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## Supporting Information

2 **CRISPR/Cas13a-based single-nucleotide polymorphism detection for**  
3 **reliable determination of ABO blood group genotypes**

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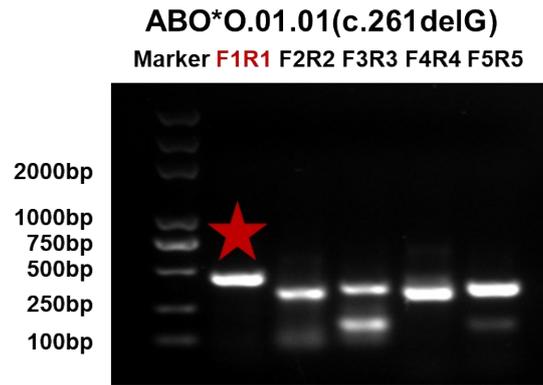
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15 **Table S1. The sequences of Amplicons, PCR primers and crRNAs**

DNA	Detail of sequences 5`-3`
<b>Amplicon of Exon 6-261</b>	<b>GTCGCATTTGCCTCTGGTTGGTTTCCCGGGGAAGGGCGGCTGCCTCTGG AAGGGTGGTCAGAGGAGGCAGAAGCTGAGTGGAGTTTCCAGGTGGGGGC GGCCGTGTGCCAGAGGCGCATGTGGGTGGCACCCCTGCCAGCTCCATGTGA CCGCACGCCTCTCTCCATGTGCAGTAGGAAGGATGTCTCTCGTGGT</b> <b>NACCC CTTGGCTGGCTCCCATTGTCTGGGAGGGCACATTCAACATCGACATCCTC AACGAGCAGTTCAGGCTCCAGAACCACCATTGGGTAACTGTGTTTGC CATCAAGAAGTAAGTCAGTGAGGTGGCCGAGGGTAGAGACCCAGGCAGT GGCAGTGACTGTGGACATTGAGGTCTCTCCTTGTGTTCAAGACAGAGT GGGGTG</b>
<b>ABO-6-T7-F1</b>	<b>GAAATTAATACGACTCACTATAGGGGTCGCATTTGCCTCTGGTTG</b>
ABO-6-T7-F2	GAAATTAATACGACTCACTATAGGGGGCAGAAGCTGAGTGGAGTT
ABO-6-T7-F3	GAAATTAATACGACTCACTATAGGGTCCAATGTTGAGGGAGGGCT
ABO-6-T7-F4	GAAATTAATACGACTCACTATAGGGGGGAATGATTTGCCCGGTTG
ABO-6-T7-F5	GAAATTAATACGACTCACTATAGGGGTTCCCGCAGGTCCAATGTT
<b>ABO-6-R1</b>	<b>CACCCCACTCTGTCTTGAACA</b>
ABO-6-R2	AATGTCCACAGTCACTCGCC
ABO-6-R3	CCCAATGGTGGTGTCTGGA
ABO-6-R4	GAGCCTGAACTGCTCGTTGA
ABO-6-R5	ACTGCTCGTTGAGGATGTCG
6-261-G1	GT <b>G</b> ACCCCTTGGCTGGCTCCCATTGTCT
<b>6-261-G2</b>	<b>ATGTCCTCGTGGT</b> <b>G</b> ACCCCTTGGCTGGC
6-261-G3	GCAGTAGGAAGGATGTCCTCGTGGT <b>G</b> AC
6-261-D1	GT <b> </b> ACCCCTTGGCTGGCTCCCATTGTCTG
6-261-D2	GATGTCCTCGTGGT <b> </b> ACCCCTTGGCTGGC
<b>6-261-D3</b>	<b>TGCAGTAGGAAGGATGTCCTCGTGGT</b> <b> </b> <b>AC</b> <b>CCGTGTCCACTACTATGTCTTACCAGCAGCNGGCCGCGGTGCCCGC</b> <b>GTGACGCTGGGGACCGTCCGCAGCTGTCAGTGCTGGAGGTGNGCGCCT</b> <b>ACAAGCGCTGGCAGGACGTGTCCATGCGCCGCATGGAGATGATCAGTGA</b> <b>CTTCTGCGAGCGGCGCTTCTCAGCGAGGTGGATTACCTGGTGTGCGTGG</b> <b>ACGTGGACATGGAGNTCCGCGACCACGTGGGCGTGGAGATCCTGACTCC</b> <b>GCTGTTCGGCACCCCTGCACCCCGGCTTCTACGGAAGCAGCCGGGAGGCCT</b> <b>TCACCTACGAGCGCCGCCCCAGTCCCAGGCCTACATCCCCAAGGACGAG</b> <b>GGCGATTTCTACTAC</b> <b>N</b> <b>TGGGGGNGTTCTTCGGGGGGTTCGGTGAAGAGGT</b> <b>GCAGCGGCTCACCAGGGCCTGCCACCAGGCCATGATGGTTCGACCAGGCC</b> <b>AACGGCATCGAGGCCGTGTGGCACGACGAGAGCCACCTGAACAAGTACC</b> <b>TGCTGCGCCACAAACCCACCAAGGTGCTCTCCCCGAGTACTTGTGG</b>
<b>Amplicon of Exon 7-796</b>	
<b>ABO-7-T7-F1</b>	<b>GAAATTAATACGACTCACTATAGGGCCGTGTCCACTACTATGTCTTACC</b>
ABO-7-T7-F2	GAAATTAATACGACTCACTATAGGGCCTGCCTTGCAGATACGTGG
ABO-7-T7-F3	GAAATTAATACGACTCACTATAGGGTCTAAGCCTTCCAATGGCCG
ABO-7-T7-F4	GAAATTAATACGACTCACTATAGGGGATGAAGTGAATCGCAGCCC
ABO-7-T7-F5	GAAATTAATACGACTCACTATAGGGCCTTACCTACGAGCGCC
ABO-7-T7-F6	GAAATTAATACGACTCACTATAGGGTGCTGCTCTAAGCCTTCCAAT
ABO-7-T7-F7	GAAATTAATACGACTCACTATAGGGGGCGTGGAGATCCTGACT
ABO-7-T7-F8	GAAATTAATACGACTCACTATAGGGATCCTGACTCCGCTGTTCG

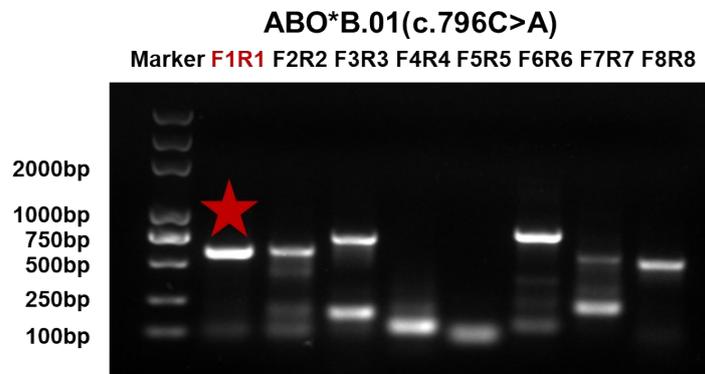
<b>ABO-7-R1</b>	<b>CCACAAGTACTCGGGGGAG</b>
ABO-7-R2	TTG TTCAGGTGGCTCTCGTC
ABO-7-R3	AGCACCTTGGTGGGTTTGTG
ABO-7-R4	ACAACAGGACGGACAAAGGAAACAG
ABO-7-R5	ACTTG TTCAGGTGGCTCTCG
ABO-7-R6	CACAAGTACTCGGGGGAGAG
ABO-7-R7	CACAACAGGACGGACAAAGGA
ABO-7-R8	TTCTGCTAAAACCAAGGGCG
<b>7-796-C1</b>	<b>CAAGGACGAGGGCGATTTCTACTACCTG</b>
7-796-C2	CAAGGACGAGGGCGATTTCTACT <u>IC</u> CTG
7-796-C3	AGGACGAGGGCGATTTCTACTACCTGGG
7-796-A1	CAAGGACGAGGGCGATTTCTACTACATG
7-796-A2	CAAGGACGAGGGCGATTTCTACT <u>IC</u> ATG
<b>7-796-A3</b>	<b>AGGACGAGGGCGATTTCTACTACATGGG</b>
FAM-Reporter	FAM-UUUUU-BHQ1

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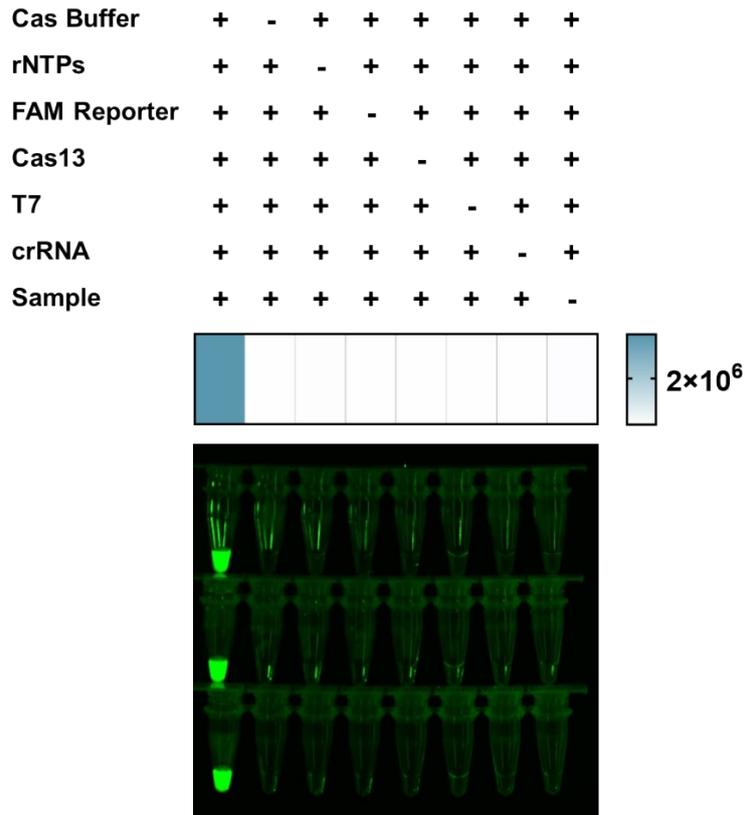
18 **Figure S1.** Five pairs of PCR primers were screened for ABO\*O.01.01(c.261delG)  
 19 locus, the best of which was F1R1.



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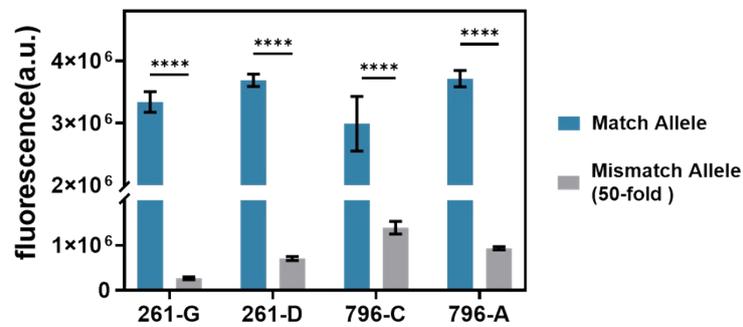
21 **Figure S2.** Eight pairs of PCR primers were screened for ABO\*B.01(c.796C>A) locus,  
 22 the best of which was F1R1.

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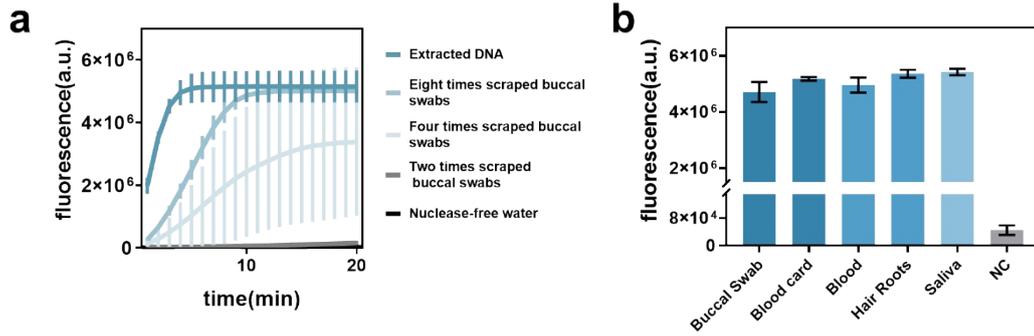
25 **Figure S3.** Verification of the effect of each component of the Cas13 trans-cleavage  
 26 activity showed that each component is indispensable.



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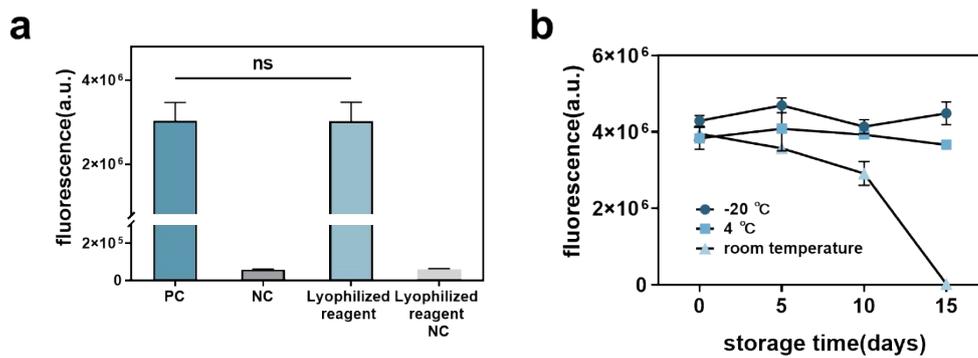
28 **Figure S4.** The discrimination capacity validation experiment. The four screened  
 29 crRNAs demonstrated good discrimination capacity in samples with matched  
 30 sequences and 50-fold concentrations of mismatched sequences with SNP.

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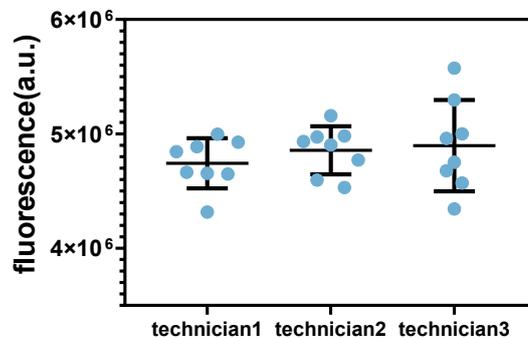
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33 **Figure S5.** Extraction-free sample pretreatment was tested. (a) There is no significant  
 34 difference between positive control (extracted DNA) and directly treated buccal swabs  
 35 when scraped times outnumber eight. (b) Different samples including buccal swabs,  
 36 blood cards, peripheral blood, hair roots, and saliva tested show the adaptivity of the  
 37 method.



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39 **Figure S6.** (a) Lyophilization reagent techniques were tested, which is also compatible  
 40 with this assay. (b) The end-point fluorescence intensity at 20 minutes using the  
 41 lyophilized reagents stored at  $-20\text{ }^{\circ}\text{C}$ ,  $4\text{ }^{\circ}\text{C}$ , and room temperature for 0, 5, 10, and 15  
 42 days.



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44 **Figure S7.** The reproducibility of our method was validated by three different lab  
 45 technicians conducting eight experiments.