

Figure S1 Flowchart of the EEG signal feature extraction procedure for Auto-Neo-EEG. The boxes in yellow, orange, blue and green represent features related to amplitude, rEEG, spectral density and connectivity, respectively. EEG, electroencephalography; FIR, finite-impulse response; PSD, power spectral density; FD, fractal dimension; CSD, cross-PSD; rEEG, range EEG.

Table S1 Thirty findings for each EEG recording clinical report

Finding class	General description for the findings
Sleep-wake cycling	Abnormal sleep-wake cycling Sleep cycling can be divided into AS and QS period Sleep cycling cannot be divided into AS and QS state
Background	Tracé discontinu pattern in sleep state Tracé alternant pattern in sleep state Continuous pattern in sleep state Tracé discontinu pattern in awake state Tracé alternant pattern in awake state Continuous pattern in awake state Burst suppression Abnormal symmetry and synchrony Hemisphere asymmetry/asynchronous $\leq 50\%$ Hemisphere asymmetry/asynchronous $> 50\%$ Borderline low voltage Abnormally low voltage Dysmaturity
Seizures	No obvious discharge Seizure
Waves	Spike waves rhythmic discharges Sharp waves rhythmic discharges Low amplitude fast wave rhythmic discharges Sleep state sharp wave Sleep state sharp-slow wave complex Sleep state spike wave Sleep state spike-slow wave complex Awake state abnormal wave Awake state sharp wave Awake state sharp-slow wave complex Awake state spike wave Awake state spike-slow wave complex

EEG, electroencephalography; AS, active sleep; QS, quiet sleep.

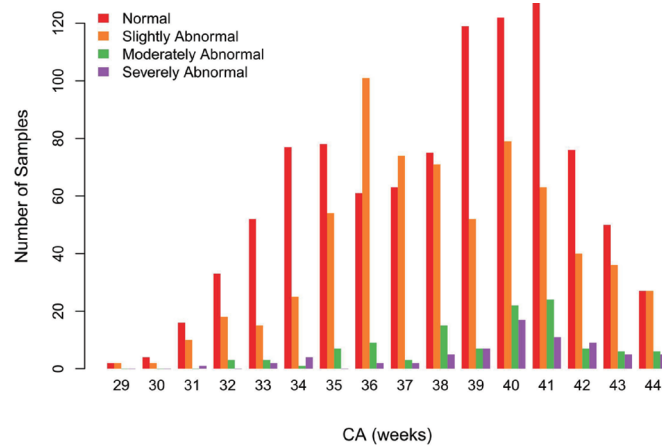


Figure S2 Detailed CA distribution for all samples in the neonatal EEG dataset. Sample numbers in each CA week are divided into four groups according to the EEG report conclusion label. EEG, electroencephalography; CA, conceptional age.

Table S2 Relationship between clinical systems and EEG report conclusion

Disease Classification	Normal (N=992)	Slightly abnormal (N=674)	Moderately abnormal (N=115)	Severely abnormal (N=70)
HIE	71 (50%, OR =0.93, P=0.7)	40 (28%, OR =0.77, P=0.2)	19 (13%, OR =2.2, P=0.006)	12 (8%, OR =2.2, P=0.02)
Cardiopulmonary disease	62 (56%, OR =1, P=0.8)	44 (40%, OR =1.1, P=0.6)	5 (5%, OR =0.73, P=0.7)	0 (0%, OR =0, P=0.03)
Central nervous system infection	43 (41%, OR =0.76, P=0.1)	41 (39%, OR =1.1, P=0.8)	13 (12%, OR =2, P=0.05)	9 (8%, OR =2.2, P=0.04)
Intracranial hemorrhage	81 (63%, OR =1.2, P=0.3)	32 (25%, OR =0.68, P=0.06)	11 (9%, OR =1.4, P=0.4)	5 (4%, OR =1, P=0.8)
Congenital metabolic disease	7 (24%, OR =0.45, P=0.05)	11 (38%, OR =1, P=0.9)	3 (10%, OR =1.7, P=0.4)	8 (28%, OR =7.3, P=7e-05)
Temporary metabolic disorder	131 (72%, OR =1.3, P=0.02)	44 (24%, OR =0.66, P=0.02)	5 (3%, OR =0.44, P=0.09)	2 (1%, OR =0.29, P=0.09)
Perinatal stroke	5 (38%, OR =0.72, P=0.6)	6 (46%, OR =1.3, P=0.6)	1 (8%, OR =1.2, P=0.6)	1 (8%, OR =2, P=0.4)
Premature	482 (53%, OR =0.99, P=0.9)	372 (41%, OR =1.1, P=0.1)	35 (4%, OR =0.62, P=0.02)	22 (2%, OR =0.64, P=0.07)
Genetic factors/syndrome	32 (50%, OR =0.93, P=0.8)	21 (33%, OR =0.9, P=0.8)	8 (12%, OR =2, P=0.08)	3 (5%, OR =1.2, P=0.7)
Unexplained convulsions	31 (35%, OR =0.65, P=0.05)	38 (43%, OR =1.2, P=0.4)	14 (16%, OR =2.5, P=0.005)	6 (7%, OR =1.8, P=0.2)
Other	47 (63%, OR =1.2, P=0.4)	25 (33%, OR =0.92, P=0.8)	1 (1%, OR =0.21, P=0.1)	2 (3%, OR =0.71, P=1)

Each box contains the number of patients, percentage of patients in this clinical system, OR compared to background and P value (P) by Fisher's exact test. EEG, electroencephalography; HIE, hypoxic ischemic encephalopathy; OR, odds ratio.

Table S3 Interrater agreement for report conclusion between two experts for 96 EEG recording subjects

Patient	CA	Gender	Outcome by expert 1	Outcome by expert 2
1	42+4	Male	Severely abnormal	Severely abnormal
2	40+3	Male	Normal	Normal
3	34+2	Male	Slightly abnormal	Slightly abnormal
4	43+4	Female	Severely abnormal	Severely abnormal
5*	41+3	Male	Severely abnormal	Moderately abnormal
6	39+6	Male	Normal	Normal
7	40+2	Male	Normal	Normal
8*	39+1	Female	Slightly abnormal	Normal
9	41+5	Female	Slightly abnormal	Slightly abnormal
10	35+2	Male	Moderately abnormal	Moderately abnormal
11	41+6	Male	Normal	Normal
12	40	Male	Normal	Normal
13	44+4	Male	Moderately abnormal	Moderately abnormal
14	40+1	Female	Moderately abnormal	Moderately abnormal
15	42	Female	Moderately abnormal	Moderately abnormal
16	41	Female	Severely abnormal	Severely abnormal
17	39+5	Female	Severely abnormal	Severely abnormal
18	39	Male	Normal	Normal
19	40+4	Male	Normal	Normal
20	37+3	Female	Slightly abnormal	Slightly abnormal
21	37+2	Male	Moderately abnormal	Moderately abnormal
22	41+5	Male	Normal	Normal
23	41+1	Female	Normal	Normal
24	44	Male	Severely abnormal	Severely abnormal
25	34+3	Male	Normal	Normal
26	41+1	Male	Normal	Normal
27	42+5	Female	Slightly abnormal	Slightly abnormal
28	40+5	Male	Normal	Normal
29	34+4	Female	Normal	Normal
30	37+3	Female	Normal	Normal
31	33+2	Female	Normal	Normal
32	38+4	Female	Slightly abnormal	Slightly abnormal
33	36+2	Female	Slightly abnormal	Slightly abnormal
34	35+2	Male	Normal	Normal
35	37	Male	Slightly abnormal	Slightly abnormal
36	39	Male	Normal	Normal
37	34+3	Male	Normal	Normal
38	39+6	Female	Slightly abnormal	Slightly abnormal
39	41+1	Male	Slightly abnormal	Slightly abnormal
40	36+3	Female	Normal	Normal
41	41+3	Male	Normal	Normal
42	35+3	Female	Normal	Normal
43	38+2	Female	Slightly abnormal	Slightly abnormal
44	32+1	Male	Normal	Normal
45	33+4	Female	Slightly abnormal	Slightly abnormal
46	42+1	Male	Normal	Normal
47	39	Male	Normal	Normal
48*	37+5	Female	Slightly abnormal	Normal
49	32+3	Male	Slightly abnormal	Slightly abnormal
50	39	Female	Slightly abnormal	Slightly abnormal
51	31+4	Male	Slightly abnormal	Slightly abnormal
52	34+5	Male	Slightly abnormal	Slightly abnormal
53	37+2	Female	Slightly abnormal	Slightly abnormal
54	38+1	Female	Slightly abnormal	Slightly abnormal
55	38+3	Male	Moderately abnormal	Moderately abnormal
56	33+3	Female	Slightly abnormal	Slightly abnormal
57	35+3	Male	Moderately abnormal	Moderately abnormal
58*	37+3	Female	Normal	Slightly abnormal
59	35+1	Male	Normal	Normal
60	34+3	Male	Normal	Normal
61	36+5	Male	Normal	Normal
62	39+6	Male	Normal	Normal
63*	36+4	Male	Normal	Slightly abnormal
64	37+6	Male	Normal	Normal
65	34+4	Male	Normal	Normal
66*	40+1	Male	Slightly abnormal	Moderately abnormal
67	41+3	Male	Normal	Normal
68	33+1	Male	Slightly abnormal	Slightly abnormal
69	38+4	Male	Normal	Normal
70	39+1	Female	Slightly abnormal	Slightly abnormal
71	33+6	Male	Normal	Normal
72	35+2	Female	Normal	Normal
73	42+1	Female	Moderately abnormal	Moderately abnormal
74	41+4	Male	Normal	Normal
75	34+6	Female	Normal	Normal
76	35	Male	Normal	Normal
77	40+6	Female	Moderately abnormal	Moderately abnormal
78	43+4	Male	Moderately abnormal	Moderately abnormal
79	40+3	Male	Severely abnormal	Severely abnormal
80	40+2	Male	Severely abnormal	Severely abnormal
81	39+5	Female	Moderately abnormal	Moderately abnormal
82	39+6	Male	Severely abnormal	Severely abnormal
83	40+1	Male	Moderately abnormal	Moderately abnormal
84	41+4	Male	Severely abnormal	Severely abnormal
85	40+3	Female	Severely abnormal	Severely abnormal
86	35+5	Male	Severely abnormal	Severely abnormal
87	42+2	Male	Moderately abnormal	Moderately abnormal
88	38+1	Female	Moderately abnormal	Moderately abnormal
89	40+1	Male	Moderately abnormal	Moderately abnormal
90	40+2	Female	Moderately abnormal	Moderately abnormal
91	39+2	Female	Severely abnormal	Severely abnormal
92*	40+1	Male	Moderately abnormal	Severely abnormal
93*	38+3	Female	Moderately abnormal	Severely abnormal
94	41+5	Male	Severely abnormal	Severely abnormal
95	43+2	Male	Severely abnormal	Severely abnormal
96*	40+4	Male	Moderately abnormal	Slightly abnormal

*, subjects inconsistent for report conclusion level between two experts. EEG, electroencephalography.

Table S4 The performance of Auto-Neo-EEG in predicting report conclusions by absolute CA difference

Strategy	Dataset	Predicted label	Original label				TP	TN	FP	FN	Sensitivity	Specificity	Accuracy
			Normal	Slightly abnormal	Moderately abnormal	Severely abnormal							
Predicted by absolute CA difference	Model-developing dataset	Normal (0, 2.9]	352	129	19	5	352	593	153	493	41.66%	79.49%	59.4%
		Slightly abnormal (2.9, 7.6]	321	177	16	12	177	658	349	407	30.31%	65.34%	52.48%
		Moderately abnormal (7.6, 14.5]	152	172	27	8	27	1161	332	71	27.55%	77.76%	74.67%
		Severely abnormal (14.5, Inf]	20	106	36	39	39	1365	162	25	60.94%	89.39%	88.25%
	Validation dataset	Normal (0, 2.9]	42	21	2	1	42	89	24	105	28.57%	78.76%	50.38%
		Slightly abnormal (2.9, 7.6]	51	25	2	0	25	117	53	65	27.78%	68.82%	54.62%
		Moderately abnormal (7.6, 14.5]	35	25	5	1	5	182	61	12	29.41%	74.9%	71.92%
		Severely abnormal (14.5, Inf]	19	19	8	4	4	208	46	2	66.67%	81.89%	81.54%

EEG, electroencephalography; CA, conceptional age; TP, true positive, TN, true negative, FP, false positive, FN, false negative.

Table S5 The performance of Auto-Neo-EEG in predicting report conclusions in each pair-wise comparison

Prediction strategy	Dataset	Prediction Label	Original label				Sensitivity	Specificity	Accuracy	AUC (95% CI)
			Normal	Slightly abnormal	Moderately abnormal	Severely abnormal				
Severely abnormal vs. others	Model-developing dataset	F		1,507		0	100%	98.69%	98.74%	1.000 (0.999–1.000)
		T		20		64				
	Validation dataset	F		244		0	100%	96.06%	96.15%	0.984 (0.970–0.999)
		T		10		6				
Moderately abnormal vs. slightly abnormal + normal	Model-developing dataset	F	1,245		14	–	85.71%	87.12%	87.03%	0.919 (0.885–0.955)
		T	184		84	–				
	Validation dataset	F	210		6	–	64.71%	88.61%	87.01%	0.857 (0.741–0.973)
		T	27		11	–				
Slightly abnormal vs. normal	Model-developing dataset	F	692	221	–	–	62.16%	81.89%	73.83%	0.784 (0.759–0.808)
		T	153	363	–	–				
	Validation dataset	F	107	48	–	–	46.67%	72.79%	62.87%	0.647 (0.576–0.718)
		T	40	42	–	–				
Abnormal vs. normal	Model-developing dataset	F	728		160		78.55%	86.15%	82.59%	0.906 (0.892–0.921)
		T	117		586					
	Validation dataset	F	109		45		60.18%	74.15%	68.08%	0.713 (0.648–0.777)
		T	38		68					
Total number	Model-developing dataset		845	584	98	64	–	–	–	–
	Validation dataset		147	90	17	6	–	–	–	–

EEG, electroencephalography; AUC, area under the curve; CI, confidence interval; T, true, F, false.