

Supplementary Table S3. Statistical analysis of the sequencing data of the degradome.

Sample	BC15T (Count)	BC15T (Percent)	BC25T (Count)	BC25T (Percent)
Raw Reads	40035871	/	63921661	/
Reads < 15 nt after removing 3 adaptor	279303	0.70%	365364	0.57%
Mappable Reads	39756568	99.30%	63556297	99.43%
Unique Raw Reads	6681964	/	11443406	/
Unique reads < 15 nt after removing 3 adaptor	49769	0.74%	69339	0.61%
Unique Mappable Reads	6632195	99.26%	11374067	99.39%
Transcript Mapped Reads	11632183	29.05%	16611634	25.99%
Unique Transcript Mapped Reads	1945301	29.11%	3320813	29.02%
Number of Input Transcript	194182	/	194182	/
Number of Covered Transcript	56560	29.13%	84673	43.60%