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	PAT. NO.	Title
1	11,261,485	Multiplex amplification methods
2	11,118,216	Nucleic acid analysis by joining barcoded polynucleotide probes
3	11,009,462	Methods, systems and apparatuses for testing and calibrating fluorescent scanners
4	10,977,478	Methods and devices for reading microarrays
5	10,920,269	Amplification and analysis of selected targets on solid supports
6	10,872,681	Differential filtering of genetic data
7	10,832,796	System, method, and computer software product for genotype determination using probe array data
8	10,822,659	Analysis of methylation using nucleic acid arrays
9	10,822,649	Multiplex targeted amplification using flap nuclease
10	10,822,642	Methods of analysis of methylation
11	10,648,030	Methods of determining the presence or absence of a plurality of target polynucleotides in a sample
12	10,648,021	DNA ligation on RNA template
13	10,586,095	Methods and devices for reading microarrays
14	10,535,420	Systems and methods for probe design to detect the presence of simple and complex indels
15	10,515,447	Analysis of data obtained from microarrays
16	10,422,750	Methods, systems and apparatuses for testing and calibrating fluorescent scanners
17	10,407,717	Methods of analysis of methylation
18	10,329,600	Locus specific amplification using array probes
19	10,316,136	Polyfluoreno[4,5-cde]oxepine conjugates and their use in methods of analyte detection
20	10,303,922	Methods and devices for reading microarrays
21	10,275,568	Methods and computer software for detecting splice variants
22	10,229,244	Methods for identifying DNA copy number changes using hidden markov model based estimations
23	10,213,762	Silane mixtures
24	10,155,976	Methods for genotyping selected polymorphism

25 10,119,162 T Multiplex targeted amplification using flap nuclease
26 10,108,777 T Methods for identifying DNA copy number changes
27 10,019,620 T Methods and devices for reading microarrays
28 9,982,293 T Amplification and analysis of selected targets on solid supports
29 9,933,365 T Methods, systems and apparatuses for testing and calibrating fluorescent scanners
30 9,932,636 T Array-based translocation and rearrangement assays
31 9,896,538 T Polyfluoreno[4,5-cde]oxepine polymers and conjugates thereof
32 9,845,494 T Enzymatic methods for genotyping on arrays
33 9,845,490 T Complexity management of genomic DNA
34 9,828,640 T Analysis of methylation using nucleic acid arrays
35 9,816,134 T Multiplex amplification methods
36 9,798,855 T Differential filtering of genetic data
37 9,767,342 T Methods and devices for reading microarrays
38 9,760,675 T System, method, and computer software product for genotype determination using probe array data
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41 9,663,822 T Multiplex capture of nucleic acids
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45 9,523,122 T Multiplex targeted amplification using flap nuclease
46 9,445,025 T System, method, and product for imaging probe arrays with small feature sizes
47 9,388,461 T Methods for genotyping selected polymorphism
48 9,388,460 T Complexity management of genomic DNA
49 9,388,459 T Methods for genotyping
50 9,388,457 T Locus specific amplification using array probes

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