#### **1** Supplemental Material:

#### 2 Supplemental Materials and Methods:

#### 3 Media conditions for cell culture

4 KCs were propagated in M154 medium supplemented with human KC growth supplement (Life 5 Technologies/Thermo Fisher Scientific, Waltham, MA), 1,000 x gentamycin/amphotericin B solution 6 (Life Technologies/Thermo Fisher), and 0.07 mM CaCl<sub>2</sub> (low calcium). Confluent KC monolayers were induced to differentiate by the addition of 1.2 mM CaCl<sub>2</sub> (high calcium) in M154 in the presence 7 8 of human KC growth supplement and gentamycin/amphotericin B solution. MCs were cultured in 9 OptiMEM (Life Technologies/Thermo Fisher) containing 1% penicillin/streptomycin (Corning, 10 Corning, NY), 5% fetal bovine serum (Millipore Sigma, St. Louis, MO), 10 ng/ml bFGF (ConnStem Inc., Cheshire, CT), 1 ng/ml heparin (Millipore Sigma), 0.1 mM N<sup>6</sup>, 2'-O-dibutyryladenosine 3:5-11 12 cyclic monophosphate (dbcAMP; Millipore Sigma), and 0.1 mM 3-isobutyl-1-methyl xanthine 13 (IBMX; Millipore Sigma).

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#### 15 Fixation and processing of organotypic cultures for imaging

Organotypic cultures fixed in 10% neutral buffered formalin were embedded in paraffin blocks and 16 cut into 4 µm sections. For indirect immunofluorescence microscopy, slides were baked at 60°C, de-17 paraffinized using xylene, dehydrated with ethanol, rehydrated in PBS and permeabilized by 0.5% 18 Triton X-100 in PBS. Antigen retrieval was performed by incubation in 0.01 M Citrate buffer (pH 19 20 6.0) at 95°C for 15 minutes. Sections were blocked in 1% BSA/2% Normal Goat Serum/PBS for 30 minutes at 37°C. Primary antibody incubation was carried out overnight at 4°C in 1% BSA/2% 21 22 Normal Goat Serum/PBS followed by washing in PBS. Secondary antibody incubation was carried 23 out at 37°C for 45 minutes followed by washing in PBS. Sections were stained with 4',6-Diamidino-2-phenylindole (DAPI – Millipore Sigma) at a final concentration of 5 ng/µl at room temperature for 24 25 2 minutes followed by washing in PBS. Cover slips were mounted on the sections with ProLong Gold 26 Antifade Reagent (Life Technologies/Thermo Fisher). 27 For whole mount imaging: Six days after lifting to the air-liquid interface, the epidermal equivalent

28 layer was removed from the collagen plug and fixed in 4% paraformaldehyde in PBS for 15 min on

- 29 ice. Samples were then washed three times in PBS for 5 min each at room temperature.
- 30 Subsequently, samples were incubated in blocking buffer (1% Triton-X 100 with 5% goat serum in
- PBS) for 1 hr at 37°C followed by incubation overnight at 37°C with S100 (ab52642, anti-S100
- 32 beta; Abcam, Cambridge, UK) diluted 1:100 in blocking buffer. Samples were washed 3 times for
- 10 min each with PBS at room temperature and then incubated overnight at 37°C with Alexa Fluor 1

34 conjugated secondary antibodies (1:250) in blocking buffer that included DAPI (2  $\mu$ g/ml). Samples

35 were washed 3 times for 10 min each with PBS at room temperature and mounted onto glass slides

36 with Prolong Gold Antifade Reagent.

#### 37 Quantitative real-time PCR

38 RNA was isolated from KCs and MCs using the RNeasy Mini Kit (Qiagen, Valencia, CA),

- 39 according to the manufacturer's instructions. MCs were grown in monoculture prior to being
- 40 incubated for 7 days with either MC media alone or with a 1:1 mixture of MC media and
- 41 conditioned media. Total RNA concentrations were equalized between samples and cDNA was
- 42 prepared using the Superscript III First Strand Synthesis Kit (Life Technologies/Thermo Fisher).
- 43 Quantitative PCR was performed using SYBR Green PCR master mix (Life Technologies/Thermo
- 44 Fisher) and gene-specific primers (Supplemental Table 1) in a StepOnePlus instrument (Thermo
- 45 Fisher Scientific Applied Biosystems). Calculations for relative mRNA levels were performed using
- 46 the  $\Delta\Delta$ Ct method, normalized to GAPDH.
- 47

#### 48 MTT assay and cell counting

- 49 MCs were plated at 5000 cells/well in 96 well plates and the next day conditioned media diluted 1:1
- 50 in MC media was added. The MTT assay protocol (Abcam), which was followed according to the
- 51 manufacturer's instructions, is based on the conversion of water soluble MTT (3-(4,5-
- 52 dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) compound to an insoluble formazan
- 53 product by viable cells. MC viability was assayed 1 day and 7 days after initiating incubation in
- 54 conditioned media diluted 1:1 in MC media. On days 1 and 7, MCs were incubated for 3 hours with
- 55 MTT reagent and 15 minutes with MTT solvent and absorbance at 590nm was measured with an
- 56 ELx800 microplate reader (Bio-Tek Instruments, Inc). Cells were also counted on days 0, 1, and 3

57 after initiating incubation in conditioned media using a hemocytometer.

58

#### 59 Immunoblot analysis of proteins

Whole cell lysates were collected from confluent monolayers of KCs or 80% confluent MCs in urea-SDS buffer (8 M urea/1% SDS/60 mM Tris (pH 6.8)/5% ß-mercaptoethanol/10% glycerol) and sonicated. Samples separated by SDS-PAGE were transferred to nitrocellulose, blocked in 5% milk/PBS, and incubated with primary antibodies in milk/PBS for 1 hour at room temperature or overnight at 4°C. After PBS washes, secondary antibodies diluted in milk/PBS were added to blots. Protein bands were visualized using exposure to X-ray film. Densitometric analyses were performed

66 of scanned immunoblots using ImageJ software.

### 68 Antibodies

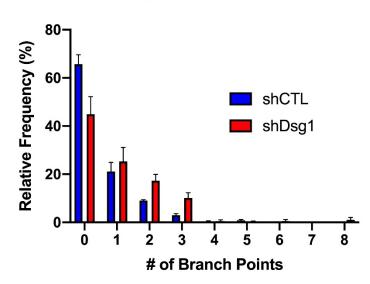
- 69 Mouse monoclonal antibodies: P124 (anti-Dsg1 extracellular domain; Progen, Heidelberg,
- 70 Germany) and 27B2 (anti-Dsg1 cytodomain; Life Technologies/Thermo Fisher). Rabbit monoclonal
- 71 antibody EP1576Y (ab52642, anti-S100 beta; Abcam, Cambridge, UK). Rabbit polyclonal
- 72 antibodies: Flag (Cell Signaling Technologies), TYRP1 (ab83774; Abcam), and GAPDH (G9545,
- 73 glyceraldehyde-3-phosphate dehydrogenase; Millipore Sigma). Secondary antibodies for
- 74 immunoblotting were goat anti-mouse and goat anti-rabbit peroxidase (SeraCare Life Sciences,
- 75 Milford, MA). Secondary antibodies for immunofluorescence were goat anti-mouse and goat-anti-
- rabbit linked to fluorophores of 488 and 568 nm (Alexa Fluor; Life Technologies/Thermo Fisher).

# 78 Supplemental Table 1. Primers used in this study.

Dsg1F: 5'-TCCATAGTTGATCGAGAGGTCAC-3' R: 5'-CTGCGTCAGTAGCATTGAGTATC-3'Dsg3F: 5'-ATCAATGCAACAGATGCAGATGCAGATGA-3' R: 5'-TGTCAAAGTGTAGCTGCTGTGT-3'IL1 $\alpha$ F: 5'-AGTAGCAACCAACGGGAAGG-3' R: 5'-TGGTTGGTCTTCATCTTGGG-3'IL1 $\beta$ F: 5'-GCAAGGGCTTCAGGCAGGCCGCG-3' R: 5'-GGTCATTCTCCTGGAAGGTCTGTG-3'IL2F: 5'-GTCACAAACAGTGCACCTAC-3' R: 5'-CCCTGGGTCTTAAGTGAAAG-3'IL4F: 5'-ACATTTGAACAGCCTCACAGAG-3' R: 5'-TTGGAGGCAGCAGCAACGAGAGAG-3'IL4F: 5'-ACATTTGAACAGCCTCACAGAG-3' R: 5'-CCATCTTTTCAGCCATCTTT -3'IL6F: 5'- ACAGCCACTCCACCTCTTCAG -3' R: 5'- TCCTTGGCAAAACTGCACCT -3'IL10F: 5'- AGGCCATTCCAAGCTGGCCGT -3' R: 5'- TCCTTGGCAAAACTGCACCT -3'IL19F: 5'- GGTTGCCAAGCCTTGTCTGA -3' R: 5'- CTTGGTCACAGCAGCACACAT -3'IL23F: 5'- GCTTCAAAATCCTTCGCAG -3' R: 5'- TATCTGAGTGCCATCCTTGAG -3' R: 5'- TATCTGAGTGCCATCCTTGAG -3' R: 5'- ATGAGTCATAGCCACAC -3' R: 5'- ATGGGAGTTATAGCCACAC -3' R: 5'- ATGGGAGTCATAGCCACAC -3' R: 5'- ATGGGCATTGTCAAAGCATGATCC -3'TNF $\alpha$ F: 5'- ATGAGCACTGAAAGCATGACTGACTGA-3' R: 5'- CTAATTATTCGGTAACTGACTGACTGA -3' R: 5'- ATGGGCATTAGCAAGAAGGATC -3'POMCF: 5'- AAGGCACTGCTGGCATCCTTGGA -3' R: 5'- ACAGTTCAGCCATCCATGGAACTGACTGACACAC -3'	<b>Target</b>	<u>Primers</u>
Dsg3F: 5'-ATCAATGCAACAGATGCAGATGA-3' R: 5'-TGTCAAAGTGTAGCTGCTGTGT-3'IL1αF: 5'-AGTAGCAACCAACGGGAAGG-3' R: 5'-TGGTTGGTCTTCATCTTGGG-3'IL1βF: 5'-GCAAGGGCTTCAGGCAGGCCGCG-3' R: 5'-GGTCATTCTCCTGGAAGGTCTGTG-3'IL2F: 5'-GTCACAAACAGTGCACCTAC-3' R: 5'-CCCTGGGTCTTAAGTGAAAG-3'IL4F: 5'-ACTTTGAACAGCCTCACAGAG-3' R: 5'-TTGGAGGCAGCAGCCAGCT-3'IL6F: 5'-ACAGCCACTCACCTCTTCAG -3' R: 5'-CCATCTTTCAGCCATCTT -3'IL8F: 5'- ATGACTTCCAAGCTGGCCGT -3' R: 5'- TCCTTGGCAAAACTGCACCT -3'IL10F: 5'- GGTTGCCAAGCCTTGTCTGA -3' R: 5'- CTTGGTCACGCAGCAAGCAAGATGACA -3' R: 5'- CTTGGTCACGCAGCAAGCAAGATGACA -3' R: 5'- TATCTGAGTGCCATCCTTGAG -3'IL23F: 5'- GAGCCATCCAAGCCTAGGCCAG -3' R: 5'- TATCTGAGTGCCATCCTTGAG -3' R: 5'- ATGAGCTCATAGCCACACACACACACACACACACACACAC	Dsg1	F: 5'-TCCATAGTTGATCGAGAGGTCAC-3'
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IL10F: 5'- GGTTGCCAAGCCTTGTCTGA -3' R: 5'- AGGGAGTTCACATGCGCCT -3'IL19F: 5'- GAGCCATCCAAGCTAAGGACA -3' R: 5'- CTTGGTCACGCAGCACACAT -3'IL23F: 5'- GCTTCAAAATCCTTCGCAG -3' R: 5'- TATCTGAGTGCCATCCTTGAG -3'CXCL1F: 5'- AACCGAAGTCATAGCCACAC -3' R: 5'- GTTGGATTTGTCACTGTTCAGC -3'TNF $\alpha$ F: 5'- ATGAGCACTGAAAGCATGATCC -3' R: 5'- GAGGGCTGATTAGAGAGAGGTC -3'IFN $\gamma$ F: 5'- CTAATTATTCGGTAACTGACTTGA -3' R: 5'- ACAGTTCAGCCATCATTGGA -3'	IL8	F: 5'- ATGACTTCCAAGCTGGCCGT -3'
R: 5'- AGGGAGTTCACATGCGCCT -3'IL19F: 5'- GAGCCATCCAAGCTAAGGACA -3'R: 5'- CTTGGTCACGCAGCACACAT -3'IL23F: 5'- GCTTCAAAATCCTTCGCAG -3'R: 5'- TATCTGAGTGCCATCCTTGAG -3'CXCL1F: 5'- AACCGAAGTCATAGCCACAC -3'R: 5'- GTTGGATTTGTCACTGTTCAGC -3'TNF $\alpha$ F: 5'- ATGAGCACTGAAAGCATGATCC -3'R: 5'- GAGGGCTGATTAGAGAGAGGTC -3'IFN $\gamma$ F: 5'- CTAATTATTCGGTAACTGACTTGA -3'R: 5'- ACAGTTCAGCCATCATTGGA -3'		R: 5'- TCCTTGGCAAAACTGCACCT -3'
IL19       F: 5'- GAGCCATCCAAGCTAAGGACA -3'         R: 5'- CTTGGTCACGCAGCACACAT -3'         IL23       F: 5'- GCTTCAAAATCCTTCGCAG -3'         R: 5'- TATCTGAGTGCCATCCTTGAG -3'         CXCL1       F: 5'- AACCGAAGTCATAGCCACAC -3'         R: 5'- GTTGGATTTGTCACTGTTCAGC -3'         R: 5'- GTTGGATTTGTCACTGTTCAGC -3'         TNFα       F: 5'- ATGAGCACTGAAAGCATGATCC -3'         IFNγ       F: 5'- CTAATTATTCGGTAACTGACTTGA -3'         R: 5'- ACAGTTCAGCCATCACTTGGA -3'	IL10	F: 5'- GGTTGCCAAGCCTTGTCTGA -3'
R: 5'- CTTGGTCACGCAGCACACAT -3'IL23F: 5'- GCTTCAAAATCCTTCGCAG -3'R: 5'- TATCTGAGTGCCATCCTTGAG -3'CXCL1F: 5'- AACCGAAGTCATAGCCACAC -3'R: 5'- GTTGGATTTGTCACTGTTCAGC -3'TNF $\alpha$ F: 5'- ATGAGCACTGAAAGCATGATCC -3'R: 5'- GAGGGCTGATTAGAGAGAGGTC -3'IFN $\gamma$ F: 5'- CTAATTATTCGGTAACTGACTTGA -3'R: 5'- ACAGTTCAGCCATCACTTGGA -3'		R: 5'- AGGGAGTTCACATGCGCCT -3'
IL23       F: 5'- GCTTCAAAATCCTTCGCAG -3'         R: 5'- TATCTGAGTGCCATCCTTGAG -3'         CXCL1       F: 5'- AACCGAAGTCATAGCCACAC -3'         R: 5'- GTTGGATTTGTCACTGTTCAGC -3'         TNFα       F: 5'- ATGAGCACTGAAAGCATGATCC -3'         R: 5'- GAGGGCTGATTAGAGAGAGGTC -3'         IFNγ       F: 5'- CTAATTATTCGGTAACTGACTTGA -3'         R: 5'- ACAGTTCAGCCATCACTTGGA -3'	IL19	F: 5'- GAGCCATCCAAGCTAAGGACA -3'
R: 5'- TATCTGAGTGCCATCCTTGAG -3'CXCL1F: 5'- AACCGAAGTCATAGCCACAC -3'R: 5'- GTTGGATTTGTCACTGTTCAGC -3'TNF $\alpha$ F: 5'- ATGAGCACTGAAAGCATGATCC -3'R: 5'- GAGGGCTGATTAGAGAGAGGTC -3'IFN $\gamma$ F: 5'- CTAATTATTCGGTAACTGACTTGA -3'R: 5'- ACAGTTCAGCCATCACTTGGA -3'		R: 5'- CTTGGTCACGCAGCACACAT -3'
CXCL1         F: 5'- AACCGAAGTCATAGCCACAC -3'           R: 5'- GTTGGATTTGTCACTGTTCAGC -3'         R: 5'- ATGAGCACTGAAAGCATGATCC -3'           TNFα         F: 5'- ATGAGGCTGATTAGAGAGAGGTC -3'           IFNγ         F: 5'- CTAATTATTCGGTAACTGACTTGA -3'           R: 5'- ACAGTTCAGCCATCACTTGGA -3'	IL23	F: 5'- GCTTCAAAATCCTTCGCAG -3'
R: 5'- GTTGGATTTGTCACTGTTCAGC -3'TNFαF: 5'- ATGAGCACTGAAAGCATGATCC -3'R: 5'- GAGGGCTGATTAGAGAGAGGTC -3'IFNγF: 5'- CTAATTATTCGGTAACTGACTTGA -3'R: 5'- ACAGTTCAGCCATCACTTGGA -3'		R: 5'- TATCTGAGTGCCATCCTTGAG -3'
TNFαF: 5'- ATGAGCACTGAAAGCATGATCC -3'R: 5'- GAGGGCTGATTAGAGAGAGGTC -3'IFNγF: 5'- CTAATTATTCGGTAACTGACTTGA -3'R: 5'- ACAGTTCAGCCATCACTTGGA -3'	CXCL1	F: 5'- AACCGAAGTCATAGCCACAC -3'
R: 5'- GAGGGCTGATTAGAGAGAGGTC -3'IFNγF: 5'- CTAATTATTCGGTAACTGACTTGA -3'R: 5'- ACAGTTCAGCCATCACTTGGA -3'		R: 5'- GTTGGATTTGTCACTGTTCAGC -3'
IFNγ       F: 5'- CTAATTATTCGGTAACTGACTTGA -3'         R: 5'- ACAGTTCAGCCATCACTTGGA -3'	ΤΝFα	F: 5'- ATGAGCACTGAAAGCATGATCC -3'
R: 5'- ACAGTTCAGCCATCACTTGGA -3'		R: 5'- GAGGGCTGATTAGAGAGAGGTC -3'
	IFNγ	F: 5'- CTAATTATTCGGTAACTGACTTGA -3'
POMC F: 5'- AGGCACTTGCTGGATTCTCC -3'		R: 5'- ACAGTTCAGCCATCACTTGGA -3'
	POMC	F: 5'- AGGCACTTGCTGGATTCTCC -3'

	R: 5'- GCCCTTCTTGTAGGCGTTCT -3'
KITL	F: 5'- TCGATGACCTTGTGGAGTGC -3'
	R: 5'- TGCTGTCATTCCTAAGGGAGC -3'
END-1	F: 5'- GACATCATTTGGGTCAACACTC -3'
	R: 5'- GGCATCTATTTTCACGGTCTGT-3'
MITF	F: 5'- TTATAGTACCTTCTCTTTGCCAGTCC -3'
	R: 5'- CTTATAAAATCCCTCTTTTTCACAGTTGGA -3'
MC1R	F: 5'- ACTTCTCACCAGCAGTCGTG -3'
	R: 5'- CATTGGAGCAGACGGAGTGT -3'
TYRP1	F: 5'- GTGCCACTGTTGAGGCTTTG -3'
	R: 5'- ATGGGGATACTGAGGGCTGT -3'
GAPDH	F: 5'- ACCACAGTCCATGCCATCAC -3'
	R: 5'- TCCACCACCCTGTTGCTGTA -3'

Dsg1, Desmoglein 1; Dsg3, Desmoglein 3; IL, Interleukin, CXCL1, Chemokine ligand 1; TNF,
Tumor Necrosis Factor; IFN, Interferon; POMC, Pro-opiomelanocortin; MITF, Melanogenesis
associated transcription factor; MC1R, Melanocortin 1 receptor; TYRP1, Tyrosinase related protein
1; KITL, Kit ligand; END-1, Endothelin-1; GAPDH, Glyceraldehyde 3-phosphate dehydrogenase.

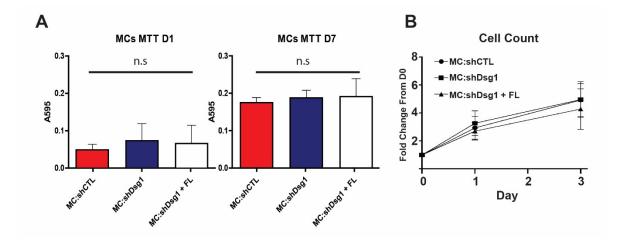


## **Histogram of Branch Points**

Supplemental Figure 1. MCs in Dsg1-deficient organotypic cultures exhibit increased dendrite branching compared to controls. Histogram of the frequency of occurrence of 0, 1, 2, 3, 4, 5, 6, 7, and 8 branch points per cell. The average number of branch points per cell are presented graphically in Figure 1F. There were fewer occurrences of 0 MC dendrite branch points and more occurrences of 2 and 3 branch points in the Dsg1-depleted organotypic cultures compared to the controls. A total of 239 MCs were analyzed for shCTL samples and 204 MCs for shDsg1 samples across all images (around 50 cells per organotypic culture). Data are presented as mean (N=4 independent experiments) and SEM.

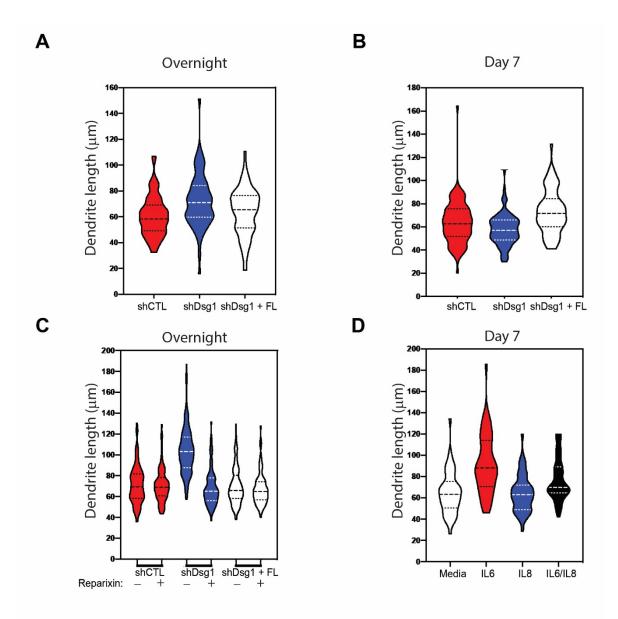
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116 Supplemental Figure 2. MCs are viable and have similar growth curves when grown in 117 conditioned media from control, Dsg1-deficient, and Dsg1-deficient rescued with full length Dsg1 KCs. (A) The MTT assay was conducted on MCs 1 day or 7 days after incubation in 1:1 mixture of 118 119 MC media: KC conditioned media (MC:shCTL, MC:shDsg1, MCLshDsg1 + FL). No significant changes were observed in cell viability at either time point regardless of type of conditioned media 120 121 (One Way ANOVA, repeated measures, Tukey's post-hoc test, N=3 MC isolates, 1 KC conditioned media, bar graph represents mean and SEM). (B) MCs (3 MC isolates) were counted using a 122 hemocytometer on days 1, 2, and 3 after incubation in 1:1 MC:KC conditioned media. Fold changes 123 124 from day 0 were calculated and represented as the mean and SEM of the 3 MC isolates treated with 125 1 set of KC conditioned media (MC:shCTL, MC:shDsg1, MC:shDsg1 + FL).



Supplemental Figure 3. Dsg1-deficient KCs change MC dendricity, partially dependent upon 127 128 cytokine/chemokine signaling. A) Violin plots of the spread of dendrite lengths of all cells after overnight incubation in media conditioned by KCs infected with shCTL, shDsg1, or shDsg1 + FL-129 expressing viruses. Incubation with media from KCs depleted of Dsg1 resulted in dendrite 130 lengthening compared to shCTL, as shown in Figure 4B. N = 5 independent MC:KC pairs. B) Violin 131 132 plots as in A except after 7 days of incubation in media conditioned by KCs expressing shCTL, shDsg1, or shDsg1 + FL. Incubation with media from KCs depleted of Dsg1 resulted in dendrite 133 shortening compared to the other two treatment groups, as shown in Figure 4C. N = 5 independent 134 135 MC:KC pairs. C) Violin plots of dendrite lengths after overnight incubation in media conditioned by

- 136 KCs expressing shCTL or shDsg1 with and without treatment with the CXCL1/IL8 receptor inhibitor,
- 137 reparixin. While media from Dsg1-depleted KCs resulted in lengthened MC dendrites as in A,
- treatment with reparixin blocked the effect, as shown in Figure 4D. N = 3 independent experiments.
- 139 D) Violin plots of dendrite lengths from MCs treated for 7 days with media spiked with recombinant
- 140 IL6, IL8, or the combination of IL6/IL8. Exposure to IL6 resulted in increased dendrite length as
- shown in Figure 4E. N = 3 independent experiments. Dotted lines inside violin plots represent means
- and SEM of all data points.