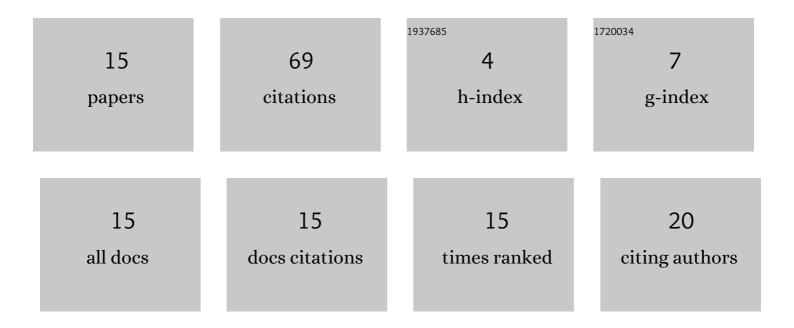
## T Galluccio

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9569717/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Identification of the novel <scp>HLAâ€DPA1</scp> allele, <scp><i>DPA1</i></scp> * <i>02:53</i> by nextâ€generation sequencing. Hla, 2022, 99, 149-150.	0.6	3
2	Identification of the novel <scp>HLAâ€B</scp> allele, <scp><i>HLAâ€B*44:532</i></scp> by nextâ€generation sequencing. Hla, 2022, 99, 210-211.	0.6	3
3	Characterization of the novel <i>HLAâ€DQA1*05:49</i> allele by sequencingâ€based typing. Hla, 2022, 99, 140-141.	0.6	3
4	Characterization of the novel <scp><i>HLAâ€DPA1*01:42</i></scp> allele by sequencingâ€based typing. Hla, 2021, 97, 93-94.	0.6	3
5	<i>HLA-Cw6</i> and other HLA-C alleles, as well as <i>MICB-DT, DDX58,</i> and <i>TYK2</i> genetic variants associate with optimal response to anti-IL-17A treatment in patients with psoriasis. Expert Opinion on Biological Therapy, 2021, 21, 259-270.	3.1	22
6	Identification of the novel <i><scp>HLAâ€DPB1</scp>*1149:01</i> . Hla, 2021, 97, 468-469.	0.6	3
7	Identification of the novel HLAâ€DRB1*03:01:32 in an Italian patient. Hla, 2021, 98, 179-180.	0.6	4
8	Identification of the novel <scp>HLAâ€DPA1</scp> , <i><scp>DPA1</scp>*01:56</i> by nextâ€generation sequencing. Hla, 2021, 98, 502-503.	0.6	3
9	Characterization of the novel <i>HLAâ€DPA1*02:26</i> allele by sequencingâ€based typing. Hla, 2020, 95, 160-161.	0.6	2
10	Characterization of the novel HLAâ€ÐRB3*01:86 allele by sequencingâ€based typing. Hla, 2020, 96, 535-537.	0.6	6
11	Characterization of the novel <scp><i>HLAâ€DQA1*03:15</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 236-237.	0.6	6
12	Characterization of the novel <scp><i>HLAâ€DRB3*02:142</i></scp> allele by sequencingâ€based typing. Hla, 2020, 95, 581-582.	0.6	2
13	Characterization of the novel <i>HLAâ€DQA1*01:25</i> allele by sequencingâ€based typing. Hla, 2019, 94, 174-175.	0.6	2
14	Characterization of the novel <i>HLAâ€DRB3*03:01:07</i> allele by sequencingâ€based typing. Hla, 2019, 93, 240-241.	0.6	2
15	Identification of the novel HLAâ€A*01:289 allele in an Italian patient. Hla, 2019, 93, 484-485.	0.6	5