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Signature	Representative sample profile	Description	Potential mechanism
scSig1	10	(A) Mainly small and medium-sized DNA segments (0-5Mb); (B) Large sized (>5Mb) CN amplification (mostly 5 copies); (C) The signature is associated with ecDNA.	EcDNA
scSig2	2	 (A) Large sized (>5Mb) CN amplification (mostly 4 copies); (B) Adjacent CN change is within 2 copies; (C) The signature is associated with WGD. 	WGD
scSig3	3 2 1 0	(A) Long DNA segments (>10Mb) without copy number alteration (CN = 2).	Normal genome
scSig4	5 4 3 2 2 1 0	(A) Mainly large sized (>5Mb) DNA segment (CN = 3);(B) Adjacent CN change is within 2;	Triploid chromosome
scSig5	5	 (A) Many localized high CN amplification (copies ≥ 5, size<5Mb); (B) Adjacent CN change is irregular; (C) The associated with chromothripsis. 	Chromothripsis
scSig6	5 4 3 2 1 0	(A) Mainly large sized (>5Mb) DNA segment localized amplification and deletions; (B) The associated with LST (HRD).	LST (HRD)
scSig7	4 3 2 1 0	 (A) Mainly small DNA segments (0-5Mb) with localized low amplification (CN = 3); (B) Adjacent CN change is within 2; 	Unknown

Abbreviations: EcDNA: Extrachromosomal circular DNA; WGD: Whole Genome Duplication; HRD: Homologous Recombination Deficiency; LST: Large-scale State Transitions