

Table S4. Analysis of migratory DFC phenotypes in integrin morphants at 80-90% epiboly

Marker	Injection stage	Morpholino	mRNA	n_T	n_{exp}	WT	Mutant	Absent
cas	1 to 4 cell	Uninjected	-	124	5	77.9±6.9%	19.9±6.8%	2.2±2.5%
		1.75 ng $\alpha V1miss$	-	51	2	75.7±6.1%	17.1±4.0%	7.1±10.1%
		None	200 pg $m\alpha V$	19	2	63.3±4.7%	26.7±9.4%	16.1±8.6%
		1.25 ng $\alpha V1$	-	69	4	17.6±6.3%	69.3±9.5%	13.2±7.3%
		1.25 ng $\alpha V1$	200 pg $m\alpha V$	23	2	56.3±1.1%	43.7±1.1%	0%
		1.75 ng $\alpha V2$	-	59	4	23.9±6.8%	67.9±9.3%	8.2±3.4%
		1.75 ng $\alpha V2$	200 pg $m\alpha V$	19	2	52.2±11.0%	42.8±18.1%	5.0±7.1%
		5.00 ng $\alpha VE110$	-	23	2	47.3±15.5%	52.7±15.5%	0%
		0.75 ng $\beta 1b1$	-	40	3	21.1±18.3%	72.2±18.0%	6.7±8.2%
5.00 ng $\beta 1bE110$	-	23	2	48.5±21.4%	51.5±21.4%	0%		
sox17	1 to 4 cell	Uninjected	-	53	3	68.8±7.5%	22.9±7.9%	8.3±4.8%
		1.75 ng $\alpha V1miss$	-	17	2	70.8±4.2%	12.5±12.5%	16.7±16.7%
		1.25 ng $\alpha V1$	-	27	2	19.2±4.2%	53.3±13.3%	27.5±19.2%
		1.75 ng $\alpha V2$	-	28	2	32.5±7.5%	47.5±27.5%	20.0±20.0%
		5.00 ng $\alpha VE110$	-	12	1	58.3%	33.3%	8.3%
		0.75 ng $\beta 1b1$	-	18	2	37.0±8.4%	33.1±24.0%	29.9±15.6%
		5.00 ng $\beta 1bE110$	-	21	1	38.1%	47.6%	14.3%
ntl	1 to 4 cell	Uninjected	-	42	3	79.0±1.4%	16.5±3.9%	4.5±4.5%
		1.75 ng $\alpha V1miss$	-	25	3	68.1±3.7%	8.3±8.3%	23.6±6.1%
		1.25 ng $\alpha V1$	-	53	4	29.2±7.6%	60.9±2.0%	9.9±5.8%
		5.00 ng $\alpha VE110$	-	46	2	36.9±0.1%	40.4±7.0%	22.7±6.9%
		0.75 ng $\beta 1b1$	-	31	3	34.2±4.6%	58.6±3.8%	8.3±8.3%
		5.00 ng $\beta 1bE110$	-	29	2	41.4±0.2%	58.6±0.2%	0%
cas	512-1000 cell	Uninjected	-	63	2	80.6±10.3%	11.1±1.5%	8.3±11.8%
		2.50 ng $\alpha V1miss$	-	47	3	75.5±3.6%	19.2±7.1%	5.3±3.6%
		1.25 ng $\alpha V1$	-	48	3	36.8±5.3%	41.5±8.8%	21.7±3.8%
		1.75 ng $\alpha V2$	-	12	1	41.7%	41.7%	16.7%
		0.75 ng $\beta 1b1$	-	39	3	36.1±9.0%	42.6±6.0%	21.3±14.7%
		5.00 ng $\beta 1bE110$	-	25	2	23.4±7.3%	64.0±0.5%	12.7±7.8%