

研究成果の刊行に関する一覧表

雑誌

発表者氏名	論文タイトル名	発表誌名	巻号	ページ	出版年
Shigemizu D, Mitsumori R, Akiyama S, Miyashita A, Morizono T, Higaki S, Asanomi Y, Hara N, Tamiya G, Kinoshita K, Ikeuchi T, Niida S, <u>Ozaki K</u> .	Ethnic and trans-ethnic genome-wide association studies identify new loci influencing Japanese Alzheimer's disease risk.	Translational Psychiatry	11	151	2021
Shigemizu D, Akiyama S, Higaki S, Sugimoto T, Sakurai T, Boroevich KA, Sharma A, Tsunoda T, Ochiya T, Niida S, <u>Ozaki K</u> .	Prognosis prediction model for conversion from mild cognitive impairment to Alzheimer's disease created by integrative analysis of multi-omics data.	Alzheimer's Research & Therapy	12	145	2020
Johnson T, Mashimo Y, Wu JY, Yoon D, Hata A, Kubo M, Takahashi A, Tsunoda T, <u>Ozaki K</u> , Tanaka T, Ito K, Suzuki H, Hamada H, Kobayashi T, Hara T, Chen CH, Lee YC, Liu YM, Chang LC, Chang CP, Hong YM, Jang G, Yun S, Yu J, Lee KY, Kim JJ, Park T, Lee JK, Chen YT, <u>O no u c h i Y</u> .	Published Online 26 October (2020). doi: 10.1038/s10038-020-00864-z. Association of an IGHV3-66 gene variant with Kawasaki disease.	Journal of Human Genetics	Published Online		2020
Koyama S, Ito K, Terao C, Akiyama M, Horikoshi M, Momozawa Y, Matsunaga H, Ieki H, <u>Ozaki K</u> , Onouchi Y, et al.	Population-specific and trans-ancestry genome-wide analyses identify distinct and shared genetic risk loci for coronary artery disease.	Nature Genetics	52	1169-1177	2020
Ishigaki K, Akiyama M, Kanai M, Takahashi A, Kawakami E, Sugishita H, Sakaue S, Matoba N, Low SK, Okada Y, Terao C, Amariuta T, Gazal S, Kochi Y, Horikoshi M, Suzuki K, Ito K, Koyama S, <u>Ozaki K</u> , Niida S. et al.	Large-scale genome-wide association study in a Japanese population identifies novel susceptibility loci across different diseases.	Nature Genetics	52 (7)	669-679	2020

<p>Matsunaga H, Akiyama M, Takahashi A, Nomura S, Ozaki K, Onouchi Y, Suni a S, Ogishima S, Yamamoto M, Satoh M, Sasaki M, Yamaji T, Iwasaki M, Tsugane S, Tanaka K, Naito M, Akai W, Tanaka H, Sakata Y, Morita H, Matsuda K, Murakami Y, Akazawa H, Kubo M, Kamatani Y, Komuro I, Ito K.</p>	<p>Transethnic meta-analysis of genome-wide association studies identifies three new loci for coronary artery disease.</p>	<p>Circulation; Genomics and Precision Medicine.</p>	<p>13(3)</p>	<p>e002670</p>	<p>2020</p>
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