6 subgroups, 10 signatures (100.0%) with fdr < 0.05 2522 confident samples Prob Purity Linear_Amplicons z-score 40 0.8 30 0.6 20 0.5 0.4 10 0.2 pboâse_mean Class Telomere_Content(log2) 154 12₀ Ploidy 2 3 5 -4 6 Sig3 rel_diff Sex **Age** 100 LOHs female Sig9 800 male 600 0.5 Sig4 60 Tumor_Stage 400 200 Sig11 Status (barplots) Sig6 confident **Fusions** Unknown 150 ambiguous Sig1 2000 100 WGD_Status 1500 Sig7 no_wgd wgd 1000 50 500 Sig2 0 HRD_Status Chromothripsis Sig10 **DELs** hrd no_hrd **2**0 200 15 Sig8 150 10 100 Age 50 **AMPs** 0 DELs **BFBs** Affected_Chrs Affected_Chrs 25 20 CNA_Burden Purity 15 Ploidy 10 LOHs ecDNAs **Fusions** Chromothripsis CNA_Burden **BFBs** ecDNAs 0.5 HRs Linear_Amplicor HRs Telomere_Content(log2) Sex Tumor_Stage WGD_Status HRD_Status