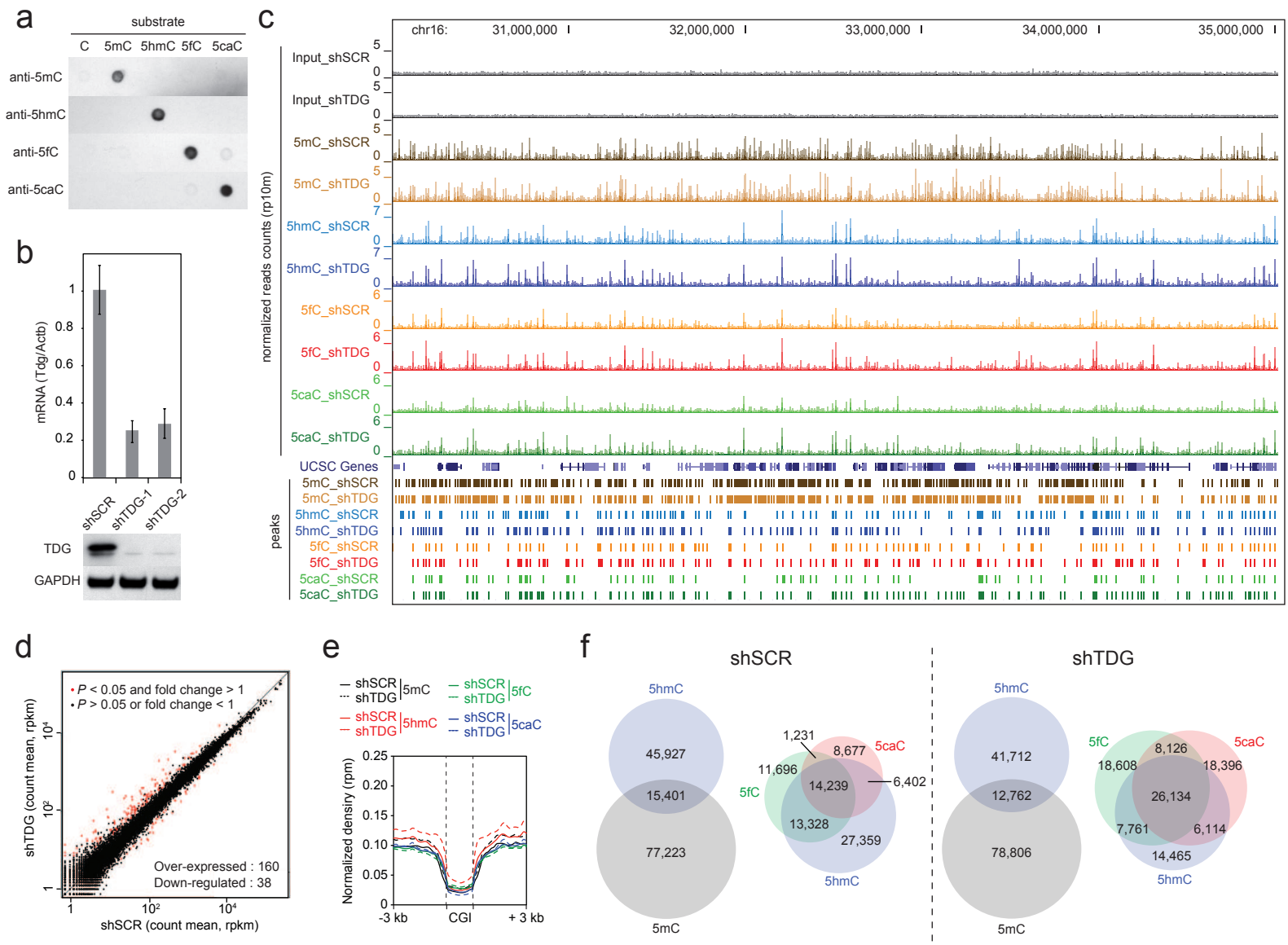

g

Family	Size (Mb)	% genome	Sub-family	Size (Mb)	% genome
LTR	273.6	10.1	IAP_{int}	119.6	4.4
			IAP_{ext}	27.9	1.2
LINEs	522.3	19.3	L1Md	172.4	6.4
			(> 5 kb)	(75.7)	(2.8)
SINEs	202.9	7.5	B1m+B2m	45.8	1.7
Simple repeats	63.4	2.3	CA repeats	19.5	0.7
Major Satellites	7	0.3			
DNA transposons	21.8	0.93			

Supplemental_Fig.1