Corrections

GENETICS

Correction for "Global analysis of *trans*-splicing in *Drosophila*," by C. Joel McManus, Michael O. Duff, Jodi Eipper-Mains, and Brenton R. Graveley, which appeared in issue 29, July 20, 2010, of *Proc Natl Acad Sci USA* (107:12975–12979; first published July 1, 2010; 10.1073/pnas.1007586107).

The authors note that, within the supporting information Web link "http://intron.ccam.uchc.edu/Graveley/Publications/Publications.html" should be removed. Tables S1–S6 have been added to the online publication. The online version has been corrected.

IMMUNOLOGY

Correction for "Association of RIG-I with innate immunity of ducks to influenza," by Megan R. W. Barber, Jerry R. Aldridge, Jr., Robert G. Webster, and Katharine E. Magor, which appeared in issue 13, March 30, 2010, of *Proc Natl Acad Sci USA* (107:5913–5918; first published March 22, 2010; 10.1073/pnas.1001755107).

The authors note that on page 5917, right column, second full paragraph, line 12 " 5′-GTG TAT GGA GGA AAA CCC TAT TTC TTA ACT-3′ " should instead appear as " 5′-GTG TAT GGA GGA AAA CCC TAT TCT TAA CT-3′ ".

MEDICAL SCIENCES

Correction for "Prolonged nerve blockade delays the onset of neuropathic pain," by Sahadev A. Shankarappa, Jonathan H. Tsui, Kristine N. Kim, Gally Reznor, Jenny C. Dohlman, Robert Langer, and Daniel S. Kohane, which appeared in issue 43, October 23, 2012, of *Proc Natl Acad Sci USA* (109:17555–17560; first published October 8, 2012; 10.1073/pnas.1214634109).

The authors note that the following statement should be added as a new Acknowledgments section: "This work was supported by National Institute of General Medical Sciences Grant GM073626 (to D.S.K.)."

BIOCHEMISTRY, ENVIRONMENTAL SCIENCES

Correction for "Proteomic analysis of skeletal organic matrix from the stony coral *Stylophora pistillata*," by Jeana L. Drake, Tali Mass, Liti Haramaty, Ehud Zelzion, Debashish Bhattacharya, and Paul G. Falkowski, which appeared in issue 10, March 5, 2013, of *Proc Natl Acad Sci USA* (110:3788–3793; first published February 19, 2013; 10.1073/pnas.1301419110).

The authors note that Table 1 appeared incorrectly. Within the Name column, "CARP8" should instead appear as "CARP4," and "CARP9" should instead appear as "CARP5." These errors do not affect the conclusions of the article.

Table 1. Thirty-six predicted proteins in *S. pistillata* SOM samples detected by LC-MS/MS and their bioinformatics analysis

Protein	Gene	Accession no.	Name	P. damicornis	A. digitifera	Favia sp.	N. vectensis	P. maxima	S. purpuratus	E. huxleyii	R. filosa	H. sapiens	T. pseudonana
P1	g11108	KC509948	Protocadherin fat-like	−	+	+*†‡	+	+	−	−	−	−	−
P2	g11187	KC493647	CARP4	+	+‡	+†‡	−	−	−	−	−	−	−
P3	g12510	KC342189	Thrombospondin	−	+	+†	+	−	−	−	−	−	−
P4	g9861	KC342190	Viral inclusion protein	+*	+	+*†	+	−	−	−	−	−	−
P5	g11674	KC150884	Hemicentin	+	+†	+‡	+‡	+	−	−	−	+‡	−
P6	g11666	KC149520	Actin	+*†	+*	+*	+*	+*	+*	+*	+*	+*	+*
P7	g4601	KC342191	Actin	+*	+*‡	+*	+*†	+*	+*	+*	+*	+*	+*
P8	g9654	KC342192	Major yolk protein	+‡	+	+†‡	−	+	+	−	−	−	−
P9	g10811	KC000002	Protocadherin fat-like	−	+‡	+†‡	+‡	−	−	−	−	−	−
P10	g11107	KC509947	Cadherin	+*	+	+†‡	+	−	−	−	−	−	−
P11	g13727	KC342193	Actin	+*	+*†	+*	+*	+*	+*	+*	+*	+*	+*
P12[3]	g2385	JX891654	—	−	−	+†‡	−	−	−	−	−	−	−
P13	g6918	KC342194	Sushi domain-containing	+	+†	+	−	−	−	−	−	−	−
P14	g9951	KC342195	Collagen - alpha	−	+	+†	−	−	−	−	−	−	−
P15	g1532	KC493648	CARP5	−	+‡	+†‡	−	−	−	−	−	−	−
P16	g11702	KC342196	—	−	+†	+*	+	−	−	−	−	−	−
P17	g12472	KC149521	Glyceraldehyde 3-phosphatase dehydrogenase	+*	+*†	+*	+*	+	+	+	+	+*	+
P18	g810	KC342197	Collagen - alpha	−	+	+†	+	−	−	−	−	−	−
P19	g20041	KC342198	Contactin-associated protein	−	+	+†‡	+	−	−	−	−	−	−
P20	g6066	KC342199	MAM domain anchor protein	+	+†‡	+*	+	−	+	+	−	+‡	−
P21	g18277	KC479163	Zona pellucida	+*†‡	+	+*‡	+	−	−	−	−	−	−
P22	g19762	KC493649	—	−	−	−	−	−	−	−	−	−	−
P23	g1057	KC000004	Protocadherin	+	+	+*	+†	+	+	−	−	−	−
P24	g15888	KC479164	Vitellogenin	−	+	+*†‡	−	−	+	−	−	−	−
P25	g11220	KC479165	Ubiquitin	+*	+*‡	+*	+*†	+*	+*	+*	+*	+*	+*
P26	g1441	KC479166	Vitellogenin	+	−	+†‡	−	−	−	−	−	−	−
P27	g18472	KC479167	Integrin - alpha	+*	+	+†‡	+	−	−	−	−	−	−
P28	g11651	KC149519	Late embryogenesis protein	+†	−	−	−	−	−	−	−	−	−
P29	g13377	KC479168	Tubulin - beta	+*	+*	+*†	+*	+*	+*	+*	+*	+*	+
P30	g11056	KC000003	Myosin regulatory light chain	+*	+	+†‡	+	+	−	−	−	−	−
P31	g20420	KC479169	Neurexin	−	+	+†‡	−	−	−	−	−	−	−
P32	g5540	KC479170	Kielin/chordin like	+*†	+‡	+†	−	−	−	−	−	+‡	−
P33	g8985	KC479171	Flagellar associated protein	+*†	+*	+*	+	−	−	−	−	−	−
P34	g1714	KC479172	MAM/LDL receptor domain containing protein	+	+‡	+*†	+	+	+	+	−	+	−
P35	g7349	EU532164.1	Carbonic anhydrase (STPCA2)	+*†	+‡	+	−	+	−	−	−	+	−
P36	g13890	KC479173	Zonadhesion-like precursor	+	+*‡	+	+*	+	+	−	−	−	−

Returned sequences with e-values ≤10^{-10} are presented in order of decreasing e-value. "Protein name" is the best BLAST hit in NCBI. "Gene" is the code number in our *S. pistillata* gene prediction model.
The "+" and "−" represent presence and absence, respectively, of similar sequences in comparison species.
*Sequence similarity is greater than 70%.
†Most similar sequence by bit score.
‡Indicates export signal.

CORRECTIONS