

	Absent or strongly reduced <i>pu.1</i> Expression, %	
wt (n>20)	0	
<i>etsrp</i> MO (n=52)	95	
<i>etsrp</i> MO + <i>scl</i> RNA (n=17)	0	
<i>scl</i> RNA (n=16)	0	
	Absent or strongly reduced <i>flk1</i> expression, %	Increased <i>flk1</i> expression, %
wt (n>20)	0	0
<i>etsrp</i> MO (n>50)	100	0
<i>etsrp</i> MO + <i>scl</i> RNA (n=20)	100	0
<i>scl</i> RNA (n=17)	0	71
	Absent or strongly reduced <i>pu.1</i> expression, %	Increased / ectopic <i>pu.1</i> expression, %
wt (n=40)	0	0
<i>scl</i> MO (n=38)	92	0
<i>etsrp</i> RNA (n=29)	10	48
<i>scl</i> MO + <i>etsrp</i> RNA (n=40)	95	0
	Absent or strongly reduced <i>pu.1</i> expression, %	Increased / ectopic <i>pu.1</i> expression, %
wt (n>40)	0	0
<i>alk8</i> MO (n=25)	100	0
<i>etsrp</i> RNA (n=31)	0	82
<i>alk8</i> MO + <i>etsrp</i> RNA (n=33)	100	0
	Some / moderate number of ectopic <i>pu.1</i> cells, %	Dramatic increase in <i>pu.1</i> cell number, %
wt (n=40)	0	0
<i>alk8</i> RNA (n=40)	0	0
<i>etsrp</i> RNA (n=51)	59	4
<i>alk8</i> + <i>etsrp</i> RNA (n=67)	55	21