

	ZNF143					ZNF143-HA	ZNF76	ICN1
	HeLa rep1	HeLa rep2	HPB-ALL	K562	293T-rex	FLP143-HA	FLP76	HPB-ALL
Number of reads	25680120	27890160	30950880	31759080	33242640	26253249	26523782	44174782
Mapped reads	20140058	22064111	24563873	24630672	26617590	19908969	20025151	33293640
Peaks	4250	2388	2823	6329	2570	4071	3134	18686
Peak at <= 2kb from TSS	2502	1632	1886	2797	1518	2016	1424	8467
Peak at > 2kb from TSS	1748	756	937	3532	1052	2055	1710	10219

	Zfp143		
	mES cells	NIH3T3	MEF
Number of reads	30617537	36603053	27806809
Mapped reads	22023766	24518391	19759166
Peaks	6119	5453	2736
Peak at <= 2kb from TSS	2150	2744	1697
Peak at > 2kb from TSS	3969	2709	1039

Table S1 : Total number of reads obtained from sequencing. Number of reads mapped on human hg19 and mouse mm9 genomes. Number of peaks obtained with MACS software (p-value of 10⁻⁵ and minimum height of 20 reads for all except ICN1; For ICN1 p-value of 10⁻⁵, FDR of 0.01 and minimum height of 50 reads).