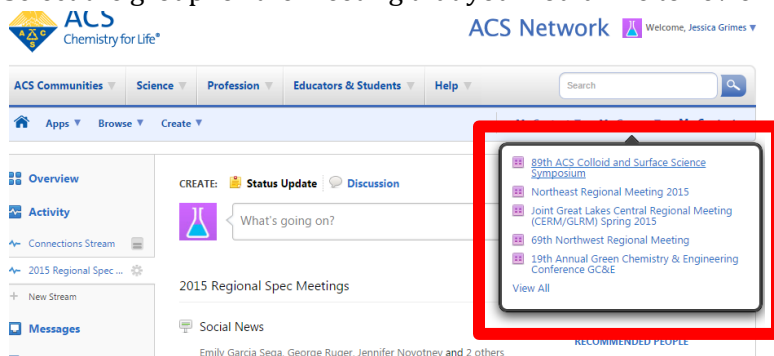


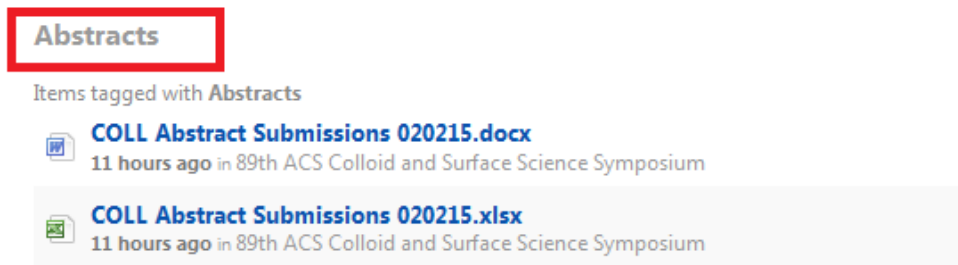
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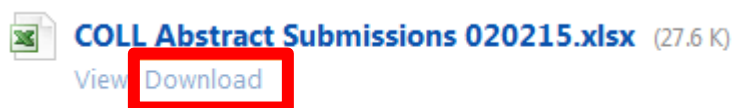
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CONTROL ID: 2194676

TITLE: [Self-assembly of linear peptide analogs and their interaction with lipid bilayers using MD simulations](#)

PRESENTER: Brian Novak

INSTITUTIONS (ALL): 1. Louisiana State University, Baton Rouge, LA, United States.

PRESENTATION TYPE: Oral Preferred

ABSTRACT BODY:

Abstract: The self-assembly and DPPC bilayer interaction of cationic linear peptide analogs (LPAs) with the formula $\text{Lys-NH}(\text{CH}_2)_n\text{CO-Arg-NH}(\text{CH}_2)_n\text{CO-Arg}$ with n varying from 4 to 13 has been studied using molecular dynamics simulations. These LPAs were designed to deliver an anionic phosphopeptide into cells, and experimental work on $n = 4, 7,$ and 11 LPAs[1-3] showed that: (i) LPAs induced changes in lipid phase behavior, (ii) the $n = 11$ LPAs caused the most structural changes to and leakage from DPPC, DPPC/DPPG, and DPPC/DPPS liposomes, and (iii) only the $n = 11$ LPAs were taken up by cells to a significant amount. Self-assembly of $n = 11$ LPAs in solution was also observed, and may be related to their activity. Our simulations showed that stable LPAs micelles formed in aqueous solution (0.1 M LPAs) for $n > 7$, LPAs in those micelles had increased β -sheet type structure consistent with experimental results, and that micelles with more than 8 molecules were cylindrical. The equilibrium depth of single LPAs in a DPPC bilayer increased with increasing n up to $n = 8$, but remained almost constant for $n > 8$. A 5 molecule, $n = 11$ LPA micelle sank into an 85% DPPC/15% DPPS bilayer, but remained intact and did not cause major disruption to the bilayer within about 350 ns. We will also present results of the behavior of $n = 11$ LPAs added to a DPPC bilayer one at a time.

[1] A. Gupta *et al.*, *Eur. Biophys. J. Biophys.* **40**, 727 (2011).

[2] G. F. Ye *et al.*, *Colloids Surf., B* **76**, 76 (2010).

[3] G. F. Ye *et al.*, *J. Med. Chem.* **50**, 3604 (2007).

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