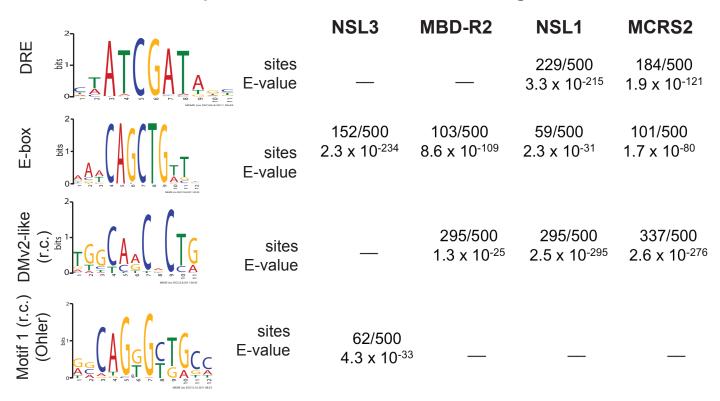
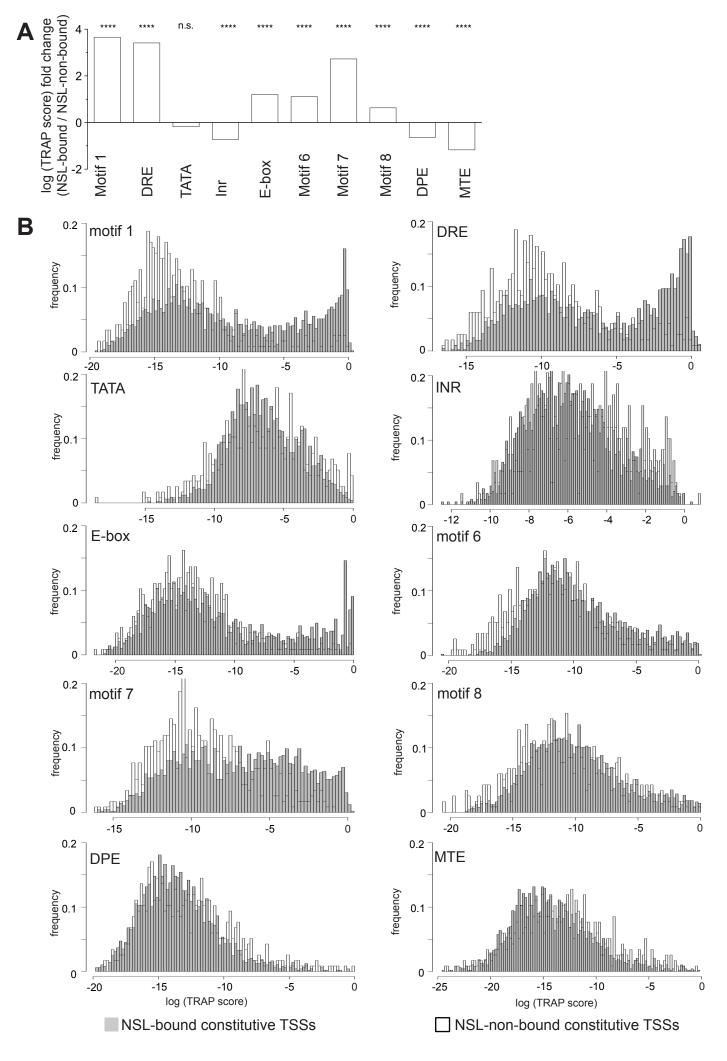


A Peak summits with tag count ranks 500 - 1,000

		NSL3	MBD-R2	NSL1	MCRS2
DRA SEE TO SEE T	sites E-value	_	_	183/500 6.5 x 10 ⁻¹⁸⁸	123/500 3.4 x 10 ⁻¹⁰⁴
WIND THE REPORT OF THE REPORT	sites E-value	117/500 3.4 x 10 ⁻¹¹⁵	83/500 1.3 x 10 ⁻⁵⁴	77/500 1.7 x 10 ⁻⁶¹	90/500 8.3 x 10 ⁻⁹¹
DMA2-like Company of the control of	sites E-value	_	_	263/500 3.9 x 10 ⁻¹⁸⁹	222/500 6.4 x 10 ⁻¹⁴⁶
Motif 7 (r.c.) Ohler Ohler The state of t	sites E-value	64/500 4.2 x 10 ⁻³⁵	63/500 1.7 x 10 ⁻⁴²		_

B Peak summits of peaks associated with constitutive genes





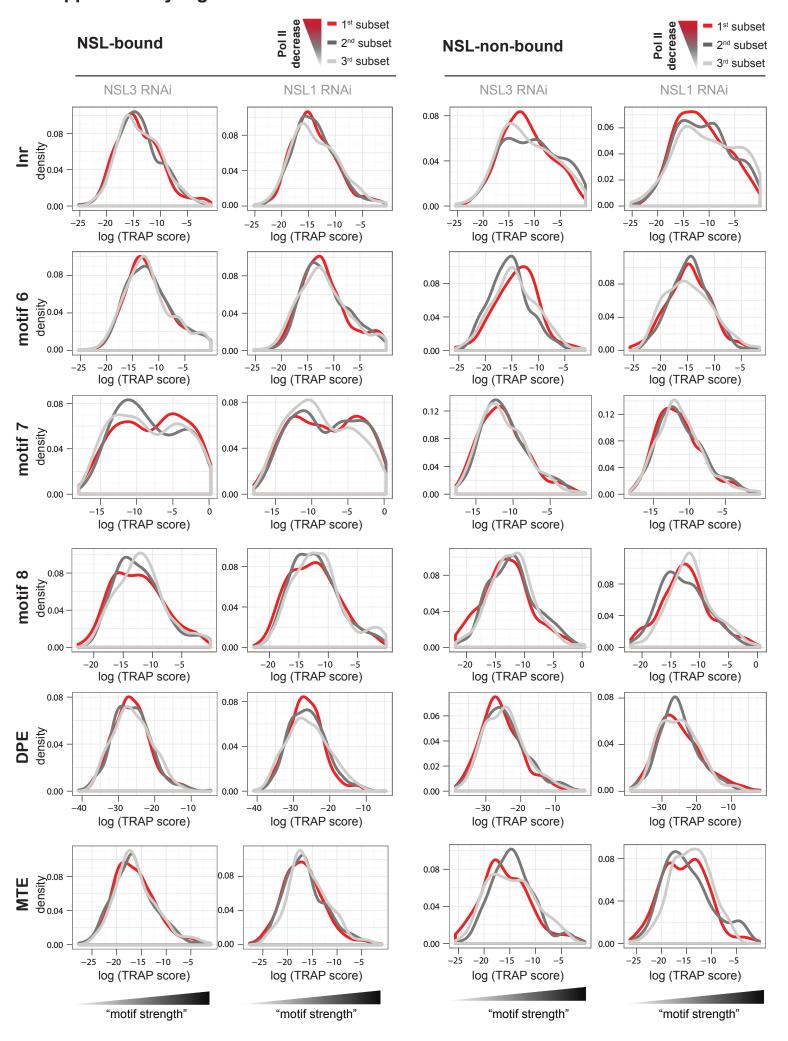


Table S1

		NSL1	MCRS2	NSL3	MBD-R2
peaks (MACS)		3,541	3,733	4,244	3,399
median peak size		1,129 bp	1,314 bp	2,227 bp	2,676 bp
(sub)peaks (PeakSplitter)		11,797	12,234	9,409	9,957
subpeaks per peak (median)		3	3	2	2
median peak size		350 bp	379 bp	1,110 bp	1,040 bp
TSSs bound	general	28.7 %	28.9 %	36.8 %	36.1 %
	active	55.7 %	54.1 %	68.3 %	65.9 %
	constitutive	63.2 %	63.2 %	77.9 %	75.2 %