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Catalog	Organism	ArrayFormat	Comments
GT_CAT_78	Solanum lycopersicum (Tomato) SNP	1x244K	Tilling probes designed from all available ESTs & mRNAs from NCBI & TIGR. Probe length 25 mer
GT_CAT_79	Eucalyptus and Populus	1x244K	1 Probe per transcript for Eucalyptus and Poppulas ESTs & mRNAs from NCBI
GT_CAT_80	Cucumis melo (Melon)	2x105K	3 Probe per transcript for Eucalyptus and Poppulas ESTs & mRNAs from NCBI
GT_CAT_81	Nicotiana tabacum	1x244K	3 Probe per transcript for N. tabacum ESTs & mRNAs from NCBI
GT_CAT_82	Cucumis sativus (Cucumber)	4x44K	2 Probe per transcript for Cummber ESTs & mRNAs from NCBI
GT_CAT_83	Banana	4x44K	Covers all the available ESTs NCBI's dbEST (Design in progress). Comprehensive annotation has been provided based on sequence homology to Arabidopsis & Rice.
GT_CAT_84	Brassica juncea	8x15K	1 Probe per transcript for Brassica ESTs & mRNAs from NCBI
GT_CAT_144	Arabidopsis thaliana	1x244K(2SlideSet)	Probes tiled across whole Arabidopsis genome with 212 nt average spacing. content sourced from TIGR ATH1
GT_CAT_145	Oryza (Rice)	4x44K	43,803 Rice probes represented. Content sourced from National Institute of Agrobiological Sciences (NIAS), RefSeq and UniGene
GT_CAT_146	Zea mays (Maize)	4x44K	21,000 Maize probes represented, replicated twice. Content sourced from RefSeq and GenBank
GT_CAT_147	Glycine max (Soybean)	4x44K	43,803 Soybean probes represented. Content sourced from RefSeq, UniGene, TIGR Plant TA and TIGR Gene Indices
GT_CAT_148	Nicotiana tabacum (Tobacco)	4x44K	43,803 Tobacco probes represented. Content sourced from UniGene, TIGR, IGR Plant Transcript Assemblies
GT_CAT_149	Arabidopsis thaliana	4x44K	43,803 A. thaliana probes represented. Content sourced from RefSeq, UniGene, TAIR, TIGR and ATH1
GT_CAT_150	Hordeum vulgare (Barley)	4x44K	43,803 Barley probes represented. Content sourced from RefSeq, UniGene, TIGR Transcript Assemblies and TIGR Gene Indices
GT_CAT_151	Solanum lycopersicum (Tomato)	4x44K	43,803 Tomato probes represented. Content sourced from RefSeq, UniGene, TIGR Plant Transcript Assemblies and IGR Gene Indices
GT_CAT_152	Triticum aestivum (Wheat)	4x44K	43,803 Wheat probes represented. Content sourced from RefSeq, UniGene, TIGR Plant Transcript Assemblies and TIGR Gene Indices
GT_CAT_153	Brassica napus	4x44K	43,803 Brassica probes represented. Content sourced from RefSeq, UniGene, TIGR Plant TA and TIGR Gene Indices
GT_CAT_154	Gossypium hirsutum (Cotton)	4x44K	43,803 Cotton probes represented. Content sourced from RefSeq, UniGene, TIGR Plant TA and TIGR Gene Indices
GT_CAT_155	Medicago	4x44K	43,803 Medicago probes represented. Content sourced from RefSeq, UniGene, TIGR Plant TA and TIGR Gene Indices
GT_CAT_256	Sorghum bicolor	1x244K	Sorghum bicolor gene expression array
GT_CAT_257	Solanum tuberosum (Potato)	8x60K	Potato ESTs Gene expression array
GT_CAT_258	Antirrhinum majus (snapdragon)	4x44K	Antirrhinum majus ESTs Gene expression array
GT_CAT_259	Chlamydomonas reinhardtii	4x44K	Chlamydomonas reinhardtii whole genome Gene expression array
GT_CAT_326	Arabidopsis thaliana	8x60K	Arabiopsis thaliana Whole genome including organalle specific expression profiling with 1 probe for each transcript genes along with mitochondrial genes and chloroplast genes
GT_CAT_327	Oryza sativa - Japonica	8x60K	Rice Whole genome including organalle specific expression profiling with 1 probe for each transcript genes along with mitochondrial genes and chloroplast genes
GT_CAT_404	Oryza sativa	8x60K	Oryza sativa whole genome gene expression microarray
GT_CAT_406	Curcuma longa	4x180K	Curcuma longa whole genome gene expression microarray
GT_CAT_409	Zea mays	4x180K	Zea mays whole genome gene expression microarray
GT_CAT_410	Brassica rapa	4x44K	Brassica rapa Whole genome Chip array
GT_CAT_411	Oryza sativa	4x44K	Oryza sativa Whole genome Chip array
GT_CAT_38	Chlamydomonas reinhardtii	4x44K	Average 1 probes for each ORF. Non-mRNA(structural RNAs) transcripts are included. Content source JGI.
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