

A

RNAi phenotype	33d- <i>GFP</i>	9d- <i>Smed-smg-1</i>	22d- <i>Smed-smg-1</i>	33d- <i>Smed-smg-1</i>
Normal	100% (84/84)	20.24% (17/84)		
Hyperplasia		34.52% (29/84)	7.14% (6/84)	
Hyperplasia and outgrowths		35.71% (30/84)	61.90% (52/84)	
Dead after outgrowths		9.52% (8/84)	30.95% (26/84)	100% (87/87)

B

Trunks 25dR	<i>Control RNAi</i>	<i>Smed-smg-1 RNAi</i>
Normal	100% (75/75)	8% (6/75)
Abnormal unpigmented blastemas with or without hyperplasia		38.67% (29/75)
Abnormal unpigmented blastemas, hyperplasia and/or outgrowths		9.33% (7/75)
Dead after outgrowths		44% (33/75)

C