## **Carnegie Mellon University** Materials Science & Engineering

presents

## Screening and Characterizing DNA-based Ligands for Gold Nanomaterials

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**ABSTRACT:** Oligonucleotide aptamers are single-stranded sequences that exhibit high affinity and specificity for a particular non-nucleotide target including, but not limited to small molecules, proteins, and even whole cells. Aptamers are conventionally isolated and identified using a multi-round screening approach called "Systematic Evolution of Ligands by EXponential Enrichment"

(SELEX) in which a pool of approximately  $10^9$  random candidate sequences is continuously enriched with amplified copies of "winning" sequences or adsorbates from prior selection rounds. While SELEX has revolutionized the discovery of numerous DNA and RNA-based aptamers for a variety of targets and dominated the field for two decades as a screening approach, we have developed a non-SELEX screening approach we call CompELS (Competition-Enhanced Ligand Selection) to identify single-stranded DNA aptamers for gold substrates. One of the key differences in our competition-based screening approach is the elimination of intermittent, timeintensive elution and amplification steps of random sequences that (1) can introduce undesired PCR side products (e.g. partially elongated duplexes) into the candidate pool and (2) bias the candidate pool towards early winners that may simply outnumber higher affinity aptamer candidates. Following aptamer selection against our gold-based target (e.g. planar crystalline gold, gold nanospheres, gold nanorods) we then evaluate sequences to identify base consensus motifs as well as structural elements such as hairpins, internal loops, and multibranched loops to reveal any shared patterns in the identified primary and predicted secondary structures of the ~20-40 identified aptamer sequences for a given gold target. Lastly, we have ranked our aptamer sequences for one of our gold targets in terms of their occurrence as a bound species using a high throughput sequencing method known as "deep sequencing" or next generation sequencing (NGS). As aptamers continue to be pursued as potential analogs and even substitutes for antibodies and other ligands in the broader materials community, we continue to adapt our unconventional screening approach to hopefully enable faster and easier aptamer identification for a rich range of material targets.

**BIOGRAPHY:** Valeria Milam received her B.S. in Materials Science and Engineering with Honors from the University of Florida in 1993. She specialized in metallurgy and also received a minor in Portuguese. For her doctoral studies at UIUC, she explored the phase behavior, structure and properties of nanoparticle-microsphere suspensions and reported a novel colloidal stabilization mechanism called nanoparticle "haloing." Her postdoctoral work at the University of Pennsylvania focused on DNA-mediated colloidal assembly. She joined the faculty at Georgia Tech in July 2004 as an assistant professor in the School of Materials Science & Engineering and was promoted to associate professor in July 2011. Current research efforts in the Milam group focus on oligonucleotides (1) as isothermal colloidal assembly and disassembly tools and (2) as synthetic ligands called aptamers that bind in a strong, but noncovalent manner to nonnucleotide targets. Her honors include a Georgia Cancer Coalition Distinguished Scholar Award, CETL/BP Teaching Award, AFRL Summer Faculty Fellowship, NSF CAREER, and 3M Nontenured Faculty Award.

