

Aleksander S Popel

List of Publications by Year in Descending Order

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Version: 2024-04-26

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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.

295
papers

11,080
citations

56
h-index

88
g-index

322
ext. papers

12,749
ext. citations

4.4
avg, IF

6.67
L-index

#	Paper	IF	Citations
295	Dynamic Multiscale Regulation of Perfusion Recovery in Experimental Peripheral Arterial Disease: A Mechanistic Computational Model.. <i>JACC Basic To Translational Science</i> , 2022 , 7, 28-50	8.7	1
294	Systems biology modeling of endothelial cell and macrophage signaling in angiogenesis in human diseases 2022 , 163-172		
293	Quantitative Systems Pharmacology Modeling of PBMC-Humanized Mouse to Facilitate Preclinical Immuno-oncology Drug Development. <i>ACS Pharmacology and Translational Science</i> , 2021 , 4, 213-225	5.9	2
292	Peptides that immunoactivate the tumor microenvironment. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2021 , 1875, 188486	11.2	5
291	A data-driven computational model enables integrative and mechanistic characterization of dynamic macrophage polarization. <i>IScience</i> , 2021 , 24, 102112	6.1	9
290	Quantitative systems pharmacology model predictions for efficacy of atezolizumab and nab-paclitaxel in triple-negative breast cancer 2021 , 9,		4
289	A systems biology model of junctional localization and downstream signaling of the Ang-Tie signaling pathway. <i>Npj Systems Biology and Applications</i> , 2021 , 7, 34	5	2
288	Protocol for simulating macrophage signal transduction and phenotype polarization using a large-scale mechanistic computational model. <i>STAR Protocols</i> , 2021 , 2, 100739	1.4	1
287	Predictive models of response to neoadjuvant chemotherapy in muscle-invasive bladder cancer using nuclear morphology and tissue architecture. <i>Cell Reports Medicine</i> , 2021 , 2, 100382	18	1
286	Integrating single cell sequencing with a spatial quantitative systems pharmacology model spQSP for personalized prediction of triple-negative breast cancer immunotherapy response. <i>Immunoinformatics</i> , 2021 , 1-2,		4
285	Systems biology of angiogenesis signaling: Computational models and omics.. <i>WIREs Mechanisms of Disease</i> , 2021 , e1550	0.3	1
284	Regulation of the tumor immune microenvironment and vascular normalization in TNBC murine models by a novel peptide. <i>Onc Immunology</i> , 2020 , 9, 1760685	7.2	5
283	A Quantitative Systems Pharmacology Model of T Cell Engager Applied to Solid Tumor. <i>AAPS Journal</i> , 2020 , 22, 85	3.7	9
282	Cytokines secreted by stromal cells in TNBC microenvironment as potential targets for cancer therapy. <i>Cancer Biology and Therapy</i> , 2020 , 21, 560-569	4.6	8
281	Conducting a Virtual Clinical Trial in HER2-Negative Breast Cancer Using a Quantitative Systems Pharmacology Model With an Epigenetic Modulator and Immune Checkpoint Inhibitors. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020 , 8, 141	5.8	17
280	Immunoactivating the tumor microenvironment enhances immunotherapy as predicted by integrative computational model. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 4447-4449	11.5	13
279	Mathematical Models of Transport Phenomena in Normal and Neoplastic Tissue 2020 , 169-183		1

278	Simulations of Combination Therapy in Hepatocellular Carcinoma Using a Quantitative Systems Pharmacology Model. <i>FASEB Journal</i> , 2020 , 34, 1-1	0.9	
277	QSP-IO: A Quantitative Systems Pharmacology Toolbox for Mechanistic Multiscale Modeling for Immuno-Oncology Applications. <i>CPT: Pharmacometrics and Systems Pharmacology</i> , 2020 , 9, 484-497	4.5	11
276	Suppression of Ocular Vascular Inflammation through Peptide-Mediated Activation of Angiopoietin-Tie2 Signaling. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	4
275	Digital Pathology Analysis Quantifies Spatial Heterogeneity of CD3, CD4, CD8, CD20, and FoxP3 Immune Markers in Triple-Negative Breast Cancer. <i>Frontiers in Physiology</i> , 2020 , 11, 583333	4.6	12
274	Combination therapy with T cell engager and PD-L1 blockade enhances the antitumor potency of T cells as predicted by a QSP model 2020 , 8,		12
273	Multiscale Agent-Based and Hybrid Modeling of the Tumor Immune Microenvironment. <i>Processes</i> , 2019 , 7,	2.9	61
272	Mechanistic Computational Models of MicroRNA-Mediated Signaling Networks in Human Diseases. <i>International Journal of Molecular Sciences</i> , 2019 , 20,	6.3	13
271	simulation of a clinical trial with anti-CTLA-4 and anti-PD-L1 immunotherapies in metastatic breast cancer using a systems pharmacology model. <i>Royal Society Open Science</i> , 2019 , 6, 190366	3.3	26
270	Dynamic Changes in Microvascular Flow Conductivity and Perfusion After Myocardial Infarction Shown by Image-Based Modeling. <i>Journal of the American Heart Association</i> , 2019 , 8, e011058	6	7
269	Tumor Ensemble-Based Modeling and Visualization of Emergent Angiogenic Heterogeneity in Breast Cancer. <i>Scientific Reports</i> , 2019 , 9, 5276	4.9	12
268	Mechanistically detailed systems biology modeling of the HGF/Met pathway in hepatocellular carcinoma. <i>Npj Systems Biology and Applications</i> , 2019 , 5, 29	5	10
267	Anisotropic poly(lactic-co-glycolic acid) microparticles enable sustained release of a peptide for long-term inhibition of ocular neovascularization. <i>Acta Biomaterialia</i> , 2019 , 97, 451-460	10.8	9
266	A QSP Model for Predicting Clinical Responses to Monotherapy, Combination and Sequential Therapy Following CTLA-4, PD-1, and PD-L1 Checkpoint Blockade. <i>Scientific Reports</i> , 2019 , 9, 11286	4.9	34
265	A Computational Model of Neoadjuvant PD-1 Inhibition in Non-Small Cell Lung Cancer. <i>AAPS Journal</i> , 2019 , 21, 79	3.7	25
264	Angiopoietin-Tie Signaling Pathway in Endothelial Cells: A Computational Model. <i>iScience</i> , 2019 , 20, 497-511		19
263	A collagen IV-derived peptide disrupts $\beta 1$ integrin and potentiates Ang2/Tie2 signaling. <i>JCI Insight</i> , 2019 , 4,	9.9	23
262	A mechanistic integrative computational model of macrophage polarization: Implications in human pathophysiology. <i>PLoS Computational Biology</i> , 2019 , 15, e1007468	5	20
261	Deciphering microvascular changes after myocardial infarction through 3D fully automated image analysis. <i>Scientific Reports</i> , 2018 , 8, 1854	4.9	11

260	Human expression patterns: qualitative and quantitative analysis of thrombospondin-1 under physiological and pathological conditions. <i>Journal of Cellular and Molecular Medicine</i> , 2018 , 22, 2086-2097	5.6	22
259	Biomimetic peptide display from a polymeric nanoparticle surface for targeting and antitumor activity to human triple-negative breast cancer cells. <i>Journal of Biomedical Materials Research - Part A</i> , 2018 , 106, 1753-1764	5.4	14
258	Computer Simulation of TSP1 Inhibition of VEGF-Akt-eNOS: An Angiogenesis Triple Threat. <i>Frontiers in Physiology</i> , 2018 , 9, 644	4.6	20
257	Computational modeling of synergistic interaction between α 5 β 1 integrin and VEGFR2 in endothelial cells: Implications for the mechanism of action of angiogenesis-modulating integrin-binding peptides. <i>Journal of Theoretical Biology</i> , 2018 , 455, 212-221	2.3	12
256	Simultaneous blockade of IL-6 and CCL5 signaling for synergistic inhibition of triple-negative breast cancer growth and metastasis. <i>Breast Cancer Research</i> , 2018 , 20, 54	8.3	41
255	Modeling triple-negative breast cancer heterogeneity: Effects of stromal macrophages, fibroblasts and tumor vasculature. <i>Journal of Theoretical Biology</i> , 2018 , 452, 56-68	2.3	30
254	Quantitative Characterization of CD8+ T Cell Clustering and Spatial Heterogeneity in Solid Tumors. <i>Frontiers in Oncology</i> , 2018 , 8, 649	5.3	17
253	Three-Dimensional Transport Model for Intravitreal and Suprachoroidal Drug Injection 2018 , 59, 5266-5276		17
252	Tyrosine kinase blocking collagen IV-derived peptide suppresses ocular neovascularization and vascular leakage. <i>Science Translational Medicine</i> , 2017 , 9,	17.5	29
251	Gene delivery nanoparticles to modulate angiogenesis. <i>Advanced Drug Delivery Reviews</i> , 2017 , 119, 20-43	8.5	46
250	Transcriptional and Post-Transcriptional Regulation of Thrombospondin-1 Expression: A Computational Model. <i>PLoS Computational Biology</i> , 2017 , 13, e1005272	5	13
249	Computational investigation of sphingosine kinase 1 (SphK1) and calcium dependent ERK1/2 activation downstream of VEGFR2 in endothelial cells. <i>PLoS Computational Biology</i> , 2017 , 13, e1005332	5	17
248	A computational multiscale agent-based model for simulating spatio-temporal tumour immune response to PD1 and PDL1 inhibition. <i>Journal of the Royal Society Interface</i> , 2017 , 14,	4.1	64
247	An agent-based model of triple-negative breast cancer: the interplay between chemokine receptor CCR5 expression, cancer stem cells, and hypoxia. <i>BMC Systems Biology</i> , 2017 , 11, 68	3.5	26
246	Inhibition of VEGFR2 Activation and Its Downstream Signaling to ERK1/2 and Calcium by Thrombospondin-1 (TSP1): Investigation. <i>Frontiers in Physiology</i> , 2017 , 8, 48	4.6	17
245	Abstract 4531: Systems pharmacology to predict cellular biomarkers and optimize mono- and combination-therapy regimens: Focusing on immune checkpoint targets PD-1, PD-L1 and CTLA-4 2017 ,		2
244	Crosstalk between stromal components and tumor cells of TNBC via secreted factors enhances tumor growth and metastasis. <i>Oncotarget</i> , 2017 , 8, 60210-60222	3.3	27
243	Therapeutic potential of an anti-angiogenic multimodal biomimetic peptide in hepatocellular carcinoma. <i>Oncotarget</i> , 2017 , 8, 101520-101534	3.3	5

242	Effects of endothelial cell proliferation and migration rates in a computational model of sprouting angiogenesis. <i>Scientific Reports</i> , 2016 , 6, 36992	4.9	80
241	A multiscale computational model predicts distribution of anti-angiogenic isoform VEGF in peripheral arterial disease in human and mouse. <i>Scientific Reports</i> , 2016 , 6, 37030	4.9	16
240	Multiscale Modeling in the Clinic: Drug Design and Development. <i>Annals of Biomedical Engineering</i> , 2016 , 44, 2591-610	4.7	36
239	Endothelial cells decode VEGF-mediated Ca ²⁺ signaling patterns to produce distinct functional responses. <i>Science Signaling</i> , 2016 , 9, ra20	8.8	56
238	Multi-scale Modeling in Clinical Oncology: Opportunities and Barriers to Success. <i>Annals of Biomedical Engineering</i> , 2016 , 44, 2626-41	4.7	48
237	The Angiogenic Secretome in VEGF overexpressing Breast Cancer Xenografts. <i>Scientific Reports</i> , 2016 , 6, 39460	4.9	15
236	Crosstalk between cancer cells and blood endothelial and lymphatic endothelial cells in tumour and organ microenvironment. <i>Expert Reviews in Molecular Medicine</i> , 2015 , 17, e3	6.7	55
235	Analysis of gene expression of secreted factors associated with breast cancer metastases in breast cancer subtypes. <i>Scientific Reports</i> , 2015 , 5, 12133	4.9	30
234	Vasculature-specific MRI reveals differential anti-angiogenic effects of a biomimetic peptide in an orthotopic breast cancer model. <i>Angiogenesis</i> , 2015 , 18, 125-36	10.6	8
233	A 3D Fractal-Based Approach towards Understanding Changes in the Infarcted Heart Microvasculature. <i>Lecture Notes in Computer Science</i> , 2015 , 173-180	0.9	2
232	Computational systems biology approaches to anti-angiogenic cancer therapeutics. <i>Drug Discovery Today</i> , 2015 , 20, 187-97	8.8	36
231	PADPIN: protein-protein interaction networks of angiogenesis, arteriogenesis, and inflammation in peripheral arterial disease. <i>Physiological Genomics</i> , 2015 , 47, 331-43	3.6	12
230	Pharmacokinetics of Anti-VEGF Agent Aflibercept in Cancer Predicted by Data-Driven, Molecular-Detailed Model. <i>CPT: Pharmacometrics and Systems Pharmacology</i> , 2015 , 4, 641-9	4.5	14
229	Computational Model of MicroRNA Control of HIF-VEGF Pathway: Insights into the Pathophysiology of Ischemic Vascular Disease and Cancer. <i>PLoS Computational Biology</i> , 2015 , 11, e1004612	5.2	25
228	Computational drug repositioning for peripheral arterial disease: prediction of anti-inflammatory and pro-angiogenic therapeutics. <i>Frontiers in Pharmacology</i> , 2015 , 6, 179	5.6	6
227	Heterogeneity of chemokine cell-surface receptor expression in triple-negative breast cancer. <i>American Journal of Cancer Research</i> , 2015 , 5, 1295-307	4.4	14
226	In Vitro and in Vivo Analyses of the Effects of Sunitinib on Endothelial Cell-Surface Vascular Endothelial Growth Factor Receptor-2. <i>FASEB Journal</i> , 2015 , 29, 780.4	0.9	1
225	Image-based Characterization of Functional and Structural Heterogeneity of Tumor Xenografts using Blood Flow modeling, Oxygenation Modeling and Multivariate Analysis. <i>FASEB Journal</i> , 2015 , 29, 787.11	0.9	1

224	Inhibition of breast cancer growth and metastasis by a biomimetic peptide. <i>Scientific Reports</i> , 2014 , 4, 7139	4.9	31
223	Lymphatic endothelial cells support tumor growth in breast cancer. <i>Scientific Reports</i> , 2014 , 4, 5853	4.9	37
222	Extracellular regulation of VEGF: isoforms, proteolysis, and vascular patterning. <i>Cytokine and Growth Factor Reviews</i> , 2014 , 25, 1-19	17.9	198
221	Breast cancer cells condition lymphatic endothelial cells within pre-metastatic niches to promote metastasis. <i>Nature Communications</i> , 2014 , 5, 4715	17.4	108
220	An agent-based model of cancer stem cell initiated avascular tumour growth and metastasis: the effect of seeding frequency and location. <i>Journal of the Royal Society Interface</i> , 2014 , 11, 20140640	4.1	22
219	A systems biology view of blood vessel growth and remodelling. <i>Journal of Cellular and Molecular Medicine</i> , 2014 , 18, 1491-508	5.6	92
218	A bioimage informatics based reconstruction of breast tumor microvasculature with computational blood flow predictions. <i>Microvascular Research</i> , 2014 , 91, 8-21	3.7	52
217	A biomimetic collagen derived peptide exhibits anti-angiogenic activity in triple negative breast cancer. <i>PLoS ONE</i> , 2014 , 9, e111901	3.7	9
216	Angiogenesis interactome and time course microarray data reveal the distinct activation patterns in endothelial cells. <i>PLoS ONE</i> , 2014 , 9, e110871	3.7	14
215	Antiangiogenic cancer drug sunitinib exhibits unexpected proangiogenic effects on endothelial cells. <i>OncoTargets and Therapy</i> , 2014 , 7, 1571-82	4.4	13
214	Quantitative fluorescent profiling of VEGFRs reveals tumor cell and endothelial cell heterogeneity in breast cancer xenografts. <i>Cancer Medicine</i> , 2014 , 3, 225-44	4.8	41
213	Pre-treatment of mice with tumor-conditioned media accelerates metastasis to lymph nodes and lungs: a new spontaneous breast cancer metastasis model. <i>Clinical and Experimental Metastasis</i> , 2014 , 31, 67-79	4.7	25
212	Synergy between a collagen IV mimetic peptide and a somatotropin-domain derived peptide as angiogenesis and lymphangiogenesis inhibitors. <i>Angiogenesis</i> , 2013 , 16, 159-70	10.6	16
211	Inhibition of lymphangiogenesis and angiogenesis in breast tumor xenografts and lymph nodes by a peptide derived from transmembrane protein 45A. <i>Neoplasia</i> , 2013 , 15, 112-24	6.4	40
210	Long-term suppression of ocular neovascularization by intraocular injection of biodegradable polymeric particles containing a serpin-derived peptide. <i>Biomaterials</i> , 2013 , 34, 7544-51	15.6	41
209	Computational model of VEGFR2 pathway to ERK activation and modulation through receptor trafficking. <i>Cellular Signalling</i> , 2013 , 25, 2496-510	4.9	30
208	Endothelial cell-by-cell profiling reveals the temporal dynamics of VEGFR1 and VEGFR2 membrane localization after murine hindlimb ischemia. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2013 , 304, H1085-93	5.2	51
207	Compartment model predicts VEGF secretion and investigates the effects of VEGF trap in tumor-bearing mice. <i>Frontiers in Oncology</i> , 2013 , 3, 196	5.3	34

206	Effect of tumor microenvironment on tumor VEGF during anti-VEGF treatment: systems biology predictions. <i>Journal of the National Cancer Institute</i> , 2013 , 105, 802-11	9.7	69
205	Computational Model of Gab1/2-Dependent VEGFR2 Pathway to Akt Activation. <i>PLoS ONE</i> , 2013 , 8, e67438	3.7	26
204	Identification of morphological and hemodynamic biomarkers for tumor vascular perfusion through mathematical modeling and high-resolution imaging. <i>FASEB Journal</i> , 2013 , 27, 685.12	0.9	
203	Structure-activity relationship study of collagen-derived anti-angiogenic biomimetic peptides. <i>Chemical Biology and Drug Design</i> , 2012 , 80, 27-37	2.9	12
202	Multiscale imaging and computational modeling of blood flow in the tumor vasculature. <i>Annals of Biomedical Engineering</i> , 2012 , 40, 2425-41	4.7	40
201	Predicting the effects of anti-angiogenic agents targeting specific VEGF isoforms. <i>AAPS Journal</i> , 2012 , 14, 500-9	3.7	45
200	Serpin-derived peptides are antiangiogenic and suppress breast tumor xenograft growth. <i>Translational Oncology</i> , 2012 , 5, 92-7	4.9	18
199	Blood-plasma separation in Y-shaped bifurcating microfluidic channels: a dissipative particle dynamics simulation study. <i>Physical Biology</i> , 2012 , 9, 026010	3	64
198	Computational models of VEGF-associated angiogenic processes in cancer. <i>Mathematical Medicine and Biology</i> , 2012 , 29, 85-94	1.3	18
197	Constructing the angiome: a global angiogenesis protein interaction network. <i>Physiological Genomics</i> , 2012 , 44, 915-24	3.6	25
196	Collagen IV and CXC chemokine-derived antiangiogenic peptides suppress glioma xenograft growth. <i>Anti-Cancer Drugs</i> , 2012 , 23, 706-12	2.4	14
195	Effects of fiber type and size on the heterogeneity of oxygen distribution in exercising skeletal muscle. <i>PLoS ONE</i> , 2012 , 7, e44375	3.7	26
194	Expression of VEGF receptors on endothelial cells in mouse skeletal muscle. <i>PLoS ONE</i> , 2012 , 7, e44791	3.7	45
193	Applications of Network Bioinformatics to Cancer Angiogenesis		1
192	Simulating Therapeutics Using Multiscale Models of the VEGF Receptor System in Cancer		2
191	Novel peptide-specific quantitative structure-activity relationship (QSAR) analysis applied to collagen IV peptides with antiangiogenic activity. <i>Journal of Medicinal Chemistry</i> , 2011 , 54, 6492-500	8.3	17
190	Small peptides derived from somatotropin domain-containing proteins inhibit blood and lymphatic endothelial cell proliferation, migration, adhesion and tube formation. <i>International Journal of Biochemistry and Cell Biology</i> , 2011 , 43, 1812-21	5.6	19
189	Analysis of VEGF--a regulated gene expression in endothelial cells to identify genes linked to angiogenesis. <i>PLoS ONE</i> , 2011 , 6, e24887	3.7	21

188	Theoretical models for coronary vascular biomechanics: progress & challenges. <i>Progress in Biophysics and Molecular Biology</i> , 2011 , 104, 49-76	4.7	51
187	Quantification and cell-to-cell variation of vascular endothelial growth factor receptors. <i>Experimental Cell Research</i> , 2011 , 317, 955-65	4.2	76
186	Angiogenesis-associated crosstalk between collagens, CXC chemokines, and thrombospondin domain-containing proteins. <i>Annals of Biomedical Engineering</i> , 2011 , 39, 2213-22	4.7	22
185	Module-based multiscale simulation of angiogenesis in skeletal muscle. <i>Theoretical Biology and Medical Modelling</i> , 2011 , 8, 6	2.3	53
184	Pharmacokinetics and pharmacodynamics of VEGF-neutralizing antibodies. <i>BMC Systems Biology</i> , 2011 , 5, 193	3.5	51
183	Formation of VEGF isoform-specific spatial distributions governing angiogenesis: computational analysis. <i>BMC Systems Biology</i> , 2011 , 5, 59	3.5	52
182	Development of a biomimetic peptide derived from collagen IV with anti-angiogenic activity in breast cancer. <i>Cancer Biology and Therapy</i> , 2011 , 12, 808-17	4.6	12
181	A two-compartment model of VEGF distribution in the mouse. <i>PLoS ONE</i> , 2011 , 6, e27514	3.7	29
180	Anti-angiogenic peptides for cancer therapeutics. <i>Current Pharmaceutical Biotechnology</i> , 2011 , 12, 1101-16		119
179	Blood flow and cell-free layer in microvessels. <i>Microcirculation</i> , 2010 , 17, 615-28	2.9	168
178	Quantifying the proteolytic release of extracellular matrix-sequestered VEGF with a computational model. <i>PLoS ONE</i> , 2010 , 5, e11860	3.7	59
177	VEGF and soluble VEGF receptor-1 (sFlt-1) distributions in peripheral arterial disease: an in silico model. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2010 , 298, H2174-91	5.2	33
176	Increase of plasma VEGF after intravenous administration of bevacizumab is predicted by a pharmacokinetic model. <i>Cancer Research</i> , 2010 , 70, 9886-94	10.1	65
175	A systems biology perspective on sVEGFR1: its biological function, pathogenic role and therapeutic use. <i>Journal of Cellular and Molecular Medicine</i> , 2010 , 14, 528-52	5.6	135
174	In Silico Modeling of Angiogenesis at Multiple Scales: From Nanoscale to Organ System 2010 , 287-320		2
173	Pentastatin-1, a collagen IV derived 20-mer peptide, suppresses tumor growth in a small cell lung cancer xenograft model. <i>BMC Cancer</i> , 2010 , 10, 29	4.8	37
172	Systems biology of pro-angiogenic therapies targeting the VEGF system. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , 2010 , 2, 694-707	6.6	70
171	Gene therapy from the perspective of systems biology. <i>Current Opinion in Molecular Therapeutics</i> , 2010 , 12, 570-7		22

170	Computational models of personalized medicine: leading indicators for angiogenesis therapies. <i>FASEB Journal</i> , 2010 , 24, 593.8	0.9	
169	A peptide derived from type 1 thrombospondin repeat-containing protein WISP-1 inhibits corneal and choroidal neovascularization 2009 , 50, 3840-5		17
168	A compartment model of VEGF distribution in humans in the presence of soluble VEGF receptor-1 acting as a ligand trap. <i>PLoS ONE</i> , 2009 , 4, e5108	3.7	54
167	Computational analysis of the tether-pulling experiment to probe plasma membrane-cytoskeleton interaction in cells. <i>Physical Review E</i> , 2009 , 80, 041905	2.4	13
166	Computational kinetic model of VEGF trapping by soluble VEGF receptor-1: effects of transendothelial and lymphatic macromolecular transport. <i>Physiological Genomics</i> , 2009 , 38, 29-41	3.6	17
165	The presence of VEGF receptors on the luminal surface of endothelial cells affects VEGF distribution and VEGF signaling. <i>PLoS Computational Biology</i> , 2009 , 5, e1000622	5	45
164	Computational fluid dynamics of aggregating red blood cells in postcapillary venules. <i>Computer Methods in Biomechanics and Biomedical Engineering</i> , 2009 , 12, 385-97	2.1	3
163	Elongation, proliferation & migration differentiate endothelial cell phenotypes and determine capillary sprouting. <i>BMC Systems Biology</i> , 2009 , 3, 13	3.5	93
162	Multiscale models of angiogenesis. <i>IEEE Engineering in Medicine and Biology Magazine</i> , 2009 , 28, 14-31		116
161	Modeling of growth factor-receptor systems from molecular-level protein interaction networks to whole-body compartment models. <i>Methods in Enzymology</i> , 2009 , 467, 461-497	1.7	13
160	Effects of erythrocyte deformability and aggregation on the cell free layer and apparent viscosity of microscopic blood flows. <i>Microvascular Research</i> , 2009 , 77, 265-72	3.7	157
159	Hemorrhagic shock and nitric oxide release from erythrocytic nitric oxide synthase: a quantitative analysis. <i>Microvascular Research</i> , 2009 , 78, 107-18	3.7	15
158	Peptides derived from type IV collagen, CXC chemokines, and thrombospondin-1 domain-containing proteins inhibit neovascularization and suppress tumor growth in MDA-MB-231 breast cancer xenografts. <i>Neoplasia</i> , 2009 , 11, 1285-91	6.4	47
157	Nitric oxide production pathways in erythrocytes and plasma. <i>Biorheology</i> , 2009 , 46, 107-19	1.7	26
156	Integration of angiogenesis modules at multiple scales: from molecular to tissue. <i>Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing</i> , 2009 , 316-27	1.3	13
155	Investigating transient and prolonged VEGF signaling through regulation of intracellular calcium. <i>FASEB Journal</i> , 2009 , 23, 767.15	0.9	
154	Pro-angiogenic and anti-angiogenic therapies targeting the VEGF-VEGFR system. <i>FASEB Journal</i> , 2009 , 23, 592.27	0.9	
153	Modeling Skeletal Muscle Angiogenesis from the Molecular to the Tissue Level. <i>FASEB Journal</i> , 2009 , 23, 592.24	0.9	

152	Effects of an anti-VEGF administration on whole-body VEGF distribution assessed by a molecularly-detailed pharmacokinetic model: a cancer study. <i>FASEB Journal</i> , 2009 , 23, 592.22	0.9	
151	A Systems Model of HIF1 α Dynamics in Ischemia. <i>FASEB Journal</i> , 2009 , 23, 767.6	0.9	
150	Experimental and theoretical studies of oxygen gradients in rat pial microvessels. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 2008 , 28, 1597-604	7.3	45
149	A compartment model of VEGF distribution in blood, healthy and diseased tissues. <i>BMC Systems Biology</i> , 2008 , 2, 77	3.5	99
148	Nitric oxide in the vasculature: where does it come from and where does it go? A quantitative perspective. <i>Antioxidants and Redox Signaling</i> , 2008 , 10, 1185-98	8.4	137
147	Nitric oxide from nitrite reduction by hemoglobin in the plasma and erythrocytes. <i>Nitric Oxide - Biology and Chemistry</i> , 2008 , 18, 47-60	5	53
146	Modeling the mechanics of tethers pulled from the cochlear outer hair cell membrane. <i>Journal of Biomechanical Engineering</i> , 2008 , 130, 031007	2.1	8
145	Trans-scleral delivery of antiangiogenic proteins. <i>Journal of Ocular Pharmacology and Therapeutics</i> , 2008 , 24, 70-9	2.6	27
144	A systematic methodology for proteome-wide identification of peptides inhibiting the proliferation and migration of endothelial cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 13775-80	11.5	61
143	INTEGRATION OF ANGIOGENESIS MODULES AT MULTIPLE SCALES: FROM MOLECULAR TO TISSUE 2008 ,		3
142	Novel anti-angiogenic peptides derived from ELR-containing CXC chemokines. <i>Journal of Cellular Biochemistry</i> , 2008 , 104, 1356-63	4.7	16
141	Red blood cell aggregation and dissociation in shear flows simulated by lattice Boltzmann method. <i>Journal of Biomechanics</i> , 2008 , 41, 47-55	2.9	197
140	Systems biology of vascular endothelial growth factors. <i>Microcirculation</i> , 2008 , 15, 715-38	2.9	118
139	Reactive oxygen species regulate hypoxia-inducible factor 1 α differentially in cancer and ischemia. <i>Molecular and Cellular Biology</i> , 2008 , 28, 5106-19	4.8	138
138	Nitric oxide release from nitric oxide synthase on erythrocytes during hemorrhagic shock. <i>FASEB Journal</i> , 2008 , 22, 749.7	0.9	
137	Towards a systematic design of novel endogenous peptide-based antiangiogenic approaches. <i>FASEB Journal</i> , 2008 , 22, 925.15	0.9	
136	An immersed boundary lattice Boltzmann approach to simulate deformable liquid capsules and its application to microscopic blood flows. <i>Physical Biology</i> , 2007 , 4, 285-95	3	139
135	Application of Chimera grid to modelling cell motion and aggregation in a narrow tube. <i>International Journal for Numerical Methods in Fluids</i> , 2007 , 53, 105-128	1.9	24

134	Dimerization of VEGF receptors and implications for signal transduction: a computational study. <i>Biophysical Chemistry</i> , 2007 , 128, 125-39	3.5	90
133	A computational model of oxygen delivery by hemoglobin-based oxygen carriers in three-dimensional microvascular networks. <i>Journal of Theoretical Biology</i> , 2007 , 248, 657-74	2.3	45
132	Where is VEGF in the body? A meta-analysis of VEGF distribution in cancer. <i>British Journal of Cancer</i> , 2007 , 97, 978-85	8.7	197
131	Three autocrine feedback loops determine HIF1 alpha expression in chronic hypoxia. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2007 , 1773, 1511-25	4.9	26
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