

Sleep Disturbances are Associated with Brain Structural Covariance Networks in Preadolescence

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Beyond isolated brain regions:

Are sleep disturbances reflected in how the developing brain is organized?



Background

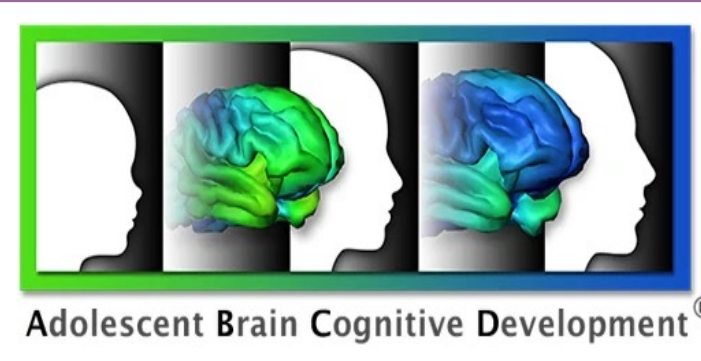
- Sleep disturbances (SD) affect 30–40% of children and are associated with adverse developmental outcomes [1].
- Preadolescence is marked by reorganization of brain morphology, reflected in coordinated structural variation across distributed brain systems [2].
- Existing group-level evidence has linked SD to regional brain structural alterations; however, little is known about how SD relate to individual-specific deviations in brain structural covariance organization and their developmental course.

Research Aim

- Characterize individual-specific alterations in brain structural covariance organization in children with SD.
- Determine whether these individual-specific alterations form a shared pattern across children with SD.
- Examine the developmental course of the shared structural covariance pattern according to sleep-disturbance status.

Sample Overview

ABCD	
Subjects	10,447
Age	9–10 years old at baseline
Follow-up	2-year
Sleep	Sleep Disturbance Scale for Children (SDSC)
Imaging	T1-weighted MRI
Covariates	Age, Sex, BMI, Puberty Stage, Site, Total Intracranial Volume (TIV)



Data Preparation

Baseline **1-y FU** **2-y FU**

SDSC:  **T1:** 

Inclusion criteria: at least one available T1-weighted MRI

FU: follow-up

N = 10,477 **N = 9692**

Sleep Phenotyping

Clinical cut-off: SDSC Total Score ≥ 39

Group	SDSC Total Score	Baseline: n	2-y FU: n
Stable Good sleepers (Never SD)	< 39	4420	3995
With SD history	≥ 39 at least at one time point	6057	5697

Time Point	Current SD	Current non-SD
Baseline	n = 3182	n = 2875
2-y FU	n = 2850	n = 2847

MRI Processing

T1-Weighted MRI $\rightarrow$ **CAT12 Morphometry** $\rightarrow$ **Parcellation** (Schaefer (200) + Tian (43)) $\rightarrow$ **Residualized GMVs** (Subjects $\times$ Parcels)

Covariate adjustment (age, Sex, BMI, TIV, site)

GMV: gray matter volume

Aim 1

Individual Differences in Structural Covariance Organization

Approach

- Compute covariance matrix
- Recompute covariance matrix
- Compute differences
- Edge wise Z-score and Threshold

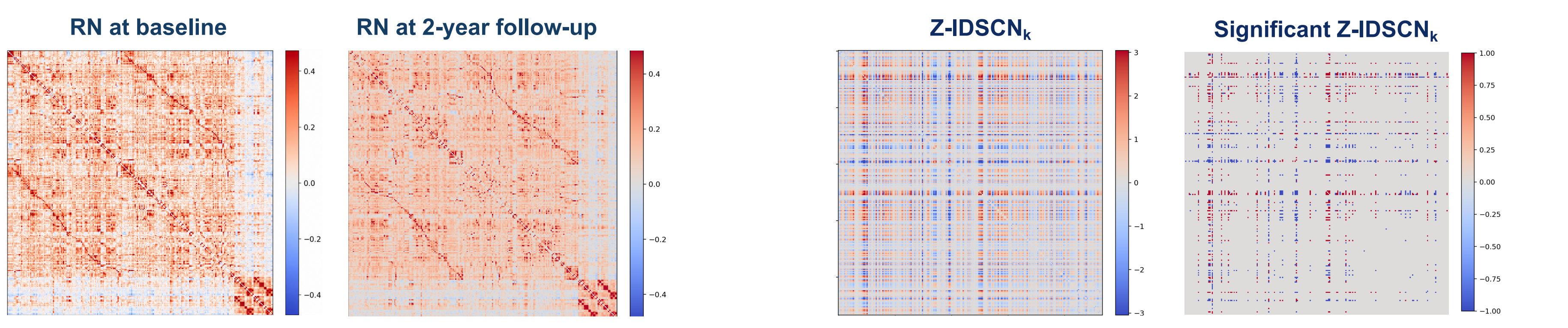
Good sleepers **Good sleepers + SD Subject_k**

RN **PN** **IDSCN = PN - RN**

RN: reference covariance network **PN:** perturbed covariance network **IDSCN:** individual differential structural covariance network

Key findings

- Reference networks showed high stability across resampling procedures
- Individual structural covariance deviations were reliably detectable beyond normative perturbation levels, demonstrating that IDSCN can capture inter/intra individual differences among children with SD.



Aim 2:

Shared Structural Covariance Alterations Across Children with SD

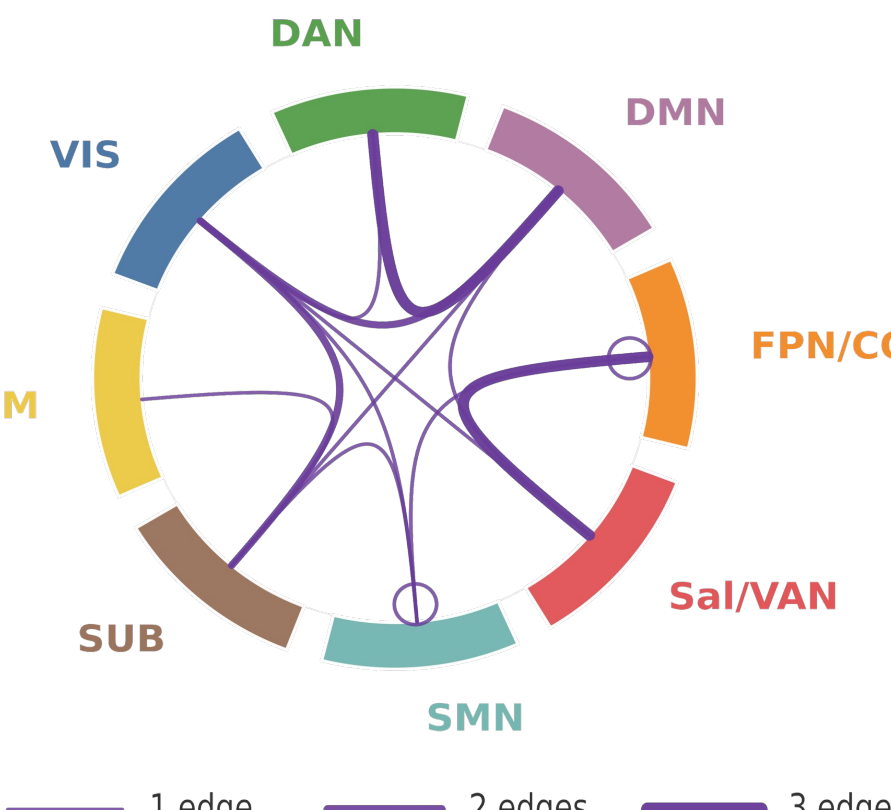
Approach

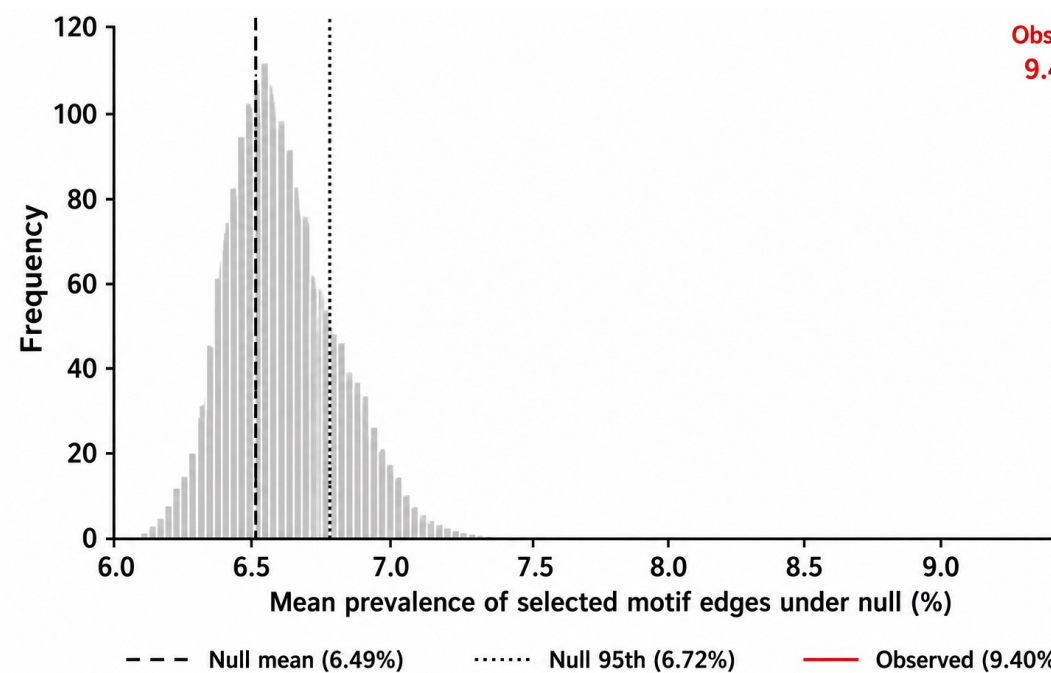
Significant Z-IDSCNs $\rightarrow$ **Frequency Map** $\rightarrow$ **Count across subjects** $\rightarrow$ **Top-20 edges** $\rightarrow$ **Permutation test** ($p < 0.05$)

The shared pattern was characterized by distributed cortical–subcortical systems rather than localized to a single network.

Key findings

- Despite substantial inter-individual variability, a shared pattern of structural covariance deviations emerged across children with SD. The pattern was defined by the 20 most prevalent deviation edges and was observed significantly more often than expected by chance.

Parcer i – Parcel j	Prevalence	Network Pair	Network Contribution
IDefaultB_PFCv_2 $\rightarrow$ ISalVentAttnB_IPL_1	10.01%	DMN – Sal/VAN	
IDefaultA_pCunPCC_1 $\rightarrow$ IVisPeri_StriCal_1	9.77%	DMN – VIS	
ISomMotB_Cent_2 $\rightarrow$ rSomMotA_9	9.53%	SMN – SMN	
ISomMotA_1 $\rightarrow$ IVisPeri_StriCal_1	9.53%	SMN – VIS	
ISalVentAttnB_IPL_1 $\rightarrow$ rDorsAttnA_IPS_2	9.53%	Sal/VAN – FPN/CON	
IDefaultA_pCunPCC_1 $\rightarrow$ rDorsAttnA_TempOcc_1	9.45%	DMN – DAN	
rVisCent_ExStr_3 $\rightarrow$ subcort_THA-DP-rh	9.45%	VIS – SUB	
IContA_PFCi_2 $\rightarrow$ ISalVentAttnB_IPL_1	9.45%	FPN/CON – Sal/VAN	
rDefaultA_PFCm_1 $\rightarrow$ rDorsAttnB_PostC_4	9.45%	DMN – DAN	
IContA_pCun_2 $\rightarrow$ rContB_Temp_2	9.45%	FPN/CON – FPN/CON	
IDefaultA_PFCi_2 $\rightarrow$ ISomMotB_S2_2	9.29%	FPN/CON – SMN	
rVisCent_ExStr_3 $\rightarrow$ subcort_aGP-rh	9.29%	VIS – SUB	
IDefaultC_IPL_1 $\rightarrow$ IVisPeri_StriCal_1	9.29%	DMN – VIS	
IContA_PFCi_2 $\rightarrow$ ISalVentAttnB_PFCi_1	9.29%	FPN/CON – Sal/VAN	
ISomMotA_4 $\rightarrow$ subcort_mAMY-rh	9.29%	SMN – SUB	
IVisPeri_StriCal_1 $\rightarrow$ rDorsAttnA_TempOcc_1	9.21%	VIS – DAN	
ISalVentAttnB_PFCi_1 $\rightarrow$ IVisCent_ExStr_5	9.21%	Sal/VAN – VIS	
IDefaultB_PFCd_3 $\rightarrow$ rDorsAttnB_PostC_4	9.21%	DMN – DAN	
ILimbicA_TempPole_2 $\rightarrow$ subcort_mAMY-rh	9.13%	LIM – SUB	
rDefaultA_PFCm_2 $\rightarrow$ subcort_pPUT-rh	9.13%	DMN – SUB	



Aim 3:

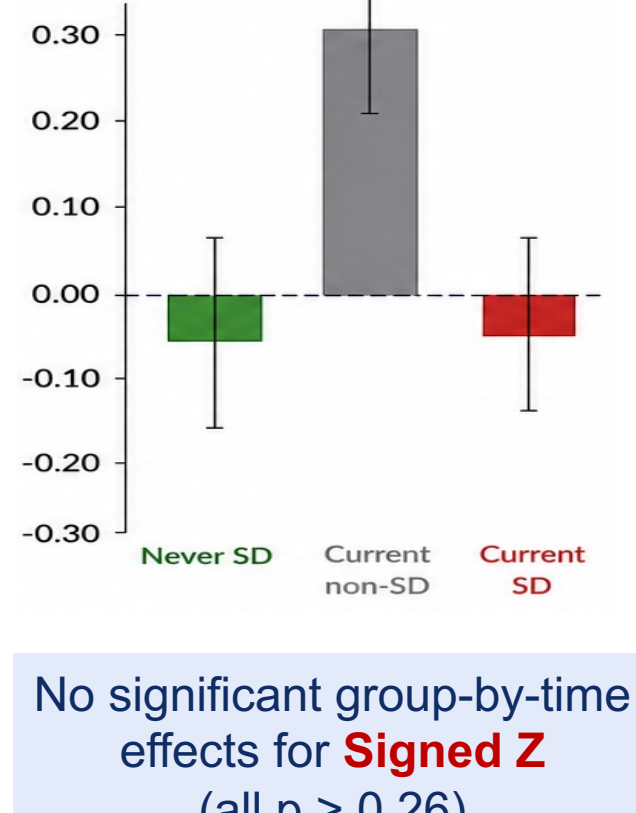
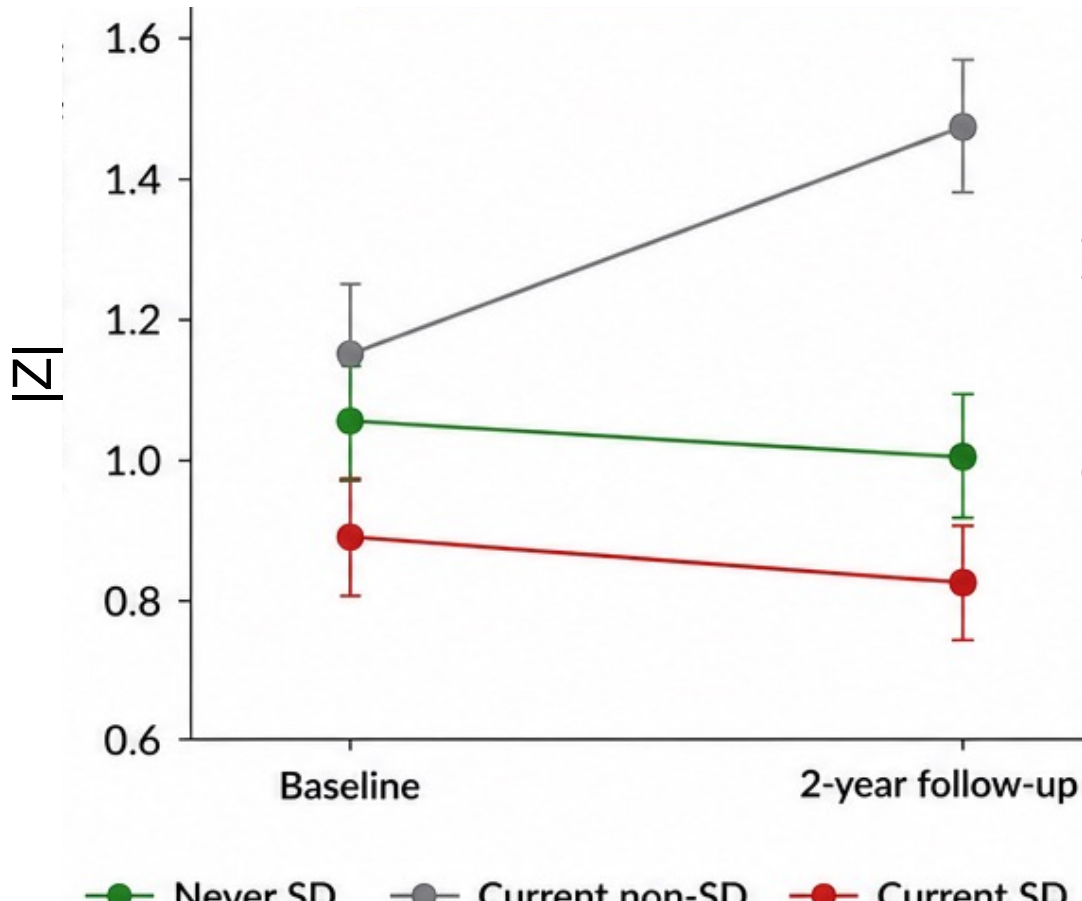
Longitudinal changes of the SD-related structural covariance pattern

Model : $|Z| \sim \text{Group} \times \text{Time} + \text{Pubertal stage} + (1|\text{subject}) + (1|\text{edge})$

Linear mixed-effect model with random intercepts for subject and edge

Longitudinal change in $ Z $	β	p
Current SD (vs never SD)	-0.003	0.86
Current non- SD (vs never SD)	0.037	0.06
Current non- SD vs Current SD	-0.041	0.006

Pubertal stage $\rightarrow$ -0.027 < 0.001



No significant group-by-time effects for **Signed Z** (all $p > 0.26$)

The longitudinal changes of the shared deviation pattern differed according to current SD status

Conclusions

- Distributed alterations in brain structural covariance organization support a network-level rather than region-specific view of SD in preadolescence.
- SD history may matter: longitudinal differences in structural covariance alteration were not fully captured by current symptom status alone.

Reference

- Lewien, C., et al., Sleep-related difficulties in healthy children and adolescents. BMC Pediatrics, 2021. 21(1): p. 82.
- Bethlehem, R.A.I., et al., Brain charts for the human lifespan. Nature, 2022. 604(7906): p. 525-533.

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