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• Several eukaryotic cells have been shown to contain two histone fold structures that are homologous to a portion of the bacteriophage T4 integrase. A major goal of this project is to elucidate the structural and functional relationship between these histone fold proteins. How do they function in chromosomal

organization and gene regulation? This question will be explored in two different systems: (1) integration of the T4 bacteriophage genome into the *E. coli* chromosome, and (2) transcriptional regulation of specific genes in *E. coli* and *S. typhimurium*. The integration reaction and its regulation will be studied using fusion

proteins that contain both the bacteriophage integrase and a specific regulatory protein. By analyzing the structure of the various proteins as a function of time and temperature, a mechanism for the integration reaction and its regulation will be determined. Using antibodies to unique peptides of the bacterial

proteins, the presence of the histone fold proteins will be studied during both the integration and transcriptional regulation reactions. A new histone fold protein will be isolated from rat liver. The amino acid sequence will be determined and antibody against the protein will be prepared. The NMR structure will be established using standard

techniques. The primary sequence of the protein will be used to search for related proteins in other organisms. A 6d1f23a050

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