

Dysregulation of the Wnt/β-catenin signaling pathway via Rnf146 upregulation in a VPA-induced mouse model of autism spectrum disorder

Seoyeon Kim^{1,2,15}, Gaeun Park^{3,4,15}, Wooyoung Eric Jang^{5,15}, Edson Luck Gonzales⁶, Jungeun Ji^{1,2}, Seunghwan Choi⁷, Yujin Kim^{1,2}, Ji Hwan Park⁸, Hazara Begum Mohammad⁸, Geul Bang⁹, Minkyung Kang^{3,4}, Soobin Kim^{3,4}, Se Jin Jeon⁶, Jin Young Kim⁹, Kwang Pyo Kim^{5,10}, Chan Young Shin^{6*}, Joon-Yong An^{1,2,7*}, Min-Sik Kim^{8,11,12*} and Yong-Seok Lee^{3,4,13,14*}

¹Department of Integrated Biomedical and Life Science, Korea University, Seoul 02841, Republic of Korea. ²BK21FOUR R&E Center for Learning Health Systems, Korea University, Seoul 02841, Republic of Korea. ³Department of Biomedical Science, Seoul National University College of Medicine, Seoul 03080, Republic of Korea. ⁴Department of Physiology, Seoul National University College of Medicine, Seoul 03080, Republic of Korea. ⁵Department of Applied Chemistry, Institute of Natural Science, Global Center for Pharmaceutical Ingredients Materials, Kyung Hee University, Yongin 17104, Republic of Korea. ⁶School of Medicine and Center for Neuroscience Research, Konkuk University, Seoul 05029, Republic of Korea. ⁷School of Biosystem and Biomedical Science, College of Health Science, Korea University, Seoul 02841, Republic of Korea. ⁸Department of New Biology, DGIST, Daegu 42988, Republic of Korea. ⁹Research Center for Bioconvergence Analysis, Korea Basic Science Institute, Ochang 28119, Republic of Korea. ¹⁰Department of Biomedical Science and Technology, Kyung Hee Medical Science Research Institute, Kyung Hee University, Seoul 02447, Republic of Korea. ¹¹New Biology Research Center, DGIST, Daegu 42988, Republic of Korea. ¹²Center for Cell Fate Reprogramming and Control, DGIST, Daegu 42988, Republic of Korea. ¹³Neuroscience Research Institute, Seoul National University College of Medicine, Seoul 03080, Republic of Korea. ¹⁴Wide River Institute of Immunology, Seoul National University, Hongcheon 25159, Republic of Korea. ¹⁵These authors contributed equally: Seoyeon Kim, Gaeun Park, Wooyoung Eric Jang. *Correspondence: chanyshin@kku.ac.kr; joonan30@korea.ac.kr; mkim@dgist.ac.kr; yongseok7@snu.ac.kr



ABSTRACT

Maternal exposure to valproic acid (VPA) during pregnancy is linked to autism spectrum disorder (ASD). This study used high-resolution mass spectrometry to analyze protein expression changes in the prefrontal cortex (PFC) of mice exposed to VPA in utero. The analysis revealed differential expression of proteins associated with ASD risk genes, particularly in the Wnt/β-catenin signaling pathway. Overexpression of Rnf146 in the PFC of adult mice resulted in regulatory abnormalities and social behavior defects in the Wnt/β-catenin pathway, along with increased excitatory synaptic metastasis in PFC neurons. We further examined the expression profile of Rnf146 in mouse forebrain. The findings suggest that Rnf146 disrupts the Wnt/β-catenin signaling pathway, contributing to ASD development following fetal exposure to VPA.

METHODS

Pregnant mice received E10 subcutaneous injections of VPA or saline, followed by social behavior tests at 6–7 weeks post-birth. PFC samples from VPA-exposed and control mice were collected for proteomic and western blot analyses, using high-resolution mass spectrometry and MaxQuant software. Enrichment analyses was performed in VPA-exposed mice, followed by functional mapping and protein-protein interaction network analysis. PFC samples from Rnf146-overexpressing mice underwent RNA sequencing, processed with Salmon software. Weighted gene coexpression network analysis (WGCNA) was applied for functional topology interpretation. Neuronal Rnf146 expression was achieved through an adeno-associated viral (AAV) vector, followed by targeted AAV injection via surgical procedures..

RESULTS

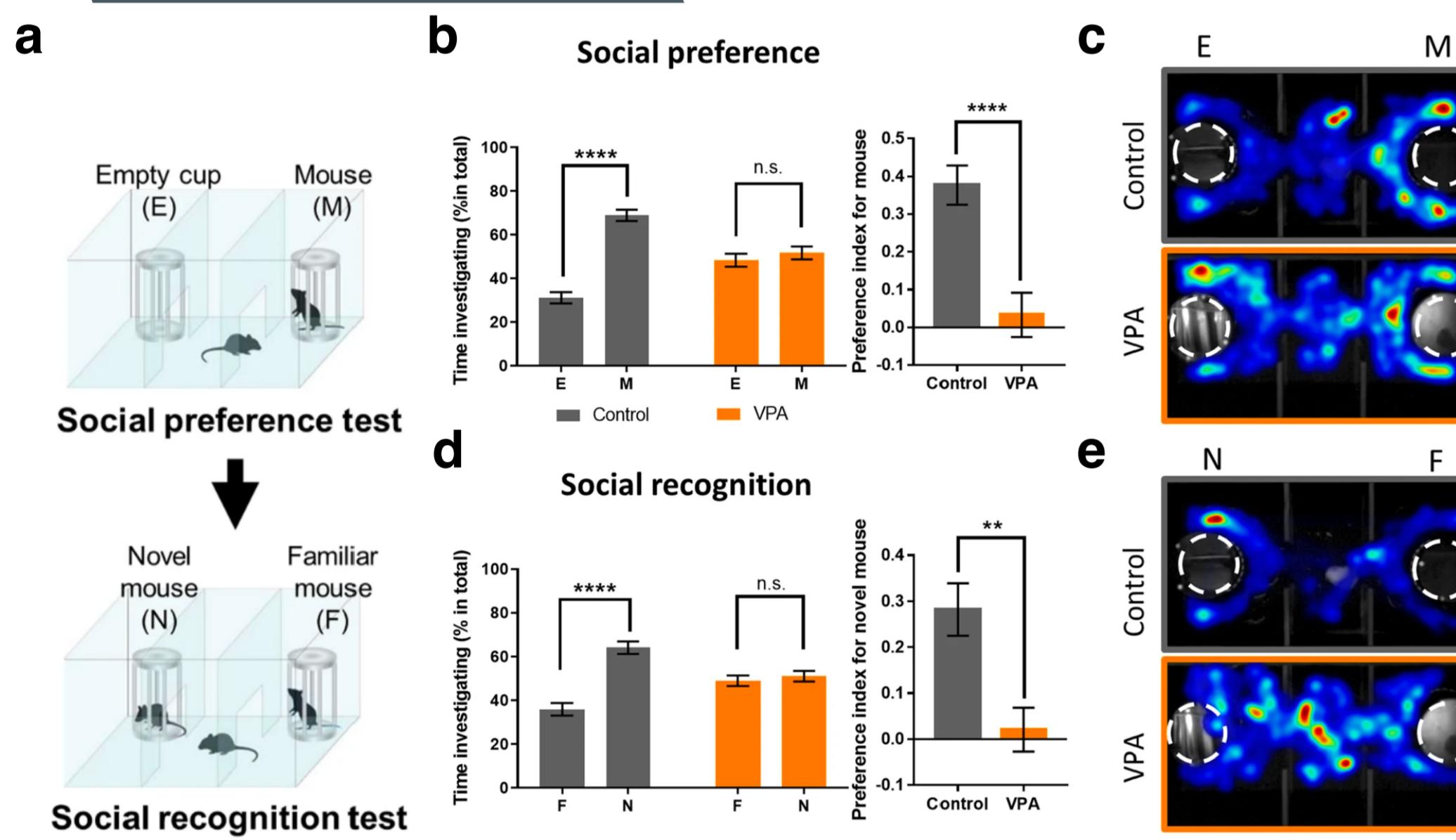


Figure 1. Social deficits in VPA-exposed mice. **a.** Schematic diagram of the three chamber social behavior test. **b.** Social preference test and preference index between VPA-exposed mice ($n = 19$) and control mice ($n = 20$). **c.** Representative heatmap images of the social preference test of control and VPA-exposed mice. **d.** Social recognition test result and preference index between VPA. **e.** Representative heatmap images of the social recognition test of control and VPA-exposed mice.



Figure 2. VPA-exposed mice show upregulation of the Wnt signaling pathway. **a.** The overall design of the proteomic experiment. **b.** Volcano plot showing DEPs between VPA-exposed and control mice. **c.** Enrichment analysis of VPA-DEPs with mouse brain celltype markers. **d.** Enrichment analysis of VPA-DEPs with disorder risk genes. Autism spectrum disorder (ASD), developmental disorder (DD), epilepsy (EP), and schizophrenia (SCZ). **e.** Functional annotations for VPA-DEPs showing significantly enriched biological pathways ($FDR \leq 0.05$). **f.** Protein-protein interaction network of VPA-DEPs related to Wnt signaling and neuronal development.

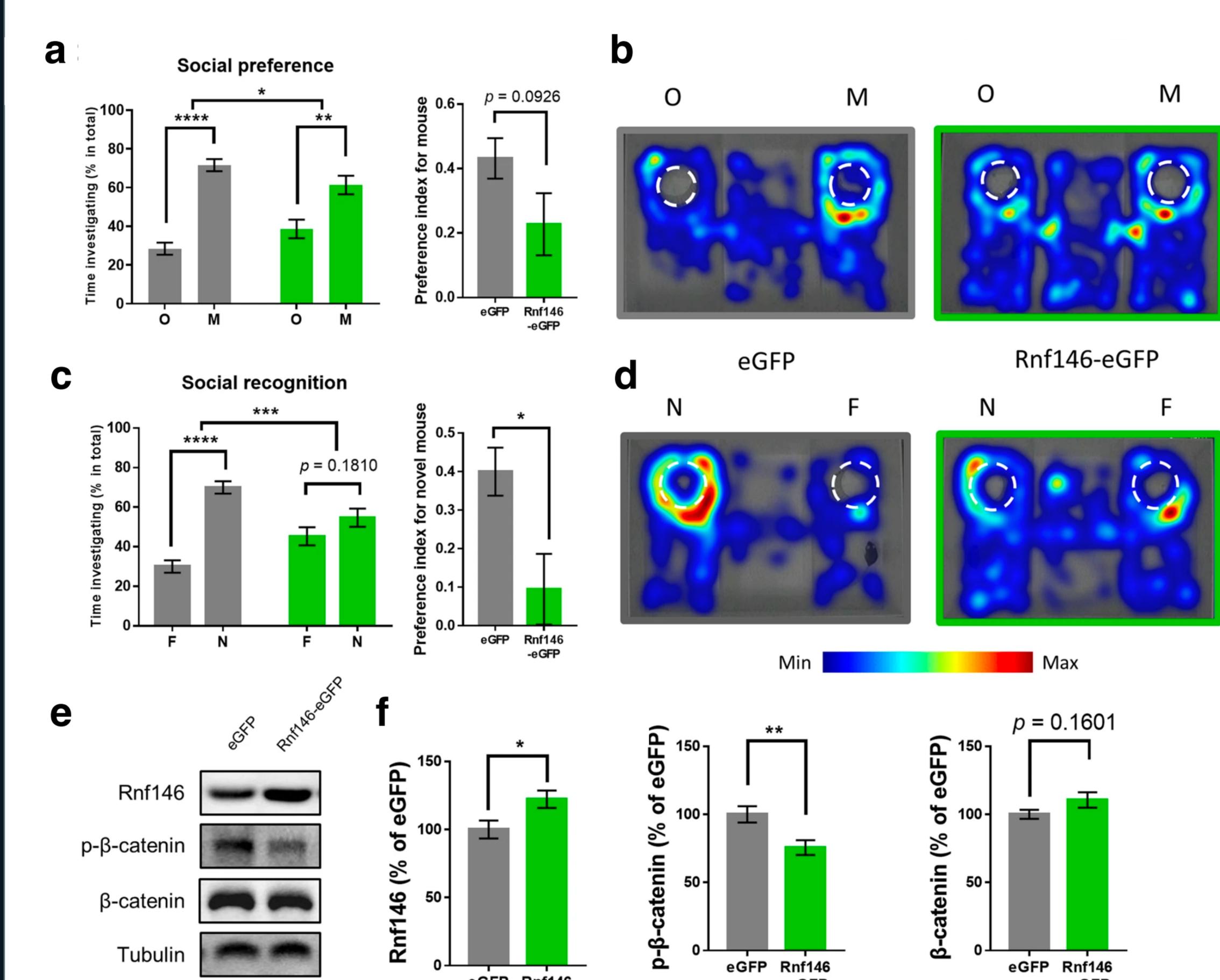


Figure 3. Rnf146 overexpression in the PFC causes social deficits in mice. **a.** Social preference test and preference index between Rnf146-overexpressing mice compared to eGFP-overexpressing control mice. **b.** Representative heatmap images of the social preference test. **c.** Social recognition test result and preference index between Rnf146- and eGFP-overexpressing mice. **d.** Representative heatmap images of the social recognition test. **e.** Western blot analysis result. **f.** Changes of Rnf146, p-β-catenin, and β-catenin levels in Rnf146 overexpression.

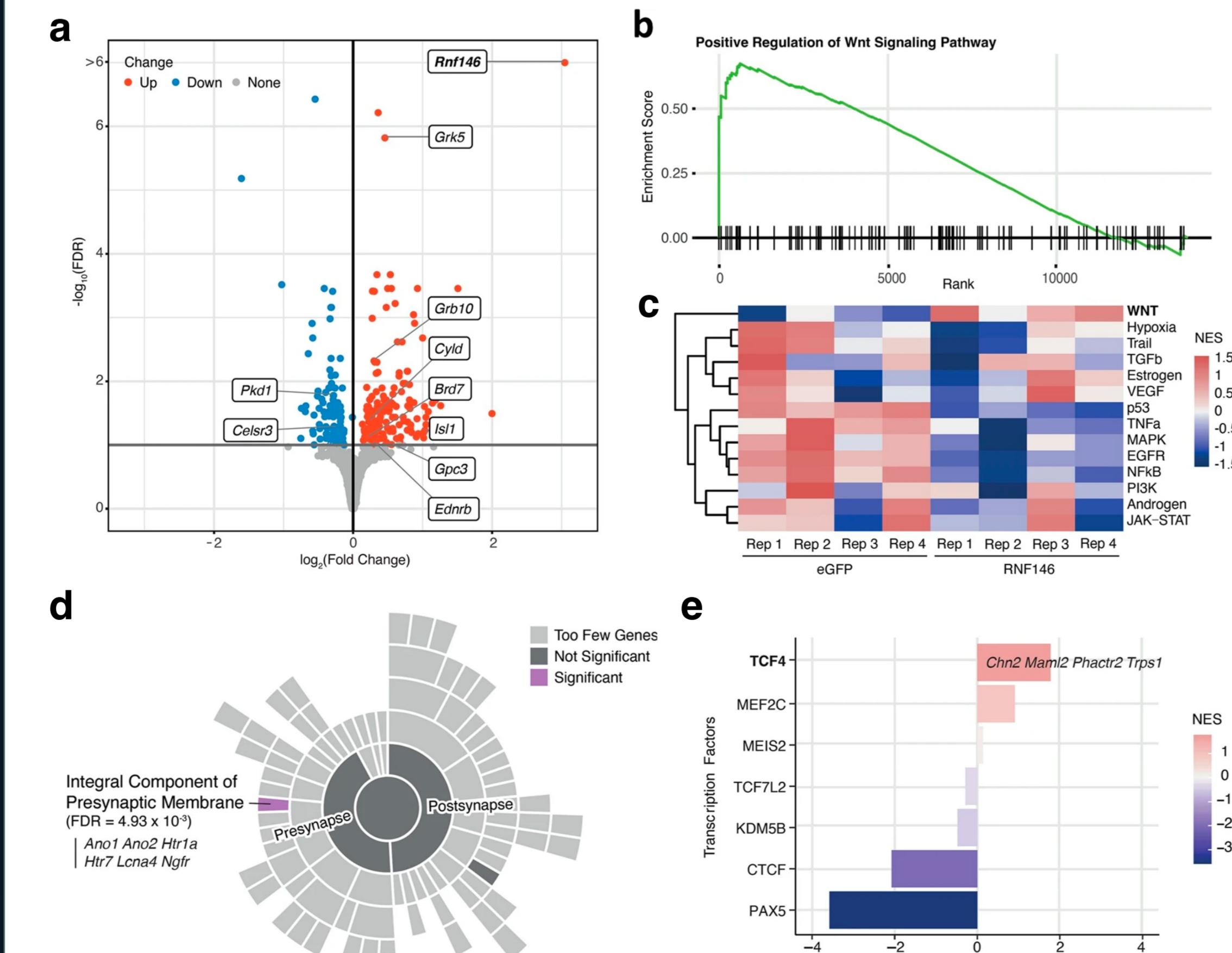


Figure 4. The Wnt signaling pathway is promoted by Rnf146 overexpression. **a.** Volcano plot showing DEGs between Rnf146- and eGFP-overexpressing mice. **b.** Enrichment plot depicting the elevated positive regulation of Wnt signaling pathway in Rnf146-overexpression. **c.** Heatmap of the pathway activity inferred by PROGENY. **d.** Enrichment of upregulated Rnf146-DEGs with synaptic locations. **e.** Enrichment with inferred transcription factor activities associated with ASD.

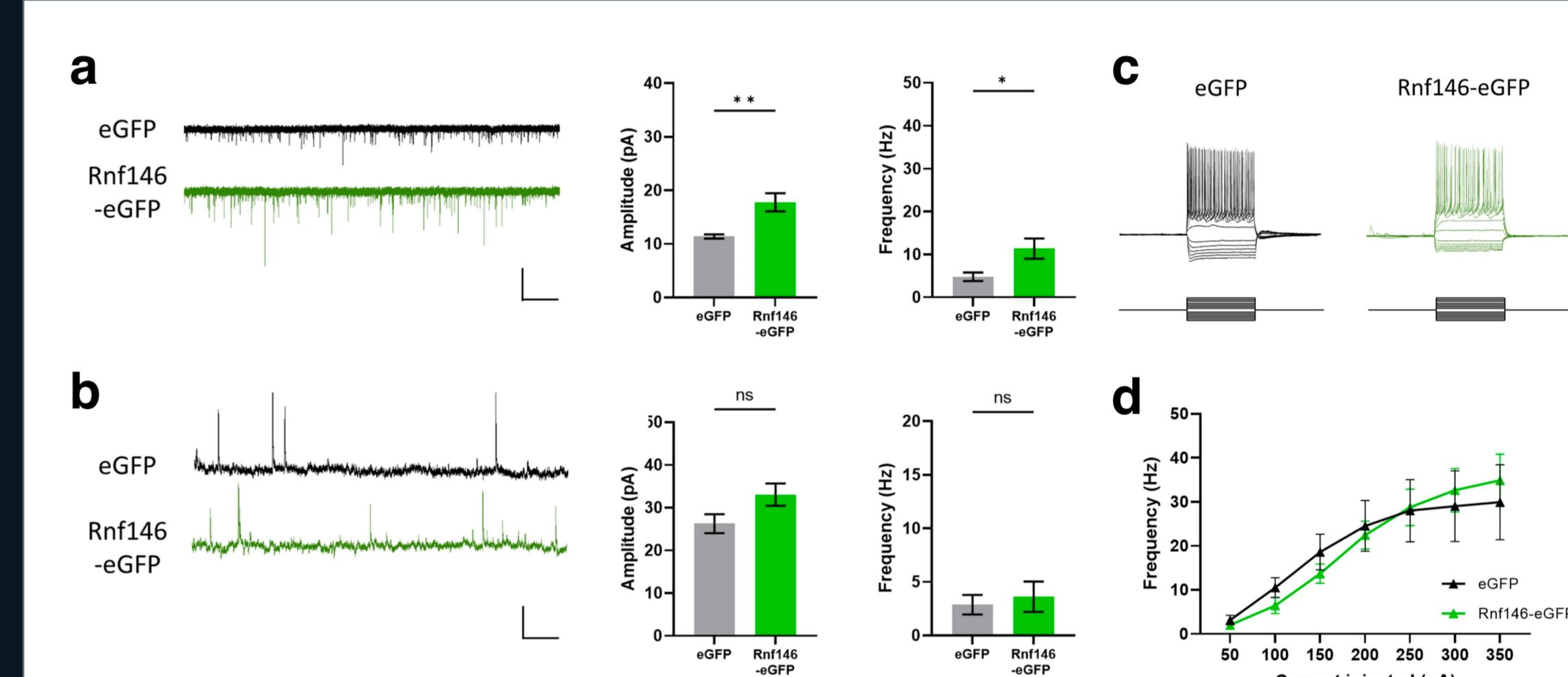


Figure 5. Rnf146 overexpression increases excitatory synaptic transmission in prefrontal pyramidal neurons. **a, b.** Spontaneous excitatory and inhibitory postsynaptic current (sEPSC, iEPSC) traces of prefrontal neurons in Rnf146- and eGFP-overexpressing mice. **c.** Representative traces of voltage responses of prefrontal neurons in Rnf146- or eGFP-overexpressing mice. **d.** Summary data of the number of action potentials evoked in response to 300 pA current steps.

DISCUSSIONS

Prenatal exposure to valproic acid (VPA) induces ASD-like social deficits, linked to increased Rnf146 expression activating the Wnt/β-catenin pathway. Rnf146 overexpression leads to impaired social behavior through heightened excitatory synaptic transmission. Dysregulation of the Wnt signaling is identified as a contributor to ASD etiology, impacting development and synaptic functions. These findings offer new insights into the molecular mechanism of VPA-associated ASD, suggesting the Rnf146-Wnt/β-catenin pathway as a potential target to alleviate social deficits in ASD.

REFERENCES

- H. S. Go et al., Prenatal exposure to valproic acid increases the neural progenitor cell pool and induces macrocephaly in rat brain via a mechanism involving the GSK-3beta/beta-catenin pathway. *Neuropharmacology* 63, 1028–1041 (2012).
- T. Chomiak, N. Turner, B. Hu, What We Have Learned about Autism Spectrum Disorder from Valproic Acid. *Patholog Res Int* 2013, 712758 (2013).
- C. Nicolini, M. Fahnestock, The valproic acid-induced rodent model of autism. *Experimental Neurology* 299, 217–227 (2018)
- H. O. Kalkman, A review of the evidence for the canonical Wnt pathway in autism spectrum disorders. *Mol Autism* 3, 10 (2012).

ACKNOWLEDGEMENT

This study was supported by grants from the National Research Foundation of Korea (NRF-2017M3C7A1026942 to Y.-S.L. and M.-S.K.; NRF-2018R1A5A2025964 to Y.-S.L.; NRF-2019M3E5D3073568 and NRF-2020R1C1C1003426 to J.-Y.A.; NRF-2022R1A2C2013377 to M.-S.K.; and NRF-2018H1A2A1061381 to G.P.) and Korea University (to J.-Y.A.) and Korea Basic Science Institute research program (C270100 to J.Y.K.).