

Use of DNA thermodynamics for low-abundance mutation detection by DNA hybridization

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Problem Statement

diagnostics, hybridization-based sensors In cancer detect mutant DNA by matching probes. The presence of a large amount of wild-type DNA lowers the detection sensitivity, as wild-type also binds to the probe.

Mutant (minority)

Wild-type (majority)

Mutant matching probe:



Sensor Design & Depletion

The OWL sensor consists of two DNA adaptor strands (R & P) and a universal molecular beacon (UMB). The 'fragile' structural rigidity of the P-strand provides **highly selective** binding to mutant DNA [1].



The sensor differentiates mutant DNA at room temperature in a wide temperature range. It has recently been equiped with additional strands, capable of unfolding secondary structures of sample DNA [2].

> Interactions are described by Langmuir theory [3]:



Blocker strands selectively

 c_{mut}

 $c_{\rm tot}$



Wild-type binding and experimental noise determines limit of detection. This limit is improved by the **technique of depletion**:

reduce DNA concentrations: Wild-type matching $c^*_{
m mut}$ Blocker **C***

Enhanced mutant ratio improves limit of detection

To test the OWL sensor and the effect of depletion, experiments were conducted on lung cancer biomarker EGFR. Results show that depletion greatly improves the limit of detection. As yet, the sensor allows discrimitation of mixtures containing more than 0.07% mutant DNA, reaching the accuracy of clinical methods. We are currently exploring the kinetics of the OWL sensor, which could provide a new detection method for mutant DNA.

mutant content c_{mut}/c_{tot}

Figure: Surface-based sensors show depletion results in a \sim 10-fold improvement in limit of detection [4].

References & Acknowledgement

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