

The Cancer Genome Atlas Pilot Project
Broad Institute of MIT & Harvard
Raw Data Description

Platforms: Affymetrix SNP 6.0 Array & Affymetrix HTA Array

Raw Data

The CEL file stores the results of the intensity calculations on the pixel values of the DAT file. This includes an intensity value, standard deviation of the intensity, the number of pixels used to calculate the intensity value, a flag to indicate an outlier as calculated by the algorithm and a user defined flag indicating the feature should be excluded from future analysis. The file stores the previously stated data for each feature on the probe array.