

SUPPLEMENTAL MATERIAL

Boulter et al., <http://www.jem.org/cgi/content/full/jem.20121657/DC1>**Table S1.** Genotyping primer list

Primer	Sequence
A	5'-GATAGACGGGAGTATTAGCGAGGC-3'
B	5'-CTCATGGTGCCTGCAGAAACGG-3'
C	5'-GGACATGTTACGGGATCGCCAGGCG-3'
D	5'-GCATAACCAAGTAAACAGCATTGCTG-3'
F	5'-CAGGGTTCTGTGTATGTGGGCGG-3'

Table S2. QPCR primers

Primer	Sequence
mJag1 F	5'-TCTCTGACCCTGCCATAAC-3'
mJag1 R	5'-TTGAATCCATTACCAGATCC-3'
mHes1F	5'-ACACCGGACAAACCAAAGAC-3'
mHes1R	5'-CGCCTCTTCTCCATGATAGG-3'
mShh F	5'-CAAGTACGCATGCTGGCTC-3'
mShh R	5'-AAGGTGAGGAAGTCGCTGTA-3'
mPtc2 F	5'-GGTCCTCCGCACCTCATATC-3'
mPtc2 R	5'-GCGCAGTCTGAATCAACATC-3'
mGli F	5'-AAGCCTGAGCCTGAGTCTGT-3'
mGli R	5'-GGTCACTGGCATTGCTAAAG-3'
mLef1 F	5'-TGAGTGACGCTAAAGGAGA-3'
mLef1 R	5'-ATAATTGTCTCGCGTGACC-3'
mGAPDH F	5'-ACCACAGTCCATGCCATCAC-3'
mGAPDH R	5'-TCCACCACCCTGTTGCTGTA-3'

Table S3. LC-ESI-MSMS protein identification

Peptide number	m/z	$\Delta m/z$	Range	Sequence
1	532.7500	-75.791	63-71	R.SEIYGLVQR.C
2	599.1700	-217.346	463-471	R.LYIQTMQER.V
3	337.7300	43.158	572-577	R.IGFLPK.I
4	584.2700	-70.99	808-816	K.VLDQVFYQR.V
5	550.8400	70.905	820-829	R.EGILSPSELR.K
6	539.2500	-6.734	950-957	K.YTEWPPER.E
7	685.7500	-76.123	1273-1283	K.STQEDWQSF.SR.Y