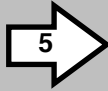
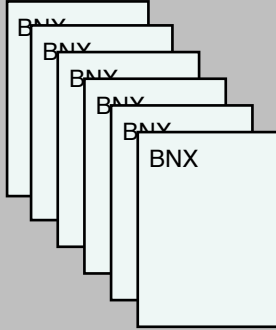
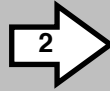
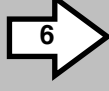


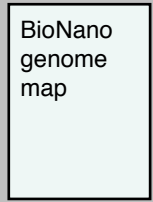
Autodetect: Converts Irys tiff files to BNX files



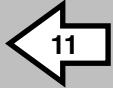
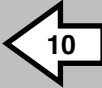
AssembleIrysCluster: uses *in silico* genome map to prep assemblies



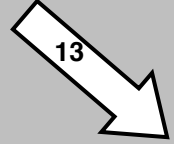
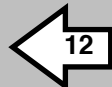
Assembler: assembles molecules



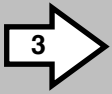
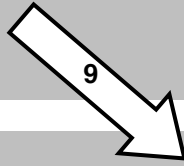
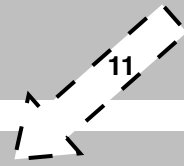
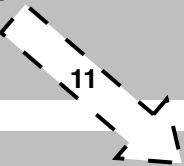
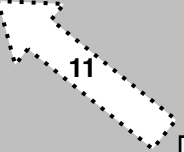
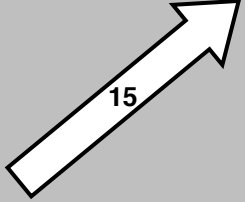
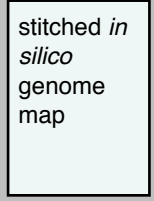
RefAligner: aligns *in silico* and BioNano genome maps



stitch: uses XMAP, scaffold and *in silico* genome maps to super scaffold



fa2cmap_multi: *in silico* nicks stitched NGS scaffolds



fa2cmap_multi: The genome scaffolds are *in silico* labeled

