Robert F Hillary

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

| 37 papers | 493 | 11 | 22 |
|-------------|--------------------|---------|-----------|
| | citations | h-index | g-index |
| 46 | 892 ext. citations | 10 | 3.95 |
| ext. papers | | avg, IF | L-index |

| # | Paper | IF | Citations |
|----|--|------|-----------|
| 37 | Blood-based epigenome-wide analyses of cognitive abilities <i>Genome Biology</i> , 2022 , 23, 26 | 18.3 | 1 |
| 36 | Epigenetic scores for the circulating proteome as tools for disease prediction ELife, 2022, 11, | 8.9 | 2 |
| 35 | Genome- and epigenome-wide studies of plasma protein biomarkers for Alzheimer's disease implicate TBCA and TREM2 in disease risk <i>Alzheimers</i> and Dementia: Diagnosis, Assessment and Disease Monitoring, 2022 , 14, e12280 | 5.2 | |
| 34 | The genetic and epigenetic profile of serum \$\frac{1}{2}100 In the Lothian Birth Cohort 1936 and its relationship to Alzheimer's disease Wellcome Open Research, 2021 , 6, 306 | 4.8 | |
| 33 | Birth weight associations with DNA methylation differences in an adult population. <i>Epigenetics</i> , 2021 , 16, 783-796 | 5.7 | 7 |
| 32 | Meta-analysis of genome-wide DNA methylation identifies shared associations across neurodegenerative disorders. <i>Genome Biology</i> , 2021 , 22, 90 | 18.3 | 6 |
| 31 | Epigenetic predictors of lifestyle traits applied to the blood and brain. <i>Brain Communications</i> , 2021 , 3, fcab082 | 4.5 | 1 |
| 30 | An epigenetic predictor of death captures multi-modal measures of brain health. <i>Molecular Psychiatry</i> , 2021 , 26, 3806-3816 | 15.1 | 31 |
| 29 | Structural brain correlates of serum and epigenetic markers of inflammation in major depressive disorder. <i>Brain, Behavior, and Immunity</i> , 2021 , 92, 39-48 | 16.6 | 14 |
| 28 | Creating and Validating a DNA Methylation-Based Proxy for Interleukin-6. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2021 , 76, 2284-2292 | 6.4 | 1 |
| 27 | The neuropathology of autism: A systematic review of post-mortem studies of autism and related disorders. <i>Neuroscience and Biobehavioral Reviews</i> , 2021 , 129, 35-62 | 9 | 5 |
| 26 | Identification of plasma proteins relating to brain neurodegeneration and vascular pathology in cognitively normal individuals. <i>Alzheimers</i> and <i>Dementia: Diagnosis, Assessment and Disease Monitoring</i> , 2021 , 13, e12240 | 5.2 | 0 |
| 25 | Multi-method genome- and epigenome-wide studies of inflammatory protein levels in healthy older adults. <i>Genome Medicine</i> , 2020 , 12, 60 | 14.4 | 9 |
| 24 | DNA methylation outlier burden, health, and ageing in Generation Scotland and the Lothian Birth Cohorts of 1921 and 1936. <i>Clinical Epigenetics</i> , 2020 , 12, 49 | 7.7 | 8 |
| 23 | MethylDetectR: a software for methylation-based health profiling. <i>Wellcome Open Research</i> , 2020 , 5, 283 | 4.8 | 2 |
| 22 | Characterisation of an inflammation-related epigenetic score and its association with cognitive ability. <i>Clinical Epigenetics</i> , 2020 , 12, 113 | 7.7 | 15 |
| 21 | Epigenetic measures of ageing predict the prevalence and incidence of leading causes of death and disease burden. <i>Clinical Epigenetics</i> , 2020 , 12, 115 | 7.7 | 40 |

| 20 | MethylDetectR: a software for methylation-based health profiling. <i>Wellcome Open Research</i> , 2020 , 5, 283 | 4.8 | 1 |
|-------------------|---|------|-------|
| 19 | Age-related clonal haemopoiesis is associated with increased epigenetic age. <i>Current Biology</i> , 2019 , 29, R786-R787 | 6.3 | 20 |
| 18 | Genome and epigenome wide studies of neurological protein biomarkers in the Lothian Birth Cohort 1936. <i>Nature Communications</i> , 2019 , 10, 3160 | 17.4 | 21 |
| 17 | An epigenome-wide association study of sex-specific chronological ageing. <i>Genome Medicine</i> , 2019 , 12, 1 | 14.4 | 43 |
| 16 | A meta-analysis of genome-wide association studies of epigenetic age acceleration. <i>PLoS Genetics</i> , 2019 , 15, e1008104 | 6 | 38 |
| 15 | Childhood intelligence attenuates the association between biological ageing and health outcomes in later life. <i>Translational Psychiatry</i> , 2019 , 9, 323 | 8.6 | 8 |
| 14 | A lifetime of stress: ATF6 in development and homeostasis. <i>Journal of Biomedical Science</i> , 2018 , 25, 48 | 13.3 | 77 |
| 13 | Stem Cells to Inform the Neurobiology of Mental Illness. <i>Current Topics in Behavioral Neurosciences</i> , 2018 , 40, 13-43 | 3.4 | 4 |
| 12 | Epigenetic signatures of starting and stopping smoking. <i>EBioMedicine</i> , 2018 , 37, 214-220 | 8.8 | 36 |
| | | | |
| 11 | Epigenetic prediction of complex traits and death. <i>Genome Biology</i> , 2018 , 19, 136 | 18.3 | 77 |
| 10 | Epigenetic prediction of complex traits and death. <i>Genome Biology</i> , 2018 , 19, 136 The genetic and epigenetic profile of serum 15100 In the Lothian Birth Cohort 1936 and its relationship to Alzheimer disease. <i>Wellcome Open Research</i> , 6, 306 | 18.3 | 77 |
| | The genetic and epigenetic profile of serum \$1000n the Lothian Birth Cohort 1936 and its | | 77 |
| 10 | The genetic and epigenetic profile of serum[\$100[in the Lothian Birth Cohort 1936 and its relationship to Alzheimer disease. Wellcome Open Research, 6, 306 Childhood intelligence attenuates the association between biological ageing and health outcomes | | |
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| 10 9 8 7 6 | The genetic and epigenetic profile of serum[\$100[in the Lothian Birth Cohort 1936 and its relationship to Alzheimer® disease. Wellcome Open Research, 6, 306 Childhood intelligence attenuates the association between biological ageing and health outcomes in later Creating and validating a DNA methylation-based proxy for Interleukin-6 A comparison of blood and brain-derived ageing and inflammation-related DNA methylation signatures and their association with microglial burdens Characterisation of an inflammation-related epigenetic score and its association with cognitive ability | | 2 2 1 |

Birth weight associations with psychiatric and physical health, cognitive function, and DNA methylation differences in an adult population

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Blood-based epigenome-wide analyses of cognitive abilities

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