

This module was created for the Cancer Proteogenomics Workshop held at the Rajiv Gandhi Center for Biotechnology in India during Sep 26-30, 2016.

The modules capture analysis algorithms for analysing proteome, phosphoproteome and genome data, specifically those used for the Mertins, Mani, et. al., *Nature* paper. See workshop hands-on material for use and documentation.

Contact D. R. Mani (manidr@broadinstitute.org) with questions or comments.