## John W Snape

List of Publications by Year in descending order

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IOHN W SNADE

#	Article	IF	CITATIONS
1	Genetic variation in wheat grain quality is associated with differences in the galactolipid content of flour and the gas bubble properties of dough liquor. Food Chemistry: X, 2020, 6, 100093.	4.3	12
2	Natural Selection Towards Wild-Type in Composite Cross Populations of Winter Wheat. Frontiers in Plant Science, 2019, 10, 1757.	3.6	15
3	Increased pericarp cell length underlies a major quantitative trait locus for grain weight in hexaploid wheat. New Phytologist, 2017, 215, 1026-1038.	7.3	103
4	Mapping of quantitative trait loci for root hair length in wheat identifies loci that co-locate with loci for yield components. Journal of Experimental Botany, 2016, 67, 4535-4543.	4.8	35
5	Comparative analysis of performance and stability among composite cross populations, variety mixtures and pure lines of winter wheat in organic and conventional cropping systems. Field Crops Research, 2015, 183, 235-245.	5.1	77
6	Genetic Dissection of Grain Size and Grain Number Trade-Offs in CIMMYT Wheat Germplasm. PLoS ONE, 2015, 10, e0118847.	2.5	88
7	Identification and independent validation of a stable yield and thousand grain weight QTL on chromosome 6A of hexaploid wheat (Triticum aestivum L.). BMC Plant Biology, 2014, 14, 191.	3.6	161
8	Molecular marker-based characterization of a set of wheat genotypes adapted to Central Europe. Cereal Research Communications, 2014, 42, 189-198.	1.6	4
9	Analysis of the Genetic Structure of a Barley Collection Using DNA Diversity Array Technology (DArT). Plant Molecular Biology Reporter, 2013, 31, 280-288.	1.8	17
10	Exploitation of interspecific diversity for monocot crop improvement. Heredity, 2013, 110, 475-483.	2.6	39
11	Identifying wheat genomic regions for improving grain protein concentration independently of grain yield using multiple inter-related populations. Molecular Breeding, 2013, 31, 587-599.	2.1	49
12	Control of late maturity alpha-amylase in wheat by the dwarfing gene Rht-D1b and genes on the 1B/1R translocation. Molecular Breeding, 2013, 32, 425-436.	2.1	23
13	Effect of wheat dwarfing genes on nitrogen-use efficiency. Journal of Agricultural Science, 2012, 150, 3-22.	1.3	66
14	Location of genes for common bunt resistance in the European winter wheat cv. Trintella. Euphytica, 2012, 186, 257-264.	1.2	20
15	Simulation of environmental and genotypic variations of final leaf number and anthesis date for wheat. European Journal of Agronomy, 2012, 42, 22-33.	4.1	56
16	Meta-QTL analysis of the genetic control of crop height in elite European winter wheat germplasm. Molecular Breeding, 2012, 29, 159-171.	2.1	127
17	Evidence of selective changes in winter wheat in middle-European environments reflected by allelic diversity at loci affecting plant height and photoperiodic response. Journal of Agricultural Science, 2011, 149, 313-326.	1.3	5
18	Relationships between carbon isotope discrimination and grain yield in winter wheat under well-watered and drought conditions. Journal of Agricultural Science, 2011, 149, 257-272.	1.3	22

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19	Identification of traits to improve the nitrogen-use efficiency of wheat genotypes. Field Crops Research, 2011, 123, 139-152.	5.1	243
20	Anthesis date mainly explained correlations between post-anthesis leaf senescence, grain yield, and grain protein concentration in a winter wheat population segregating for flowering time QTLs. Journal of Experimental Botany, 2011, 62, 3621-3636.	4.8	193
21	An SSR-based genetic linkage map of the model grass Brachypodium distachyon. Genome, 2010, 53, 1-13.	2.0	55
22	Effects of reduced height (Rht) and photoperiod insensitivity (Ppd) alleles on yield of wheat in contrasting production systems. Euphytica, 2010, 172, 169-181.	1.2	42
23	Effects of specific Rht and Ppd alleles on agronomic traits in winter wheat cultivars grown in middle Europe. Euphytica, 2010, 172, 221-233.	1.2	14
24	Mapping quantitative trait loci for resistance against Russian wheat aphid (Diuraphis noxia) in wheat (Triticum aestivum L.). Crop and Pasture Science, 2010, 61, 970.	1.5	17
25	A Genetic Framework for Grain Size and Shape Variation in Wheat Â. Plant Cell, 2010, 22, 1046-1056.	6.6	397
26	Raising yield potential in wheat. Journal of Experimental Botany, 2009, 60, 1899-1918.	4.8	508
27	Wheat Antixenosis, Antibiosis, and Tolerance to Infestation by <i>Delphacodes kuscheli</i> (Hemiptera: Delphacidae), a Vector of "Mal de RÃo Cuarto―in Argentina. Journal of Economic Entomology, 2009, 102, 1801-1807.	1.8	0
28	Reduced height (Rht) and photoperiod insensitivity (Ppd) allele associations with establishment and early growth of wheat in contrasting production systems. Euphytica, 2009, 166, 249.	1.2	60
29	Inheritance and QTL mapping of leaf rust resistance in the European winter wheat cultivar †Beaver'. Euphytica, 2009, 169, 253-261.	1.2	23
30	A consensus map of rye integrating mapping data from five mapping populations. Theoretical and Applied Genetics, 2009, 118, 793-800.	3.6	46
31	Meta-QTL analysis of the genetic control of ear emergence in elite European winter wheat germplasm. Theoretical and Applied Genetics, 2009, 119, 383-395.	3.6	225
32	Development of consistently crossable wheat genotypes for alien wheat gene transfer through fine-mapping of the Kr1 locus. Theoretical and Applied Genetics, 2009, 119, 1371-1381.	3.6	13
33	A protocol for Agrobacterium-mediated transformation of Brachypodium distachyon community standard line Bd21. Nature Protocols, 2009, 4, 638-649.	12.0	129
34	A novel transcriptomic approach to identify candidate genes for grain quality traits in wheat. Plant Biotechnology Journal, 2009, 7, 401-410.	8.3	18
35	Intronâ€mediated enhancement as a method for increasing transgene expression levels in barley. Plant Biotechnology Journal, 2009, 7, 856-866	8.3	38
36	Barley Transformation Using Agrobacterium-Mediated Techniques. Methods in Molecular Biology, 2009, 478, 137-147.	0.9	78

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37	Mapping of a gene (Vir) for a non-glaucous, viridescent phenotype in bread wheat derived from Triticum dicoccoides, and its association with yield variation. Euphytica, 2008, 159, 333-341.	1.2	32
38	Mapping quantitative trait loci for growth responses to exogenously applied stress induced hormones in wheat. Euphytica, 2008, 164, 719-727.	1.2	11
39	Genetic mapping of a new flowering time gene on chromosome 3B of wheat. Euphytica, 2008, 164, 779-787.	1.2	15
40	Susceptibility to Fusarium head blight is associated with the Rht-D1b semi-dwarfing allele in wheat. Theoretical and Applied Genetics, 2008, 116, 1145-1153.	3.6	101
41	Agrobacterium-mediated transformation of the temperate grass Brachypodium distachyon (genotype) Tj ETQq1 1	0.784314 8.3	⊦rgβT /Over
42	Corrigendum. Plant Biotechnology Journal, 2008, 6, 941-941.	8.3	3
43	Effects of drought and the presence of the 1BL/1RS translocation on grain vitreosity, hardness and protein content in winter wheat. Journal of Cereal Science, 2008, 47, 457-468.	3.7	50
44	Identification and genetic mapping of variant forms of puroindoline b expressed in developing wheat grain. Journal of Cereal Science, 2008, 48, 722-728.	3.7	51
45	High-throughput Agrobacterium-mediated barley transformation. Plant Methods, 2008, 4, 22.	4.3	163
46	PAPER PRESENTED AT INTERNATIONAL WORKSHOP ON INCREASING WHEAT YIELD POTENTIAL, CIMMYT, OBREGON, MEXICO, 20–24 MARCH 2006 Genetic progress in yield potential in wheat: recent advances and future prospects. Journal of Agricultural Science, 2007, 145, 17-29.	1.3	136
47	The pCLEAN Dual Binary Vector System for <i>Agrobacterium</i> -Mediated Plant Transformation. Plant Physiology, 2007, 145, 1211-1219.	4.8	69
48	Identifying physiological traits associated with improved drought resistance in winter wheat. Field Crops Research, 2007, 103, 11-24.	5.1	213
49	A Pseudo-Response Regulator is misexpressed in the photoperiod insensitive Ppd-D1a mutant of wheat (Triticum aestivum L.). Theoretical and Applied Genetics, 2007, 115, 721-733.	3.6	691
50	Dissecting geneÂ×Âenvironmental effects on wheat yields via QTL and physiological analysis. Euphytica, 2007, 154, 401-408.	1.2	125
51	Development and genetic mapping of sequence-tagged microsatellites (STMs) in bread wheat (Triticum) Tj ETQq1	1.0.7843 3.6	14 rgBT /O∨
52	Major Genetic Changes in Wheat with Potential to Affect Disease Tolerance. Phytopathology, 2006, 96, 680-688.	2.2	34
53	Identification and characterization of quantitative trait loci related to lodging resistance and associated traits in bread wheat. Plant Breeding, 2005, 124, 234-241.	1.9	70
54	Mapping antixenosis genes on chromosome 6A of wheat to greenbug and to a new biotype of Russian wheat aphid. Plant Breeding, 2005, 124, 229-233.	1.9	33

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55	A comparison of transgenic barley lines produced by particle bombardment and Agrobacterium-mediated techniques. Plant Cell Reports, 2005, 23, 780-789.	5.6	238
56	Deletion analysis of genes regulating cold- and PEG-induced carbohydrate accumulation in hydroponically raised wheat seedlings. Acta Agronomica Hungarica: an International Multidisciplinary Journal in Agricultural Science, 2005, 53, 359-370.	0.2	2
57	The Distribution of Transgene Insertion Sites in Barley Determined by Physical and Genetic Mapping. Genetics, 2004, 167, 1371-1379.	2.9	36
58	Strategies for precise quantification of transgene expression levels over several generations in rice. Journal of Experimental Botany, 2004, 55, 1307-1313.	4.8	13
59	Mapping quantitative trait loci in wheat for resistance against greenbug and Russian wheat aphid. Plant Breeding, 2004, 123, 361-365.	1.9	25
60	Development of a standard operating procedure (SOP) for the precise quantification of transgene expression levels in rice plants. Physiologia Plantarum, 2004, 120, 650-656.	5.2	6
61	Effects of a photoperiod-response gene Ppd-D1 on yield potential and drought resistance in UK winter wheat. Euphytica, 2004, 135, 63-73.	1.2	85
62	Mapping quantitative trait loci for flag leaf senescence as a yield determinant in winter wheat under optimal and drought-stressed environments. Euphytica, 2004, 135, 255-263.	1.2	217
63	The effect of additional virulence genes on transformation efficiency, transgene integration and expression in rice plants using the pGreen/pSoup dual binary vector system. Transgenic Research, 2004, 13, 593-603.	2.4	29
64	A new approach to extending the wheat marker pool by anchored PCR amplification of compound SSRs. Theoretical and Applied Genetics, 2004, 108, 733-742.	3.6	19
65	A large-scale study of rice plants transformed with different T-DNAs provides new insights into locus composition and T-DNA linkage configurations. Theoretical and Applied Genetics, 2004, 109, 815-826.	3.6	80
66	Transgene behaviour in populations of rice plants transformed using a new dual binary vector system: pGreen/pSoup. Theoretical and Applied Genetics, 2003, 107, 210-217.	3.6	74
67	Mapping genes affecting flowering time and frost resistance on chromosome 5B of wheat. Theoretical and Applied Genetics, 2003, 107, 509-514.	3.6	165
68	Developmental responses to vernalization in wheat deletion lines for chromosomes 5A and 5D. Plant Breeding, 2003, 122, 35-39.	1.9	15
69	Analysis of the expression patterns of the Arabidopsis thaliana tubulin-1 and Zea mays ubiquitin-1 promoters in rice plants in association with nematode infection. Physiological and Molecular Plant Pathology, 2002, 60, 197-205.	2.5	26
70	RFLP mapping of a Hordeum bulbosum gene highly expressed in pistils and its relationship to homoeologous loci in other Gramineae species. Theoretical and Applied Genetics, 2002, 105, 271-276.	3.6	6
71	Transgene behaviour across two generations in a large random population of transgenic rice plants produced by particle bombardment. Theoretical and Applied Genetics, 2002, 105, 878-889.	3.6	70
72	Utility of barley and wheat simple sequence repeat (SSR) markers for genetic analysis of Hordeum chilense and tritordeum. Theoretical and Applied Genetics, 2002, 104, 735-739.	3.6	39

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73	The relationship between homozygous and hemizygous transgene expression levels over generations in populations of transgenic rice plants. Theoretical and Applied Genetics, 2002, 104, 553-561.	3.6	62
74	Use of the firefly luciferase gene in a barley ( Hordeum vulgare ) transformation system. Plant Cell Reports, 2002, 21, 320-326.	5.6	17
75	Constructing plant radiation hybrid panels. Plant Journal, 2002, 31, 223-228.	5.7	41
76	A core genetic map of Hordeum chilense and comparisons with maps of barley (Hordeum vulgare) and wheat (Triticum aestivum). Theoretical and Applied Genetics, 2001, 102, 1259-1264.	3.6	63
77	Microsatellites and RFLP probes from maize are efficient sources of molecular markers for the biomass energy crop Miscanthus. Theoretical and Applied Genetics, 2001, 102, 616-622.	3.6	66
78	An RFLP map of diploid Hordeum bulbosum L. and comparison with maps of barley (H. vulgare L.) and wheat (Triticum aestivum L.). Theoretical and Applied Genetics, 2001, 103, 869-880.	3.6	26
79	Title is missing!. Euphytica, 2001, 119, 173-177.	1.2	11
80	Mapping genes for flowering time and frost tolerance in cereals using precise genetic stocks. Euphytica, 2001, 120, 309-315.	1.2	73
81	Title is missing!. Euphytica, 2001, 121, 265-271.	1.2	5
82	Title is missing!. Euphytica, 2001, 122, 309-317.	1.2	135
83	Title is missing!. Euphytica, 2000, 111, 67-76.	1.2	31
84	A skeletal linkage map of Hordeum bulbosum L. and comparative mapping with barley (H. vulgare L.). Euphytica, 2000, 115, 115-120.	1.2	9
85	Matrix attachment regions increase transgene expression levels and stability in transgenic rice plants and their progeny. Plant Journal, 1999, 18, 233-242.	5.7	93
86	â€~Green revolution' genes encode mutant gibberellin response modulators. Nature, 1999, 400, 256-261.	27.8	1,876
87	Luciferase as a reporter gene for transformation studies in rice ( Oryza sativa L.). Plant Cell Reports, 1999, 18, 715-720.	5.6	17
88	Physical mapping of the Vrn-A1 and Fr1 genes on chromosome 5A of wheat using deletion lines. Theoretical and Applied Genetics, 1999, 99, 199-202.	3.6	74
89	Induction and Characterization of Ph1 Wheat Mutants. Genetics, 1999, 153, 1909-1918.	2.9	132
90	The green fluorescent protein (GFP) as a vital screenable marker in rice transformation. Theoretical and Applied Genetics, 1998, 96, 164-169.	3.6	79

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91	Expression of an engineered cysteine proteinase inhibitor (Oryzacystatin-IΔD86) for nematode resistance in transgenic rice plants. Theoretical and Applied Genetics, 1998, 96, 266-271.	3.6	130
92	A simple PCR-based method for scoring the ph1b deletion in wheat. Theoretical and Applied Genetics, 1998, 96, 371-375.	3.6	54
93	Comparative mapping of the wheat chromosome 5A Vrn-A1 region with rice and its relationship to QTL for flowering time. Theoretical and Applied Genetics, 1998, 97, 103-109.	3.6	52
94	Title is missing!. Euphytica, 1997, 94, 335-340.	1.2	17
95	Comparative RFLP mapping of the chlorotoluron resistance gene (Su1) in cultivated wheat (Triticum) Tj ETQq1	1 0.784314	rgßT /Overlo
96	Location of a gene regulating cold-induced carbohydrate production on chromosome 5A of wheat. Theoretical and Applied Genetics, 1997, 95, 265-270.	3.6	54
97	The genetic and molecular characterization of pollen-derived plant lines from octoploid triticale � wheat hybrids. Theoretical and Applied Genetics, 1996, 92, 811-816.	3.6	3
98	QTL analysis: unreliability and bias in estimation procedures. Molecular Breeding, 1995, 1, 273-282.	2.1	85
99	Transformation studies in Hordeum vulgare using a highly regenerable microspore system. Euphytica, 1995, 85, 113-118.	1.2	30
100	Spontaneous wheat/rye translocations from female meiotic products of hybrids between octoploid triticale and wheat. Euphytica, 1995, 81, 265-270.	1.2	10
101	A comparison of male and female recombination frequency in wheat using RFLP maps of homoeologous group 6 and 7 chromosomes. Theoretical and Applied Genetics, 1995, 91, 744-746.	3.6	55
102	RFLP mapping of the vernalization (Vrn1) and frost resistance (Fr1) genes on chromosome 5A of wheat. Theoretical and Applied Genetics, 1995, 90, 1174-1179.	3.6	329
103	A partial genome assay for quantitative trait loci in wheat (Triticum aestivum) using different analytical techniques. Theoretical and Applied Genetics, 1994, 89, 735-741.	3.6	25
104	Genetic analysis of a photoperiod response gene on the short arm of chromosome 2(2H) of Hordeum vulgare (barley). Heredity, 1994, 72, 619-627.	2.6	139
105	Stability of transgenes and presence of N6 methyladenine DNA in transformed wheat cells. Plant Journal, 1994, 5, 429-436.	5.7	16
106	The genetic characterisation of novel multi-addition doubled haploid lines derived from triticale x wheat hybrids. Theoretical and Applied Genetics, 1993, 87, 531-536.	3.6	13
107	The high-molecular-weight glutenin subunit compositions of Chinese bread wheat varieties and their relationship with bread-making quality. Euphytica, 1993, 68, 205-212.	1.2	16
108	Assignment of the denso Dwarfing Gene to the Long Arm of Chromosome 3(3H) of Barley by Use of RFLP Markers. Plant Breeding, 1993, 111, 198-203.	1.9	61

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109	A Similar Metabolism of Chlorotoluron in Cell Suspension Cultures from Near-Isogenic Susceptible and Tolerant Lines of Wheat. Pesticide Biochemistry and Physiology, 1993, 47, 51-59.	3.6	4
110	The location and effects of genes modifying the response of wheat to the herbicide difenzoquat. Journal of Agricultural Science, 1992, 118, 9-15.	1.3	0
111	The relationship between in vitro performance of haploid embryos and the agronomic performance of the derived doubled haploid lines in barley. Theoretical and Applied Genetics, 1992, 84, 118-122.	3.6	6
112	Herbicide response polymorphisms in wild emmer wheat: ecological and isozyme correlations. Theoretical and Applied Genetics, 1992, 84, 209-216.	3.6	10
113	The location of major genes and associated quantitative trait loci on chromosome arm 5BL of wheat. Theoretical and Applied Genetics, 1992, 85-85, 197-204.	3.6	29
114	Herbicide response polymorphism in wild populations of emmer wheat. Heredity, 1991, 66, 251-257.	2.6	16
115	The cytological and genetic characterisation of doubled haploid lines derived from triticale×wheat hybrids. Theoretical and Applied Genetics, 1991, 81, 369-375.	3.6	8
116	The assessment of in vitro characters and their influence on the success rates of doubled haploid production in barley. Euphytica, 1991, 58, 137-144.	1.2	4
117	The agronomic performance of wheat doubled haploid lines derived from wheat x maize crosses. Theoretical and Applied Genetics, 1990, 79, 813-816.	3.6	21
118	The resistance of Hordeum bulbosum and its hybrids with H. vulgare to common fungal pathogens. Euphytica, 1989, 41, 273-276.	1.2	29
119	Location of a gene for frost resistance on chromosome 5A of wheat. Euphytica, 1989, 42, 41-44.	1.2	113
120	Genetic analysis of anther culture response in wheat using aneuploid, chromosome substitution and translocation lines. Theoretical and Applied Genetics, 1989, 77, 7-11.	3.6	97
121	Tests for the presence of gametoclonal variation in barley and wheat doubled haploids produced using the Hordeum bulbosum system. Theoretical and Applied Genetics, 1988, 75, 509-513.	3.6	34
122	The detection and estimation of linkage using doubled haploid or single seed descent populations. Theoretical and Applied Genetics, 1988, 76, 125-128.	3.6	15
123	Studies of the Genetic Relationship between Anther Culture and Somatic Tissue Culture Abilities in Wheat. Plant Breeding, 1988, 100, 26-33.	1.9	55
124	The chromosomal locations in wheat of genes conferring differential response to the wild oat herbicide, difenzoquat. Journal of Agricultural Science, 1987, 108, 543-548.	1.3	14
125	Factors affecting haploid production in wheat using the Hordeum bulbosum system. 1. Genotypic and environmental effects on pollen grain germination, pollen tube growth and the frequency of fertilization. Euphytica, 1987, 36, 483-496.	1.2	40
126	Factors affecting haploid production in wheat using the Hordeum bulbosum system. 2. The effect of the timing of pollination. Euphytica, 1987, 36, 497-504.	1.2	10

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127	Factors affecting haploid production in wheat using the Hordeum bulbosum system. 3. Post-fertilization effects on embryo survival. Euphytica, 1987, 36, 763-773.	1.2	9
128	The utilisation of doubled haploid lines in quantitative genetics. Bulletin De La Société Botanique De France Actualités Botaniques, 1986, 133, 59-66.	0.0	2
129	Doubled haploid production in winter wheat and triticale genotypes, using the Hordeum bulbosum system. Euphytica, 1986, 35, 1045-1051.	1.2	19
130	Genetical analysis of chromosome 5A of wheat and its influence on important agronomic characters. Theoretical and Applied Genetics, 1985, 71, 518-526.	3.6	113
131	Intrachromosomal mapping of crossability genes in wheat (Triticum aestivum). Theoretical and Applied Genetics, 1985, 70, 309-314.	3.6	78
132	Intrachromosomal mapping of the nucleolar organiser region relative to three marker loci on chromosome 1B of wheat (Triticum aestivum). Theoretical and Applied Genetics, 1985, 69, 263-270.	3.6	88
133	Methods for estimating gene numbers for quantitative characters using doubled haploid lines. Theoretical and Applied Genetics, 1984, 67, 143-148.	3.6	43
134	The use of irradiated pollen for differential gene transfer in wheat (Triticum aestivum). Theoretical and Applied Genetics, 1983, 65, 103-111.	3.6	45
135	Predicting the frequencies of transgressive segregants for yield and yield components in wheat. Theoretical and Applied Genetics, 1982, 62, 127-134.	3.6	29
136	The genetical expectations of doubled haploid lines derived from different filial generations. Theoretical and Applied Genetics, 1981, 60, 123-128.	3.6	66
137	The detection of homologous chromosome variation in wheat using backcross reciprocal monosomic lines. Heredity, 1980, 45, 187-200.	2.6	22
138	Genetical analysis of chromosome substitution lines of bread wheat using second generation hybrids. Heredity, 1979, 42, 247-258.	2.6	6
139	The crossabilities of wheat varieties with Hordeum bulbosum. Heredity, 1979, 42, 291-298.	2.6	157
140	The genetical relationship between height and yield in wheat. Heredity, 1978, 40, 133-151.	2.6	115
141	Whole chromosome analysis of height in wheat. Heredity, 1977, 38, 25-36.	2.6	36
142	Effects of linkage and interaction in a comparison of theoretical populations derived by diploidized haploid and single seed descent methods. Theoretical and Applied Genetics, 1977, 49, 111-115.	3.6	30
143	A theoretical comparison of diploidised haploid and single seed descent populations. Heredity, 1976, 36, 275-277.	2.6	38
144	Chromosome variation for loci controlling ear emergence time on chromosome 5A of wheat. Heredity, 1976, 37, 335-340.	2.6	44

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145	A method for the detection of epistasis in chromosome substitution lines of hexaploid wheat. Heredity, 1975, 34, 297-303.	2.6	9
146	Genetical consequences of single seed descent in the breeding of self-pollinating crops. Heredity, 1975, 35, 211-219.	2.6	41
147	The breeding system of Arabidopsis thaliana. Heredity, 1971, 27, 299-302.	2.6	49