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# Diversity Arrays Technology (DarT) in genetics and breeding of wheat and barley

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Diversity Arrays Technology (DarT) complements the molecular toolkit for whole genome-based studies. DarT has been developed for over 20 plant species representing a wide range of genome size, ploidy level and breeding systems, including a number of cereal crops. DarT is a sequence-independent, highthroughput technology, discovering and typing hundreds of markers in a single hybridisation-based assay. DarT markers are mostly based on Single Nucleotide Polymorphisms, but detect also Insertion/Deletion and methylation polymorphisms. DarT has been effectively used in characterisation and management of genetic diversity in domesticated species and their relatives, association mapping and pedigree-based mapping, rapid creation of linkage maps for QTL analysis, marker-trait associations through Bulk Segregant Analysis, accelerated introgression from wild germplasm (e.g. AB QTL). DarT is commercially delivered for wheat and barley through Triticarte service ([www.triticarte.com.au](http://www.triticarte.com.au)). Since its establishment Triticarte has rapidly increased the numbers of samples in 2006/2007. Currently used arrays contain approximately 2000 markers for both barley and wheat, with 4000 markers target for wheat in early 2007. Average number of polymorphic markers reported to customers in 2005 was 472 for barley and 63 for wheat, with average price per datapoint at 7 US cents. Such Throughput and cost enables utilisation of DarT-based whole genome profiles in breeding.