



March 18, 2015

Dear ASSEDA User:

For the past 2 years, we have asked registered users to support the Automated Splice Site and Exon Definition Server (ASSEDA) through a voluntary contribution. Some did subscribe ([link](#)), but many others simply kept renewing trial access under a series of aliases. If you are a repeat trial users of ASSEDA, please read on.

To generate the sequences you analyzed, you had to purchase reagents, pay technologists, equip a lab or pay a third party to sequence DNA or cDNA. ASSEDA is now being managed and updated by professional bioinformatics software developers who provide a valuable service to you. Their work cannot be delivered free of charge. We announced this change in service to all registrants by direct email and web marketing late last year.

ASSEDA has proven itself. After 10 years, and many updates and improvements in the webservice, we have summarized the considerable body of literature that has been published based on information theory in a recent acclaimed review article ([Caminsky et al. 2014](#)). By any reasonable set of criteria, the utility of this software has been amply demonstrated, and its value is proven for mutation analysis.

Trial access to ASSEDA is no longer available, now that the MutationForecaster system, which integrates this software, has been released. It is a part of commercial web service includes a suite of tools to help researchers quickly examine millions of DNA/mRNA variants simultaneously. These tools include ASSEDA, the Shannon splicing mutation pipeline, and Veridical (for mutation validation), VEP for coding sequence analysis, the Cytognomix User Variant Database, and our Genome Visual Analytics Decision Support Tool.

You may obtain a subscription (fee for either 2 months or 1 year) MutationForecaster at <http://mutationforecaster.com/signup.php>. Once you login, you will see an Account menu at the top of the window, where you can subscribe to Mutation Forecaster. Select this menu and you will be presented with either invoicing or PayPal options.

Upon acceptance of our terms of use and receipt of the subscription fee, your account will be activated. All of the components of the MutationForecaster system are already well documented, and the documentation is publicly available on the website. The website will be updated as new documentation is created. If you are a subscriber, we will alert you to these changes when they are introduced.

We plow all subscription funding back into the development of new algorithms and software components. For example, we are currently in the process of developing a completely automated workflow for genome scale coding and splicing mutation analysis and reporting.

We are confident that these and other capabilities that we will be introducing in new releases of the software will justify your Institute, Department, University or company investment in our product. We look forward to providing you with access to this new service, and we hope it will suit your growing needs for genome-scale variant analysis.

Kind regards,

A handwritten signature in black ink, appearing to read "Peter K. Rogan".

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