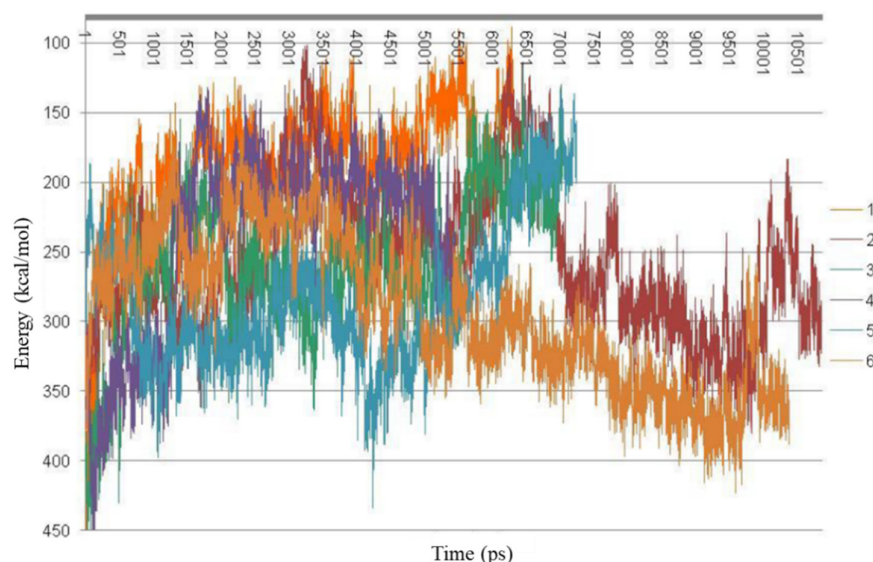


Article

# SpAD Biofunctionalized Cellulose Acetate Scaffolds Inhibit *Staphylococcus aureus* Adherence in a Coordinating Function with the von Willebrand A1 Domain (vWF A1)

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**Figure S1.** Comparison of binding energies for complexes 1-6. The low energy value for complex 6, which is also the most likely conformation for the interaction, is evident. The value of the final energy is very close to the energy provided by the zdock server (-370 kcal/mol).