Table S224 putative Wnt target genes in common between Jackson et al., 2014 and this work. Ofthe 239 putative Wnt pathway regulated genes from seam cells and VPCs identified here by mRNA tagging,24 genes were also identified by Jackson et al., 2014 in a non-tissue specific analysis of Wnt upregulatedgenes.

Gene WB ID G	
••••••••	ene
WBGene00000615 co	ol-38
WBGene00000626 co	ol-49
WBGene00000647 co	ol-71
WBGene00000930 da	ao-4
WBGene00001984 h	og-1
WBGene00004202 pl	ry-1
WBGene00006509 ta	ng-164
WBGene00006539 tb	ob-6
WBGene00008472 E	03H4.4
WBGene00009898 de	od-23
WBGene00010470 co	dr-4
WBGene00011077 R	07B1.5
WBGene00011594 7	07G12.3
WBGene00011624 7	08G5.3
WBGene00012046 72	26E4.4
WBGene00012783 Y	43C5A.3
WBGene00013754 fb	oxa-116
WBGene00015442 C	04F1.1
WBGene00016058 ns	spd-3
WBGene00017485 F	15E6.4
WBGene00018295 04	ac-29
WBGene00018416 re	etr-1
WBGene00019234 ug	gt-8
WBGene00043147 ns	spd-6