

Axiom™ Analysis Suite (AxAS) v5.4

APT version

AxAS 5.4 is associated with **APT 2.12**.

New analysis features and changes

- When adding CEL files to a new analysis, Array type will now auto-populate for arrays that have only one version of library file package.
- AxAS Best Practices and Genotyping Analysis now includes **remoteCN** when the library package includes **remoteCN** genotyping.
- Cluster Plot **CopyID(s)** now copies **probeset_id** in addition to Sample Filename.

Known issues

Known issues and behaviors when using the General Analysis **Inbred File | Value** option include:

- An Inbred file is still accepted even if it contains an invalid value.
- If a Value and a File are supplied for Inbred, the File is used even if the **Value** check box was checked.
- If the **Inbred** check box was checked prior to restoring an Analysis configuration, it remains checked.
- **Percent Concordance** is displayed with full (5 decimal) precision, not 3 decimals (its default).
- In certain Windows 10 environments, the **Export IGV | Output File** Prefix field may be partially hidden after sample attributes are loaded.