

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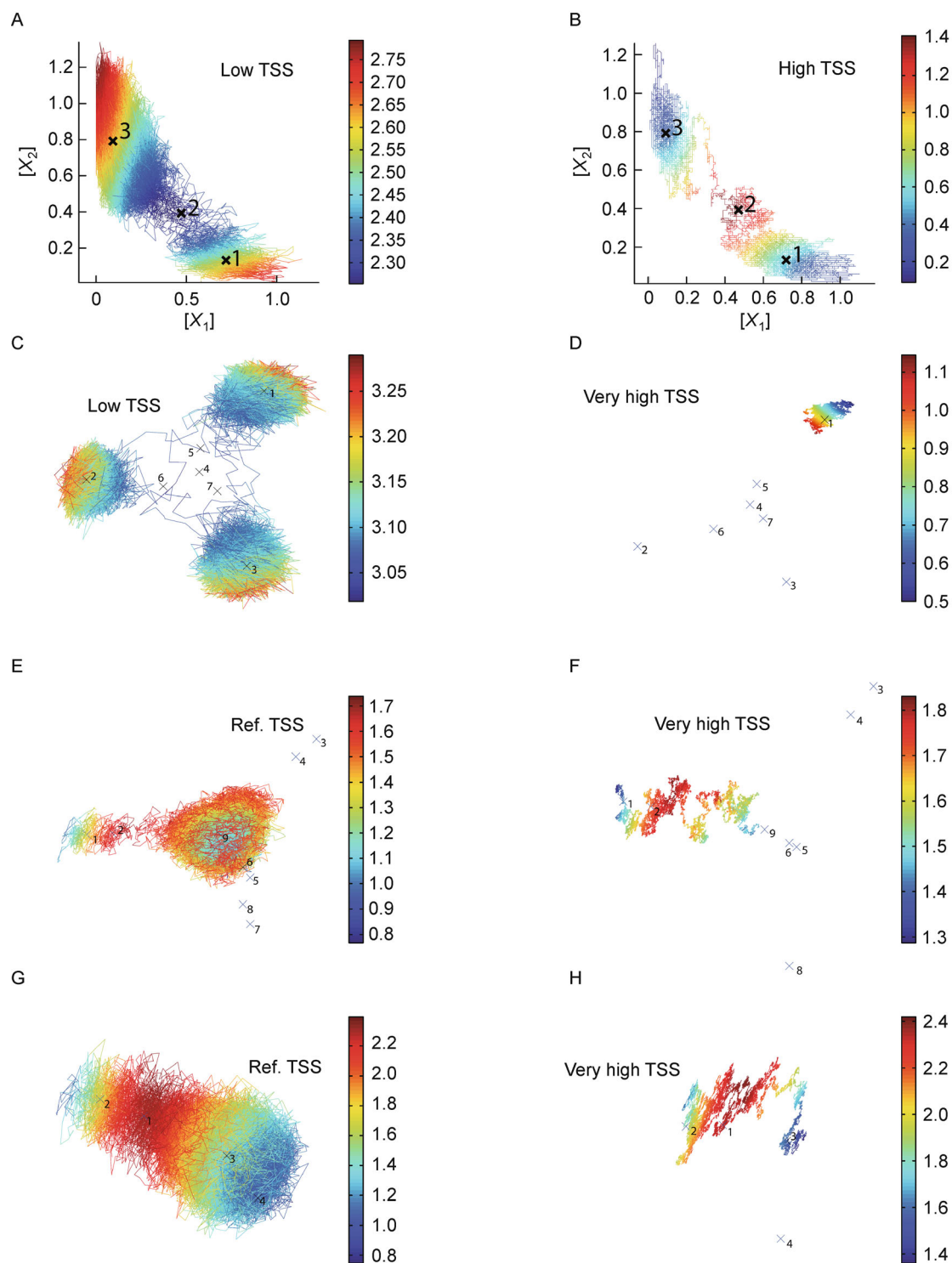
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The original version of this article contained an error in panel D of Figure 4. The original panel D erroneously duplicated panel C. The author apologize for this oversight. The corrected Figure 4 can be found below.



**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.