The successful candidate for this position will work with interdisciplinary teams of Researchers and  
Scientists on projects utilizing next generation high-throughput DNA sequencing. Duties include  
providing bioinformatics training, consulting with clients, providing applications support, as well  
as the design, development, and implementation of software to be employed in these endeavors. Due to the  
need to engage in multiple collaborations simultaneously, it is critical that this person have excellent  
organizational skills, manage commitments to projects effectively, and  
be able to negotiate concrete  
deliverables that will be completed in a timely manner. The above  
statements describe the general  
nature and level of work being performed by individuals assigned to  
this classification. This is not  
intended to be an exhaustive list of all responsibilities and duties  
required of personnel so  
classified.

Work requires at least 5 years of experience in bioinformatics. Experience with web-based bioinformatics tools, public domain biological databases and software tools for sequence, domain and structural analysis. Extensive familiarity with and development of computational tools in biology that use genomic data to generate biological hypotheses. Experience with a procedural language, proficient in Java, Perl, �C�, web design, DNA genome informatics, proteomics informatics, statistics, and computer science. Experience with relational databases and SQL helpful. OR AN EQUIVALENT COMBINATION OF RELEVANT