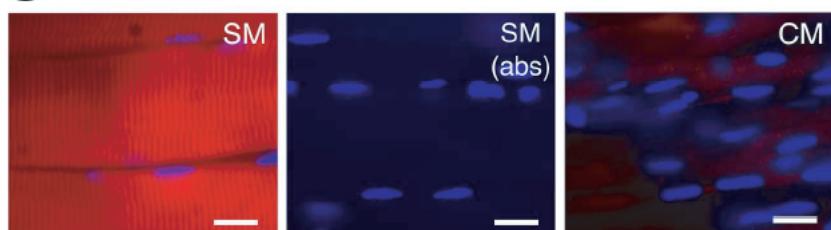
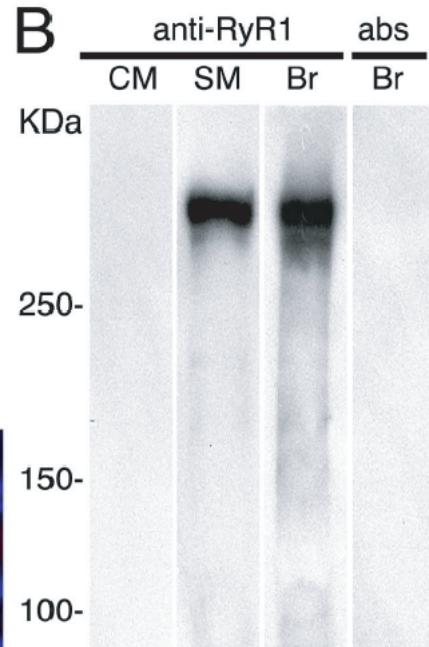
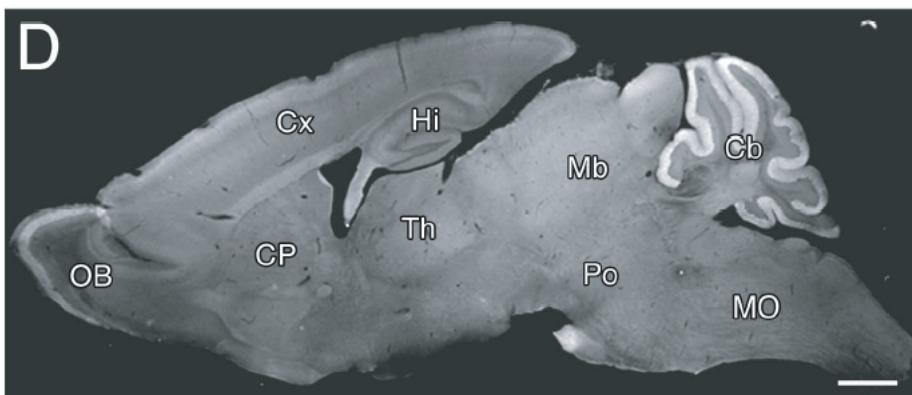


A

RyR1 (1313-1347) SPGLQPPAEDEARAAEPDTDYENLRRSAGGWGEAE
 * * * * * * * * * * * *
 RyR2 (1326-1357) --GFYGPKS-D-LEDFDVDSDFEVLMKTAHGLV-PD

RyR1 (1348-1382) GGKDGTAKEGTPGGTAQAGVEAQPARAENEKDSTT
 ** * * * * * * * * * *
 RyR2 (1358-1382) RIDKD--KETPKPEFNNHKDYAQ-----EKPS--

RyR1 (1383-1417) EKNKKRGFLFKAKKVAMMTQPPSTPALPRLPRDV
 * * * * * * * * * * * *
 RyR2 (1383-1409) --RLKQRFLLRRTK-----PDYSTGHSARLTED-V

C**B****D****E****Supplementary Figure 5 (Kakizawa et al.)**