Federico De Masi

List of Publications by Year in descending order

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#	Article	lF	CITATIONS
1	High-resolution DNA-binding specificity analysis of yeast transcription factors. Genome Research, 2009, 19, 556-566.	5.5	365
2	Genomic and drug target evaluation of 90 cardiovascular proteins in 30,931 individuals. Nature Metabolism, 2020, 2, 1135-1148.	11.9	327
3	Systematic Discovery of New Recognition Peptides Mediating Protein Interaction Networks. PLoS Biology, 2005, 3, e405.	5.6	310
4	A Multiparameter Network Reveals Extensive Divergence between C. elegans bHLH Transcription Factors. Cell, 2009, 138, 314-327.	28.9	242
5	ATAF1 transcription factor directly regulates abscisic acid biosynthetic gene <i>NCED3</i> in <i>Arabidopsis thaliana</i> . FEBS Open Bio, 2013, 3, 321-327.	2.3	182
6	Ibuprofen alters human testicular physiology to produce a state of compensated hypogonadism. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E715-E724.	7.1	88
7	A DNA-binding-site landscape and regulatory network analysis for NAC transcription factors in <i>Arabidopsis thaliana</i> . Nucleic Acids Research, 2014, 42, 7681-7693.	14.5	84
8	Using a structural and logics systems approach to infer bHLH–DNA binding specificity determinants. Nucleic Acids Research, 2011, 39, 4553-4563.	14.5	73
9	High throughput production of mouse monoclonal antibodies using antigen microarrays. Proteomics, 2005, 5, 4070-4081.	2.2	71
10	High-throughput sequencing enhanced phage display enables the identification of patient-specific epitope motifs in serum. Scientific Reports, 2015, 5, 12913.	3.3	62
11	miRandola 2017: a curated knowledge base of non-invasive biomarkers. Nucleic Acids Research, 2018, 46, D354-D359.	14.5	61
12	Using protein design algorithms to understand the molecular basis of disease caused by protein–DNA interactions: the Pax6 example. Nucleic Acids Research, 2010, 38, 7422-7431.	14.5	55
13	Predicting and elucidating the etiology of fatty liver disease: A machine learning modeling and validation study in the IMI DIRECT cohorts. PLoS Medicine, 2020, 17, e1003149.	8.4	47
14	Four groups of type 2 diabetes contribute to the etiological and clinical heterogeneity in newly diagnosed individuals: An IMI DIRECT study. Cell Reports Medicine, 2022, 3, 100477.	6.5	39
15	Control of lysosomal biogenesis and Notch-dependent tissue patterning by components of the TFEB-V-ATPase axis in <i>Drosophila melanogaster</i> . Autophagy, 2016, 12, 499-514.	9.1	34
16	High-throughput immuno-profiling of mamba (Dendroaspis) venom toxin epitopes using high-density peptide microarrays. Scientific Reports, 2016, 6, 36629.	3.3	33
17	Genetic studies of abdominal MRI data identify genes regulating hepcidin as major determinants of liver iron concentration. Journal of Hepatology, 2019, 71, 594-602.	3.7	23
18	Discovery of biomarkers for glycaemic deterioration before and after the onset of type 2 diabetes: descriptive characteristics of the epidemiological studies within the IMI DIRECT Consortium. Diabetologia, 2019, 62, 1601-1615.	6.3	22

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19	Profiles of Glucose Metabolism in Different Prediabetes Phenotypes, Classified by Fasting Glycemia, 2-Hour OGTT, Glycated Hemoglobin, and 1-Hour OGTT: An IMI DIRECT Study. Diabetes, 2021, 70, 2092-2106.	0.6	17
20	Cross-recognition of a pit viper (Crotalinae) polyspecific antivenom explored through high-density peptide microarray epitope mapping. PLoS Neglected Tropical Diseases, 2017, 11, e0005768.	3.0	17
21	Processes Underlying Glycemic Deterioration in Type 2 Diabetes: An IMI DIRECT Study. Diabetes Care, 2021, 44, 511-518.	8.6	16
22	The role of physical activity in metabolic homeostasis before and after the onset of type 2 diabetes: an IMI DIRECT study. Diabetologia, 2020, 63, 744-756.	6.3	12
23	Whole blood co-expression modules associate with metabolic traits and type 2 diabetes: an IMI-DIRECT study. Genome Medicine, 2020, 12, 109.	8.2	8
24	The governance structure for data access in the DIRECT consortium: an innovative medicines initiative (IMI) project. Life Sciences, Society and Policy, 2018, 14, 20.	3.2	7
25	Post-load glucose subgroups and associated metabolic traits in individuals with type 2 diabetes: An IMI-DIRECT study. PLoS ONE, 2020, 15, e0242360.	2.5	7
26	Genome-Wide Association Analysis of Pancreatic Beta-Cell Glucose Sensitivity. Journal of Clinical Endocrinology and Metabolism, 2021, 106, 80-90.	3.6	5
27	Dietary metabolite profiling brings new insight into the relationship between nutrition and metabolic risk: An IMI DIRECT study. EBioMedicine, 2020, 58, 102932.	6.1	3
28	The Interplay of Non-coding RNAs and X Chromosome Inactivation in Human Disease. RNA Technologies, 2018, , 229-238.	0.3	0
29	Title is missing!. , 2020, 17, e1003149.		0
30	Title is missing!. , 2020, 17, e1003149.		0
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