

% <i>Golli</i> / <i>Gapdh</i> in spinal cord of mice										
	P7		P14		P21		P30		P90	
Mouse line	% ± SEM	n(F:M)	% ± SEM	n(F:M)	% ± SEM	n(F:M)	% ± SEM	n(F:M)	% ± SEM	n(F:M)
WT	100 ± 3%	3:3	100 ± 3%	3:3	100 ± 4%	3:3	100 ± 11%	3:3	100 ± 3%	3:3
M3KO	7 ± 0% **	3:3	7 ± 0% **	2:2	8 ± 0% **	3:3	9 ± 1% **	2:2	11 ± 1% **	2:4
M3(225)KO	-	-	90 ± 8%	3:3	-	-	141 ± 5% **	3:3	66 ± 4% **	0:5
M4KO	-	-	97 ± 5%	2:2	-	-	114 ± 10%	2:2	94 ± 4%	8:3
M5KOΔ3.6kb	96 ± 3%	3:3	95 ± 2%	3:3	101 ± 4%	1:4	112 ± 6%	3:3	109 ± 6%	2:3
M5KOΔ1kb	-	-	90 ± 2%	3:3	-	-	-	-	-	-
M3M5KO	7 ± 0% **	3:3	7 ± 0% **	2:2	10 ± 0% **	3:3	8 ± 0% **	3:3	12 ± 0% **	3:3
M1EM3M5KO	-	-	7 ± 0% **	3:2	8 ± 1% **	4:1	9 ± 0% **	4:2	-	-

S3 Table. Relative *Golli* mRNA accumulation in spinal cord of enhancer knock-out mice at P7, P14, P21, P30 and P90. The values are presented as % ± standard error of the mean. “***” represent p-values ≤ 0.01. n(F:M) represents number of Female and Male mice analyzed.