

Supplementary table S3

F	a_0	a_1	a_2	a_3	b_0	b_1	b_2	n_1	n_2	T	γ_1	γ_2
10	0.01	0.25	1	1	0.5	0.25	1	2	1	20	1	20
								P	τ			
								0.2	inner cell	150		
								1	outer cell			

Table S3: Parameters used in the simulations of the gene network in the position-based trophoctoderm formation model. The parameters in the top table are the same as in polarity-based model.