

# Johannes P Langedijk

## List of Publications by Year in descending order

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48  
papers

4,633  
citations

172457

29  
h-index

197818

49  
g-index

50  
all docs

50  
docs citations

50  
times ranked

7755  
citing authors

#	ARTICLE	IF	CITATIONS
1	Universal stabilization of the influenza hemagglutinin by structure-based redesign of the pH switch regions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	7
2	Stabilizing the closed SARS-CoV-2 spike trimer. <i>Nature Communications</i> , 2021, 12, 244.	12.8	139
3	Ad26.COVS.S protects Syrian hamsters against G614 spike variant SARS-CoV-2 and does not enhance respiratory disease. <i>Npj Vaccines</i> , 2021, 6, 39.	6.0	38
4	Safety and immunogenicity of two heterologous HIV vaccine regimens in healthy, HIV-uninfected adults (TRAVERSE): a randomised, parallel-group, placebo-controlled, double-blind, phase 1/2a study. <i>Lancet HIV</i> , 2020, 7, e688-e698.	4.7	58
5	Ad26 vector-based COVID-19 vaccine encoding a prefusion-stabilized SARS-CoV-2 Spike immunogen induces potent humoral and cellular immune responses. <i>Npj Vaccines</i> , 2020, 5, 91.	6.0	286
6	Single-shot Ad26 vaccine protects against SARS-CoV-2 in rhesus macaques. <i>Nature</i> , 2020, 586, 583-588.	27.8	765
7	Automated Design by Structure-Based Stabilization and Consensus Repair to Achieve Prefusion-Closed Envelope Trimers in a Wide Variety of HIV Strains. <i>Cell Reports</i> , 2020, 33, 108432.	6.4	32
8	Adenovector 26 encoded prefusion conformation stabilized RSV-F protein induces long-lasting Th1-biased immunity in neonatal mice. <i>Npj Vaccines</i> , 2020, 5, 49.	6.0	24
9	Structure-Based Design of Prefusion-Stabilized Filovirus Glycoprotein Trimers. <i>Cell Reports</i> , 2020, 30, 4540-4550.e3.	6.4	46
10	HIV-1 anchor inhibitors and membrane fusion inhibitors target distinct but overlapping steps in virus entry. <i>Journal of Biological Chemistry</i> , 2019, 294, 5736-5746.	3.4	24
11	Transient opening of trimeric prefusion RSV F proteins. <i>Nature Communications</i> , 2019, 10, 2105.	12.8	71
12	Primary resistance mechanism of the canine distemper virus fusion protein against a small-molecule membrane fusion inhibitor. <i>Virus Research</i> , 2019, 259, 28-37.	2.2	10
13	A Universal Approach to Optimize the Folding and Stability of Prefusion-Closed HIV-1 Envelope Trimers. <i>Cell Reports</i> , 2018, 23, 584-595.	6.4	93
14	Structural basis for recognition of the central conserved region of RSV G by neutralizing human antibodies. <i>PLoS Pathogens</i> , 2018, 14, e1006935.	4.7	50
15	Therapeutic efficacy of a respiratory syncytial virus fusion inhibitor. <i>Nature Communications</i> , 2017, 8, 167.	12.8	58
16	Peptide mimetics of immunoglobulin A (IgA) and FcγRI block IgA-induced human neutrophil activation and migration. <i>European Journal of Immunology</i> , 2017, 47, 1835-1845.	2.9	16
17	HIV-1 Escape from a Peptidic Anchor Inhibitor through Stabilization of the Envelope Glycoprotein Spike. <i>Journal of Virology</i> , 2016, 90, 10587-10599.	3.4	18
18	Molecular mechanism of respiratory syncytial virus fusion inhibitors. <i>Nature Chemical Biology</i> , 2016, 12, 87-93.	8.0	121

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19	Sequential Conformational Changes in the Morbillivirus Attachment Protein Initiate the Membrane Fusion Process. <i>PLoS Pathogens</i> , 2015, 11, e1004880.	4.7	35
20	A highly stable prefusion RSV F vaccine derived from structural analysis of the fusion mechanism. <i>Nature Communications</i> , 2015, 6, 8143.	12.8	248
21	Identification of Amino Acid Substitutions with Compensational Effects in the Attachment Protein of Canine Distemper Virus. <i>Journal of Virology</i> , 2014, 88, 8057-8064.	3.4	17
22	A gp41 MPER-specific Llama VHH Requires a Hydrophobic CDR3 for Neutralization but not for Antigen Recognition. <i>PLoS Pathogens</i> , 2013, 9, e1003202.	4.7	64
23	Structure-Based Design for High-Hanging Vaccine Fruits. <i>Advances in Immunology</i> , 2012, 114, 33-50.	2.2	7
24	Structural Rearrangements of the Central Region of the Morbillivirus Attachment Protein Stalk Domain Trigger F Protein Refolding for Membrane Fusion. <i>Journal of Biological Chemistry</i> , 2012, 287, 16324-16334.	3.4	63
25	A sweet surprise for HIV broadly neutralizing antibodies. <i>Nature Medicine</i> , 2012, 18, 1616-1617.	30.7	3
26	Epitope Mapping of Broadly Neutralizing HIV-2 Human Monoclonal Antibodies. <i>Journal of Virology</i> , 2012, 86, 12115-12128.	3.4	27
27	Peptides based on the presenilin-1 APP binding domain inhibit APP processing and A $\beta$ production through interfering with the APP transmembrane domain. <i>FASEB Journal</i> , 2012, 26, 3765-3778.	0.5	11
28	A Neutralizing Antibody Selected from Plasma Cells That Binds to Group 1 and Group 2 Influenza A Hemagglutinins. <i>Science</i> , 2011, 333, 850-856.	12.6	1,092
29	Helical peptide arrays for lead identification and interaction site mapping. <i>Analytical Biochemistry</i> , 2011, 417, 149-155.	2.4	11
30	Resistance of Human Immunodeficiency Virus Type 1 to a Third-Generation Fusion Inhibitor Requires Multiple Mutations in gp41 and Is Accompanied by a Dramatic Loss of gp41 Function. <i>Journal of Virology</i> , 2011, 85, 10785-10797.	3.4	66
31	Canine Distemper Virus Infects Canine Keratinocytes and Immune Cells by Using Overlapping and Distinct Regions Located on One Side of the Attachment Protein. <i>Journal of Virology</i> , 2011, 85, 11242-11254.	3.4	31
32	Analysis of Memory B Cell Responses and Isolation of Novel Monoclonal Antibodies with Neutralizing Breadth from HIV-1-Infected Individuals. <i>PLoS ONE</i> , 2010, 5, e8805.	2.5	405
33	Identification of Key Residues in Virulent Canine Distemper Virus Hemagglutinin That Control CD150/SLAM-Binding Activity. <i>Journal of Virology</i> , 2010, 84, 9618-9624.	3.4	32
34	Detailed Mechanistic Insights into HIV-1 Sensitivity to Three Generations of Fusion Inhibitors. <i>Journal of Biological Chemistry</i> , 2009, 284, 26941-26950.	3.4	71
35	Heterologous Stacking of Prion Protein Peptides Reveals Structural Details of Fibrils and Facilitates Complete Inhibition of Fibril Growth. <i>Journal of Biological Chemistry</i> , 2009, 284, 12809-12820.	3.4	5
36	Selection of T1249-Resistant Human Immunodeficiency Virus Type 1 Variants. <i>Journal of Virology</i> , 2008, 82, 6678-6688.	3.4	76

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37	SUMO Assay with Peptide Arrays on Solid Support: Insights into SUMO Target Sites. <i>Journal of Biochemistry</i> , 2008, 144, 39-49.	1.7	12
38	Polyanion induced fibril growth enables the development of a reproducible assay in solution for the screening of fibril interfering compounds, and the investigation of the prion nucleation site. <i>Amyloid: the International Journal of Experimental and Clinical Investigation: the Official Journal of the International Society of Amyloidosis</i> , 2007, 14, 205-219.	3.0	3
39	Evidence of a Potential Receptor-Binding Site on the Nipah Virus G Protein (NiV-G): Identification of Globular Head Residues with a Role in Fusion Promotion and Their Localization on an NiV-G Structural Model. <i>Journal of Virology</i> , 2006, 80, 7546-7554.	3.4	47
40	Activation of human microglia by fibrillar prion protein-related peptides is enhanced by amyloid-associated factors SAP and C1q. <i>Neurobiology of Disease</i> , 2005, 19, 273-282.	4.4	21
41	Measles Virus (MV) Hemagglutinin: Evidence that Attachment Sites for MV Receptors SLAM and CD46 Overlap on the Globular Head. <i>Journal of Virology</i> , 2004, 78, 9051-9063.	3.4	79
42	An in vitro screening assay based on synthetic prion protein peptides for identification of fibril-interfering compounds. <i>Analytical Biochemistry</i> , 2004, 333, 372-380.	2.4	25
43	Vaccine-Induced Immunopathology during Bovine Respiratory Syncytial Virus Infection: Exploring the Parameters of Pathogenesis. <i>Journal of Virology</i> , 2003, 77, 12067-12073.	3.4	85
44	Translocation Activity of C-terminal Domain of Pestivirus Erns and Ribotoxin L3 Loop. <i>Journal of Biological Chemistry</i> , 2002, 277, 5308-5314.	3.4	57
45	Novel Strategy for Inhibiting Viral Entry by Use of a Cellular Receptor-Plant Virus Chimera. <i>Journal of Virology</i> , 2002, 76, 4412-4419.	3.4	35
46	Structural and Functional Relationship between the Receptor Recognition and Neuraminidase Activities of the Newcastle Disease Virus Hemagglutinin-Neuraminidase Protein: Receptor Recognition Is Dependent on Neuraminidase Activity. <i>Journal of Virology</i> , 2001, 75, 1918-1927.	3.4	77
47	Solution Structure of the Immunodominant Region of Protein G of Bovine Respiratory Syncytial Virus. <i>Biochemistry</i> , 1996, 35, 14684-14688.	2.5	38
48	A Hidden Region in the Third Variable Domain of HIV-1 IIIB gp120 Identified by a Monoclonal Antibody. <i>AIDS Research and Human Retroviruses</i> , 1993, 9, 605-612.	1.1	23