## **Supplementary Material**

## Discriminating Unalike Single Nucleobase Mismatches using a Molecularly Resolved, Label-free, Interfacial LNA-based Assay

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Fig. S1: RAIR spectra of DNA and LNA layers on gold(111) surface acquired for the same probe concentration  $0.1\mu M$  and the same incubation time 4 h.

Assignment of the primary IR frequencies of that differentiate between DNA and LNA cases

ASSIGNMENT	DNA	LNA
2997 <sup>1</sup>		V(C-H)aromatic, present in nucleobases
2965-2968 <sup>1</sup>	Vasym(CH <sub>3</sub> ), present in thymine bases	Vasym(CH <sub>3</sub> ), present in thymine bases, <b>peak is more intense than in</b> <b>case of DNA</b>
2937 <sup>1</sup>		Vasym(CH <sub>2</sub> ), present in hexyl spacer and locked sugar moieties
28811		Vsym(CH <sub>2</sub> )
1734-1743 <sup>2,3</sup>	v(C=O) nucleobase, present in thymine and guanine	v(C=O), present in thymine and guanine, <b>peak much sharper and</b> <b>intense</b>
1607-1654 <sup>2,3</sup>	v(C=O) nucleobase, present in cytosine and adenine	v(C=O), present in cytosine and adenine, <b>peak much sharper and</b> <b>intense</b>
1614 <sup>4</sup>		$\delta(-NH_2)$ , v(C=N), present in nucleobases
1544 <sup>4</sup>		v(C=C), v(C=N), present in nucleobases
1458 <sup>4</sup>		purine imidazolic ring vib.
1416 <sup>5-7</sup>		sugar vib. (C3'- endo)
1275 <sup>8</sup>		thymidine (N3–H bending vib.)
11919		sugar phosphate backbone vib. and N-type sugar conformation
1039-1096 <sup>10</sup>	v(PO <sub>2</sub> ), sugar-phosphate backbone vibration	v(PO <sub>2</sub> ) <sup>-</sup> , sugar-phosphate backbone vibration, <b>peak much sharper and</b> <b>intense</b>

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Fig. S2: Representative plots for on-surface melting temperature of LNA-DNA duplex (A) for fully matched LNA-DNA duplex,  $T_m = 52.1$  °C; and [B-G] for single mismatch at the central position (B) G: G,  $T_m = 31$  °C; (C) G: T,  $T_m = 31.4$  °C; (D) T: G,  $T_m = 33$  °C; (E) A: G,  $T_m = 30$  °C; (F) A: C,  $T_m = 31.1$  °C; (G) C: A,  $T_m = 34.3$  °C; (H) C: T,  $T_m = 33.8$  °C; (I) C: C,  $T_m = 33.4$  °C; (J) A: A,  $T_m = 30.4$  °C; (K) T: T,  $T_m = 31.8$  °C.



Fig. S3: Unbinding force values for fully matched as well as the mismatched combinations presented including the standard errors from mean.