

**Table S5 Tissue-specific RNAi of BAF and PBAF subunits**

RNAi	Genotype	% Gon <sup>a</sup> +/- SD	n
none	wild-type	0	>200
	<i>hnd-1::rde-1</i> <sup>b</sup>	0 +/- 0	184
<i>swns-4</i>	wild-type	Emb	169
	<i>hnd-1::rde-1</i> <sup>b</sup>	99.0 +/- 1.1	99
<i>pbrm-1</i>	wild-type	Lvl	105
	<i>hnd-1::rde-1</i> <sup>b,c</sup>	10.2 +/- 3.3	137
<i>let-526</i>	wild-type	Emb	78
	<i>hnd-1::rde-1</i> <sup>b</sup>	100 +/- 0	379

<sup>a</sup> Gonadogenesis defects were assessed using a dissecting microscope. The average penetrance and standard deviation (SD) are reported. Embryonic (Emb) or Larval (Lvl) lethality is indicated.

<sup>b</sup> genotype is *unc-119(ed3)*; *rde-1(ne219)*; *rdls7* [*hnd-1::rde-1*]

<sup>c</sup> also includes *ccls4444* [*arg-1::GFP*]