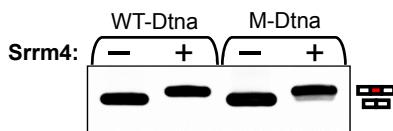


Figure S11

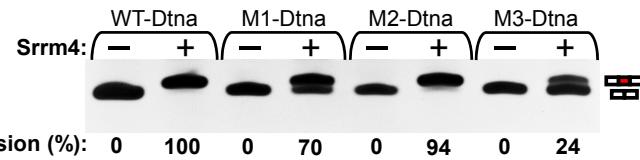
A

Dtna pre-mRNA (mutated region)
 WT: UGCUGCCAAAAG-**UGAUACUUG**-GUAAGUGAUGAAC
 M: UGCUGCCAAAAG-**UGCCGGUG**-GUAAGUGAUGAAC



B

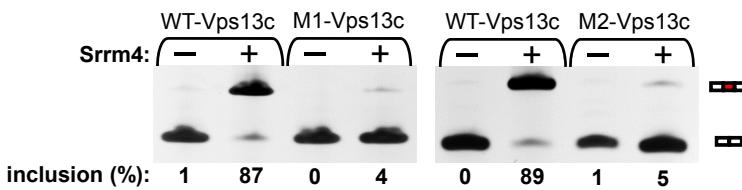
Dtna pre-mRNA (mutated region)
 WT: UGCUGCCAAAAG-exon
 M1: **UAC**UGCCAAAAG-exon
 M2: UGC**UAC**CCAAAAG-exon
 M3: **UACUAC**CCAAAAG-exon



inclusion (%): 0 100 0 70 0 94 0 24

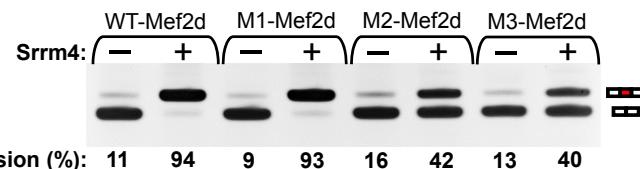
C

Vps13c pre-mRNA (mutated region)
 WT: UGUUUCCACUGAGAAC-exon
 M1: **UAC**UUCCACUGAGAAC-exon
 M2: UGUUUCCACUGAGAAC-exon



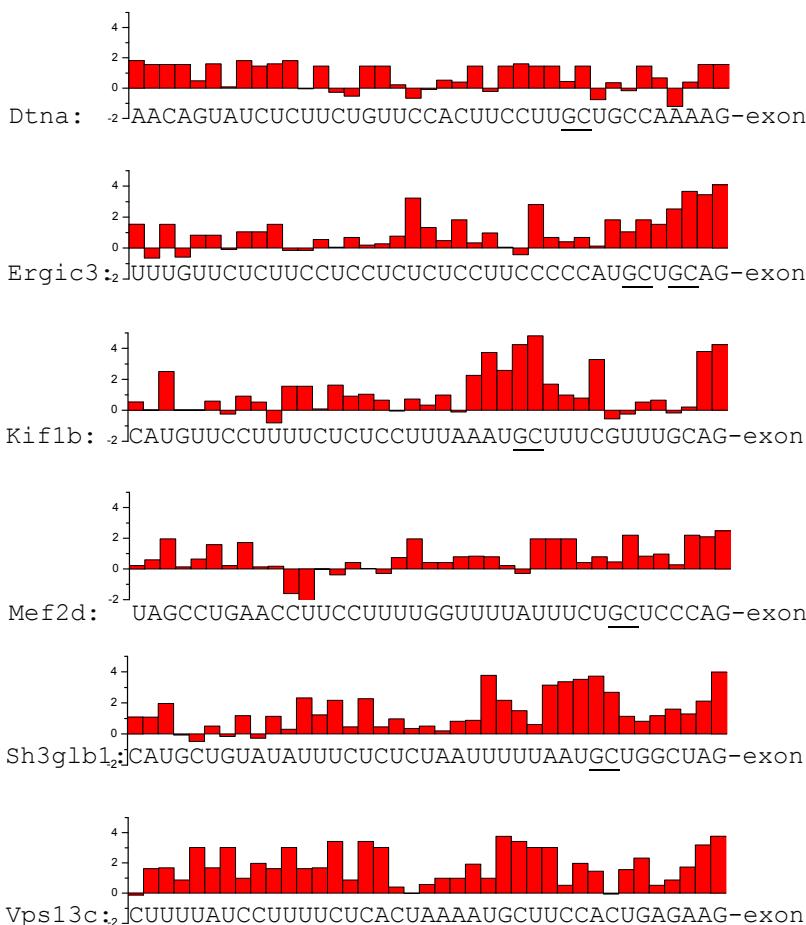
inclusion (%): 1 87 0 4 0 89 1 5

Mef2d pre-mRNA (mutated region)
 WT: UGGUUUUAUUUUCUGCUCCCAG-exon
 M1: **UAA**UUUUAUUUUCUGCUCCCAG-exon
 M2: UGGUUUUAUUU**C**UCCCAG-exon
 M3: **UAA**UUUUAUUU**C**UCCCAG-exon

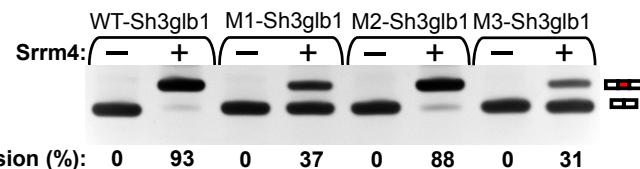


inclusion (%): 11 94 9 93 16 42 13 40

D

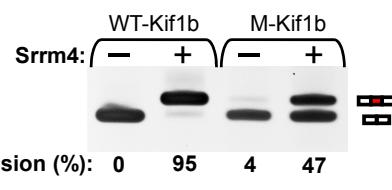


Sh3glb1 pre-mRNA (mutated region)
 WT: UGCUGGUAG-exon
 M1: **UAC**UGGUAG-exon
 M2: UGC**UAC**UAG-exon
 M3: **UACUAC**UAG-exon



inclusion (%): 0 93 0 37 0 88 0 31

Kif1b pre-mRNA (mutated region)
 WT: UGCUUUCGUUUGCAG-exon
 M: **UAC**UUUCGUUUGCAG-exon



inclusion (%): 0 95 4 47