

# Ofer Sarig

## List of Publications by Year in descending order

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75  
papers

1,965  
citations

257357

24  
h-index

276775

41  
g-index

76  
all docs

76  
docs citations

76  
times ranked

2984  
citing authors

#	ARTICLE	IF	CITATIONS
1	Desmoglein 1 deficiency results in severe dermatitis, multiple allergies and metabolic wasting. <i>Nature Genetics</i> , 2013, 45, 1244-1248.	9.4	289
2	Severe dermatitis, multiple allergies, and metabolic wasting syndrome caused by a novel mutation in the N-terminal plakin domain of desmoplakin. <i>Journal of Allergy and Clinical Immunology</i> , 2015, 136, 1268-1276.	1.5	103
3	Variant <i>PADI3</i> in Central Centrifugal Cicatricial Alopecia. <i>New England Journal of Medicine</i> , 2019, 380, 833-841.	13.9	102
4	Stabilizing mutations of KLHL24 ubiquitin ligase cause loss of keratin 14 and human skin fragility. <i>Nature Genetics</i> , 2016, 48, 1508-1516.	9.4	101
5	Population-Specific Association between a Polymorphic Variant in ST18, Encoding a Pro-Apoptotic Molecule, and Pemphigus Vulgaris. <i>Journal of Investigative Dermatology</i> , 2012, 132, 1798-1805.	0.3	98
6	CEDNIK syndrome results from loss-of-function mutations in SNAP29. <i>British Journal of Dermatology</i> , 2011, 164, no-no.	1.4	69
7	A Mutation in LIPN, Encoding Epidermal Lipase N, Causes a Late-Onset Form of Autosomal-Recessive Congenital Ichthyosis. <i>American Journal of Human Genetics</i> , 2011, 88, 482-487.	2.6	62
8	The Genetics of Pemphigus Vulgaris. <i>Frontiers in Medicine</i> , 2018, 5, 226.	1.2	60
9	Loss-of-function mutations in caspase recruitment domain-containing protein 14 (CARD14) are associated with a severe variant of atopic dermatitis. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 143, 173-181.e10.	1.5	60
10	Short Stature, Onychodysplasia, Facial Dysmorphism, and Hypotrichosis Syndrome Is Caused by a POC1A Mutation. <i>American Journal of Human Genetics</i> , 2012, 91, 337-342.	2.6	59
11	Autosomal dominant inheritance of central centrifugal cicatricial alopecia in black South Africans. <i>Journal of the American Academy of Dermatology</i> , 2014, 70, 679-682.e1.	0.6	54
12	Pyoderma gangrenosum, acne and ulcerative colitis in a patient with a novel mutation in the <i>PSTPIP1</i> gene. <i>Clinical and Experimental Dermatology</i> , 2015, 40, 367-372.	0.6	53
13	Identification of a Functional Risk Variant for Pemphigus Vulgaris in the ST18 Gene. <i>PLoS Genetics</i> , 2016, 12, e1006008.	1.5	53
14	A phenotype combining hidradenitis suppurativa with Dowling-Degos disease caused by a founder mutation in <i>PSENEN</i> . <i>British Journal of Dermatology</i> , 2018, 178, 502-508.	1.4	48
15	Olmsted Syndrome Caused by a Homozygous Recessive Mutation in TRPV3. <i>Journal of Investigative Dermatology</i> , 2014, 134, 1752-1754.	0.3	44
16	Treatment of epidermolysis bullosa pruriginosa-associated pruritus with dupilumab. <i>British Journal of Dermatology</i> , 2020, 182, 1495-1497.	1.4	41
17	Infantile mitochondrial hepatopathy is a cardinal feature of MEGDEL syndrome (Methylglutaconic) Tj ETQq1 1 0.784314 rgBT /Over mutations in <i>SERAC1</i> . <i>American Journal of Medical Genetics, Part A</i> , 2013, 161, 2204-2215.	0.7	39
18	Filaggrin 2 Deficiency Results in Abnormal Cell-Cell Adhesion in the Cornified Cell Layers and Causes Peeling Skin Syndrome Type A. <i>Journal of Investigative Dermatology</i> , 2018, 138, 1736-1743.	0.3	37

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19	Non-syndromic autosomal recessive congenital ichthyosis in the Israeli population. <i>Clinical and Experimental Dermatology</i> , 2013, 38, 911-916.	0.6	34
20	Segmental basal cell naevus syndrome caused by an activating mutation in <i>smoothed</i> . <i>British Journal of Dermatology</i> , 2016, 175, 178-181.	1.4	33
21	Mutations in TSPEAR, Encoding a Regulator of Notch Signaling, Affect Tooth and Hair Follicle Morphogenesis. <i>PLoS Genetics</i> , 2016, 12, e1006369.	1.5	32
22	Establishment of Two Mouse Models for CEDNIK Syndrome Reveals the Pivotal Role of SNAP29 in Epidermal Differentiation. <i>Journal of Investigative Dermatology</i> , 2016, 136, 672-679.	0.3	31
23	Loss-of-Function Mutations in SERPINB8 Linked to Exfoliative Ichthyosis with Impaired Mechanical Stability of Intercellular Adhesions. <i>American Journal of Human Genetics</i> , 2016, 99, 430-436.	2.6	27
24	Analysis of CARD14 Polymorphisms in Pityriasis Rubra Pilaris: Activation of NF- $\kappa$ B. <i>Journal of Investigative Dermatology</i> , 2015, 135, 1905-1908.	0.3	24
25	SAM syndrome is characterized by extensive phenotypic heterogeneity. <i>Experimental Dermatology</i> , 2018, 27, 787-790.	1.4	22
26	ST18 Enhances PV-IgG-Induced Loss of Keratinocyte Cohesion in Parallel to Increased ERK Activation. <i>Frontiers in Immunology</i> , 2019, 10, 770.	2.2	20
27	Calpain 12 Function Revealed through the Study of an Atypical Case of Autosomal Recessive Congenital Ichthyosis. <i>Journal of Investigative Dermatology</i> , 2017, 137, 385-393.	0.3	19
28	Treatment of hereditary hypotrichosis simplex of the scalp with topical gentamicin. <i>British Journal of Dermatology</i> , 2020, 183, 114-120.	1.4	19
29	<i>RBM</i> 28, a protein deficient in <i>ANE</i> syndrome, regulates hair follicle growth via miR-203 and p63. <i>Experimental Dermatology</i> , 2015, 24, 618-622.	1.4	17
30	<i>SVEP</i> 1 plays a crucial role in epidermal differentiation. <i>Experimental Dermatology</i> , 2017, 26, 423-430.	1.4	17
31	The Role of Desmoglein 1 in Gap Junction Turnover Revealed through the Study of SAM Syndrome. <i>Journal of Investigative Dermatology</i> , 2020, 140, 556-567.e9.	0.3	17
32	The Molecular Revolution in Cutaneous Biology: Era of Next-Generation Sequencing. <i>Journal of Investigative Dermatology</i> , 2017, 137, e79-e82.	0.3	16
33	IGFBP7 as a Potential Therapeutic Target in Psoriasis. <i>Journal of Investigative Dermatology</i> , 2011, 131, 1767-1770.	0.3	14
34	Loss-of-Function Variants in SERPINA12 Underlie Autosomal Recessive Palmoplantar Keratoderma. <i>Journal of Investigative Dermatology</i> , 2020, 140, 2178-2187.	0.3	14
35	Molecular Analysis of a Series of Israeli Families with Com $\bar{1}$ -Netherton Syndrome. <i>Dermatology</i> , 2014, 228, 183-188.	0.9	12
36	Coagulation Factor XIII-A Subunit Missense Mutation in the Pathobiology of Autosomal Dominant Multiple Dermatofibromas. <i>Journal of Investigative Dermatology</i> , 2020, 140, 624-635.e7.	0.3	12

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37	Loss-of-function variants in C3ORF52 result in localized autosomal recessive hypotrichosis. <i>Genetics in Medicine</i> , 2020, 22, 1227-1234.	1.1	12
38	Papillon-Loefgren syndrome: report of six patients and identification of a novel mutation. <i>International Journal of Dermatology</i> , 2016, 55, 898-902.	0.5	11
39	Identification of a recurrent mutation in <i>ATP2C1</i> demonstrates that papular acantholytic dyskeratosis and Hailey-Hailey disease are allelic disorders. <i>British Journal of Dermatology</i> , 2018, 179, 1001-1002.	1.4	11
40	Semidominant Inheritance in Epidermolytic Ichthyosis. <i>Journal of Investigative Dermatology</i> , 2013, 133, 2626-2628.	0.3	10
41	Recessive epidermolytic ichthyosis results from loss of keratin 10 expression, regardless of the mutation location. <i>Clinical and Experimental Dermatology</i> , 2018, 43, 187-190.	0.6	10
42	Molecular epidemiology of non-syndromic autosomal recessive congenital ichthyosis in a Middle-Eastern population. <i>Experimental Dermatology</i> , 2021, 30, 1290-1297.	1.4	10
43	Evidence for cutaneous dysbiosis in dystrophic epidermolysis bullosa. <i>Clinical and Experimental Dermatology</i> , 2021, 46, 1223-1229.	0.6	10
44	Extensive lentigo simplex, linear epidermolytic naevus and epidermolytic naevus comedonicus caused by a somatic mutation in <i>KRT10</i> . <i>British Journal of Dermatology</i> , 2015, 173, 293-296.	1.4	9
45	Epidermolysis bullosa simplex due to biallelic <i>DST</i> mutations: Case series and review of the literature. <i>Pediatric Dermatology</i> , 2021, 38, 436-441.	0.5	9
46	Non-keratinocyte <i>SNAP29</i> influences epidermal differentiation and hair follicle formation in mice. <i>Experimental Dermatology</i> , 2016, 25, 647-649.	1.4	8
47	Striate palmoplantar keratoderma resulting from a missense mutation in <i>DSG1</i> . <i>British Journal of Dermatology</i> , 2018, 179, 755-757.	1.4	8
48	PLACK syndrome shows remarkable phenotypic homogeneity. <i>Clinical and Experimental Dermatology</i> , 2019, 44, 580-583.	0.6	8
49	A novel homozygous deletion in <i>EXPH5</i> causes a skin fragility phenotype. <i>Clinical and Experimental Dermatology</i> , 2016, 41, 915-918.	0.6	7
50	ST18 affects cell-cell adhesion in pemphigus vulgaris in a tumour necrosis factor-1-dependent fashion*. <i>British Journal of Dermatology</i> , 2021, 184, 1153-1160.	1.4	7
51	Vorinostat, a histone deacetylase inhibitor, as a potential novel treatment for psoriasis. <i>Experimental Dermatology</i> , 2022, 31, 567-576.	1.4	7
52	Coexistence of pachyonychia congenita and hidradenitis suppurativa: more than a coincidence. <i>British Journal of Dermatology</i> , 2022, 187, 392-400.	1.4	7
53	ARCI7 Revisited and Repositioned. <i>Journal of Investigative Dermatology</i> , 2017, 137, 970-972.	0.3	6
54	Immune-regulatory genes as possible modifiers of familial pityriasis rubra pilaris – lessons from a family with <i>PRP</i> and psoriasis. <i>Journal of the European Academy of Dermatology and Venereology</i> , 2018, 32, e389-e392.	1.3	6

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55	A Mutation in TP63 Causing a Mild Ectodermal Dysplasia Phenotype. <i>Journal of Investigative Dermatology</i> , 2014, 134, 2277-2280.	0.3	5
56	Somatic Mosaicism for a Lethal GJB2 Mutation Results in a Patterned Form of Spiny Hyperkeratosis without Eccrine Involvement. <i>Pediatric Dermatology</i> , 2016, 33, 322-326.	0.5	5
57	Punctate palmoplantar keratoderma: an unusual mutation causing an unusual phenotype. <i>British Journal of Dermatology</i> , 2018, 178, 1455-1457.	1.4	5
58	Palmoplantar keratoderma caused by a missense variant in CTSB encoding cathepsin B. <i>Clinical and Experimental Dermatology</i> , 2021, 46, 103-108.	0.6	5
59	Identification of clinically useful predictive genetic variants in pachyonychia congenita. <i>Clinical and Experimental Dermatology</i> , 2021, 46, 867-873.	0.6	5
60	Autosomal dominant cutis laxa resulting from an intronic mutation in ELN. <i>Experimental Dermatology</i> , 2015, 24, 885-887.	1.4	4
61	Glutathione S-transferase polymorphisms in patients with photosensitive and non-photosensitive drug eruptions. <i>Photodermatology Photoimmunology and Photomedicine</i> , 2019, 35, 214-220.	0.7	4
62	Neonatal inflammatory skin and bowel disease type 1 caused by a complex genetic defect and responsive to combined anti-tumour necrosis factor- $\alpha$ and interleukin-12/23 blockade. <i>British Journal of Dermatology</i> , 2022, 186, 1026-1029.	1.4	4
63	Novel POFUT 1 mutation associated with hidradenitis suppurativa "Dowling-Degos disease firm up a role for Notch signalling in the pathogenesis of this disorder: reply from the authors. <i>British Journal of Dermatology</i> , 2018, 178, 986-986.	1.4	3
64	Loss-of-function variants in KLF4 underlie autosomal dominant palmoplantar keratoderma. <i>Genetics in Medicine</i> , 2022, 24, 1085-1095.	1.1	3
65	Guanine polynucleotides are self-antigens for human natural autoantibodies and are significantly reduced in the human genome. <i>Immunology</i> , 2015, 146, 401-410.	2.0	2
66	Phenotypic suppression of acral peeling skin syndrome in a patient with autosomal recessive congenital ichthyosis. <i>Experimental Dermatology</i> , 2020, 29, 742-748.	1.4	2
67	Identification of a founder mutation in KRT14 associated with Naegeli-Franceschetti-Jadassohn syndrome. <i>British Journal of Dermatology</i> , 2020, 183, 756-757.	1.4	2
68	A unique skin phenotype resulting from a large heterozygous deletion spanning six keratin genes. <i>British Journal of Dermatology</i> , 2022, 187, 773-777.	1.4	2
69	A case for diagnosis. <i>Clinical and Experimental Dermatology</i> , 2015, 40, 697-699.	0.6	1
70	Molecular epidemiology of pachyonychia congenita in the Israeli population. <i>Clinical and Experimental Dermatology</i> , 2021, 46, 663-668.	0.6	1
71	Epidermolytic epidermal nevus caused by a somatic mutation in KRT2. <i>Pediatric Dermatology</i> , 2021, 38, 538-540.	0.5	1
72	Up-regulation of ST18 in pemphigus vulgaris drives a self-amplifying p53-dependent pathomechanism resulting in decreased desmoglein 3 expression. <i>Scientific Reports</i> , 2022, 12, 5958.	1.6	1

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73	Heterozygous variants in the integrin subunit beta 4 gene (ITGB4) cause autosomal dominant nail dystrophy. <i>British Journal of Dermatology</i> , 2022, 187, 826-828.	1.4	1
74	Atypical presentation of laryngo-onycho-cutaneous syndrome resulting from novel mutations in LAMA3A. <i>Clinical and Experimental Dermatology</i> , 2021, 46, 990-992.	0.6	0
75	Acute Respiratory Distress Syndrome in a Carrier of an Interleukin-36 Receptor Antagonist Mutation With Generalized Pustular Psoriasis. <i>Journal of Psoriasis and Psoriatic Arthritis</i> , 2022, 7, 9-12.	0.3	0