

molecular and biochemical markers associated with salt ... - the identified molecular and biochemical markers could introduce a great benefit for breeding programs to select the salt tolerant individuals. The detection of salt tolerance genes like *badh* could be adding a molecular base add great value to the application of *badh* gene in plant salinity tolerance in **molecular markers associated with salt tolerance in ...** - growing tolerant genotypes on soils with salinity reflects the shift to a strategy of "tailoring the plants to fit the soil"™ in contrast to the older strategy of "tailoring the soil to fit the plants".

molecular markers assisted genetic characterization of ... - mahmood et al., j. anim. plant sci. 23(5):2013 1441 molecular markers assisted genetic characterization of different salt tolerant plant species **molecular profiling using protein markers for salt ...** - protein for determining salt tolerance in sugarcane. khaled and teixeira da silva fw, dw and plant height decreased. they concluded that these traits could be used as indicators for the effect of salt **physiological and molecular markers for salt tolerance in ...** - european scientific journal january 2014 edition vol.10, no.3 issn: 1857-7881 (print) e - issn 1857-7431 253 plants respond to salt stress through modifications of their **bulked segregant analysis to develop molecular markers for ...** - arab j. biotech., vol. 9, no. (1) jan. (2006): 129-142. bulked segregant analysis to develop molecular markers for salt tolerance in egyptian cotton **tagging and validation of ssr markers to salinity ...** - these markers could be useful in molecular mapping and marker assisted selections. keywords-qtls, saltol, ssr markers, salt tolerance, breeding populations. i. introduction so far, conventional breeding methods for salt tolerance have been found ineffective due to the strong environmental effects on genotypic expression and the low narrow sense heritability of salt tolerance [8,9] which ... **identification of ssr markers for salt-tolerance in rice ...** - identification of ssr markers for salt-tolerance in rice variety csr10 by selective genotyping poonam rana1, sunita jain2*, sheetal yadav 1, navinder saini1 and r k jain1 1department of biotechnology and molecular biology, 2department of biochemistry, ccs haryana agricultural university, hisar 125 004, india a population of 171 f 3 genotypes derived from a cross between csr10 (salt tolerant ... **development of est-ssr markers related to salt tolerance ...** - 3733 genetics and molecular research 13 2: 3732-3746 2014 funpec-rp funpecrp development of cotton salt-tolerance ssr marker the objective of this research was to develop simple sequence repeat **molecular mapping and marker-assisted selection for salt ...** - molecular mapping and marker-assisted selection for salt tolerance in rice (*oryza sativa* L.) omonrice 16 (2008) 51 detected under salt tolerance **genetic studies of salinity tolerance in wheat** - genetic studies of salinity tolerance in wheat by nawar jalal shamaya b., m. a thesis submitted for the degree of doctor of philosophy school of agriculture, food and wine **evaluation of wheat landrace genotypes for salinity ...** - sity for salt tolerance in crop species as mentioned above. however, during the last decade however, during the last decade microsatellites or ssr markers have been extensively used for genetic diversity study, genome **chapter 15 molecular markers and their role in producing ...** - molecular markers and their role in producing salt-tolerant crop plants sagar satish datir abstract besides drought and temperature, soil salinity is a severe abiotic environmental constraint to world agriculture, widely affecting yield and quality of crops. increasing world population and simultaneous depletion of agricultural land created an alarming situation to meet the global food ... **hybrid identification for glycine max and glycine soja ...** - hybrid identification for glycine max and glycine soja with ssr markers and analysis of salt tolerance fayuan li 1, *, xun liu,*, shengyan wu1, qingyun luo2 and

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