BaseSpace Variant Interpreter Beta Release Notes

BaseSpace Variant Interpreter Beta v1.0.12289

October 25, 2016

Introduction

These Release Notes detail new features, known issues, and recently resolved issues for the BaseSpace Variant Interpreter Beta Testing software. For details on how to operate BaseSpace Variant Interpreter Beta Testing software, see the online help, which is available from the Help icon in the application.

NEW FEATURES

- General user interface updates
  - Additional metadata, including IP Address and Workgroup fields, has been added to the audit log.

RESOLVED ISSUES

Issue Key	Issue Category	Description
BSVI-16925	Add New Analysis Result	A user-friendly error message displays when a duplicate subject is used.
BSKN-2194	Add New Association	Publication ID allows nonintegers to be used and creates an erroneous link to the PubMed website.
BSVI-17696	Analysis Result Metadata	User is able to save with sample class "Tumor-Normal."
BSVI-9300	Annotation	BaseSpace Variant Interpreter Beta does not validate the search criteria of a chromosome as a user enters an actual position.
BSVI-9320	Annotation	When the custom field is left blank for a given row, the system still annotates the given VID.
BSVI-17842	BSSH	Users can select a workgroup in BaseSpace Sequence Hub and redirect to VI upon successful confirmation.
BSVI-16168	Filters	Users can use the Saved Filters drop-down menu to apply a saved filter from a different reference genome.
BSVI-15146	Interpretation	The Interpretation History table is updated with the interpretation Modified Date.
BSVI-17861	Manifest	If a user ingests a manifest referencing BAM path and the AR is still in processing, the link is not available.
BSVI-17531	Registry	When using the sample metadata sheet to upload 40 or more cases, the software might freeze.
BSVI-11052	Report	The inheritance mode content extends off the report page.
BSVI-17837	Variant Grid	An error banner appears while the Variant Grid is loading.
BSVI-16351	Workgroups	If a workgroup is removed, users in the workgroup are defaulted to their personal workgroup.

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Issue Key	Issue Category	Description
BSVI-17814	Workgroups	Users may experience issues when connecting to BaseSpace Sequence Hub for the first time. When prompted, the user should select their Personal workgroup.
BSVI-17822	Workgroups	After changing workgroups, the user may need to perform a browser refresh to update the data.

KNOWN ISSUES

Issue Key	Issue Category	Description
BSKN-2898	Add New Association	Clinical Trial open and close dates can appear one day earlier depending on the time zone the association was created in.
BSVI-16523	Filters	When analyzing a hg19 sample, a saved hg38 region-based filter can be incorrectly viewed in the saved filter drop-down menu.
BSVI-12422	Import	If a user tries to upload a VCF file that is not properly formatted, the upload fails with an ambiguous error message. The file shows a status of fail, and the user can then delete it. Consequently, the variant grid does not appear.
BSVI-16408	Import	Uploading of multiple Grch38 .vcf files occasionally results in an error. The error can be cleared by deleting the failed upload and retrying.
BSVI-16384	Variant Details	ClinVar data for the same variants can appear to differ between the variant summary in the grid (which reports both SCV and RCV entries) and the variant detail page (which reports only RCV).
BSVI-15768	Variant Grid	For variants with multiple dbSNP IDs, only 2 of the IDs are displayed in the variant grid.
BSVI-18409	Variant Grid	The validation message for applying Tumor-Normal specific filters to a Germline analysis result is not displayed.
ON-345	Variant Grid	Autocomplete box shows results marked as 'obsolete' by nomenclature authority.
ON-432	Variant Grid	Expansion to related phenotypes sometimes yields results that are too distant.
ON-516	Variant Grid	Results in autocomplete pop-up might be sorted inconveniently (subjective).
WAC-546	Workgroups	Invited users who are not currently registered are sent an invitation email and marked as pending, but their status is not visible in the user interface. To confirm that the invitation has been received, users can contact the invitee directly, independent of BaseSpace Variant Interpreter (Beta).

Issue Key	Issue Category	Description
WAC-730	Workgroups	In some instances, the bulk edit of user permissions checkboxes does not work. Log out/log in to resolve this, or edit users individually.