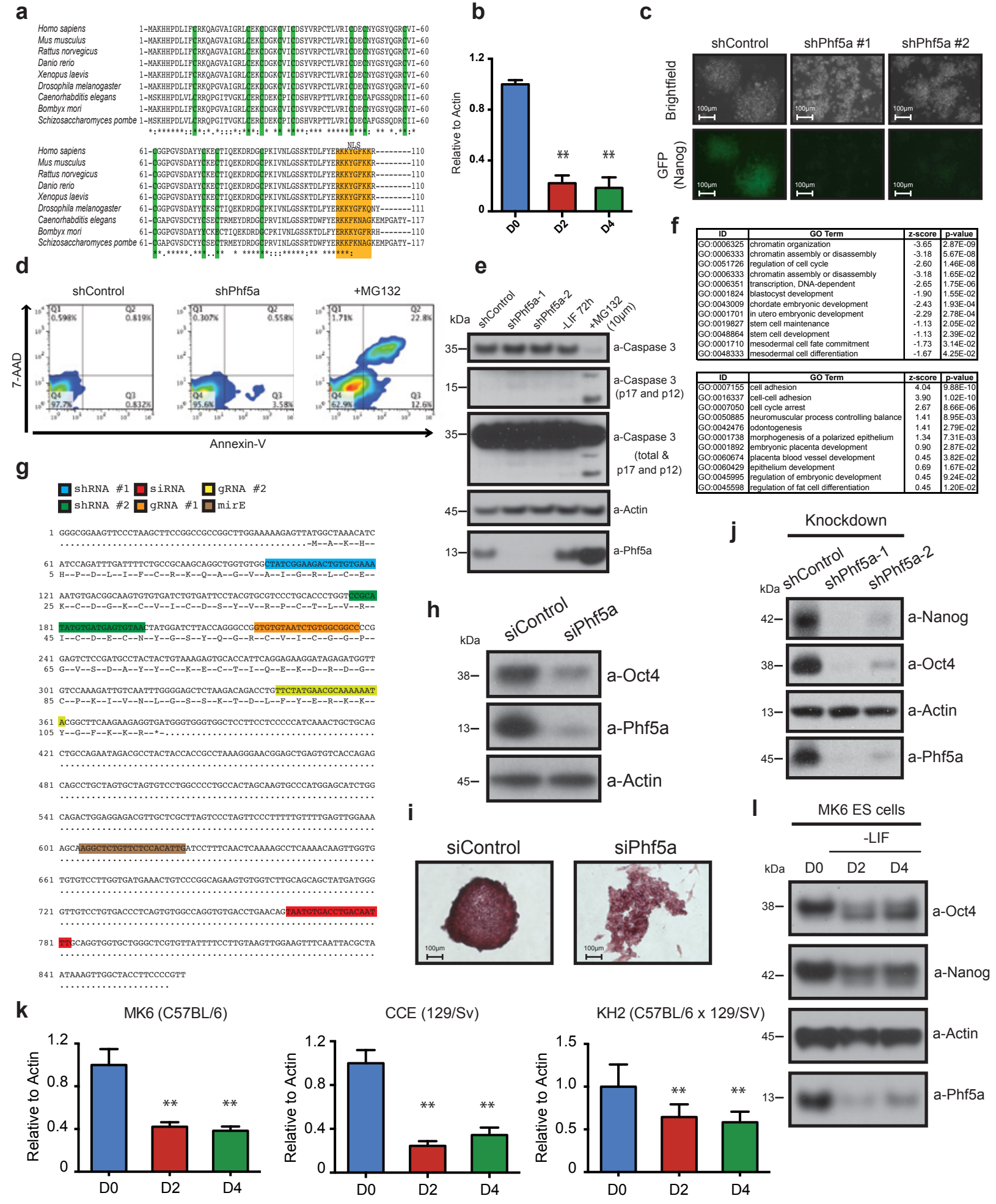
Supplementary Figure 1 - Strikoudis et al.



ID	GO Term	z-score	p-value
GO:0006325	chromatin organization	-3.65	2.87E-09
GO:0006333	chromatin assembly or disassembly	-3.18	5.67E-08
GO:0051726	regulation of cell cycle	-2.60	1.46E-08
GO:0006333	chromatin assembly or disassembly	-3.18	1.65E-02
GO:0006351	transcription, DNA-dependent	-2.65	1.75E-06
GO:0001824	blastocyst development	-1.90	1.55E-02
GO:0043009	chordate embryonic development	-2.43	1.93E-04
GO:0001701	in utero embryonic development	-2.29	2.78E-04
GO:0019827	stem cell maintenance	-1.13	2.05E-02
GO:0048864	stem cell development	-1.13	2.39E-02
GO:0001710	mesodermal cell fate commitment	-1.73	3.14E-02
GO:0048333	mesodermal cell differentiation	-1.67	4.25E-02

ID	GO Term	z-score	p-value
GO:0007155	cell adhesion	4.04	9.88E-10
GO:0016337	cell-cell adhesion	3.90	1.02E-10
GO:0007050	cell cycle arrest	2.67	8.66E-06
GO:0050885	neuromuscular process controlling balance	1.41	8.95E-03
GO:0042476	odontogenesis	1.41	2.79E-02
GO:0001738	morphogenesis of a polarized epithelium	1.34	7.31E-03
GO:0001892	embryonic placenta development	0.90	2.87E-02
GO:0060674	placenta blood vessel development	0.45	3.82E-02
GO:0060429	epithelium development	0.69	1.67E-02
GO:0045995	regulation of embryonic development	0.45	9.24E-02
GO:0045598	regulation of fat cell differentiation	0.45	1.20E-02