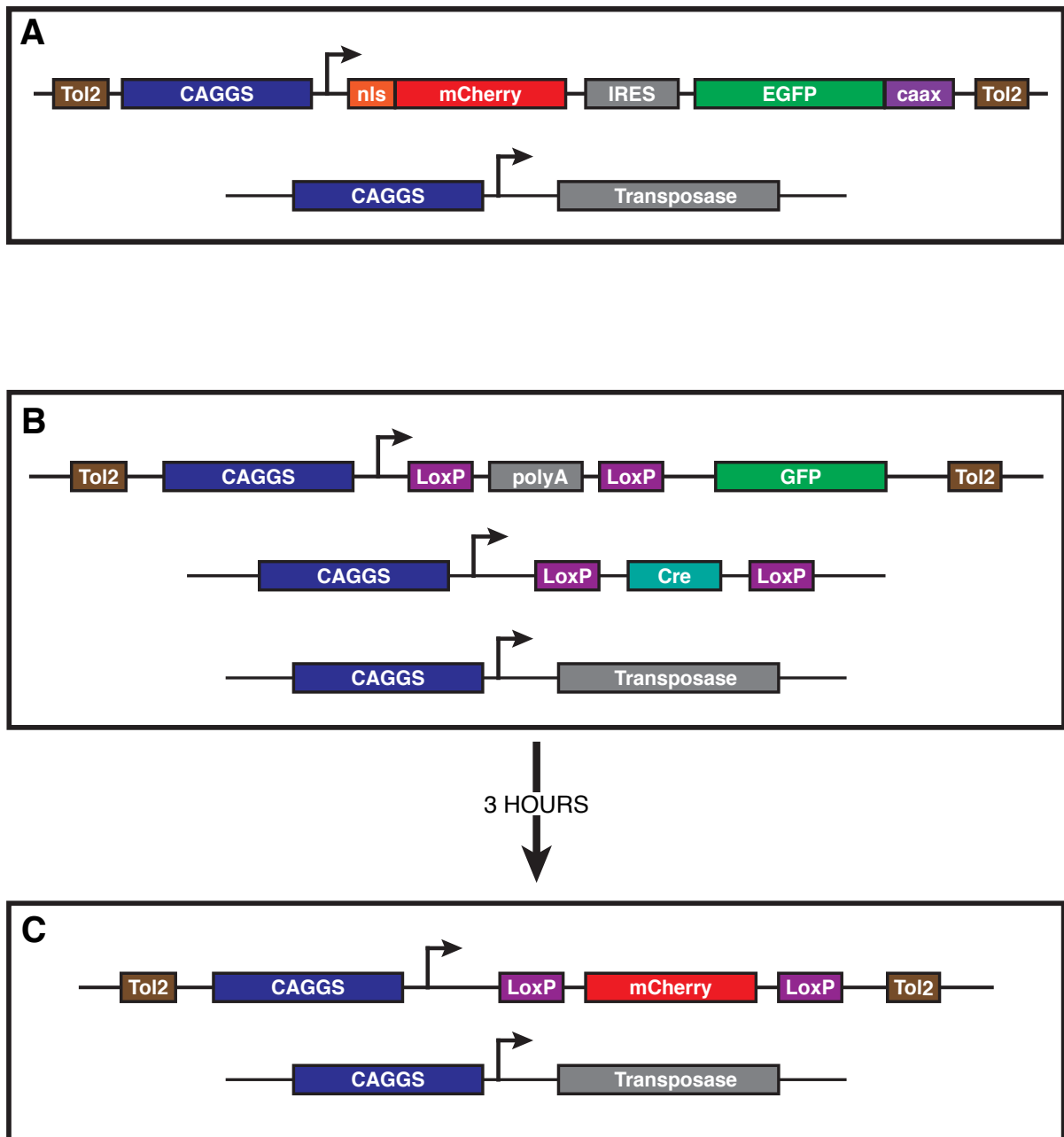


### Supplementary Figure 1.

Dorsal and ventral muscle masses of a single E5.5 chick embryo forelimb co-electroporated at E2.5 in the VLL of somites in the forelimb region with EGFP (in green) and a nuclear form of mCherry (in red). In blue, immuno-staining against MyHC. This shows that the distribution of plurinucleated myofibers is similar in both muscle masses (longest fibers in the central domain, shortest at the periphery), while a cloud of mononucleated progenitors is present at the distal leading edge of both muscle masses.



**Supplementary Figure 2.**

(A): Vectors utilized to count the number of nuclei per fiber. (B,C): Description of the double-electroporation protocol aimed at mosaically labelling a cell population with two fluorochromes. B: First co-electroporation with two plasmids, that encode a self-excising Cre and GFP, downstream of a floxed poly(A) signal. When both plasmids are co-expressed, epithelial cells are GFP-positive. If only one plasmid is expressed, cells are not fluorescent. (C): 4 hours later, the DML is electroporated once more with a plasmid encoding a floxed mCherry. DML cells express the mCherry fluorochrome, unless they were previously electroporated with Cre, which inactivates mCherry. The self-excising Cre ensures that the Cre protein is only temporarily present; if mosaically-labeled cells do not fuse, myocytes will be either red or green, if they fuse, a proportion of them will be yellow.

**Supplementary Table 1.**

Count of nuclei per fiber observed at the indicated times after DML, dorsal dermomyotome and limb electroporation.

	E3.5	E4	E4.5	E5	E5.5	E6
<b>DML EP</b>						
	n=189		n=353		n=336	
<b>1N</b>	176		265		68	
<b>2N</b>	12		75		98	
<b>3N</b>	1		10		80	
<b>4N</b>			3		44	
<b>5N</b>					33	
<b>6N</b>					11	
<b>7N</b>					2	
<b>Mean</b>	1.1		1.3		2.8	
<b>Dm EP</b>						
			n=186		n=223	
<b>1N</b>			80		40	
<b>2N</b>			71		54	
<b>3N</b>			24		58	
<b>4N</b>			10		41	
<b>5N</b>			1		21	
<b>6N</b>					7	
<b>7N</b>					1	
<b>8N</b>					1	
<b>Mean</b>			1.8		2.9	
<b>LIMB EP</b>						
	n=115	n=59	n=207	n=238	n=128	
<b>1N</b>	103	47	37	4	6	
<b>2N</b>	9	11	50	21	7	
<b>3N</b>	3	1	30	24	9	
<b>4N</b>			32	31	15	
<b>5N</b>			15	25	15	
<b>6N</b>			15	36	14	
<b>7N</b>			11	25	12	
<b>8N</b>			4	18	9	
<b>9N</b>			5	21	7	
<b>10N</b>			3	12	7	
<b>11N</b>			1	8	12	
<b>12N</b>				2	12	
<b>13N</b>				1	2	
<b>14N</b>					1	
<b>15N</b>				1	1	
<b>16N</b>			1	1		
<b>17N</b>						1
<b>Mean</b>	1.1	1.2	3.6	6.1	6.7	

**Supplementary Table 2.**

GFP (green), mCherry (red) and mixed (yellow) fibers after electroporation of the indicated somite sub-populations.

<b>DML/DML 10hpe no Cre</b>				
	<b>Green</b>	<b>Red</b>	<b>Yellow</b>	
<b>Fibers Counted</b>	48	26	101	
<b>Percentage Mean</b>	26.7	14.7	58.6	
<b>Standard Deviation</b>	8.1	6.3	13.0	
<b>DML/DML 10hpe control</b>				
	<b>Green</b>	<b>Red</b>	<b>Yellow</b>	
<b>Fibers Counted</b>	226	215	24	
<b>Percentage Mean</b>	46.9	49.7	3.4	
<b>Standard Deviation</b>	18.2	18.8	4.7	
<b>DML/DML 72hpe</b>				
	<b>Green</b>	<b>Red</b>	<b>Yellow</b>	
<b>Fibers Counted</b>	193	150	49	
<b>Percentage Mean (Adj.)</b>	49.7	42.2	8.2	
<b>Standard Deviation</b>	7.6	13.2	6.6	
<b>Dm/Dm 72hpe</b>				
	<b>Green</b>	<b>Red</b>	<b>Yellow</b>	
<b>Fibers Counted</b>	83	69	88	
<b>Percentage Mean (Adj.)</b>	36.7	30.2	33.1	
<b>Standard Deviation</b>	4.6	7.3	6.4	
<b>DML/Dm 72hpe</b>				
	<b>Green</b>	<b>Red</b>	<b>Yellow</b>	
<b>Fibers Counted</b>	91	77	28	
<b>Percentage Mean (Adj.)</b>	48.0	42.7	9.3	
<b>Standard Deviation</b>	8.1	8.2	11.2	
<b>DML/AL 72hpe</b>				
	<b>Green</b>	<b>Red</b>	<b>Yellow</b>	
<b>Fibers Counted</b>	78	66	42	
<b>Percentage Mean (Adj.)</b>	43.3	36.6	20.1	
<b>Standard Deviation</b>	11.3	8.8	7.8	
<b>DML/PL 72hpe</b>				
	<b>Green</b>	<b>Red</b>	<b>Yellow</b>	
<b>Fibers Counted</b>	78	64	38	
<b>Percentage Mean (Adj.)</b>	45.1	37.1	17.8	
<b>Standard Deviation</b>	5.4	5.5	5.1	