

# Philippe Fort

## List of Publications by Year in descending order

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109  
papers

8,876  
citations

57758

44  
h-index

42399

92  
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114  
all docs

114  
docs citations

114  
times ranked

8315  
citing authors

#	ARTICLE	IF	CITATIONS
1	Aspartateâ€phobia of thermophiles as a reaction to deleterious chemical transformations. <i>BioEssays</i> , 2022, 44, e2100213.	2.5	2
2	Regulation of Src tumor activity by its N-terminal intrinsically disordered region. <i>Oncogene</i> , 2022, 41, 960-970.	5.9	8
3	NOPCHAP1 is a PAQosome cofactor that helps loading NOP58 on RUVBL1/2 during box C/D snoRNP biogenesis. <i>Nucleic Acids Research</i> , 2021, 49, 1094-1113.	14.5	14
4	SHED-Dependent Oncogenic Signaling of the PEAK3 Pseudo-Kinase. <i>Cancers</i> , 2021, 13, 6344.	3.7	6
5	The atypical RhoU/Wrch1 Rho GTPase controls cell proliferation and apoptosis in the gut epithelium. <i>Biology of the Cell</i> , 2019, 111, 121-141.	2.0	11
6	New insights into the evolutionary conservation of the sole PIKK pseudokinase Tra1/TRRAP. <i>Biochemical Society Transactions</i> , 2019, 47, 1597-1608.	3.4	25
7	The RPAP3-Cterminal domain identifies R2TP-like quaternary chaperones. <i>Nature Communications</i> , 2018, 9, 2093.	12.8	59
8	PIP30/FAM192A is a novel regulator of the nuclear proteasome activator PA28Î³. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E6477-E6486.	7.1	29
9	SOX9 has distinct regulatory roles in alternative splicing and transcription. <i>Nucleic Acids Research</i> , 2018, 46, 9106-9118.	14.5	30
10	Rho signaling: An historical and evolutionary perspective. , 2018, , 3-18.		2
11	Binding site density enables paralog-specific activity of SLM2 and Sam68 proteins in <i>Neurexin2</i> AS4 splicing control. <i>Nucleic Acids Research</i> , 2017, 45, gkw1277.	14.5	16
12	The Evolutionary Landscape of Dbl-Like RhoGEF Families: Adapting Eukaryotic Cells to Environmental Signals. <i>Genome Biology and Evolution</i> , 2017, 9, 1471-1486.	2.5	47
13	A SLM2 Feedback Pathway Controls Cortical Network Activity and Mouse Behavior. <i>Cell Reports</i> , 2016, 17, 3269-3280.	6.4	21
14	STARs in the CNS. <i>Biochemical Society Transactions</i> , 2016, 44, 1066-1072.	3.4	10
15	High chlorpyrifos resistance in <i>Culex pipiens</i> mosquitoes: strong synergy between resistance genes. <i>Heredity</i> , 2016, 116, 224-231.	2.6	12
16	Neural Differentiation Modulates the Vertebrate Brain Specific Splicing Program. <i>PLoS ONE</i> , 2015, 10, e0125998.	2.5	10
17	Atypical RhoV and RhoU GTPases control development of the neural crest. <i>Small GTPases</i> , 2015, 6, 174-177.	1.6	22
18	Evolution of Proteasome Regulators in Eukaryotes. <i>Genome Biology and Evolution</i> , 2015, 7, 1363-1379.	2.5	77

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19	Stable coexistence of incompatible <i>Wolbachia</i> along a narrow contact zone in mosquito field populations. <i>Molecular Ecology</i> , 2015, 24, 508-521.	3.9	25
20	Pleiotropic. Small GTPases, 2014, 5, e27975.	1.6	14
21	Wolbachia Divergence and the Evolution of Cytoplasmic Incompatibility in <i>Culex pipiens</i> . <i>PLoS ONE</i> , 2014, 9, e87336.	2.5	48
22	Antagonistic functions of LMNA isoforms in energy expenditure and lifespan. <i>EMBO Reports</i> , 2014, 15, 529-539.	4.5	47
23	MBNL1 and RBFOX2 cooperate to establish a splicing programme involved in pluripotent stem cell differentiation. <i>Nature Communications</i> , 2013, 4, 2480.	12.8	120
24	Applying ecological and evolutionary theory to cancer: a long and winding road. <i>Evolutionary Applications</i> , 2013, 6, 1-10.	3.1	70
25	Targeting the Dbl and Dock-Family RhoGEFs. <i>The Enzymes</i> , 2013, 33 Pt A, 169-191.	1.7	3
26	The Tissue-Specific RNA Binding Protein T-STAR Controls Regional Splicing Patterns of Neurexin Pre-mRNAs in the Brain. <i>PLoS Genetics</i> , 2013, 9, e1003474.	3.5	74
27	Tissue-Specific Alternative Splicing of Tak1 Is Conserved in Deuterostomes. <i>Molecular Biology and Evolution</i> , 2012, 29, 261-269.	8.9	21
28	Fossil Rhabdoviral Sequences Integrated into Arthropod Genomes: Ontogeny, Evolution, and Potential Functionality. <i>Molecular Biology and Evolution</i> , 2012, 29, 381-390.	8.9	100
29	Using a Modified Yeast Two-Hybrid System to Screen for Chemical GEF Inhibitors. <i>Methods in Molecular Biology</i> , 2012, 928, 81-95.	0.9	6
30	Novel AChE Inhibitors for Sustainable Insecticide Resistance Management. <i>PLoS ONE</i> , 2012, 7, e47125.	2.5	26
31	Activity of the RhoU/Wrch1 GTPase is critical for cranial neural crest cell migration. <i>Developmental Biology</i> , 2011, 350, 451-463.	2.0	33
32	Multiple Wolbachia determinants control the evolution of cytoplasmic incompatibilities in <i>Culex pipiens</i> mosquito populations. <i>Molecular Ecology</i> , 2011, 20, 286-298.	3.9	46
33	MiniSOX9, a dominant-negative variant in colon cancer cells. <i>Oncogene</i> , 2011, 30, 2493-2503.	5.9	35
34	Tara up-regulates E-cadherin transcription by binding to the Trio RhoGEF and inhibiting Rac signaling. <i>Journal of Cell Biology</i> , 2011, 193, 319-332.	5.2	63
35	Atypical RhoV and RhoU GTPases control development of the neural crest. <i>Small GTPases</i> , 2011, 2, 310-313.	1.6	14
36	TC10 controls human myofibril organization and is activated by the sarcomeric RhoGEF obscurin. <i>Journal of Cell Science</i> , 2009, 122, 947-956.	2.0	23

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37	A Cell Active Chemical GEF Inhibitor Selectively Targets the Trio/RhoG/Rac1 Signaling Pathway. <i>Chemistry and Biology</i> , 2009, 16, 657-666.	6.0	91
38	Dynamic expression patterns of <i>RhoV</i> and <i>Chp</i> and <i>RhoU</i> / <i>Wrch</i> during chicken embryonic development. <i>Developmental Dynamics</i> , 2008, 237, 1165-1171.	1.8	15
39	Trio Controls the Mature Organization of Neuronal Clusters in the Hindbrain. <i>Journal of Neuroscience</i> , 2007, 27, 10323-10332.	3.6	43
40	Evolution of the Rho Family of Ras-Like GTPases in Eukaryotes. <i>Molecular Biology and Evolution</i> , 2007, 24, 203-216.	8.9	366
41	Variability and Expression of Ankyrin Domain Genes in <i>Wolbachia</i> Variants Infecting the Mosquito <i>Culex pipiens</i> . <i>Journal of Bacteriology</i> , 2007, 189, 4442-4448.	2.2	54
42	The small GTPase <i>RhoV</i> is an essential regulator of neural crest induction in <i>Xenopus</i> . <i>Developmental Biology</i> , 2007, 310, 113-128.	2.0	46
43	Influence of aging on cytoplasmic incompatibility, sperm modification and <i>Wolbachia</i> density in <i>Culex pipiens</i> mosquitoes. <i>Heredity</i> , 2007, 98, 368-374.	2.6	49
44	Expression of <i>RhoB</i> in the developing <i>Xenopus laevis</i> embryo. <i>Gene Expression Patterns</i> , 2007, 7, 282-288.	0.8	13
45	Fertilization regulates apoptosis of <i>Ciona intestinalis</i> extra-embryonic cells through thyroxine (T4)-dependent NF- $\kappa$ B pathway activation during early embryonic development. <i>Developmental Biology</i> , 2006, 289, 152-165.	2.0	17
46	Expression Profile of RhoGTPases and RhoGEFs During RANKL-Stimulated Osteoclastogenesis: Identification of Essential Genes in Osteoclasts. <i>Journal of Bone and Mineral Research</i> , 2006, 21, 1387-1398.	2.8	83
47	Identification of TRIO-GEFD1 chemical inhibitors using the yeast exchange assay. <i>Biology of the Cell</i> , 2006, 98, 511-522.	2.0	41
48	Identification of Rho GTPases implicated in terminal differentiation of muscle cells in ascidia. <i>Biology of the Cell</i> , 2006, 98, 577-588.	2.0	8
49	Hypervariable prophage WO sequences describe an unexpected high number of <i>Wolbachia</i> variants in the mosquito <i>Culex pipiens</i> . <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2006, 273, 495-502.	2.6	49
50	Transposable element polymorphism of <i>Wolbachia</i> in the mosquito <i>Culex pipiens</i> : evidence of genetic diversity, superinfection and recombination. <i>Molecular Ecology</i> , 2005, 14, 1561-1573.	3.9	72
51	The caspase family in urochordates: distinct evolutionary fates in ascidians and larvaceans. <i>Biology of the Cell</i> , 2005, 97, 857-866.	2.0	26
52	Distinct roles of Rac1/Cdc42 and Rho/Rock for axon outgrowth and nucleokinesis of precerebellar neurons toward netrin 1. <i>Development (Cambridge)</i> , 2004, 131, 2841-2852.	2.5	83
53	Insecticide resistance: a silent base prediction. <i>Current Biology</i> , 2004, 14, R552-R553.	3.9	76
54	Ascidians as a vertebrate-like model organism for physiological studies of Rho GTPase signaling. <i>Biology of the Cell</i> , 2003, 95, 295-302.	2.0	22

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55	RhoG regulates gene expression and the actin cytoskeleton in lymphocytes. <i>Oncogene</i> , 2003, 22, 330-342.	5.9	46
56	Insecticide resistance in mosquito vectors. <i>Nature</i> , 2003, 423, 136-137.	27.8	546
57	The GTP/GDP Cycling of Rho GTPase TCL Is an Essential Regulator of the Early Endocytic Pathway. <i>Molecular Biology of the Cell</i> , 2003, 14, 4846-4856.	2.1	61
58	A Dual Role of the GTPase Rac in Cardiac Differentiation of Stem Cells. <i>Molecular Biology of the Cell</i> , 2003, 14, 2781-2792.	2.1	58
59	A novel acetylcholinesterase gene in mosquitoes codes for the insecticide target and is non-homologous to theacegeneDrosophila. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2002, 269, 2007-2016.	2.6	233
60	The Human Rho-GEF Trio and Its Target GTPase RhoG Are Involved in the NGF Pathway, Leading to Neurite Outgrowth. <i>Current Biology</i> , 2002, 12, 307-312.	3.9	147
61	Participation of small GTPases Rac1 and Cdc42Hs in myoblast transformation. <i>Oncogene</i> , 2002, 21, 2901-2907.	5.9	31
62	Activation of ERK, Controlled by Rac1 and Cdc42 via Akt, Is Required for Anoikis. <i>Annals of the New York Academy of Sciences</i> , 2002, 973, 145-148.	3.8	32
63	Tail regression in <i>Ciona intestinalis</i> (Prochordate) involves a Caspase-dependent apoptosis event associated with ERK activation. <i>Development (Cambridge)</i> , 2002, 129, 3105-3114.	2.5	109
64	Tail regression in <i>Ciona intestinalis</i> (Prochordate) involves a Caspase-dependent apoptosis event associated with ERK activation. <i>Development (Cambridge)</i> , 2002, 129, 3105-14.	2.5	35
65	The gene for a new brain specific RhoA exchange factor maps to the highly unstable chromosomal region 1p36.2-1p36.3.. <i>Oncogene</i> , 2001, 20, 7307-7317.	5.9	60
66	Kinectin Is a Key Effector of RhoG Microtubule-Dependent Cellular Activity. <i>Molecular and Cellular Biology</i> , 2001, 21, 8022-8034.	2.3	73
67	Raf-MEK-Erk Cascade in Anoikis Is Controlled by Rac1 and Cdc42 via Akt. <i>Molecular and Cellular Biology</i> , 2001, 21, 6706-6717.	2.3	108
68	Extinction of Rac1 and Cdc42Hs signalling defines a novel p53-dependent apoptotic pathway. <i>Oncogene</i> , 2000, 19, 2377-2385.	5.9	34
69	Critical Activities of Rac1 and Cdc42Hs in Skeletal Myogenesis: Antagonistic Effects of JNK and p38 Pathways. <i>Molecular Biology of the Cell</i> , 2000, 11, 2513-2528.	2.1	101
70	Characterization of TCL, a New GTPase of the Rho Family related to TC10 and Cdc42. <i>Journal of Biological Chemistry</i> , 2000, 275, 36457-36464.	3.4	110
71	Cdc42Hs and Rac1 GTPases Induce the Collapse of the Vimentin Intermediate Filament Network. <i>Journal of Biological Chemistry</i> , 2000, 275, 33046-33052.	3.4	57
72	A fluorescent reporter gene as a marker for ventricular specification in ES-derived cardiac cells. <i>FEBS Letters</i> , 2000, 478, 151-158.	2.8	106

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73	The yeast exchange assay, a new complementary method to screen for Dbp-like protein specificity: identification of a novel RhoA exchange factor. <i>FEBS Letters</i> , 2000, 480, 287-292.	2.8	19
74	Signalling pathways controlled by the GTPase RhoG. <i>Biology of the Cell</i> , 1999, 91, 551-552.	2.0	0
75	Small GTPases of the Rho Family and Cell Transformation. <i>Progress in Molecular and Subcellular Biology</i> , 1999, 22, 159-181.	1.6	16
76	A Presumptive Developmental Role for a Sea Urchin Cyclin B Splice Variant. <i>Journal of Cell Biology</i> , 1998, 140, 283-293.	5.2	30
77	RhoG GTPase Controls a Pathway That Independently Activates Rac1 and Cdc42Hs. <i>Molecular Biology of the Cell</i> , 1998, 9, 1379-1394.	2.1	152
78	A Simple Luciferase Assay for Signal Transduction Activity Detection of Epidermal Growth Factor Displayed on Phage. <i>Nucleic Acids Research</i> , 1997, 25, 1585-1590.	14.5	31
79	Expression and Human Chromosomal Localization to 17q25 of the Growth-Regulated Gene Encoding the Mitochondrial Ribosomal Protein MRPL12. <i>Genomics</i> , 1997, 41, 453-457.	2.9	17
80	Structure of the Human ARHG Locus Encoding the Rho/Rac-like RhoG GTPase. <i>Genomics</i> , 1997, 42, 157-160.	2.9	22
81	Structure and Chromosomal Assignment to 22q12 and 17qter of the ras-Related Rac2 and Rac3 Human Genes. <i>Genomics</i> , 1997, 44, 242-246.	2.9	23
82	The small GTPases Cdc42Hs, Rac1 and RhoG delineate Raf-independent pathways that cooperate to transform NIH3T3 cells. <i>Current Biology</i> , 1997, 7, 629-637.	3.9	100
83	A Delayed-early Response Nuclear Gene Encoding MRPL12, the Mitochondrial Homologue to the Bacterial Translational Regulator L7/L12 Protein. <i>Journal of Biological Chemistry</i> , 1996, 271, 11468-11476.	3.4	30
84	[18] Serum induction of RhoG expression. <i>Methods in Enzymology</i> , 1995, 256, 151-162.	1.0	0
85	Growth-Regulated Expression of FKBP-59 Immunophilin in Normal and Transformed Fibroblastic Cells. <i>Experimental Cell Research</i> , 1995, 220, 152-160.	2.6	10
86	Concerted evolution in the GAPDH family of retrotransposed pseudogenes. <i>Mammalian Genome</i> , 1993, 4, 695-703.	2.2	32
87	Localization of ARHG, a Member of the RAS Homolog Gene Family, to 11p15.5-11p15.4 by Fluorescence in Situ Hybridization. <i>Genomics</i> , 1993, 16, 788-790.	2.9	4
88	S26 ribosomal protein RNA: an invariant control for gene regulation experiments in eucaryotic cells and tissues. <i>Nucleic Acids Research</i> , 1993, 21, 1498-1498.	14.5	184
89	Transduction du signal mitogénique, cytosquelette et petites protéines G : vers un réseau de protéines GAP ? <i>Medecine/Sciences</i> , 1993, 9, 59.	0.2	0
90	Worldwide migration of amplified insecticide resistance genes in mosquitoes. <i>Nature</i> , 1991, 350, 151-153.	27.8	283

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91	Nucleotide sequence of hamster glyceraldehyde-3-phosphate dehydrogenase mRNA. <i>Nucleic Acids Research</i> , 1990, 18, 3054-3054.	14.5	15
92	Versatile vectors for pulsed expression in eukaryotic cells. <i>Nucleic Acids Research</i> , 1989, 17, 2874-2874.	14.5	7
93	Requirements for c-fos mRNA down regulation in growth stimulated murine cells. <i>Oncogene</i> , 1989, 4, 881-8.	5.9	28
94	Cloning and regulation of a mRNA specifically expressed in the preadipose state. <i>Journal of Biological Chemistry</i> , 1989, 264, 10119-25.	3.4	52
95	Role of RNA structures in c-myc and c-fos gene regulations. <i>Gene</i> , 1988, 72, 287-295.	2.2	14
96	The regulatory strategies of c-myc and c-fos proto-oncogenes share some common mechanisms. <i>Biochimie</i> , 1988, 70, 877-884.	2.6	18
97	Complete Sequence of Cytochrome P450 3c cDNA and Presence of Two mRNA Species with 3' Untranslated Regions of Different Lengths. <i>DNA and Cell Biology</i> , 1988, 7, 39-46.	5.2	34
98	Sequence determinants of c-myc mRNA turn-over: influence of 3' and 5' non-coding regions. <i>Oncogene Research</i> , 1988, 3, 155-66.	1.2	20
99	Regulation of c-fos gene expression in hamster fibroblasts: initiation and elongation of transcription and mRNA degradation. <i>Nucleic Acids Research</i> , 1987, 15, 5657-5667.	14.5	241
100	Sequence of a human immunoglobulin gamma 3 heavy chain constant region gene: comparison with the other human C $\gamma$ genes. <i>Nucleic Acids Research</i> , 1986, 14, 1779-1789.	14.5	153
101	Various rat adult tissues express only one major mRNA species from the glyceraldehyde-3-phosphate-dehydrogenase multigenic family. <i>Nucleic Acids Research</i> , 1985, 13, 1431-1442.	14.5	2,147
102	Nucleotide Sequence and Complementation Analysis of a Polycistronic Sporulation Operon, spoVA, in <i>Bacillus subtilis</i> . <i>Microbiology (United Kingdom)</i> , 1985, 131, 1091-1105.	1.8	78
103	Effects of Transition Mutations in the Regulatory Locus spoIIA on the Incidence of Sporulation in <i>Bacillus subtilis</i> . <i>Microbiology (United Kingdom)</i> , 1985, 131, 959-962.	1.8	8
104	Duplicated sporulation genes in bacteria. <i>FEBS Letters</i> , 1985, 188, 184-188.	2.8	62
105	Post-transcriptional regulation of glyceraldehyde-3-phosphate-dehydrogenase gene expression in rat tissues. <i>Nucleic Acids Research</i> , 1984, 12, 6951-6963.	14.5	486
106	Characterization of the transcription products of glyceraldehyde 3-phosphate-dehydrogenase gene in HeLa cells. <i>FEBS Journal</i> , 1984, 145, 299-304.	0.2	79
107	Complete nucleotide sequence of the messenger RNA coding for chicken muscle glyceraldehyde-3-phosphate dehydrogenase. <i>Biochemical and Biophysical Research Communications</i> , 1984, 118, 767-773.	2.1	111
108	Selection of Seedlings of <i>Thymus Vulgaris</i> by Grazing Slugs. <i>Journal of Ecology</i> , 1983, 71, 299.	4.0	31

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109	A warning on the use of synthetic DNA primers for initiation of reverse transcription on RNA templates: unexpected initiation at a mismatched nucleotide. <i>Gene</i> , 1982, 19, 321-326.	2.2	8