

Supplementary Material

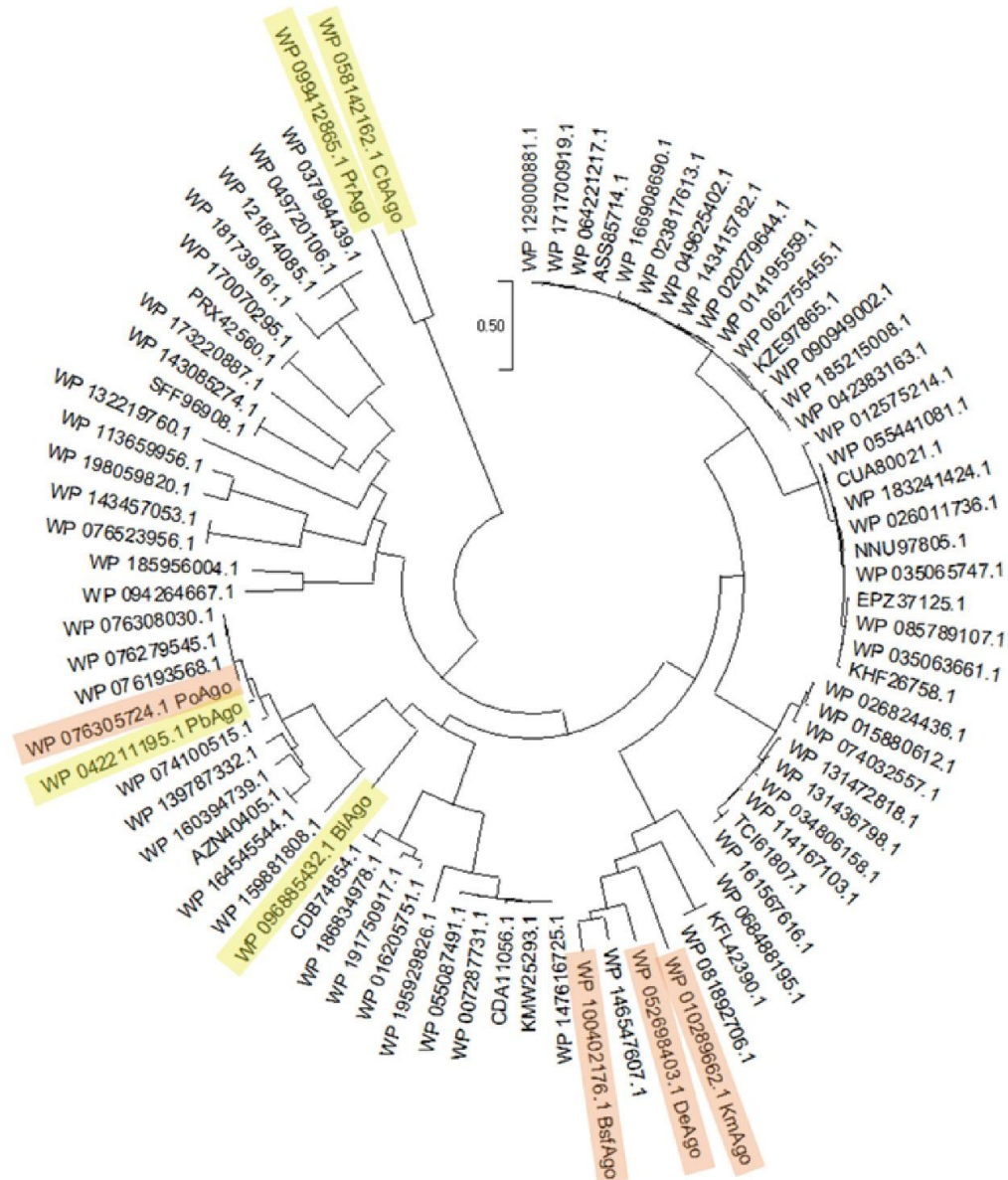
Supplementary Tables

Supplementary Table 1 The nucleotide sequences used in this research.

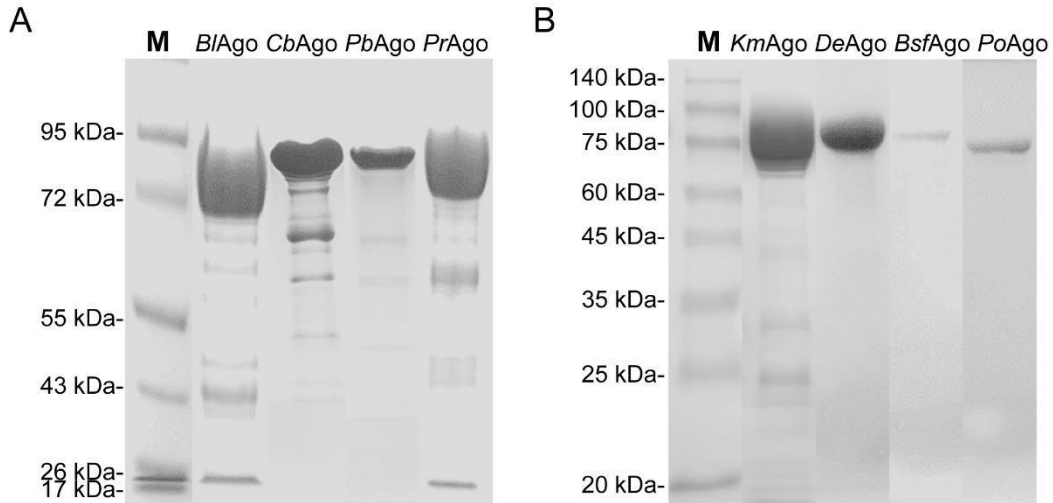
Single-stranded nucleic acids	Sequences (5'-3')
SARS-CoV-2 1b 76 nt ssDNA	TGTGTGGCGGTTCACTATATGTTAAACCAGGTGG AACCTCATCAGGAGATGCCACAACCTGCTTATGCT AATAGTGT
primary gDNA RT Guide A	P-GCAGTTGTGGCATCTCCTG
primary gDNA RT Guide B	P-CTGATGAGGTTCCACCTGG
Fluorescent Reporter	FAM-GCGACTTGACATCTCCTGATGAGGTGTGC- BHQ1
ssDNA target	ATATACTATAACAACCTACTACCTCGTATAAATTTT TAAATAAATA-FAM
ssRNA target	AUAUACUAUACAACCUACUACCUCGUAAUAAA UUUAAAUAUAAAUA-FAM
5'P DNA guide	P-TGAGGTAGTAGGTTGTATAGT
5'OH DNA guide	HO-TGAGGTAGTAGGTTGTATAGT
5'P RNA guide	P-UGAGGUAGUAGGUUGUAUAGU
5'OH RNA guide	HO-UGAGGUAGUAGGUUGUAUAGU
Inserted DNA fragment	TAATACGACTCACTATAGGGGATAAAAGTGCATT AACATTGGCCGTGACAGCTTGACAAATGTTAAAA ACACTATTAGCATAAGCAGTTGTGGCATCTCCTG ATGAGGTTCCACCTGGTTAACATATAGTGAACC GCCACACATGACCATTTCACTCAATACTTGAGCA CACTCATTAGCTAATCTATAGAAACGGTGTGACA AGCTACAACACGTTGT
FW-1b-256 bp	CAGCTATGACCATGATTACGCCAAG
RV-1b-256 bp	GGGGATCCACAACGTGTTGTAGCTT GGGGAUAAAAGUGCAUUAACAUUGGCCGUGACA GCUUGACAAAUGUUAAAAACACUAUUAGCAUAA GCAGUUGUGGCAUCUCCUGAUGAGGUUCCACCU
RNA template	GGUUUAACAUAUAGUGAACCGCCACACAUGACC AUUUCACUCAAUACUUGAGCACACUCAUUAGCU AAUCUAUAGAAACGGUGUGACAAGCUACAACAC GUUGUGGAUCCCC
RT primer	ACAACGTGTTGTAGC

Supplementary Table 2 Comparison of reported CRISPR-based and thermophilic Ago-based methods for SARS-CoV-2 detection.

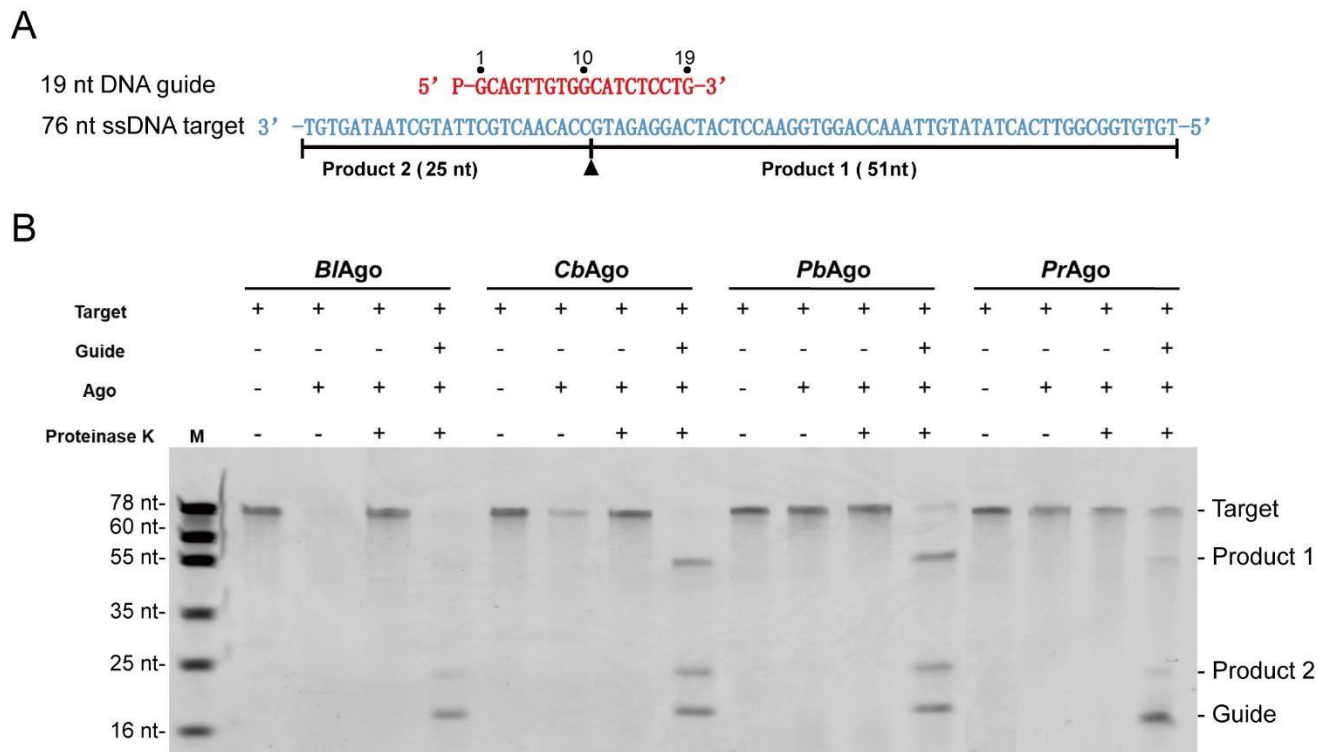
Detection methods	Nuclease	Operation	Target motif requirement	Guide oligos	Specificity	Lid opening	Portable
DETECTR (Broughton et al., 2020)	<i>LbCas12a</i>	62 °C for 20-30 min; 37 °C for 10 min	PAM	41 nt gRNAs	High	Yes	Yes
CRISPR-FDS (Huang et al., 2020)	<i>LbaCas12a</i>	42 °C for 20 min; 37 °C for 20 min	PAM	41-43 nt gRNAs	Medium	Yes	No
CRISPR/Cas12a-NER (Wang et al., 2020)	<i>FnCas12a</i>	39 °C for 30 min; 37 °C for 15 min	PAM	28 nt crRNAs	High	Yes	Yes
STOPCovid (Joung et al., 2020)	<i>AapCas12b</i>	60 °C for 1 h	PAM	111 nt crRNAs	High	No	Yes
CASdetec (Guo et al., 2020)	<i>AaCas12b</i>	42 °C for 30 min; 42 °C for 10-30 min	PAM	103-105 nt sgRNAs	High	No	Yes
SHERLOCK (Zhang et al., 2020)	<i>LwaCas13a</i>	42°C for 25 min; 37°C for 30 min	PFS	64 nt crRNAs	High	Yes	Yes
SARS-CoV-2 PAND (Wang et al., 2021)	<i>PfAgo</i>	55 °C for 10 min; 45 PCR cycles; 95 °C for 20-30 min	No	16 nt gDNAs	High	Yes	No
MULAN (Ye et al., 2022)	<i>PfAgo</i>	65 °C for 30-40 min; 95 °C for 10-15 min	No	16 nt gDNAs	High	No	Yes
MAIDEN	<i>PbAgo/KmAgo</i>	42 °C for 1 h	No	19 nt gDNAs	High	No	Yes

Supplementary Figures


Supplementary Figure 1. Phylogenetic tree analysis based on the amino acid sequences of the characterized mesophilic Agos. Candidate Agos that target DNA are highlighted in yellow. Candidates for universal Ago are highlighted in orange. Ago from *Brevibacillus laterosporus* (BlAgo, WP_096885432.1); Ago from *Clostridium butyricum* (CbAgo, WP_058142162.1); Ago from *Paenibacillus borealis* (PbAgo, WP_042211195.1); Ago from *Pseudobutyrvibrio ruminis* (PrAgo, WP_099412865.1); Ago from *Kurthia massiliensis* (KmAgo, WP_010289662.1); Ago from *Domibacillus enclensis* (DeAgo, WP_052698403.1); Ago from *Bacillus sp. FJAT-42315* (BsfAgo, WP_100402176.1); Ago from *Paenibacillus odorifer* (PoAgo, WP_076305724.1).

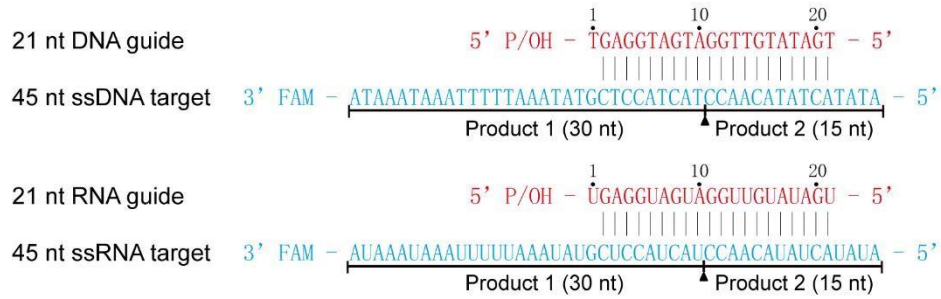


Supplementary Figure 2. SDS-PAGE analysis of purified Agos. (A) Candidate Agos that target DNA: *BlAgo*, *CbAgo*, *PbAgo*, *PrAgo*. (B) Candidates for universal Ago: *KmAgo*, *DeAgo*, *BsfAgo*, *PoAgo*.

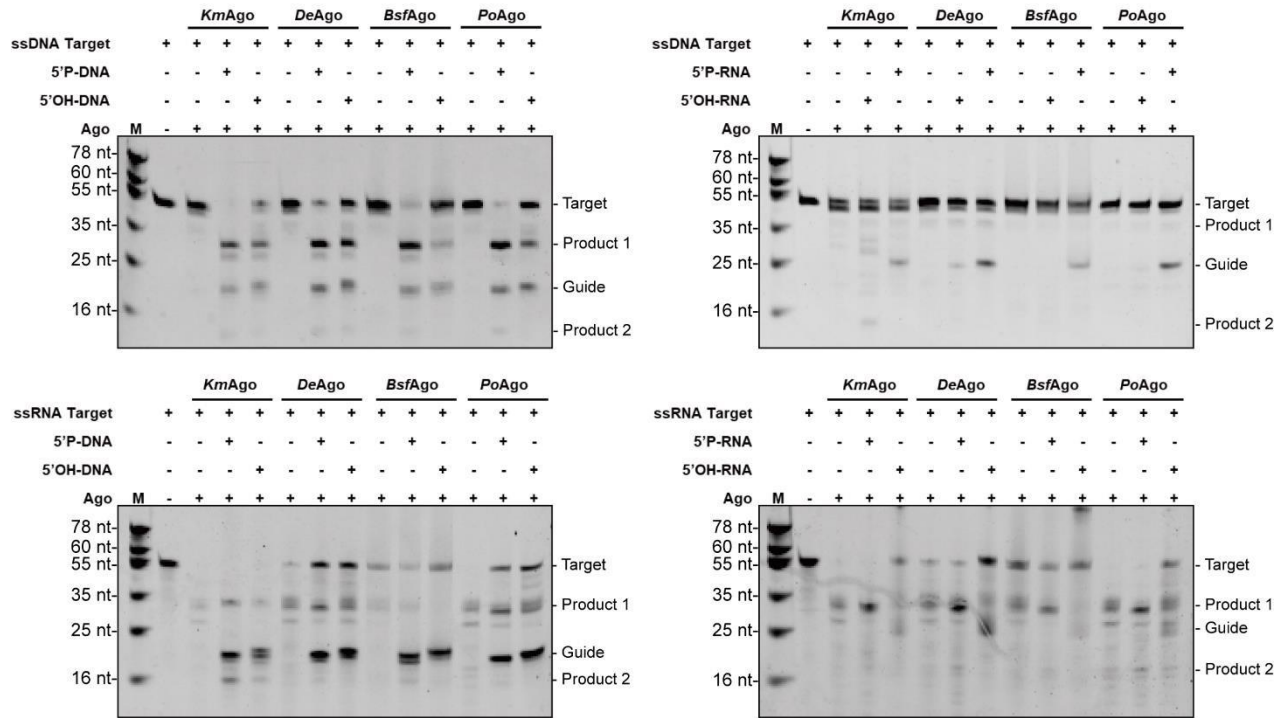


Supplementary Figure 3. *In vitro* cleavage activity determination of *BlAgo*, *CbAgo*, *PbAgo* and *PrAgo*. (A) Synthetic 5'P gDNA and ssDNA targets sequence. (B) The gel electrophoresis result of the Ago cleavage experiment.

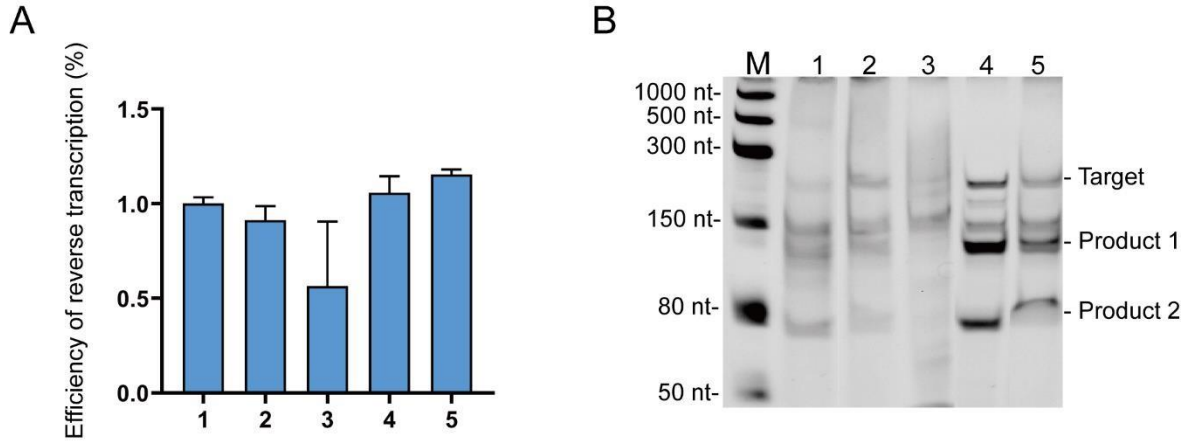
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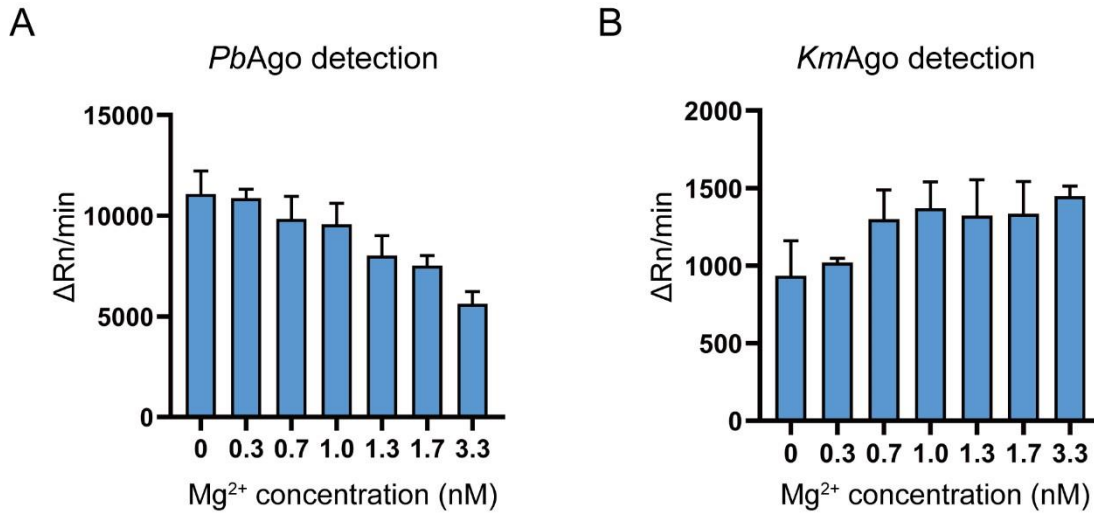
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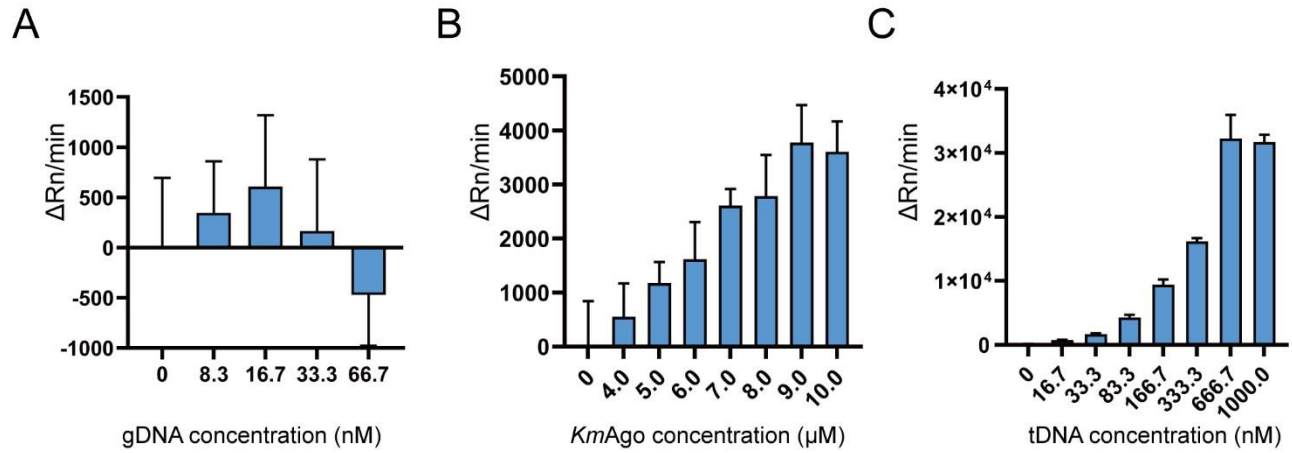
Supplementary Figure 4. *In vitro* cleavage activity determination of *KmAgo*, *DeAgo*, *BsfAgo* and *PoAgo*. (A) Synthetic 5'P and 5'OH gDNA, gRNA and ssDNA, RNA targets sequence. (B-E) The gel electrophoresis result of the Ago cleavage experiment: (B) DNA-guided cleavage of DNA target; (C) RNA-guided cleavage of DNA target; (D) DNA-guided cleavage of RNA target; (E) RNA-guided cleavage of RNA target.



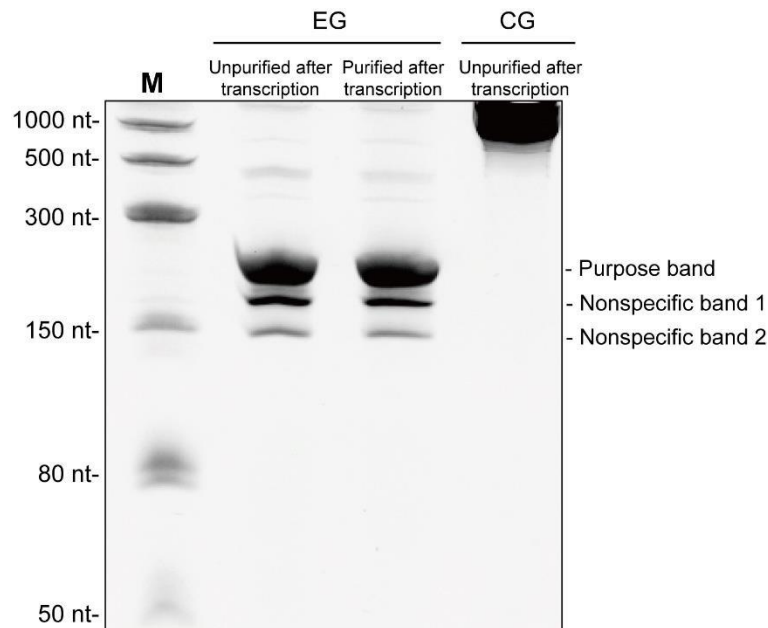
Supplementary Figure 5. Screening of enzymes for reverse transcription. (A) Transcription efficiency of the five reverse transcriptases. Error bars, standard deviation (SD) of replicates ($n = 3$). (B) Cleavage efficiency of *PbAgo* in five reverse transcription systems.



Supplementary Figure 6. Optimization of an isothermal detection reaction of mesophilic Agos. (A) Fluorescence values of *PbAgo* cascade cleavage under different concentrations of Mg²⁺. (B) Fluorescence values of *KmAgo* cascade cleavage under different concentrations of Mg²⁺. Data were collected from three independent experiments and are presented as the Mean \pm SD.



Supplementary Figure 7. Optimization of an isothermal detection reaction of *KmAgo*. (A-C) Fluorescence values of *KmAgo* cascade cleavage under different concentrations of gDNA (A), *KmAgo* (B), and tDNA (C). Data were collected from three independent experiments and are presented as the Mean ± SD.



Supplementary Figure 8. The gel electrophoresis result of transcribed RNA. EG, experimental group, including unpurified and purified 200bp RNA template after transcription. CG, control group, unpurified RNA fragment transcribed from the DNA template that comes with the transcription kit.

Supplementary References

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