Table S1. Quantification of effect of MOs and mutants								
Figure number and panel	Con	mef2d/c MO	mef2c/d MO	mef2c MO	mef2d MO	mef2c MO+mef2d MO	hoo+/tn213×hoo+/tn213	hoo+/tn213×hoo+/tn213+mef2d MO
2A	0/1178 (0%)	894/976 (92%)	298/325 (92%)	3/162 (2%)	15/204 (7%)	86/122 (70%)	0/369 (0%)	112/159 (70%)
2C	0/38 (0%)	45/47 (96%)						
2D	35/35 (100%)	19/19 (100%)						
2C,D, 3C-E, 6A, S1F, S3	0/290 (0%)	370/396 (93%)	166/181 (92%)	0/83 (0%)	6/66 (9%)	65/102 (64%)	0/161 (0%)	71/114 (62%)
2F	25/25 (100%)	28/28 (100%)						
2H 3A	8/77 (10%) 73/73 (100%)	112/120 (93%) 60/60 (100%)						
3B	78/78 (100%)	77/77 (100%)	19/19 (100%)	20/20 (100%)	21/21 (100%)	15/15 (100%)		
3F	0/22 (0%)	30/31 (97%)	13/13 (100 /0)	20/20 (100 /0)	21/21 (100/0)	13/13 (100 /0)		
3G	0/173 (0%)	70/74 (95%)						
5A	29/29 (100%)	25/25 (100%)						
5B, S4A	0/135 (0%)	214/222 (96%)		0/18 (0%)	1/22 (5%)	22/24 (92%)		
5C	0/65 (0%)	40/40 (100%)						
5D	0/38 (0%)	35/41 (85%)						
5E	103/103 (100%)	35/35 (100%)						
5F	61/61 (100%)	54/54 (100%)						
5G, S4C	146/146 (100%)	83/83 (100%)	26/26 (100%)	23/23 (100%)	24/24 (100%)	26/26 (100%)		
5H	22/22 (100%)	13/13 (100%)						
6A	0/20 (0%)	22/22 (100%)						
6B,F	0/12 (0%)	12/12 (100%)						
6C 6D	0/10 (0%) 0/18 (0%)	13/13 (100%) 17/20 (85%)						
6E,F	0/18 (0%)	19/19 (100%)						
S1A	42/42 (100%)	13/13 (100/0)		16/17 (94.1%)	0/17 (0%)			
S1B	30/30 (100%)			17/17 (100%)	0/12 (0%)			
\$1C	30/30 (100%)			0/17 (0%)	12/12 (100%)			
S1D (left)	39/39 (100%)				39/39 (100%)			
S1D (right)	19/19 (100%)			38/38 (100%)	, , , , , , , , , , , , , , , , , , , ,			
S1E,F, S3	0**/129 (0%)	192*/198 (96%)	29*/34 (85%)	0**/73 (0%)	6**/56 (11%)	40**/57 (70%)	0**/83 (0%)	10*/37 (27%) 21**/37 (57%)
S2A	0/52 (0%)							
S2B,C	0**/57 (0%)	29*/36 (81%)	18*/23 (78%)	27*/27 (100%)	0**/22 (0%)	17*/20 (85%)	13*/48 (27%) 23**/48 (48%)	5*/15 (33%) 6**/15 (40%)
S2D	0/34 (0%)			0/27 (0%)	22/22 (100%)	17/20 (85%)		
S4B	1**/32 (3%)				0**/28 (0%)		1**/32 (3%)	19*/64 (30%) 33**/64 (52%)
Con, Control. 2A Fraction of 24- to 28-hof em	hrvos with ventral curvati	ure and motility defects						
2A, Fraction of 24- to 28-hpf embryos with ventral curvature and motility defects. 2C, Fraction of embryos with co-localised slow and fast MyHC in slow fibers and absence or strong reduction of fast MyHC in fast fibres.								
2D, Fraction of Tg(acta1:GFP) embryos with normal GFP.								
2.C.E, 3.C.E, 6.A, 5.IF, 53. Fraction of 22- to 26-hpf wild-type, Tglacta1:GFP) or myogenin:GFP-injected embryos with immature slow fibres. See Fig. 3E for quantification of hoo mild and severe proportions. 21.H, Fraction of 24-hpf embryos with uprepulated myogenin mRNA expression in anterior somities.								
3A,3B,2F, Fraction of 15s embryos with normal expression of all-MyHC, smyhc1 and myhz1, respectively.								
3F, Fraction of 15s embryos with upregulated adaxial prdm/ mRNA expression in anterior somites. 3G, Fraction of 24s embryos without eng2e mRNA expression in slow or fast fibres.								
ss, raction of 24-bit emptys without eng2 a mixua expression in siow or last tipres. SA, Fraction of 24-bit embryos with normal or purequilated sin somites.								
5B, S4A, Fraction of 24-hpf embryos with reduced levels of my/n21 mRNA in somites.								
SC, Fraction of 22s embryos with reduced levels of mylz2 mRNA in somites.								
SD, Fraction of 24-hpf embryos with abolished expression of smbpc mRNA in somites. SE, Fraction of 21s embryos with normal expression of actar mRNA in somites.								
5F, Fraction of 24-hpf embryos with normal or upregulated <i>tpma</i> mRNA in somites.								
5G, 94C, Fraction of 24-hpf embryos with normal trinc mRNA in somites. 5H, Fraction of 24-hpf embryos with normal trinc mRNA in somites. 5H, Fraction of 24-hpf embryos with normal or upregulated has 9500 emRNA in somites.								
SH, Fraction of 24-hpt embryos with normal or upregulated nsp540a mRNA in somites. 6A, Fraction of 24-hpt embryos with dispersed/punctate localisation of actin.								
6B,F, Fraction of 24-hpf embryos with dispersed/punctate localisation of α-actinin.								
GC, Fraction of 24-hpf embryos with dispersed/punctate localisation of α-tropomyosin. 6D, Fraction of 24-hpf embryos with abnormal localisation of titlin.								
ou, raction of z4-ripr emptyos with undetectable My8P-C.								
S1A, Fraction of TB-stage embryos with adaxial Mef2 staining.								
518, Fraction of 15s embryos showing PSM Met2 staining. 51C, Fraction of 15s embryos showing Met2 staining in heart region nuclei.								
			s and heart region					
S1D (left), Fraction of 13-15s embryos showing normal mef2c expression in somites and heart region. S1D (right), Fraction of 13-15s embryos showing normal mef2d expression in somites.								
S1E.F. S3. Fraction of 24-hof embryos showing undetectable (*) or low (**) levels of Mef2 staining in skeletal fibre nuclei.								

S1E,F, S3, Fraction of 24-hpf embryos showing undetectable (*) or low (**) levels of Mef2 staining in skeletal fibre nuclei.

S2B,C, Fraction of 24-hpf embryos showing undetectable (*) or low (**) levels of Mef2c staining in skeletal fibre nuclei. S2D, Fraction of 24-hpf embryos showing undetectable levels of Mef2c staining in heart muscle nuclei. S4B, Fraction of 22-24s embryos with highly reduced (*) or reduced (**) levels of myhz1 mRNA in somites.

S2A, Fraction of TB-stage embryos showing Mef2c staining.