ERRATUM

Erratum to: Delineating the respective impacts of stochastic curl- and grad-forces in a family of idealized core genetic commitment circuits

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The original version of this article contained an error in panel D of Figure 4. The original panel D erroneously duplicated panel C. The author apologize for this oversight. The corrected Figure 4 can be found below.

Figure 4. Diagrams of curl sampled along one single stochastic trajectory. Propagation is displayed by sequential colored line segments joining successive locations along the stochastic track. Only a relevant part of the track is displayed; details in the text. The contribution by components 1, 2 (top row) and 1, 2, 3 (rows 2, 3 and 4) to the local curl modulus is indicated by the color ranging from each figure's minimum curl to its maximum. The curl scale is linear. Rows 1, 2, 3 and 4 are for DS1, DS2, DS3 and DS4 respectively. Left and right columns are for low TSS and high TSS respectively. In high TSS, the messenger RNA sub-manifold is essentially at rest. The X_1 , X_2 and X_3 axes are omitted on panel C through H because they would force an adversely distant viewpoint making details of the tracks imperceptible. See Table S1.