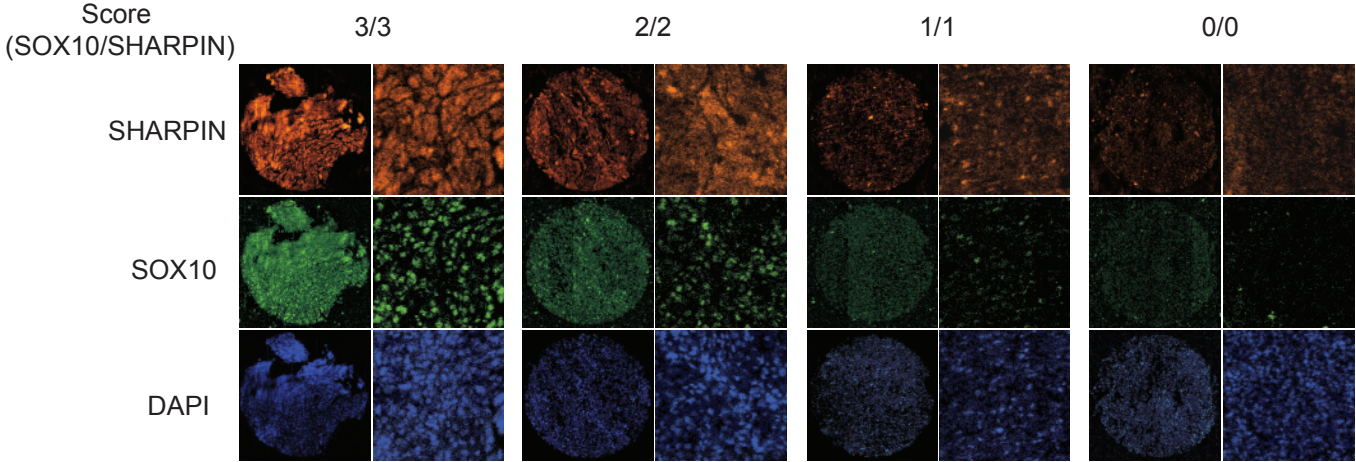


Supplementary Figure S6, Tamiya et al.



Supplementary Table S1, Tamiya et al.

Candidate protein	Control	SHARPIN
SHARPIN	0	168.33
Myosin-9 (MYH9)	2	56.33
Myosin-10 (MYH10)	0.33	29.33
Glutathione S-transferase P (GSTP1)	6	25.33
Vimentin OS=Homo sapiens (VIM)	7.33	23
Clathrin heavy chain 1 (CLTC)	0.33	22.33
Heat shock cognate 71 kDa protein (HSPA8)	10	18.67
Tubulin alpha-1A chain (TUBA1A)	3.67	13.33
60 kDa heat shock protein, mitochondrial (HSPD1)	6.33	13.33
78 kDa glucose-regulated protein (HSPA5)	7.67	13
Protein arginine N-methyltransferase 5 (PRMT5)	6.67	12.67
E3 ubiquitin-protein ligase RNF31 (RNF31)	0	11.33
X-ray repair cross-complementing protein 6 (XRCC6)	1	11.33
Serum albumin (ALB)	4	10.33
Methylosome protein 50 (WDR77)	6	9.67
Stress-70 protein, mitochondrial (HSPA9)	4.67	9.67
Myosin regulatory light chain 12A (MYL12A)	0.33	9.67
Myosin regulatory light chain 12B (MYL12B)	0.33	9.67
Glyceraldehyde-3-phosphate dehydrogenase (GAPDH)	3.67	9
Heterogeneous nuclear ribonucleoprotein U (HNRNPU)	1.67	8.67
RNA-binding protein 10 (RBM10)	4.33	8
Gelsolin (GSN)	0.33	7.67
U4/U6 small nuclear ribonucleoprotein Prp31 (PRPF31)	4.67	7.33
X-ray repair cross-complementing protein 5 (XRCC5)	0.67	7
Gamma-interferon-inducible protein 16 (IFI16)	0.33	6.33
Cytoplasmic dynein 1 heavy chain 1 (DYNC1H1)	4.33	6.33
Bifunctional polynucleotide phosphatase/kinase (PNKP)	0	6
Melanocyte protein PMEL (PMEL)	1.67	6
Poly [ADP-ribose] polymerase 1 (PARP1)	0.67	6
ADP/ATP translocase 1 (SLC25A4)	2	5.33
RanBP-type and C3HC4-type zinc finger-containing protein 1 (RBCK1)	0	5



Supplementary Table S2, Tamiya et al.

human	q-PCR primers	mouse	q-PCR primers
<i>GAPDH</i>	AGGGCTGCTTTTAACTCTGGT CCCCACTTGATTTTGGAGGGA	<i>Gapdh</i>	CATCACTGCCACCCAGAAGACTG ATGCCAGTGAGCTTCCCGTTTCAG
<i>HOIP</i>	CTTTCTTCCCCTAATCCTGCAAG CCCAAGGCTCATTTAGCATGG	<i>Sox10</i>	CAGGTGTGGCTCTGCCACG GTGTAGAGGGGCCGCTGGGA
<i>HOIL-1L</i>	AAACACCAAGCACTCTCAGC ACTGTGCAGTCTCCTTCCTG	<i>Pax3</i>	TGCCCTCAGTGAGTTCTATCAGC GCTAAACCAGACCTGCACTCGGGC
<i>SHARPIN</i>	GTGGCCCTTCCCTCACAAGTCCGAC CCCAACCCTGGTGACTGTCCCAGCA	<i>Mitf</i>	AGATTTGAGATGCTCATCCCC GATGCGTGATGTCATAGTGGA
<i>SOX10</i>	CCAGTACCCGCACCTGCAC CTTTCGTTACAGCAGCCTCCAG		
<i>PAX3</i>	GGAAGTGGAGCGTGCTTTTG GGCGGTTGCTAAACCAGAC		ChIP primers
<i>MITF</i>	CCGTCTCTCACTGGATTGGT TGGGTCTGCACCTGATAGTG	<i>ST7</i>	CCACCCAAATCTTGCTGAAT AGGATTCAGGCTCAGGGAAT
<i>SILV</i>	GCTGATCGTGGGCATCTTG AGTGACTGCTGCTATGTGG	<i>SOX10</i>	GACCCTCACCGCCTTCAC CGAGACTGACTGAGCGACTG
<i>TYR</i>	GCTGCCAATTTTCAGCTTTAGA CCGCTATCCCAGTAAGTGGA	<i>PAX3</i>	TGGGGCTGTCTCTCTCAGTT TTACCCAAAGCTTGGTCAGG
<i>DCT</i>	CTCAGACCAACTTGGCTACAGC CAACCAAAGCCACCAGTGTTCC		
<i>MLANA</i>	GGACAGCAAAGTGTCTCTTCAAG TCAGGTGTCTCGCTGGCTCTTA		
<i>ST7</i>	CCACTTGGCCTTCTCTTTC GGTCCCTACAAGTGGCTTT		
<i>RBL2</i>	GAAGTTCGTGCTGATACTGGAGG AATAGCCGCCTTCTGGTAGTGC		
<i>SKI</i>	CAGGAGCTGGAGTTCCTACG GTGACTCGTTGGCCTCTTTC		
<i>MTAP</i>	GTCATAGTGACCACAGCTTGTGG CCTCTGGCACAAGAATGACTTCC		
<i>TNFA</i>	AGGCGCTCCCCAAGAAGACA TCCTTGGCAAACCTGCACCT		
<i>IL1B</i>	ATGATGGCTTATTACAGTGGCAA AGTCGGAGATTCGTAGCTGGA		
<i>IL6</i>	ACTCACCTCTTCAGAACGAATTG CCATCTTTGGAAGGTTTCAGGTTG		
<i>PRMT5</i>	GAAGAATGGGGAAGCCAAGT ATGGGAGCCAGAAAGGAAGT		