Haploview: analysis and visualization of LD and haplotype maps

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218	Genome-wide association study of water use patterns of common bean (Phaseolus vulgaris L.) genotypes in response to drying soil.	0
217	Genomic footprints related with adaptation and fumonisins production in Fusarium proliferatum. 13,	0
216	A randomized, double-blind, placebo-controlled, pharmacogenetic study of ondansetron for treating alcohol use disorder.	0
215	QTL mapping and identification of candidate genes using a genome-wide association study for heat tolerance at anthesis in rice (Oryza sativa L.). 13,	1
214	Genomic surveillance unfolds the SARS-CoV-2 transmission and divergence dynamics in Bangladesh. 13,	1
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196	QTL Mapping of Stem Rust Resistance in Populations of Durum Wheat. **2022**, 13, 1793	0
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138	SNP - and haplotype-based single-step genomic predictions for body weight, wool, and reproductive traits in North American Rambouillet sheep.	0
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68	Targeted next-generation sequencing of genes involved in Warfarin Pharmacodynamics and pharmacokinetics pathways using the Saudi Warfarin Pharmacogenetic study (SWAP).	0
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60	Identification of QTN-by-environment interactions for yield related traits in maize under multiple abiotic stresses. 14,	0
59	Branch angle and leaflet shape are associated with canopy coverage in soybean.	0

58	Cytokinin response regulatorARR16regulates seed coat permeability in Arabidopsis natural accessions.	0
57	A new allele PEL9GG identified by genome-wide association study increases panicle elongation length in rice (Oryza sativa L.). 14,	0
56	Evidence for Epistatic Interaction between HLA-G and LILRB1 in the Pathogenesis of Nonsegmental Vitiligo. **2023**, 12, 630	1
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53	The influence of six polymorphisms of uncoupling protein 3 (UCP3) gene and childhood obesity: a case□ontrol study. **2023**, 23,	0
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48	Integrated Single-Trait and Multi-Trait GWASs Reveal the Genetic Architecture of Internal Organ Weight in Pigs. **2023**, 13, 808	0
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42	Gene-wide significant association analyses of DNMT1 genetic variants with Parkinson□ disease. 14,	0
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37	Cold-induced vasodilation response in a Japanese cohort: insights from cold-water immersion and genome-wide association studies. **2023**, 42,	0
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34	Genomic architecture of leaf senescence in sorghum (Sorghum bicolor). **2023**, 136,	0
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31	Pilot Study on the Impact of Polymorphisms Linked to Multi-Kinase Inhibitor Metabolism on Lenvatinib Side Effects in Patients with Advanced Thyroid Cancer. **2023**, 24, 5496	0
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29	Major histocompatibility complex class I-related chain A and macrophage migration inhibitory factor gene polymorphisms in a Turkish patient population with vitiligo. **2023**, 17, 11	0
28	A multi-locus linear mixed model methodology for detecting small-effect QTLs for quantitative traits in MAGIC, NAM, and ROAM populations. **2023**, 21, 2241-2252	0
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25	Association of CTNND2 gene polymorphism with schizophrenia: Two-sample case-control study in Chinese Han population. 009121742311646	0
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19	Single Nucleotide Polymorphisms in the Vitamin D Metabolic Pathway and Their Relationship with High Blood Pressure Risk. **2023**, 24, 5974	0
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17	Exploring haplotype block structure, runs of homozygosity, and effective population size among dairy cattle breeds of India. **2023**, 55,	0
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