

Table S2. Reads information of single-molecule long-read isoform sequencing (Iso-seq) data in sheep muscle

Data type	Total number	Min length	Max length	Average length
polymerase reads	442,966	51	363,026	97,901
Subreads	26,481,497	51	191,746	1,560
Circular Consensus Sequence (CCS) reads	331,820	55	15,574	2,119
Full-length Non-Concatemer (FLNC) reads	247,201	50	9,144	1,732