics*, 21(5), 382.

Shijo T, et al. (2020) Association between Polygenic Risk Score and One-Year Outcomes Following As-Needed Aflibercept Therapy for Exudative Age-Related Macular Degeneration. *Pharmaceuticals (Basel, Switzerland)*, 13(9).

Zhang J, et al. (2017) MGST2 and WNT2 are candidate genes for comitant strabismus susceptibility in Japanese patients. *PeerJ*, 5, e3935.

Sakai E, et al. (2016) TP53 mutation at early stage of colorectal cancer progression from two types of laterally spreading tumors. *Cancer science*, 107(6), 820.

Oda Y, et al. (2015) Genetic association between G protein-coupled receptor kinase 6/?-arrestin 2 and dopamine supersensitivity psychosis in schizophrenia. *Neuropsychiatric disease and treatment*, 11, 1845.

Otsuka I, et al. (2015) Association analysis of the Cadherin13 gene with schizophrenia in the Japanese population. *Neuropsychiatric disease and treatment*, 11, 1381.

Fujisawa Y, et al. (2015) Impact of a novel homozygous mutation in nicotinamide nucleotide transhydrogenase on mitochondrial DNA integrity in a case of familial glucocorticoid deficiency. *BBA clinical*, 3, 70.

Kucukkal TG, et al. (2014) Computational and experimental approaches to reveal the effects of single nucleotide polymorphisms with respect to disease diagnostics. *International journal of molecular sciences*, 15(6), 9670.

Inaguma Y, et al. (2014) SIL1, a causative cochaperone gene of Marinesco-Sjögren syndrome, plays an essential role in establishing the architecture of the developing cerebral cortex. *EMBO molecular medicine*, 6(3), 414.

Miura K, et al. (2012) An overgrowth disorder associated with excessive production of cGMP due to a gain-of-function mutation of the natriuretic peptide receptor 2 gene. *PloS one*, 7(8), e42180.

Ikegawa S, et al. (2012) A short history of the genome-wide association study: where we were and where we are going. *Genomics & informatics*, 10(4), 220.

Iida M, et al. (2012) A novel MPZ mutation in Charcot-Marie-Tooth disease type 1B with focally folded myelin and multiple entrapment neuropathies. *Neuromuscular disorders : NMD*, 22(2), 166.

De Luca D, et al. (2011) Secretory phospholipase A2 pathway in various types of lung injury in neonates and infants: a multicentre translational study. *BMC pediatrics*, 11, 101.

Cui H, et al. (2011) Association study of EP1 gene polymorphisms with suicide completers in the Japanese population. *Progress in neuro-psychopharmacology & biological psychiatry*, 35(4), 1108.

Supriyanto I, et al. (2011) Association of FKBP5 gene haplotypes with completed suicide in the Japanese population. *Progress in neuro-psychopharmacology & biological psychiatry*, 35(1), 252.

Shigeto S, et al. (2011) Improved assay for differential diagnosis between Pompe disease and acid  $\alpha$ -glucosidase pseudodeficiency on dried blood spots. *Molecular genetics and metabolism*, 103(1), 12.

Daimon M, et al. (2011) Association of the clusterin gene polymorphisms with type 2 diabetes mellitus. *Metabolism: clinical and experimental*, 60(6), 815.

Seaver LH, et al. (2011) A novel mutation in the HSD17B10 gene of a 10-year-old boy with refractory epilepsy, choreoathetosis and learning disability. *PloS one*, 6(11), e27348.

Ikeuchi T, et al. (2011) Evidence for a Common Founder and Clinical Characteristics of Japanese Families with the MAPT R406W Mutation. *Dementia and geriatric cognitive disorders extra*, 1(1), 267.

Maeda S, et al. (2010) A single nucleotide polymorphism within the acetyl-coenzyme A carboxylase beta gene is associated with proteinuria in patients with type 2 diabetes. *PLoS genetics*, 6(2), e1000842.