Release Notes

# New features

- mountcb.m: add an existing CellBase to your computer (by Tamas Laszlovszky)

- updatecb.m: allows changing CellBase preferences (by Balazs Hangya)

- mclust\_projection\_preview: plot feature data for clusters

- viewlick.m: lick-raster similar to spike raster (viewcell2b.m)

# Changes

- export\_fig.m has been updated to maintain compatibility with ghostscript

- setvalue.m now supports list of CellIDs as input argument

- Nlx converter was updated in the AddOns folder

- many functions support Open Ephys

# Bug fixes

- acg.m and ccg.m

- cellid2fnames and fnames2cellid

- MakeTrialEvents2\_gonogo.m: TTLs parsed according to their channels

- findSegs2.m contained many bugs; findSegs3.m added, with hopefully most of the bugs eliminated; findSegs3.m renders findSegs2.m obsolete

- addanalysis.m: output arguments of an added analysis do not have to be wrapped in a cell array

- many minor bug fixes