

Table S1. Summary of RNA sequencing and reads mapping.

Pooled samples:	Day 2 Control	Day 2 <i>Zbed6</i> - silenced	Day 4 Control	Day 4 <i>Zbed6</i> - silenced
Total reads (million)	49	51	53	42
Aligned reads (%)	51.7%	55.9%	57.8%	63.5%
Uniquely aligned (%)	43.2%	47.8%	48.1%	53.5%