

Element Biosciences Patents

The patents listed below are those issued in the United States. For information on patents issued outside of the United States, please contact us at legalnotices@elembio.com.

<ul style="list-style-type: none">• US10704094, Multipart reagents having increased avidity for polymerase binding• US10768173, Multivalent binding composition for nucleic acid analysis• US10876148, De novo surface preparation and uses thereof• US10982280, Multipart reagents having increased avidity for polymerase binding• US11053540, High performance fluorescence imaging module for genomic testing assay• US11060138, Nucleic acid sequencing systems• US1119812, Flow cell systems and devices• US11200446, Single-pass primary analysis• US11220707, Compositions and methods for pairwise sequencing• US11236388, Compositions and methods for pairwise sequencing• US11261489, High performance fluorescence imaging module for genomic testing assay• US11459608, High performance fluorescence imaging module for genomic testing assay• US11426732, Flow cell device and use thereof• US11427855, Compositions and methods for pairwise sequencing	<ul style="list-style-type: none">• US11408032, Tube lens design for improved depth-of-field• US11397870, Single-pass primary analysis• US11339433, Nucleic acid sequencing systems• US11365444, High performance fluorescence imaging module for genomic testing assay• US11287422, Multivalent binding composition for nucleic acid analysis• US10233490, Methods for assembling and reading nucleic acid sequences from mixed populations• US11781185, Methods and reagents for nucleic acid analysis• US11788075, Engineered polymerases with reduced sequence-specific errors• US11795504, High performance fluorescence imaging module for genomic testing assay• US11859241, Compositions and methods for pairwise sequencing• US11891651, Compositions and methods for pairwise sequencing• US11915444, Single-pass primary analysis• US11198121, Flow cell systems and devices• US12006518, Engineered polymerases with reduced sequence-specific errors
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