GRIDSS wrong_INFOtags

Legend: # samples — 3, 2, 1, 0

Column annotations: species, typeSimulations

Row labels (top to bottom):
- CLOVE max_rel_coverage_del=0.1
- CLOVE max_rel_coverage_del=0.10000000000000009
- CLOVE max_rel_coverage_del=0.6000000000000001
- CLOVE min_rel_coverage_dup=1.3
- CLOVE min_rel_coverage_dup=1.5
- CLOVE min_rel_coverage_dup=1.6
- CLOVE min_rel_coverage_dup=1.8
- GRIDSS filter_noReadPairs=False
- GRIDSS filter_noReadPairs=True
- GRIDSS filter_overlappingRepeats=False
- GRIDSS filter_overlappingRepeats=True
- GRIDSS maximum_lenght_inexactHomology=200
- GRIDSS maximum_lenght_inexactHomology=1000
- GRIDSS maximum_lenght_inexactHomology=10000000
- GRIDSS maximum_microhomology=200
- GRIDSS maximum_microhomology=1000
- GRIDSS maximum_microhomology=100000000
- GRIDSS min_Nfragments=5
- GRIDSS min_Nfragments=8
- GRIDSS min_Nfragments=10
- GRIDSS min_Nfragments=30
- GRIDSS min_af=0.05
- GRIDSS min_af=0.1
- GRIDSS min_af=0.2
- GRIDSS min_af_EitherSmallOrLargeEvent=0.05
- GRIDSS min_af_EitherSmallOrLargeEvent=0.2
- GRIDSS range_filt_DEL_breakpoints=(0, 1)
- GRIDSS range_filt_DEL_breakpoints=(50, 900)
- GRIDSS range_filt_DEL_breakpoints=(200, 700)
- GRIDSS wrong_FILTERtags=('LOW_QUAL',)
- GRIDSS wrong_FILTERtags=('NO_ASSEMBLY',)

species_simulations