

項目#	著者名	発行年	雑誌名(出版社名・ウェブ名)	タイトル	ウェブアドレス
1	リッカルド・サバティーニ Riccardo Sabatini	2016	TED	ゲノムを読んで人間を作る方法	https://www.ted.com/talks/riccardo_sabatini_how_to_read_the_genome_and_build_a_human_being?language=ja
2	厚生労働省 社会・援護局障害保健福祉部	2016	参考資料 - 厚生労働省	精神疾患を有する総患者数の推移 厚生労働省「患者調査」より	http://www.mhlw.go.jp/file/05-Shingikai-12201000-Shakaiengokyo/kyokushougai/hokenfukushibu/Kikakuka/0000108755_12.pdf
3	Ikeda M et al.	2017	Molecular Psychiatry, in press	A genome-wide association study identifies two novel susceptibility loci and trans population polygenicity associated with bipolar disorder	
3	CONVERGE consortium	2015	Nature 523, 588-591	Sparse whole-genome sequencing identifies two loci for major depressive disorder	
4	Uher R	2014	Frontiers in Psychiatry 5: 48	Gene-environment interactions in severe mental illness	
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5	Dunn DB, Smith JD	1955	Nature 175, 336-337	Occurrence of a new base in the deoxyribonucleic acid of a strain of Bacterium coli	
5	Gowher H et al.	2000	EMBO Journal 19, 6918-6923	DNA of Drosophila melanogaster contains 5-methylcytosine	
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6	Lister R et al.	2009	Nature 462, 315-322	Human DNA methylomes at base resolution show widespread epigenomic differences	
8	Goto K et al.	1994	Differentiation 56, 39-44	Expression of DNA methyltransferase gene in mature and immature neurons as well as proliferating cells in mice	
8	Feng J et al.	2005	Journal of Neuroscience Research 79, 734-746	Dynamic expression of de novo DNAmethyltransferases Dnmt3a and Dnmt3b in	

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9	岡田典弘	ラボ ウェブ	東工大 岡田研究室 Research Keywords	Aluとは・・・	
9	Hayakawa K et al.	2012	Mammalian Genome 23, 336-345	Bridging sequence diversity and tissue-specific expression by DNA methylation in genes of the mouse prolactin superfamily	
9	有馬隆博	2015	生命誌ジャーナル87号	ヒトから知るエピジェネティクスと進化	http://www.brh.co.jp/seimeishi/journal/087/research/2.html
10	Edited by Brendy TW	2017	(出版社) Academic Press	DNA modifications in the brain	
10	Szulwach KE et al.	2011	Nature Neuroscience 14, 1607-1616	5-hmC-mediated epigenetic dynamics during postnatal neurodevelopment and aging	
11	Sugawara H et al.	2011	Translational Psychiatry 1: e24	Hypermethylation of serotonin transporter gene in bipolar disorder detected by epigenome analysis of discordant monozygotic twins	
12	Nedergaard M et al.	2003	Trends in Neuroscience 26(10): 523-530.	New roles for astrocytes: redefining the functional architecture of the brain	
12	Azevedo FAC et al.	2009	The Journal of Comparative Neurology 513, 532-541	Equal numbers of neuronal and nonneuronal cells make the human brain an isometrically scaled-up primate brain	
12	Iwamoto K et al.	2011	Genome Research 21, 688-696	Neurons show distinctive DNA methylation profile and higher interindividual variations compared with non-neurons	
13	Alzheimer's Disease International	2018	World Alzheimer Report 2018		https://www.alz.co.uk/research/world-report-2018
14	Lunnon K et al.	2014	Nature Neuroscience 17, 1164-1170	Methylomic profiling implicates cortical deregulation of ANK1 in Alzheimer's disease	
14	De Jager PL et al.	2014	Nature Neuroscience 17, 1156-1163	Alzheimer's disease: early alterations in brain DNA methylation at ANK1, BIN1, RHBDF2 and other loci	
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