

Supplementary Table 1. Short tandem repeat (STR) profile, based on eight STR loci (number of repeats at each locus is indicated), of the original tissues from which the cell lines were derived*

Primary tissue	Short tandem repeat locus							
	D21S11	TH01	D3S1358	FGA	TPOX	D8S1179	vWA	D5S818
FLO-1	30, 32.2	6, 9.3	15	21	9, 11	13	16	12, 14
KYAE-1	NA	6, 9	15, 16	18	8	NA	14	10, 13
OE19	28, 30	8, 9	15, 18	23, 26	8	13, 15	17, 18	11, 14
OE33	NA	7, 8	18	22, 23	8, 11	10, 12	14, 17	11
OACM5.1	NA	6, 9.3	16, 17	22	8	13, 14	19	12
OACP4C	NA	6, 9	14, 18	20	9, 11	13, 14	16	9, 12
SK-GT-4	NA	6, 9.3	16, 17	22, 23	8, 10	NA	17	12
JH-EsoAd1*	NA	6, 7	NA	NA	8, 9	NA	18, 19	11
ESO26	30, 31.2	9	15	21	7, 9	13, 14	14, 18	11, 12
ESO51	30, 33.2	9	15, 16	21, 22	8, 11	10, 11, 12	14, 15	11, 13

*JH-EsoAd1 repeat number at additional STR loci; CSF1PO: 9, 10; Amelogenin: X, Y; D13S317: 11; D16S539: 10, 12; D7S820: 10, 12. NA = no available data due to technical failure.