

Tissue	Template	Target Enrichment	Calling Method	Majority STR Type	Targets	Informative Reads	Purpose	Refs
human blood	Bulk	multiplex PCR	Capillary Electrophoresis	hexa-	~20	--	Forensic	1
Human	Bulk	Array capture	Next Generation Sequencing	all types	7851	7%	Mutation Discovery	2
human blood	Bulk	RNA Probes	Next Generation Sequencing	tri- and longer	10764	40%	Mutation Discovery	3
Athaliana	Bulk	MIPs	Next Generation Sequencing	tri- and hexa-	102	55-64%	Evolution phylogeny	4
human leukemia	scWGA	multiplex PCR	Capillary Electrophoresis	di-	128	--	Lineage Reconstruction	5
human cancer	scWGA	Access Array	Next Generation Sequencing	di-	~2000	90%	Lineage Reconstruction	6
human cancer/normal	scWGA	duplex MIPs	Next Generation Sequencing	di-, mono-	12473	96%	Lineage Reconstruction	

#### STR capture methods summary