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*Hits **1** through **50** out of **552***

Next 50 Hits

Jump To

Refine Search AN/"Affymetrix, Inc"

	PAT. NO.	Title
1	11,118,216	Nucleic acid analysis by joining barcoded polynucleotide probes
2	11,009,462	Methods, systems and apparatuses for testing and calibrating fluorescent scanners
3	10,977,478	Methods and devices for reading microarrays
4	10,920,269	Amplification and analysis of selected targets on solid supports
5	10,872,681	Differential filtering of genetic data
6	10,832,796	System, method, and computer software product for genotype determination using probe array data
7	10,822,659	Analysis of methylation using nucleic acid arrays
8	10,822,649	Multiplex targeted amplification using flap nuclease
9	10,822,642	Methods of analysis of methylation
10	10,648,030	Methods of determining the presence or absence of a plurality of target polynucleotides in a sample
11	10,648,021	DNA ligation on RNA template
12	10,586,095	Methods and devices for reading microarrays
13	10,535,420	Systems and methods for probe design to detect the presence of simple and complex indels
14	10,515,447	Analysis of data obtained from microarrays
15	10,422,750	Methods, systems and apparatuses for testing and calibrating fluorescent scanners
16	10,407,717	Methods of analysis of methylation
17	10,329,600	Locus specific amplification using array probes
18	10,316,136	Polyfluoreno[4,5-cde]oxepine conjugates and their use in methods of analyte detection
19	10,303,922	Methods and devices for reading microarrays
20	10,275,568	Methods and computer software for detecting splice variants
21	10,229,244	Methods for identifying DNA copy number changes using hidden markov model based estimations
22	10,213,762	Silane mixtures
23	10,155,976	Methods for genotyping selected polymorphism
24	10,119,162	Multiplex targeted amplification using flap nuclease

25 10,108,777 T Methods for identifying DNA copy number changes
26 10,019,620 T Methods and devices for reading microarrays
27 9,982,293 T Amplification and analysis of selected targets on solid supports
28 9,933,365 T Methods, systems and apparatuses for testing and calibrating fluorescent scanners
29 9,932,636 T Array-based translocation and rearrangement assays
30 9,896,538 T Polyfluoreno[4,5-cde]oxepine polymers and conjugates thereof
31 9,845,494 T Enzymatic methods for genotyping on arrays
32 9,845,490 T Complexity management of genomic DNA
33 9,828,640 T Analysis of methylation using nucleic acid arrays
34 9,816,134 T Multiplex amplification methods
35 9,798,855 T Differential filtering of genetic data
36 9,767,342 T Methods and devices for reading microarrays
37 9,760,675 T System, method, and computer software product for genotype determination using probe array data
38 9,708,643 T Circulating miRNA biomaker signatures
39 9,689,867 T Assays for affinity profiling of nucleic acid binding proteins
40 9,663,822 T Multiplex capture of nucleic acids
41 9,599,561 T Methods, systems and apparatuses for testing and calibrating fluorescent scanners
42 9,556,360 T Silane mixtures
43 9,528,148 T Multiplex targeted amplification using flap nuclease
44 9,523,122 T Multiplex targeted amplification using flap nuclease
45 9,445,025 T System, method, and product for imaging probe arrays with small feature sizes
46 9,388,461 T Methods for genotyping selected polymorphism
47 9,388,460 T Complexity management of genomic DNA
48 9,388,459 T Methods for genotyping
49 9,388,457 T Locus specific amplification using array probes
50 9,359,640 T Methods for synthesizing pools of probes

Next List | Top | View Cart
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