

Toshiko Tanaka

List of Publications by Citations

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Version: 2024-04-27

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131
papers

15,047
citations

52
h-index

122
g-index

140
ext. papers

19,147
ext. citations

10.6
avg, IF

5.07
L-index

#	Paper	IF	Citations
131	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015 , 518, 197-206	50.4	2687
130	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014 , 46, 1173-86	36.3	1339
129	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015 , 518, 187-196	50.4	920
128	Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. <i>Nature Genetics</i> , 2012 , 44, 991-1005	36.3	621
127	A genome-wide approach accounting for body mass index identifies genetic variants influencing fasting glycemic traits and insulin resistance. <i>Nature Genetics</i> , 2012 , 44, 659-69	36.3	615
126	DNA methylation-based measures of biological age: meta-analysis predicting time to death. <i>Aging</i> , 2016 , 8, 1844-1865	5.6	531
125	Genetic variation in GIPR influences the glucose and insulin responses to an oral glucose challenge. <i>Nature Genetics</i> , 2010 , 42, 142-8	36.3	527
124	Epigenetic clock analysis of diet, exercise, education, and lifestyle factors. <i>Aging</i> , 2017 , 9, 419-446	5.6	317
123	Common variants at 10 genomic loci influence hemoglobin A1C levels via glycemic and nonglycemic pathways. <i>Diabetes</i> , 2010 , 59, 3229-39	0.9	314
122	Genome-wide association study of plasma polyunsaturated fatty acids in the InCHIANTI Study. <i>PLoS Genetics</i> , 2009 , 5, e1000338	6	300
121	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. <i>Nature Genetics</i> , 2017 , 49, 834-841	36.3	257
120	Seventy-five genetic loci influencing the human red blood cell. <i>Nature</i> , 2012 , 492, 369-75	50.4	257
119	Genetic loci associated with plasma phospholipid n-3 fatty acids: a meta-analysis of genome-wide association studies from the CHARGE Consortium. <i>PLoS Genetics</i> , 2011 , 7, e1002193	6	257
118	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. <i>Nature Genetics</i> , 2016 , 48, 1171-1184	36.3	251
117	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , 2015 , 47, 1294-1303	36.3	226
116	Impact of common genetic determinants of Hemoglobin A1c on type 2 diabetes risk and diagnosis in ancestrally diverse populations: A transethnic genome-wide meta-analysis. <i>PLoS Medicine</i> , 2017 , 14, e1002383	11.6	223
115	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , 2015 , 11, e1005378	6	220

114	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. <i>Nature Genetics</i> , 2014 , 46, 826-36	36.3	199
113	Genome-wide association study of vitamin B6, vitamin B12, folate, and homocysteine blood concentrations. <i>American Journal of Human Genetics</i> , 2009 , 84, 477-82	11	193
112	GWAS of longevity in CHARGE consortium confirms APOE and FOXO3 candidacy. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2015 , 70, 110-8	6.4	188
111	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. <i>Nature Communications</i> , 2016 , 7, 10495	17.4	180
110	DNA methylation signatures of chronic low-grade inflammation are associated with complex diseases. <i>Genome Biology</i> , 2016 , 17, 255	18.3	171
109	CUBN is a gene locus for albuminuria. <i>Journal of the American Society of Nephrology: JASN</i> , 2011 , 22, 555-70	17.0	170
108	Genome-wide meta-analysis identifies six novel loci associated with habitual coffee consumption. <i>Molecular Psychiatry</i> , 2015 , 20, 647-656	15.1	167
107	Plasma proteomic signature of age in healthy humans. <i>Aging Cell</i> , 2018 , 17, e12799	9.9	167
106	Genome-wide meta-analysis of observational studies shows common genetic variants associated with macronutrient intake. <i>American Journal of Clinical Nutrition</i> , 2013 , 97, 1395-402	7	161
105	A higher adherence to a Mediterranean-style diet is inversely associated with the development of frailty in community-dwelling elderly men and women. <i>Journal of Nutrition</i> , 2012 , 142, 2161-6	4.1	161
104	Measuring biological aging in humans: A quest. <i>Aging Cell</i> , 2020 , 19, e13080	9.9	158
103	Genome Analyses of >200,000 Individuals Identify 58 Loci for Chronic Inflammation and Highlight Pathways that Link Inflammation and Complex Disorders. <i>American Journal of Human Genetics</i> , 2018 , 103, 691-706	11	151
102	Novel locus including FGF21 is associated with dietary macronutrient intake. <i>Human Molecular Genetics</i> , 2013 , 22, 1895-902	5.6	134
101	Common genetic loci influencing plasma homocysteine concentrations and their effect on risk of coronary artery disease. <i>American Journal of Clinical Nutrition</i> , 2013 , 98, 668-76	7	122
100	Novel loci affecting iron homeostasis and their effects in individuals at risk for hemochromatosis. <i>Nature Communications</i> , 2014 , 5, 4926	17.4	121
99	Genetic variation at MECOM, TERT, JAK2 and HBS1L-MYB predisposes to myeloproliferative neoplasms. <i>Nature Communications</i> , 2015 , 6, 6691	17.4	120
98	FTO genetic variants, dietary intake and body mass index: insights from 177,330 individuals. <i>Human Molecular Genetics</i> , 2014 , 23, 6961-72	5.6	120
97	A genome-wide association analysis of serum iron concentrations. <i>Blood</i> , 2010 , 115, 94-6	2.2	117

96	Genome-wide association study of plasma N6 polyunsaturated fatty acids within the cohorts for heart and aging research in genomic epidemiology consortium. <i>Circulation: Cardiovascular Genetics</i> , 2014 , 7, 321-331		112
95	Mediterranean diet and mobility decline in older persons. <i>Experimental Gerontology</i> , 2011 , 46, 303-8	4.5	108
94	Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. <i>Nature Communications</i> , 2016 , 7, 10494	17.4	107
93	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. <i>Nature Communications</i> , 2017 , 8, 14977	17.4	105
92	Genome-wide physical activity interactions in adiposity - A meta-analysis of 200,452 adults. <i>PLoS Genetics</i> , 2017 , 13, e1006528	6	103
91	Multiple loci are associated with white blood cell phenotypes. <i>PLoS Genetics</i> , 2011 , 7, e1002113	6	92
90	A SERUM PROTEIN SIGNATURE OF APOE GENOTYPES IN CENTENARIANS. <i>Innovation in Aging</i> , 2019 , 3, S621-S622	0.1	78
89	52 Genetic Loci Influencing Myocardial Mass. <i>Journal of the American College of Cardiology</i> , 2016 , 68, 1435-1448	15.1	76
88	Genome-wide analyses identify a role for SLC17A4 and AADAT in thyroid hormone regulation. <i>Nature Communications</i> , 2018 , 9, 4455	17.4	75
87	Genomewide meta-analysis identifies loci associated with IGF-I and IGFBP-3 levels with impact on age-related traits. <i>Aging Cell</i> , 2016 , 15, 811-24	9.9	71
86	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. <i>Biological Psychiatry</i> , 2017 , 82, 322-329	7.9	68
85	Blood Leukocyte DNA Methylation Predicts Risk of Future Myocardial Infarction and Coronary Heart Disease. <i>Circulation</i> , 2019 , 140, 645-657	16.7	65
84	Genome-wide association analysis of total cholesterol and high-density lipoprotein cholesterol levels using the Framingham heart study data. <i>BMC Medical Genetics</i> , 2010 , 11, 55	2.1	64
83	Identification of a common variant in the TFR2 gene implicated in the physiological regulation of serum iron levels. <i>Human Molecular Genetics</i> , 2011 , 20, 1232-40	5.6	59
82	Association Between Accelerated Multimorbidity and Age-Related Cognitive Decline in Older Baltimore Longitudinal Study of Aging Participants without Dementia. <i>Journal of the American Geriatrics Society</i> , 2016 , 64, 965-72	5.6	58
81	Skeletal muscle ex vivo mitochondrial respiration parallels decline in vivo oxidative capacity, cardiorespiratory fitness, and muscle strength: The Baltimore Longitudinal Study of Aging. <i>Aging Cell</i> , 2018 , 17, e12725	9.9	57
80	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. <i>Human Molecular Genetics</i> , 2016 , 25, 358-70	5.6	54
79	Novel association to the proprotein convertase PCSK7 gene locus revealed by analysing soluble transferrin receptor (sTfR) levels. <i>Human Molecular Genetics</i> , 2011 , 20, 1042-7	5.6	51

78	Genome-wide studies of verbal declarative memory in nondemented older people: the Cohorts for Heart and Aging Research in Genomic Epidemiology consortium. <i>Biological Psychiatry</i> , 2015 , 77, 749-63	7.9	48
77	Underlying features of epigenetic aging clocks in vivo and in vitro. <i>Aging Cell</i> , 2020 , 19, e13229	9.9	37
76	Gene-Environment Interactions of Circadian-Related Genes for Cardiometabolic Traits. <i>Diabetes Care</i> , 2015 , 38, 1456-66	14.6	36
75	Effects of a behavioral intervention that emphasizes spices and herbs on adherence to recommended sodium intake: results of the SPICE randomized clinical trial. <i>American Journal of Clinical Nutrition</i> , 2015 , 102, 671-9	7	36
74	GWAS analysis of handgrip and lower body strength in older adults in the CHARGE consortium. <i>Aging Cell</i> , 2016 , 15, 792-800	9.9	33
73	Dietary fatty acids modulate associations between genetic variants and circulating fatty acids in plasma and erythrocyte membranes: Meta-analysis of nine studies in the CHARGE consortium. <i>Molecular Nutrition and Food Research</i> , 2015 , 59, 1373-83	5.9	32
72	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. <i>Nature Communications</i> , 2019 , 10, 2581	17.4	31
71	Genome-wide meta-analysis of macronutrient intake of 91,114 European ancestry participants from the cohorts for heart and aging research in genomic epidemiology consortium. <i>Molecular Psychiatry</i> , 2019 , 24, 1920-1932	15.1	30
70	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. <i>Nature Communications</i> , 2021 , 12, 24	17.4	30
69	Change in Epigenome-Wide DNA Methylation Over 9 Years and Subsequent Mortality: Results From the InCHIANTI Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2016 , 71, 1029-35	6.4	28
68	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021 , 596, 393-397	30.4	28
67	Meta-analysis of genome-wide association studies for circulating phylloquinone concentrations. <i>American Journal of Clinical Nutrition</i> , 2014 , 100, 1462-9	7	27
66	Elevated Plasma Growth and Differentiation Factor 15 Is Associated With Slower Gait Speed and Lower Physical Performance in Healthy Community-Dwelling Adults. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2020 , 75, 175-180	6.4	27
65	Adherence to a Mediterranean Diet Protects from Cognitive Decline in the Invecchiare in Chianti Study of Aging. <i>Nutrients</i> , 2018 , 10,	6.7	23
64	Association of dietary folate and vitamin B-12 intake with genome-wide DNA methylation in blood: a large-scale epigenome-wide association analysis in 5841 individuals. <i>American Journal of Clinical Nutrition</i> , 2019 , 110, 437-450	7	22
63	Genome-wide association study of breakfast skipping links clock regulation with food timing. <i>American Journal of Clinical Nutrition</i> , 2019 , 110, 473-484	7	22
62	Comparison of HapMap and 1000 Genomes Reference Panels in a Large-Scale Genome-Wide Association Study. <i>PLoS ONE</i> , 2017 , 12, e0167742	3.7	21
61	Fine-mapping, novel loci identification, and SNP association transferability in a genome-wide association study of QRS duration in African Americans. <i>Human Molecular Genetics</i> , 2016 , 25, 4350-4368	5.6	20

60	State- and trait-dependent associations of vitamin-D with brain function during aging. <i>Neurobiology of Aging</i> , 2016 , 39, 38-45	5.6	18
59	Plasma proteomic biomarker signature of age predicts health and life span. <i>ELife</i> , 2020 , 9,	8.9	18
58	Plasma proteomic signature of the risk of developing mobility disability: A 9-year follow-up. <i>Aging Cell</i> , 2020 , 19, e13132	9.9	17
57	Association of Methylation Signals With Incident Coronary Heart Disease in an Epigenome-Wide Assessment of Circulating Tumor Necrosis Factor \square <i>JAMA Cardiology</i> , 2018 , 3, 463-472	16.2	17
56	Sugar-sweetened beverage intake associations with fasting glucose and insulin concentrations are not modified by selected genetic variants in a ChREBP-FGF21 pathway: a meta-analysis. <i>Diabetologia</i> , 2018 , 61, 317-330	10.3	17
55	Cardiovascular Health Is Associated With Physical Function Among Older Community Dwelling Men and Women. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2017 , 72, 1710-1718	6.4	17
54	The complex genetics of gait speed: genome-wide meta-analysis approach. <i>Aging</i> , 2017 , 9, 209-246	5.6	16
53	Extracellular RNA profiles with human age. <i>Aging Cell</i> , 2018 , 17, e12785	9.9	16
52	Genome-wide association meta-analysis of fish and EPA+DHA consumption in 17 US and European cohorts. <i>PLoS ONE</i> , 2017 , 12, e0186456	3.7	15
51	Identification of a novel locus on chromosome 2q13, which predisposes to clinical vertebral fractures independently of bone density. <i>Annals of the Rheumatic Diseases</i> , 2018 , 77, 378-385	2.4	15
50	Vitamin D receptor and megalin gene polymorphisms are associated with central adiposity status and changes among US adults. <i>Journal of Nutritional Science</i> , 2013 , 2, e33	2.7	14
49	Genome-wide association studies identify 137 genetic loci for DNA methylation biomarkers of aging. <i>Genome Biology</i> , 2021 , 22, 194	18.3	14
48	Protein signatures of centenarians and their offspring suggest centenarians age slower than other humans. <i>Aging Cell</i> , 2021 , 20, e13290	9.9	14
47	Discovery and fine-mapping of loci associated with MUFAs through trans-ethnic meta-analysis in Chinese and European populations. <i>Journal of Lipid Research</i> , 2017 , 58, 974-981	6.3	12
46	A double blind placebo controlled randomized trial of the effect of acute uric acid changes on inflammatory markers in humans: A pilot study. <i>PLoS ONE</i> , 2017 , 12, e0181100	3.7	12
45	SPARCL1 Accelerates Symptom Onset in Alzheimer's Disease and Influences Brain Structure and Function During Aging. <i>Journal of Alzheimers Disease</i> , 2018 , 61, 401-414	4.3	12
44	Potential Interplay between Dietary Saturated Fats and Genetic Variants of the NLRP3 Inflammasome to Modulate Insulin Resistance and Diabetes Risk: Insights from a Meta-Analysis of 19,005 Individuals. <i>Molecular Nutrition and Food Research</i> , 2019 , 63, e1900226	5.9	11
43	Sex-Dependent Associations of Serum Uric Acid with Brain Function During Aging. <i>Journal of Alzheimers Disease</i> , 2017 , 60, 699-706	4.3	11

42	Associations between Common Variants in Iron-Related Genes with Haematological Traits in Populations of African Ancestry. <i>PLoS ONE</i> , 2016 , 11, e0157996	3.7	11
41	Blood Metabolite Signatures of Metabolic Syndrome in Two Cross-Cultural Older Adult Cohorts. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	10
40	PRKCZ methylation is associated with sunlight exposure in a North American but not a Mediterranean population. <i>Chronobiology International</i> , 2014 , 31, 1034-40	3.6	10
39	Proteomics in aging research: A roadmap to clinical, translational research. <i>Aging Cell</i> , 2021 , 20, e13325	9.9	10
38	Characterization of the plasma proteomic profile of frailty phenotype. <i>GeroScience</i> , 2021 , 43, 1029-1037	8.9	10
37	Genome-wide meta-analysis of muscle weakness identifies 15 susceptibility loci in older men and women. <i>Nature Communications</i> , 2021 , 12, 654	17.4	10
36	Whole-genome sequencing to understand the genetic architecture of common gene expression and biomarker phenotypes. <i>Human Molecular Genetics</i> , 2015 , 24, 1504-12	5.6	7
35	Blood Metabolite Signature of Metabolic Syndrome Implicates Alterations in Amino Acid Metabolism: Findings from the Baltimore Longitudinal Study of Aging (BLSA) and the Tsuruoka Metabolomics Cohort Study (TMCS). <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	7
34	Age-associated difference in circulating ACE2, the gateway for SARS-COV-2, in humans: results from the InCHIANTI study. <i>GeroScience</i> , 2021 , 43, 619-627	8.9	7
33	Plasma proteomic signatures predict dementia and cognitive impairment. <i>Alzheimers and Dementia: Translational Research and Clinical Interventions</i> , 2020 , 6, e12018	6	6
32	Blood DNA Methylation and Aging: A Cross-Sectional Analysis and Longitudinal Validation in the InCHIANTI Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2020 , 75, 2051-2055	6.4	6
31	Cross-Sectional and Longitudinal Effects of Genotypes on Individual Differences in Memory and Executive Function: Findings from the BLSA. <i>Frontiers in Aging Neuroscience</i> , 2017 , 9, 142	5.3	6
30	Genetic diversity is a predictor of mortality in humans. <i>BMC Genetics</i> , 2014 , 15, 159	2.6	6
29	Blood DNA methylation sites predict death risk in a longitudinal study of 12, 300 individuals. <i>Aging</i> , 2020 , 12, 14092-14124	5.6	6
28	Association Between the Multidimensional Prognostic Index and Mortality During 15 Years of Follow-up in the InCHIANTI Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2021 , 76, 1678-1685	6.4	6
27	Genome-wide association study of circulating interleukin 6 levels identifies novel loci. <i>Human Molecular Genetics</i> , 2021 , 30, 393-409	5.6	6
26	Genome-wide association study identifies novel susceptibility loci for KIT D816V positive mastocytosis. <i>American Journal of Human Genetics</i> , 2021 , 108, 284-294	11	6
25	Genome-Wide Interactions with Dairy Intake for Body Mass Index in Adults of European Descent. <i>Molecular Nutrition and Food Research</i> , 2018 , 62, 1700347	5.9	5

24	Genome-wide Association Study of Parental Life Span. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2017 , 72, 1407-1410	6.4	5
23	No evidence for genome-wide interactions on plasma fibrinogen by smoking, alcohol consumption and body mass index: results from meta-analyses of 80,607 subjects. <i>PLoS ONE</i> , 2014 , 9, e111156	3.7	5
22	Genetic analysis of dietary intake identifies new loci and functional links with metabolic traits. <i>Nature Human Behaviour</i> , 2021 ,	12.8	5
21	Association of Adherence to the Mediterranean-Style Diet with Lower Frailty Index in Older Adults. <i>Nutrients</i> , 2021 , 13,	6.7	4
20	Comparing Analytical Methods for the Gut Microbiome and Aging: Gut Microbial Communities and Body Weight in the Osteoporotic Fractures in Men (MrOS) Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2020 , 75, 1267-1275	6.4	3
19	Genetics of Energy and Macronutrient Intake in Humans. <i>Current Nutrition Reports</i> , 2014 , 3, 170-177	6	3
18	Elevated Plasma Growth and Differentiation Factor 15 Predicts Incident Anemia in Older Adults Aged 60 Years and Older. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2021 , 76, 1192-1197	6.4	3
17	Multi-trait genome-wide association meta-analysis of dietary intake identifies new loci and genetic and functional links with metabolic traits		3
16	A Plasma Proteomic Signature of Skeletal Muscle Mitochondrial Function. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	3
15	Cardiovascular Health Is Associated With Disability Among Older Community Dwelling Men and Women. <i>Journal of Aging and Health</i> , 2019 , 31, 1339-1352	2.6	2
14	Dietary Pattern Trajectories in Middle Age and Physical Function in Older Age. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2021 , 76, 513-519	6.4	2
13	A brain proteomic signature of incipient Alzheimer's disease in young ϵ carriers identifies novel drug targets. <i>Science Advances</i> , 2021 , 7, eabi8178	14.3	2
12	Adherence to the Mediterranean diet assessed by a novel dietary biomarker score and mortality in older adults: the InCHIANTI cohort study. <i>BMC Medicine</i> , 2021 , 19, 280	11.4	2
11	DNA methylation signatures reveal that distinct combinations of transcription factors specify human immune cell epigenetic identity. <i>Immunity</i> , 2021 , 54, 2465-2480.e5	32.3	2
10	Proteomics and Epidemiological Models of Human Aging. <i>Frontiers in Physiology</i> , 2021 , 12, 674013	4.6	2
9	The Mediterranean-DASH Intervention for Neurodegenerative Delay (MIND) diet is associated with physical function and grip strength in older men and women. <i>American Journal of Clinical Nutrition</i> , 2021 ,	7	1
8	DNA methylation signature of chronic low-grade inflammation and its role in cardio-respiratory diseases.. <i>Nature Communications</i> , 2022 , 13, 2408	17.4	1
7	Predicting physiological aging rates from a range of quantitative traits using machine learning. <i>Aging</i> , 2021 , 13, 23471-23516	5.6	0

6	Metabolomic Profile of Different Dietary Patterns and Their Association with Frailty Index in Community-Dwelling Older Men and Women. <i>Nutrients</i> , 2022 , 14, 2237	6.7	o
5	Interaction between Apolipoprotein E and Butyrylcholinesterase Genes on Risk of Alzheimer's Disease in a Prospective Cohort Study. <i>Journal of Alzheimer's Disease</i> , 2020 , 75, 417-427	4.3	
4	Contribution of Diet and Genes to Polyunsaturated Fatty Acid Composition. <i>Current Cardiovascular Risk Reports</i> , 2011 , 5, 45-51	0.9	
3	Caffeine, alcohol and overall nutrient adequacy are associated with longitudinal cognitive performance among US adults. <i>FASEB Journal</i> , 2013 , 27, 346.4	0.9	
2	Prior psychosocial profile and perceived impact of the COVID-19 pandemic: insights from the Baltimore Longitudinal Study of Aging.. <i>Aging Clinical and Experimental Research</i> , 2022 , 1	4.8	
1	Understanding the Human Aging Proteome Using Epidemiological Models. <i>Methods in Molecular Biology</i> , 2022 , 173-192	1.4	