

Fabio Stroppa

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8446935/publications.pdf>

Version: 2024-02-01

15
papers

161
citations

1937685

4
h-index

1372567

10
g-index

17
all docs

17
docs citations

17
times ranked

229
citing authors

#	ARTICLE	IF	CITATIONS
1	Kids (learn how to) save lives in the school with the serious game Relive. Resuscitation, 2017, 116, 27-32.	3.0	54
2	Fall detection in indoor environment with kinect sensor. , 2014, , .		39
3	RELIVE: A Markerless Assistant for CPR Training. IEEE Transactions on Human-Machine Systems, 2016, 46, 755-760.	3.5	10
4	Online adaptive assistance control in robot-based neurorehabilitation therapy. , 2017, 2017, 628-633.		7
5	Real-Time 3D Tracker in Robot-Based Neurorehabilitation. , 2018, , 75-104.		6
6	A Robust Real-Time 3D Tracking Approach for Assisted Object Grasping. Lecture Notes in Computer Science, 2014, , 400-408.	1.3	6
7	Convex polygon fitting in robot-based neurorehabilitation. Applied Soft Computing Journal, 2018, 68, 609-625.	7.2	5
8	Upper Limb Joint Angular Velocity Synergies of Human Reaching Movements. , 2018, , .		4
9	An Improved Adaptive Robotic Assistance Methodology for Upper-Limb Rehabilitation. Lecture Notes in Computer Science, 2018, , 513-525.	1.3	4
10	EasyCluster2: an improved tool for clustering and assembling long transcriptome reads. BMC Bioinformatics, 2014, 15, S7.	2.6	3
11	RELIVE Tracking for quality cardiopulmonary resuscitation training: An experimental comparison with a standard CPR training mannequin. Resuscitation, 2015, 93, e1-e2.	3.0	2
12	Teleoperated Bilateral-Arm Rehabilitation with ALEx Rehab Station. Biosystems and Biorobotics, 2019, , 185-189.	0.3	2
13	A Novel Approach to Clustering and Assembly of Large-Scale Roche 454 Transcriptome Data for Gene Validation and Alternative Splicing Analysis. Lecture Notes in Computer Science, 2012, , 641-648.	1.3	2
14	Let's play Relive! Young people may learn how to save lives with a serious game. Resuscitation, 2016, 106, e44.	3.0	1
15	Clustering and Assembling Large Transcriptome Datasets by EasyCluster2. Communications in Computer and Information Science, 2013, , 231-236.	0.5	1