**Table S3 Genes in common between van der Bent et al., 2014 and this work.** Of the 239 putative Wnt pathway regulated genes from seam cells and VPCs identified here by mRNA tagging, 35 genes were also identified by van der Bent et al., 2014 as genes that were differentially expressed between a wild type strain and a strain carrying a loss-of-function mutation in *bar-1*. The third column indicates whether expression of the gene was decreased or increased in the *bar-1(ga80)* strain relative to wild type. Genes with an asterisk (\*) were identified as downregulated upon loss of *bar-1* in van der Bent et al. 2014 and upregulated upon expression of an activated BAR-1 protein in this work and in Jackson et al. 2014.

		bar-1(ga80)
Gene WB ID	Gene	expression
WBGene00000493	che-14	decreased
WBGene00000615	col-38*	decreased
WBGene00000626	col-49*	decreased
WBGene00000647	col-71*	decreased
WBGene00000711	col-138*	decreased
WBGene00000735	col-162	decreased
WBGene00000930		decreased
	dao-4*	decreased
WBGene00001073	dpy-11*	
WBGene00001691	grd-2*	decreased
WBGene00001885	his-11	decreased
WBGene00001984	hog-1*	decreased
WBGene00002393	lpr-1	decreased
WBGene00004394	rol-1	decreased
WBGene00011077	R07B1.5*	decreased
WBGene00011594	T07G12.3*	decreased
WBGene00011665	T09F5.1	decreased
WBGene00012186	mlt-11	decreased
WBGene00013514	Y73F4A.1	decreased
WBGene00015172	B0410.3	decreased
WBGene00015442	C04F1.1*	decreased
WBGene00015453	C04G6.2	decreased
WBGene00015950	C18A11.4*	decreased
WBGene00017783	F25E5.2*	decreased
WBGene00022024	Y64H9A.2	decreased
WBGene00077489	C04G6.13	decreased
WBGene00000720	col-147	increased
WBGene00006539	tbb-6	Increased
WBGene00007297	C04F12.1	Increased
WBGene00010051	F54D5.5	Increased
WBGene00011434	T04D3.5	Increased
WBGene00011561	ttr-15	Increased
WBGene00012783	Y43C5A.3	Increased
WBGene00015865	C16E9.1	Increased
WBGene00016919	C54E4.4	Increased
WBGene00021379	Y37E11B.7	increased