Falk Zakrzewski

List of Publications by Year
in descending order

11	825	933447 10	1281871 11
papers	citations	h-index	g-index

12	12	12	1253
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	The genome of the recently domesticated crop plant sugar beet (Beta vulgaris). Nature, 2014, 505, 546-549.	27.8	569
2	DNA methylation of retrotransposons, DNA transposons and genes in sugar beet (Beta vulgaris) Tj ETQq0 0 0 rgBT /Overlock 10 T	5.7	50
3	Repeat Composition of CenH3-chromatin and H3K9me2-marked heterochromatin in Sugar Beet (Beta) Tj ETQq1 1 0.784314 rgBT /Over	3.6	32
4	Epigenetic profiling of heterochromatic satellite DNA. Chromosoma, 2011, 120, 409-422.	2.2	31
5	Diversification, evolution and methylation of short interspersed nuclear element families in sugar beet and related Amaranthaceae species. Plant Journal, 2016, 85, 229-244.	5.7	29
6	Analysis of a c0t-1 library enables the targeted identification of minisatellite and satellite families in Beta vulgaris. BMC Plant Biology, 2010, 10, 8.	3.6	28
7	Next-generation sequencing reveals differentially amplified tandem repeats as a major genome component of Northern Europeâ€™s oldest Camellia japonica. Chromosome Research, 2015, 23, 791-806.	2.2	24
8	Comparative molecular cytogenetic analyses of a major tandemly repeated DNA family and retrotransposon sequences in cultivated jute Corchorus species (Malvaceae). Annals of Botany, 2013, 112, 123-134.	2.9	23
9	The <scp>CHH</scp> motif in sugar beet satellite <scp>DNA</scp>: a modulator for cytosine methylation. Plant Journal, 2014, 78, 937-950.	5.7	17
10	Cytosine Methylation of an Ancient Satellite Family in the Wild Beet <i>Beta procumbens</i>. Cytogenetic and Genome Research, 2014, 143, 157-167.	1.1	11
11	Epigenetic Characterization of Satellite DNA in Sugar Beet (Beta vulgaris). RNA Technologies, 2017, , 445-462.	0.3	1