CCF (**ROUGH**)

Ploidy 2; Purity 0.74; n = 1927 mutations in 31 segments; 10 fragmented arms

n = 1926

QC PASS p_{PASS} = 0.932

n = 1925; Tail 18%, C1 82%

MM (25 steps; ε = 7e−05), z_{nk} > 0

n = 1627, BBin 1 100%

2021−01−31 17:15:26. evoverse pipeline for deconvolution from CCF: with fits TRUE