## **Supplementary Information**

## Mismatch detection enhancement based on silicon nanowires 3D microarray

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## Figure S1: Hairpin formation probability



 $\Delta G$  = 1.18 kcal/mole  $\Delta H$  = -16.2 kcal/mole  $\Delta S$  = -58.31 cal/ K \* mole

## Table S1: Self-dimer formation probability

Probe Sequence								
5'- CCCATTTTCCTCCCGCAATTCCTAG -3'								
Maximum Delta G: -52.23 kcal/mole								
Delta G: -5.36 kcal/mole Base Pairs: 4								
5' CTAGGAATTGCGGGAGGAAAATGGG								
: :      : :								
3' GGGTAAAAGGAGGGCGTTAAGGATC								
Delta G: -4.16 kcal/mole Base Pairs: 4								
5' CCCATTTTCCTCCCGCAATTCCTAG								
3' GATCCTTAACGCCCTCCTTTTACCC								
Delta G: -3.61 kcal/mole Base Pairs: 2								
5' CCCATTTTCCTCCCGCAATTCCTAG								
: :    : :								
3' GATCCTTAACGCCCTCCTTTTACCC								
Delta G: -3.42 kcal/mole Base Pairs: 3								
5' CCCATTTTCCTCCCGCAATTCCTAG								
3' GATCCTTAACGCCCTCCTTTTACCC								
Delta G: -3.14 kcal/mole Base Pairs: 2								
5' CCCATTTTCCTCCCGCAATTCCTAG								
:    :								
3' GATCCTTAACGCCCTCCTTTTACCC								

Figure S2: Comparison of the microarray scanned images obtained on 2D (a) and 3D (b) platforms, respectively

	1µM	ŧį.	PM	4	×,		Ŀ.	. C	>A	6	à	C .	⊳G	<i>a</i>		,	C	>T	¢.
Detailed image	25µM									0								(a	) °
	50µM	¢	0																
220 um	75µM						•	0	0	0									
300 µm	100µ3	4.																	
	150µ)	4 '	-	6. j			•	•	•		•	•				•	•		•
			PM			• • •		C	A				c>0	;	•			01	



Table S2: The hybridization thermodynamics calculated for the selected sequences, with IDTBiophysics software

Sequences	ΔG <sub>37</sub> [kcal/mole]	ΔH [kcal/mole]	ΔS [cal/(K·mole)]					
Normal	-14.78	-194.30	-578.82					
C>A	-11.73	-178.50	-537.72					
C>G	-12.46	-184.80	-555.67					
C>T	-12.24	-182.40	-548.62					