Experimental validation of 6 signalog proteins

Here we provide a detailed statistics about the experiments verifying the predicted Notch pathway membership of the *C. elegans* genes: *aqp-6*, *D1009.3*, *nsh-1*, *num-1*, *F10D7.5*, and *crb-1*. For *aqp-6*, *crb-1*, *num-1*, and *D1009.3* we analyzed loss-of-function mutations treated with *lin-12* RNAi (Supplementary Table 1.), for *nsh-1* and *F10D7.5* we applied RNAi in *lin-12(gf)* mutant animals (Supplementary Table 2. and Supplementary Figure 1.)

		Phenotype					
Genotype	RNAi	Normal vulva	Protruding vulva	Number of examined worms	P value		
N2	lin-12	82.8%	17.2%	320	_		
aqp-6(tm2407)	lin-12	74.7%	25.3%	182	0.0298		
crb-1(ok931)	lin-12	76.0%	24.0%	318	0.0359		
num-1(ok433)	lin-12	70.9%	29.1%	182	0.0018		
D1009.3(ok1349)	lin-12	70.0%	30.0%	163	0.0011		

Supplementary Table 1. Summary of the phenotypes of loss-of-function mutants treated with *lin-12* RNAi. The *N2* genotype is the control experiment. Assuming statistically independent measurements the error of the control result (N2) is ~5%. P values were computed with Chi-square probes. Compared to the control experiment in all four cases the phenotype was significantly shifted from the normal vulva towards the protruding vulva (Pvl).

		Phenotype						
Genotype	RNAi	Normal vulva	Multivulva	Number of vulvae (mean +/- std.dev.)	Number of examined worms	P value		
lin-12(gf)	-	7%	93%	3.27 +/- 0.19	118	—		
lin-12(gf)	nsh-1	17%	83%	2.75 +/- 0.16	175	0.0024		
lin-12(gf)	F10D7.5	19%	81%	2.75 +/- 0.17	176	0.0069		

Supplementary Table 2. Summary of the phenotypes of lin-12(gf) mutants. P values were computed with Chi-square probes. In both tests (performed for *nsh-1* and *F10D7.5*) the number of vulvae is reduced with a high statistical significance compared to the control case.



Supplementary Figure 1. Distribution of the number of vulvae in lin-12(gf) mutants. Observe that – compared to the lin-12(gf) control strain – both the *nsh-1* RNAi and *F10D7.5* RNAi animals were shifted towards smaller numbers of vulvae. Supplementary Table 2 displays averages of the data shown in this figure.