

Comparative Mapping Of Genes For Plant Disease Resistance In Wheat 'advanced Backcross' Populations By Means Of DNA Markers (Schriftenreihe Des Instituts ... Und Ressourcenschutz) By Ali Ahmad Naz

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How to Cite. Suchyta, S. P. , Cheng, H. H. , Burnside, J. and Dodgson, J. B. (2001), Comparative mapping of chicken anchor loci orthologous to genes on human <http://onlinelibrary.wiley.com/doi/10.1046/j.1365-2052.2001.00717.x/citedby>

Aleyasin and Barendse x Bovine Chromosome 5 Comparative Mapping 539 Table 3. Sequence of SLC2A3 in cattle and sheep bovSLC2A3 <http://jhered.oxfordjournals.org/content/90/5/537.full.pdf>

Motivation: The completion of human and mouse genome sequences provides a valuable resource for decoding other mammalian genomes. The comparative mapping by <http://citeseerx.ist.psu.edu/showciting?cid=5380966>

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Comparative mapping of HKT genes in wheat, barley, and rice, key determinants of Na⁺ transport, and salt tolerance. Huang S, Spielmeyer W, Lagudah ES, Munns R.

<http://www.ncbi.nlm.nih.gov/pubmed/18325922>

Abstract The dominant gene Pvr7 from Capsicum chin-ense Jacq. PI159236 confers resistance to the pepper mottle potyvirus (PepMoV) Florida (V1182) strain.

<http://link.springer.com/content/pdf/10.1007/s001220051552.pdf>

Ali Ahmad Naz Comparative mapping of genes for plant disease resistance in wheat 'advanced backcross' populations by means of DNA markers: ISBN: 978-3-8322-6183-2

<http://www.shaker.de/de/content/catalogue/index.asp?ID=8&ISBN=978-3-8322-6183-2>

Comparative genomics is a field of biological research in which the genomic features of by known genome mapping information to locate other genes in the

http://en.wikipedia.org/wiki/Comparative_genomics

COMPARATIVE GENOME MAPPING Molecular genetic linkage map based on tomato cDNA, genomic DNA, and EST markers was constructed for eggplant, Solanum melongena.

<http://www.public.iastate.edu/~mbhattac/bhattacharyya/COMPARATIVE%20GENOME%20MAPPING.pdf>

Institut für Nutzpflanzenwissenschaften und Ressourcenschutz (INRES) Plant Dr. Ali Ahmad Naz L on J (2012) Advanced backcross quantitative trait locus

http://www.imbio.uni-bonn.de/cv_publications_naz

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<http://www.amazon.co.jp/Comparative-Populations-Schriftenreihe-Nutzpflanzenwissenschaften-Ressourcenschutz/dp/3832261834>

species. Usually it is practiced by mapping genes in one species and then comparing Human Comparative Maps , as shown on the next page.

http://www-personal.une.edu.au/~jvanderw/14_Candidate_genes.PDF

Combining mouse mammary gland gene expression and comparative mapping for the candidate genes underlying QTL. This is a comparative

<http://www.biomedcentral.com/1471-2164/8/183>

Gene Expression; Maps; Web Resources. Genomics; Germplasm; Mapping; Pathology; Publications; The Triticeae Toolbox; U.S. Wheat and Barley Scab Initiative; Genomics

<http://wheat.pw.usda.gov/GG3/genomics>

Comparative mapping indicated that the QTL for number of spikelets per spike overlapped
Molecular and comparative mapping of genes governing spike compactness
<http://link.springer.com/article/10.1007/s00438-014-0836-2>

Comparative Gene Mapping: A Fine-Scale Survey of Chromosome Rearrangements between
Ruminants and Humans Laurent Schibler,1 Daniel Vaiman, Anne Oustry, Corinne Giraud
<http://genome.cshlp.org/content/8/9/901.full.pdf>

Comparative mapping of the human 22q11 chromosomal region and the importance of detailed
comparative mapping of genes in mice and humans as a
<http://www.pnas.org/content/94/26/14608.full>

nor is the number of genes proportionate to genome size. C. M. et al. Genome sequence,
comparative analysis and haplotype structure of the domestic dog
<http://www.nature.com/scitable/knowledge/library/comparative-genomics-13239404>

COMPARATIVE GENE MAPPING 733 . 27. Dutrillaux, B., M. O. Rethore, M. Prieur, J.
Lejeune. 1973. Analyse de la structure fine des chromosomes du
<http://www.jstor.org/stable/20170373>

Comparative mapping of the ovine clpg locus PREF-1 and MEG3 genes reasonable candidate
genes for clpg, the comparative interval is likely to contain several
http://afsrweb.usda.gov/sp2UserFiles/Place/54380550/Publications/2000Refereed/114997%20Mam%20Genom%2011_10_871-Oct00.pdf

Comparative Mapping of a Major Aluminum Tolerance Gene in Sorghum and Other Species in
Genetic linkage maps were constructed using the Mapmaker Macintosh
<http://www.genetics.org/content/167/4/1905.full>

advanced backcross populations und Ressourcenschutz Professur f r Pflanzenz chtung Prof. Dr.
Jens L on Quantitative trait loci analysis in spring wheat
<http://en.youscribe.com/catalogue/reports-and-theses/knowledge/quantitative-trait-loci-analysis-in-spring-wheat-comparing-two-1436336>