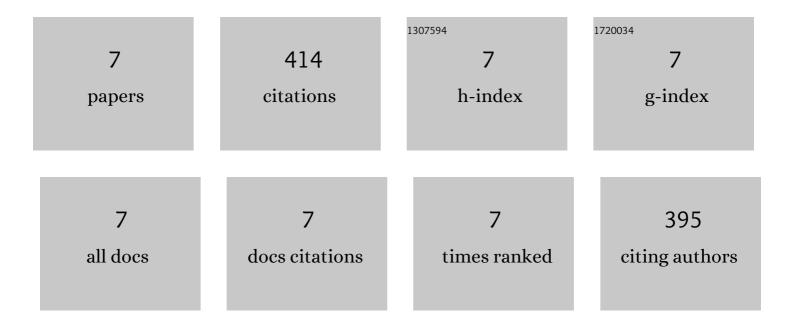
## Issa Faye

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9707036/publications.pdf Version: 2024-02-01



ISSA FAVE

#	Article	IF	CITATIONS
1	Fostered and left behind alleles in peanut: interspecific QTL mapping reveals footprints of domestication and useful natural variation for breeding. BMC Plant Biology, 2012, 12, 26.	3.6	114
2	An International Reference Consensus Genetic Map with 897 Marker Loci Based on 11 Mapping Populations for Tetraploid Groundnut (Arachis hypogaea L.). PLoS ONE, 2012, 7, e41213.	2.5	89
3	Construction of Chromosome Segment Substitution Lines in Peanut (Arachis hypogaea L.) Using a Wild Synthetic and QTL Mapping for Plant Morphology. PLoS ONE, 2012, 7, e48642.	2.5	85
4	Translational genomics for achieving higher genetic gains in groundnut. Theoretical and Applied Genetics, 2020, 133, 1679-1702.	3.6	58
5	Identification of quantitative trait loci for yield and yield related traits in groundnut (Arachis) Tj ETQq1 1 0.7843	14 rgBT /C	overlock 10 Tf
6	Improved Genetic Map Identified Major QTLs for Drought Tolerance- and Iron Deficiency Tolerance-Related Traits in Groundnut. Genes, 2021, 12, 37.	2.4	28
7	Identification of Sources of Resistance for Peanut <i>Aspergillus flavus</i> Colonization and Aflatoxin Contamination. International Journal of Agronomy, 2018, 2018, 1-7.	1.2	11