

A Review of “Dysregulation of  
Astrocyte-Secreted Pleiotrophin Contributes to  
Neuronal Structural and Functional Deficits in  
Down Syndrome” by Brandebura et al. (2025)

Reviewer 2

January 13, 2026

v1



**isitcredible.com**

## Disclaimer

This report was generated by Reviewer 2, an automated system that uses large language models to assess academic texts. It has been read and approved by a human editor on behalf of The Catalogue of Errors Ltd. The report's goal is to facilitate the discovery of knowledge by identifying errors in the existing literature. Comments can be made [here](#). Any errors will be corrected in future revisions.

I am wiser than this person; for it is likely that neither of us knows anything fine and good, but he thinks he knows something when he does not know it, whereas I, just as I do not know, do not think I know, either. I seem, then, to be wiser than him in this small way, at least: that what I do not know, I do not think I know, either.

Plato, *The Apology of Socrates*, 21d

To err is human. All human knowledge is fallible and therefore uncertain. It follows that we must distinguish sharply between truth and certainty. That to err is human means not only that we must constantly struggle against error, but also that, even when we have taken the greatest care, we cannot be completely certain that we have not made a mistake.

Karl Popper, 'Knowledge and the Shaping of Reality'

## Overview

**Citation:** Brandebura, A.N., Paumier, A., Asbell, Q.N., Tao, T., Micael, M.K.B., Sanchez, S., and Allen, N.J. (2025). Dysregulation of Astrocyte-Secreted Pleiotrophin Contributes to Neuronal Structural and Functional Deficits in Down Syndrome. *Cell Reports*, Vol. 44, 116300.

**URL:** <https://doi.org/10.1016/j.celrep.2025.116300>

**Abstract Summary:** Aberrant downregulation of the secreted molecule pleiotrophin (Ptn) contributes to neuronal phenotypes in a Down syndrome mouse model (Ts65Dn). Targeting overexpression of Ptn to astrocytes improves neuronal morphological and functional synaptic phenotypes, suggesting Ptn as a therapeutic target.

**Key Methodology:** Mouse models (Ts65Dn and Ptn KO), viral-mediated overexpression (AAV PHP.eB), smFISH, primary astrocyte and neuron cultures, western blotting, immunohistochemical synaptic analysis, ex vivo dendrite and spine analyses, and electrophysiology (fEPSP/STP).

**Research Question:** Does the dysregulation of astrocyte-secreted pleiotrophin contribute to neuronal structural and functional deficits observed in Down syndrome, and can targeting Ptn overexpression rescue these deficits?

## Summary

### Is It Credible?

The study by Brandebura et al. presents a compelling and methodologically rigorous investigation into the role of astrocyte-secreted pleiotrophin (Ptn) in Down syndrome (DS). The authors construct a logical narrative that moves from the identification of a molecular deficit—the downregulation of Ptn in Ts65Dn astrocytes—to a demonstration of functional rescue in adult mice. The central claim, that viral-mediated overexpression of Ptn can reverse established neuronal deficits in a DS mouse model, is supported by a robust body of evidence. The study is highly credible, distinguished by its willingness to acknowledge the complexities of the disease model and the limitations of the therapeutic approach. While there are statistical irregularities in secondary analyses and some variability in technical execution, these issues do not undermine the primary conclusion that Ptn is a viable target for modifying circuit aberrations in DS.

The strength of the study lies in the rigor applied to its most critical findings. When quantifying the primary structural deficits—dendrite length, complexity, and spine density—the authors adhere to a strict statistical standard, using the animal as the unit of analysis rather than the individual cell. This avoids the common pitfall of pseudoreplication and ensures that the reported rescue effects are biologically robust. The finding that Ptn overexpression fully restores dendrite morphology and spine density in both the visual cortex and the hippocampus of adult Ts65Dn mice is therefore statistically sound. Furthermore, the authors honestly grapple with the divergence between their knockout (KO) model and the disease model. They report that while Ptn KO mice share phenotypes with Ts65Dn mice, the KO deficits are transient, whereas Ts65Dn deficits are persistent. Rather than obscuring this discrepancy, the authors use it to refine their theoretical framework, correctly positing that Ptn downregulation is a contributing factor within a multifactorial disorder, rather than the sole driver of pathology.

However, the statistical rigor observed in the primary density analysis slips during the assessment of spine morphology. In determining the percentage of filopodia spines, the au-

thors treat individual dendrites as independent samples (N=40) rather than averaging per mouse (N=8), a decision that artificially inflates statistical power. While the authors acknowledge this shift in methodology, it renders the specific claims regarding spine maturity less reliable than the density data. This is particularly relevant given the finding that Ptn overexpression in wild-type mice increases the percentage of immature filopodia, suggesting a “U-shaped” dose response where excess Ptn may be detrimental. The reliance on dendrite-level statistics for this specific metric obscures whether this adverse effect is a consistent biological phenomenon or a statistical artifact.

Beyond the statistical handling of spine morphology, the study exhibits a collection of minor technical inconsistencies that reflect the challenges of integrating diverse experimental datasets. The use of different astrocyte markers (Aldh1L1, Glast, S100 $\beta$ , Sox9) across experiments and the variation in quantification methods for single-molecule fluorescence in situ hybridization (2D vs. 3D) introduce methodological noise. Similarly, the normalization of synaptic puncta data to the mean of the wild-type group within batches, while necessary to control for staining variability, imposes constraints on the variance structure of the control group. These issues are largely clerical or necessitated by experimental realities, such as the need for specific antibodies to ensure proper volume rendering. They do not appear to bias the results systematically in favor of the hypothesis, but they do reduce the seamlessness of the cross-experiment comparisons.

Ultimately, this study contributes a significant piece of knowledge to the field of neurodevelopmental disorders. It successfully demonstrates that the adult brain retains a capacity for structural remodeling that can be unlocked by restoring a single astrocyte-secreted factor. The limitations identified—specifically the statistical inflation in morphological subcategorization and the potential for adverse effects in healthy tissue—are important caveats for future translational work but do not invalidate the core discovery. The authors have provided a credible proof-of-concept that astrocyte-targeted therapies can repair circuitry after the closure of critical developmental windows.

## **The Bottom Line**

This study is a highly credible contribution to the understanding of Down syndrome pathophysiology. The authors provide robust evidence that restoring pleiotrophin levels in astrocytes can reverse neuronal structural and functional deficits in adult mice. The primary conclusions regarding dendrite length, spine density, and synaptic plasticity are supported by rigorous statistical analysis. While there is a notable statistical flaw in the secondary analysis of spine shape (pseudoreplication) and some minor inconsistencies in technical markers, these do not negate the central finding. The study is particularly strong in its honest reporting of the differences between the genetic knockout and the disease model, presenting a nuanced view of Ptn as a major, but not exclusive, contributor to the disorder.

## Specific Issues

**Pseudoreplication in spine morphology analysis:** There is a distinct inconsistency in the statistical rigor applied to spine density versus spine morphology. For spine density, the authors correctly used the average per mouse (N=6–8) as the statistical unit. However, for the analysis of spine morphology (e.g., percentage of filopodia), the authors treated individual dendrites as independent samples (N=35–40), as noted in the figure legends (pp. 7, 9) and methods (p. 25). This constitutes pseudoreplication, as dendrites from the same animal are not independent. This inflation of sample size likely overstates the statistical significance of the changes in spine maturity, particularly the increase in filopodia observed in the wild-type overexpression group and the lack of rescue in the Ts65Dn group.

**Divergent developmental trajectories:** The authors claim that Ptn KO mice display “parallel” or “overlapping” phenotypes with Ts65Dn mice (p. 5). However, the evidence shows a divergence in trajectory: Ptn KO deficits are transient and normalize by P120, whereas Ts65Dn deficits persist (pp. 5, 12). While the authors acknowledge this in the discussion, attributing the persistence in Ts65Dn to other genetic factors, the initial framing of the KO as a direct mimic of the DS condition is slightly oversimplified. This discrepancy suggests that Ptn loss is sufficient to delay development but not to maintain the deficit permanently in isolation.

**Potential adverse effects of overexpression:** The study reports that Ptn overexpression in wild-type mice leads to a significant increase in the percentage of immature filopodia spines (p. 8). Additionally, the treatment failed to reduce the elevated filopodia percentage in Ts65Dn mice. This suggests that while Ptn restores density and length, it may not restore—and could potentially disrupt—proper spine maturation when expressed at high levels. This “U-shaped” or potentially detrimental response in healthy tissue is a significant constraint on the therapeutic potential that is not fully resolved by the current data.

**Low sample size in electrophysiology:** The functional rescue claims in the hippocampus rely on a relatively small number of biological replicates. The short-term plasticity experiments utilized N=3–4 mice per group (p. 11, p. S19). While the authors correctly performed

statistics on the number of mice rather than slices, this sample size is on the lower limit of what is generally considered robust for detecting physiological interactions, increasing the risk that the results may not be generalizable.

**Methodological and reporting inconsistencies:** There are several minor technical issues that, while not fatal, accumulate to reduce the precision of the study. The authors utilized different astrocyte markers across experiments (Aldh1L1, Glast, S100 $\beta$ , Sox9) without a unified rationale for comparability (pp. 3, 25, p. S12). The quantification of smFISH data switched from 2D analysis in developmental profiles to 3D volumetric analysis for the disease model (pp. 24, 25). In vitro experiments compared effects on neurons of different developmental ages (P0–1 vs P6–7) (p. 21). Furthermore, the normalization of synaptic puncta data to the wild-type mean within batches (p. 27) constrains the variance of the control group. Finally, although the methods state both sexes were used (p. 20), the results do not explicitly report whether sex was analyzed as a factor, obscuring potential sex-dependent effects.

**Partial penetrance versus full rescue:** The authors report that deficits were “fully recovered” (pp. 7, 8) despite the viral vector achieving only partial penetrance, targeting approximately 66 percent of astrocytes (p. 6). While the authors offer a plausible explanation based on the secretion and diffusion of Ptn (p. 12), the discrepancy between partial cellular engagement and complete functional restoration warrants more detailed investigation to confirm the diffusion hypothesis.

**Partial in vitro rescue:** In the mechanistic validation, the addition of recombinant Ptn to knockout astrocyte-conditioned media resulted in only a “partial” recovery of dendrite metrics (p. 4). This contrasts with the “full” rescue observed in vivo. The authors suggest the involvement of other factors like Midkine (p. 12), but this remains a hypothesis that highlights the limitations of the single-molecule rescue model in a controlled in vitro setting.



## Future Research

**Hierarchical statistical modeling of spine morphology:** To resolve the issue of pseudoreplication in the spine morphology analysis, future analysis should employ hierarchical (nested) linear models. This approach would allow for the inclusion of all analyzed dendrites while correctly accounting for the clustering of data within individual mice. This would provide a rigorous assessment of whether the increase in filopodia spines in wild-type mice and the lack of morphological rescue in Ts65Dn mice are statistically valid biological effects or artifacts of inflated power.

**Dose-response titration studies:** Given the observation that Ptn overexpression induces immature spine phenotypes in wild-type mice, research is needed to determine the therapeutic window of Ptn. A dose-response study using tunable viral vectors or varying titers should be conducted to identify expression levels that rescue deficits in the disease model without inducing structural immaturity or aberrant plasticity in healthy neurons. This is critical for establishing the safety profile of Ptn as a therapeutic candidate.

**Combinatorial rescue experiments:** The study acknowledges that Ptn loss does not fully account for the persistent deficits in Ts65Dn mice and that in vitro rescue was only partial. Future work should investigate the co-expression of Ptn with other astrocyte-secreted factors identified as downregulated in DS, such as Midkine or Thrombospondin-1. This would test the hypothesis that a combinatorial approach is required to fully recapitulate the healthy astrocyte secretome and achieve more comprehensive rescue, potentially addressing the persistent spine immaturity.

**Sex-disaggregated analysis:** To address the reporting omission regarding biological sex, future replication of the behavioral and electrophysiological assays should explicitly power the study to detect sex differences. Analyzing males and females as separate factors is essential to determine if the Ptn rescue mechanism is equally efficacious across sexes, particularly given the known sexual dimorphism in many neurodevelopmental phenotypes.

© 2026 The Catalogue of Errors Ltd

This work is licensed under a

**Creative Commons Attribution 4.0 International License**

(CC BY 4.0)

You are free to share and adapt this material for any purpose,  
provided you give appropriate attribution.

[www.isitcredible.com](http://www.isitcredible.com)