

Supplemental Table S5. T2C sequencing and read mapping statistics (*Apol* libraries).

Library ID	Total number of read pairs	Read pairs mapped to the genome (hg18)	Read pairs mapping to the 2.8-Mbp region of interest
0min.TNF.repl1	26,806,413	15,814,978	8,257,662
0min.TNF.repl 2	27,306,144	16,217,846	8,468,038
30min.TNF.repl 1	26,479,872	14,756,133	7,472,101
30min.TNF.repl 2	26,889,575	15,114,438	7,660,360
7SKi.0min.repl1	27,916,837	18,206,522	10,661,178
7SKi.0min.repl2	28,439,924	18,739,484	10,981,808
7SKi.30min.repl1	27,999,729	19,517,813	11,318,317
7SKi.30min.repl2	28,526,014	20,096,277	11,663,259
7SKi.DRB.repl1	26,594,778	18,026,829	9,488,193
7SKi.DRB.repl2	27,045,611	18,538,760	9,760,880