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**\*Correspondence:**  
Laura Anne Lowery  
lalowery@bu.edu

†These authors have contributed  
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# Corrigendum: Wolf-Hirschhorn Syndrome-Associated Genes Are Enriched in Motile Neural Crest Cells and Affect Craniofacial Development in *Xenopus laevis*

Alexandra Mills<sup>†</sup>, Elizabeth Bearce<sup>†</sup>, Rachael Cella, Seung Woo Kim, Megan Selig, Sangmook Lee and Laura Anne Lowery\*

Biology Department, Boston College, Chestnut Hill, MA, United States

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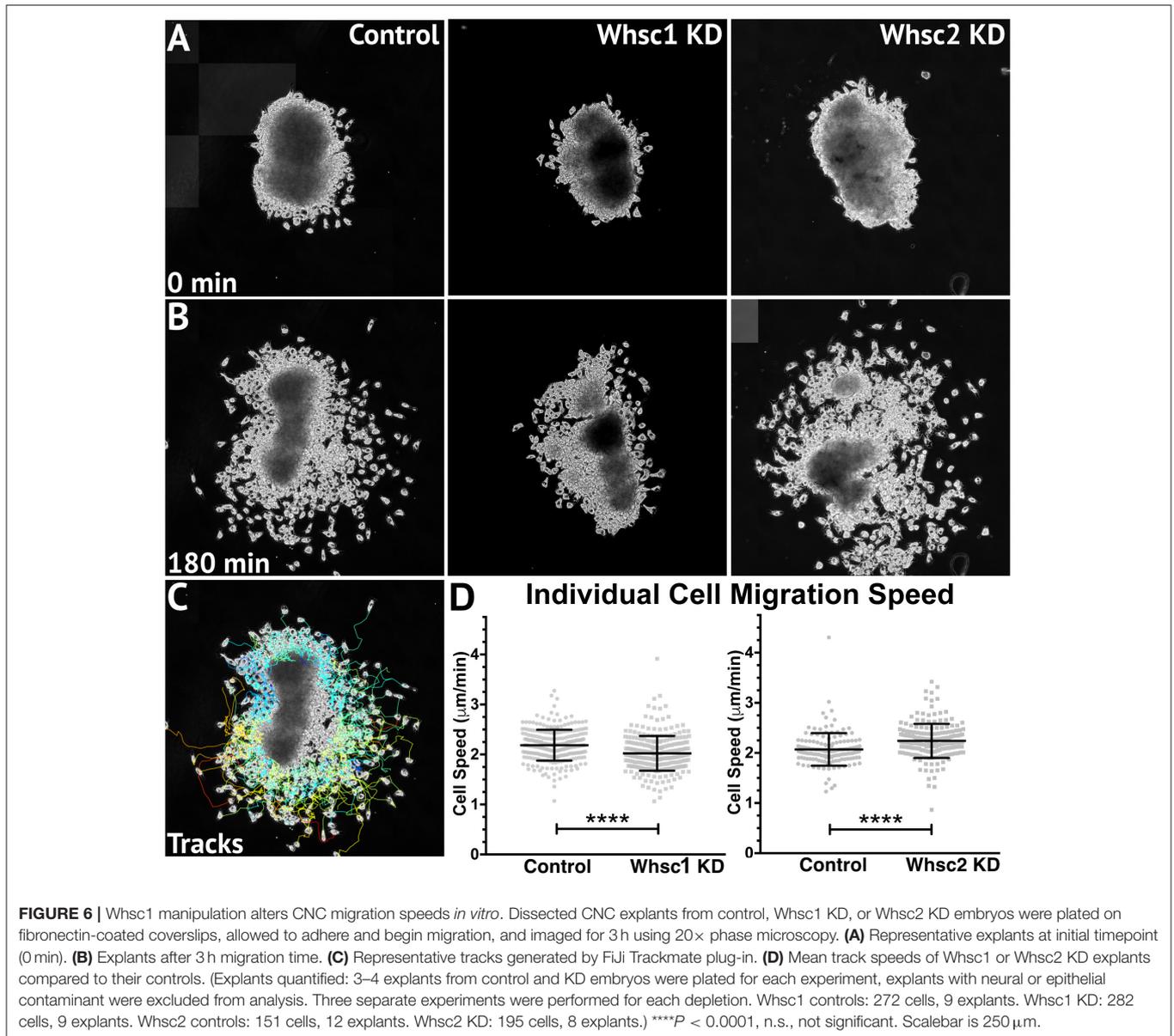
## A Corrigendum on

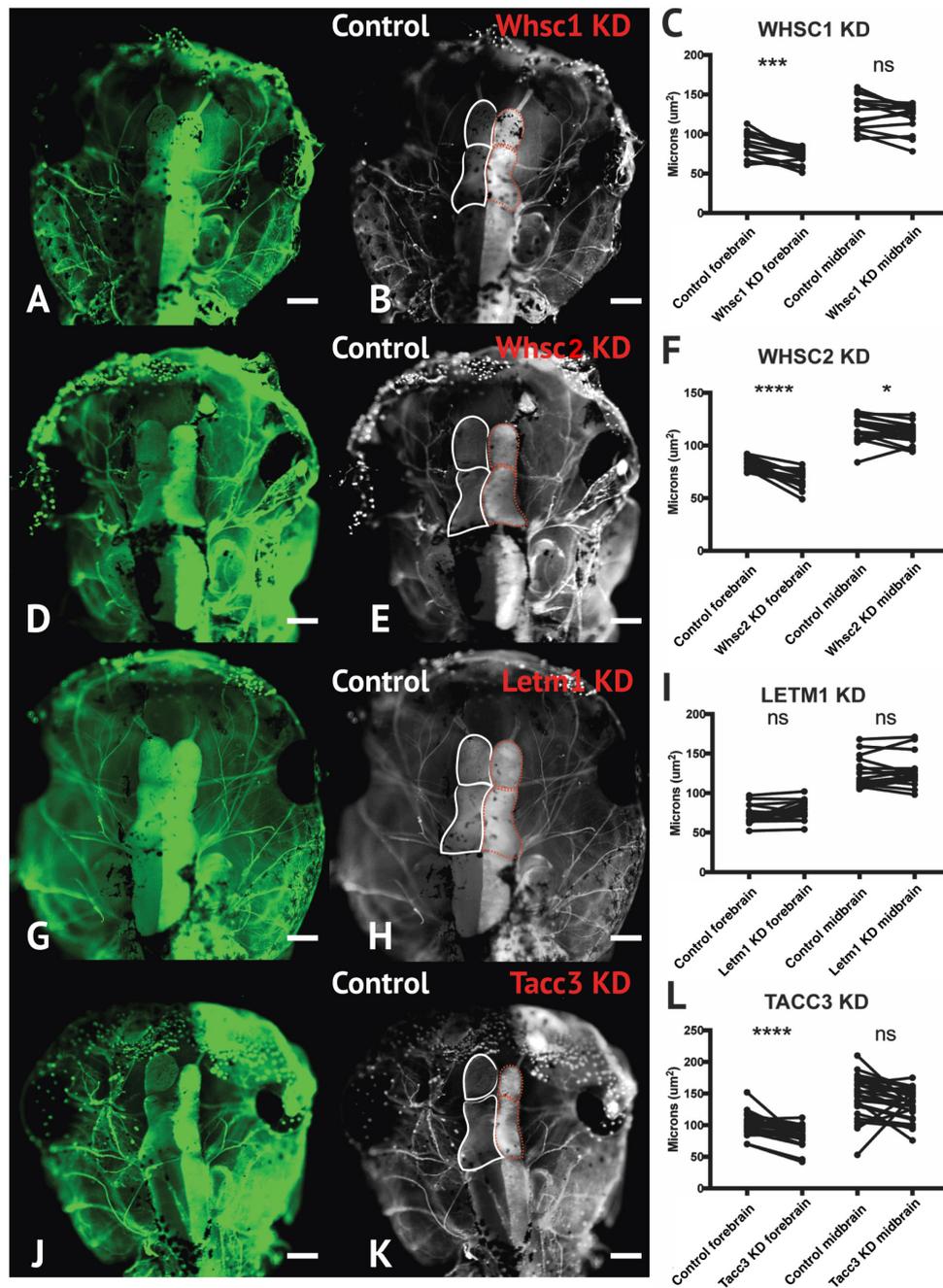
**Wolf-Hirschhorn Syndrome-Associated Genes Are Enriched in Motile Neural Crest Cells and Affect Craniofacial Development in *Xenopus laevis***

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In the original article, **Figure 6** panel D (left) was mislabeled. It has been corrected to read Whsc1 KD. In **Figure 7**, panel E was mislabeled. It has been corrected to read Whsc2 KD.

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**FIGURE 7 |** Whsc1, whsc2, and tacc3 facilitate normal forebrain development. (A,B,D,E,G,H,J,K) Dorsal view of *X. laevis* half-embryo gene depletions (6 days post-fertilization), following alpha-tubulin immunolabeling to highlight nervous system. (B,E,H,K) Dorsal view of embryos with superimposed outlines of forebrain and midbrain structures. Internal control is on left (white), depleted side is on right (dashed red). (Alpha-tubulin staining is bilateral; exogenous eGFP on KD side persisted in embryos shown, causing a unilaterally enriched green signal.) (C,F,I,L) Area of forebrain and midbrain. Whsc1 KD reduced forebrain area by 17.65%. Whsc2 KD reduced forebrain area by 17.33% and midbrain area by 4.14%. Letm1 KD caused no significant change in brain size. Tacc3 KD caused a 16.05% decrease in forebrain area. Significance determined using a student's paired *t*-test. (Embryos quantified: Whsc1 KD = 14, Whsc2 KD = 18, Letm1 KD = 12, Tacc3 KD = 26.) \*\*\*\* $P < 0.0001$ , \*\*\* $P < 0.001$ , \* $P < 0.05$ , n.s., not significant. Scalebar is 250  $\mu$ m.