

A Hierarchical Gaussian Mixture Model for Continuous High-Resolution Sleep Analysis

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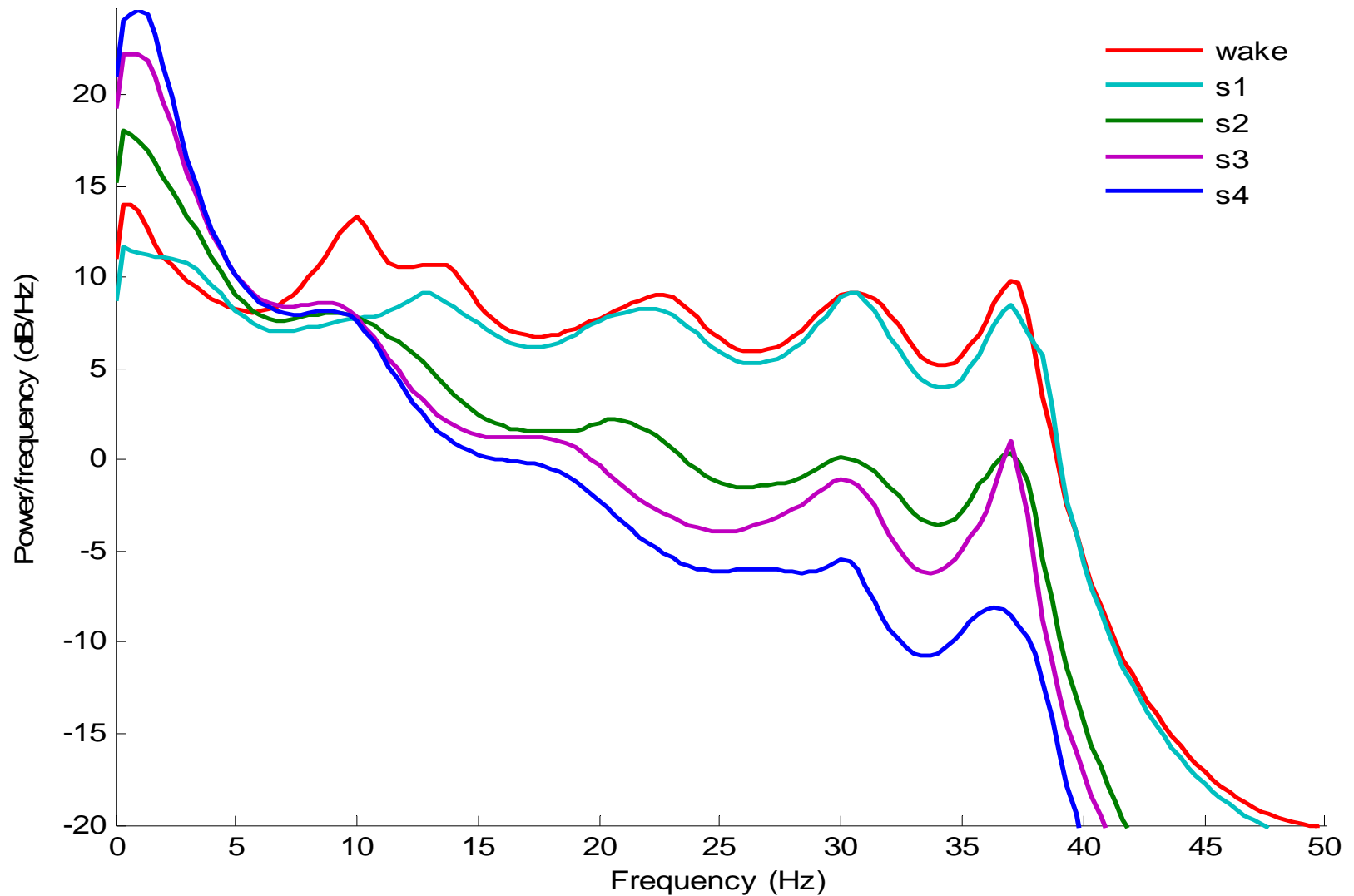
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Limits of R&K x Probabilistic Continuous Representation of Sleep

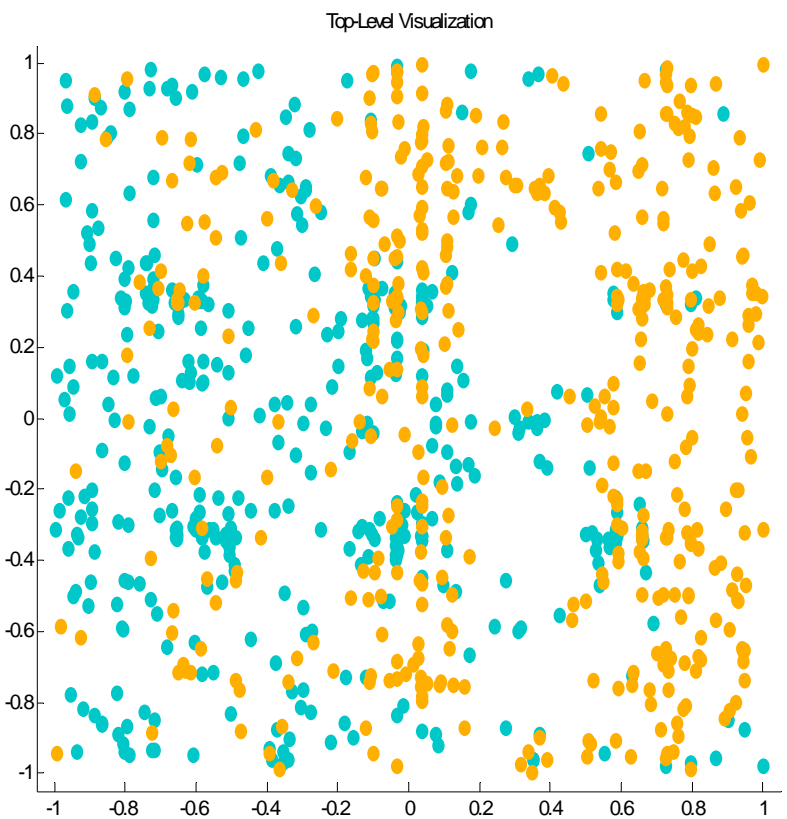
- coarse time resolution (20 or 30 sec) x 3 sec - up to the sampling frequency
- limitation to the central electrodes (C3,C4) x multi-electrodes model
- amplitude dependent delta waves rule for SWS x AR representation of the power spectra
- discrete clustering/classification x continuous probability curves of sleep stages

Example 1: mean AR(10)-based power spectrum

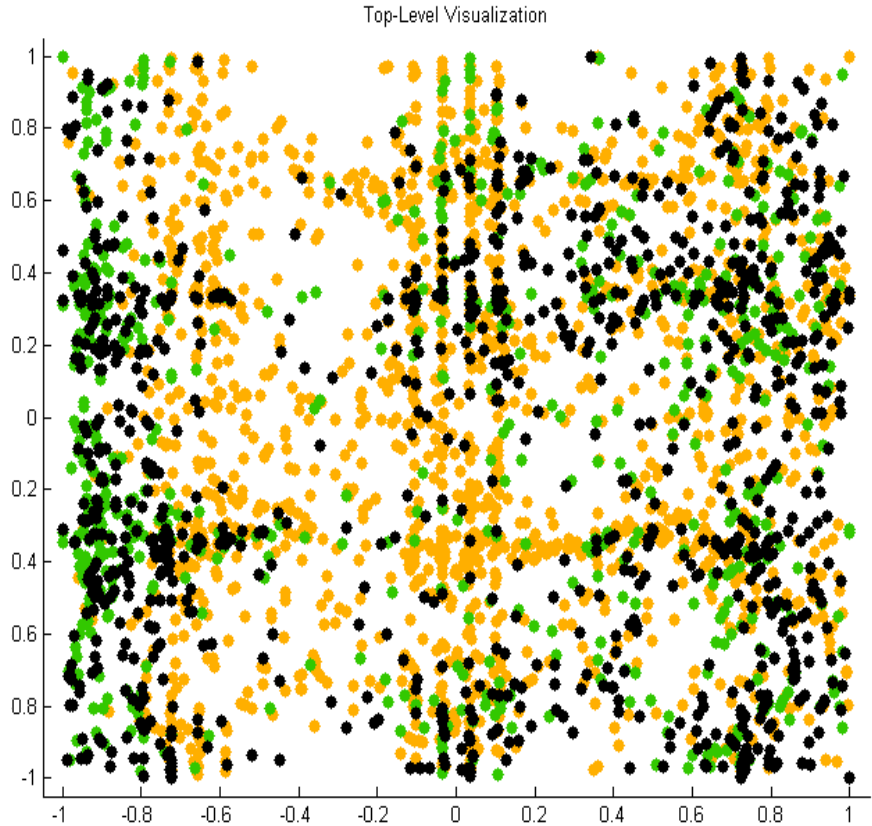


Example 2: 2-D projection of AR(10) coefficients - GTM

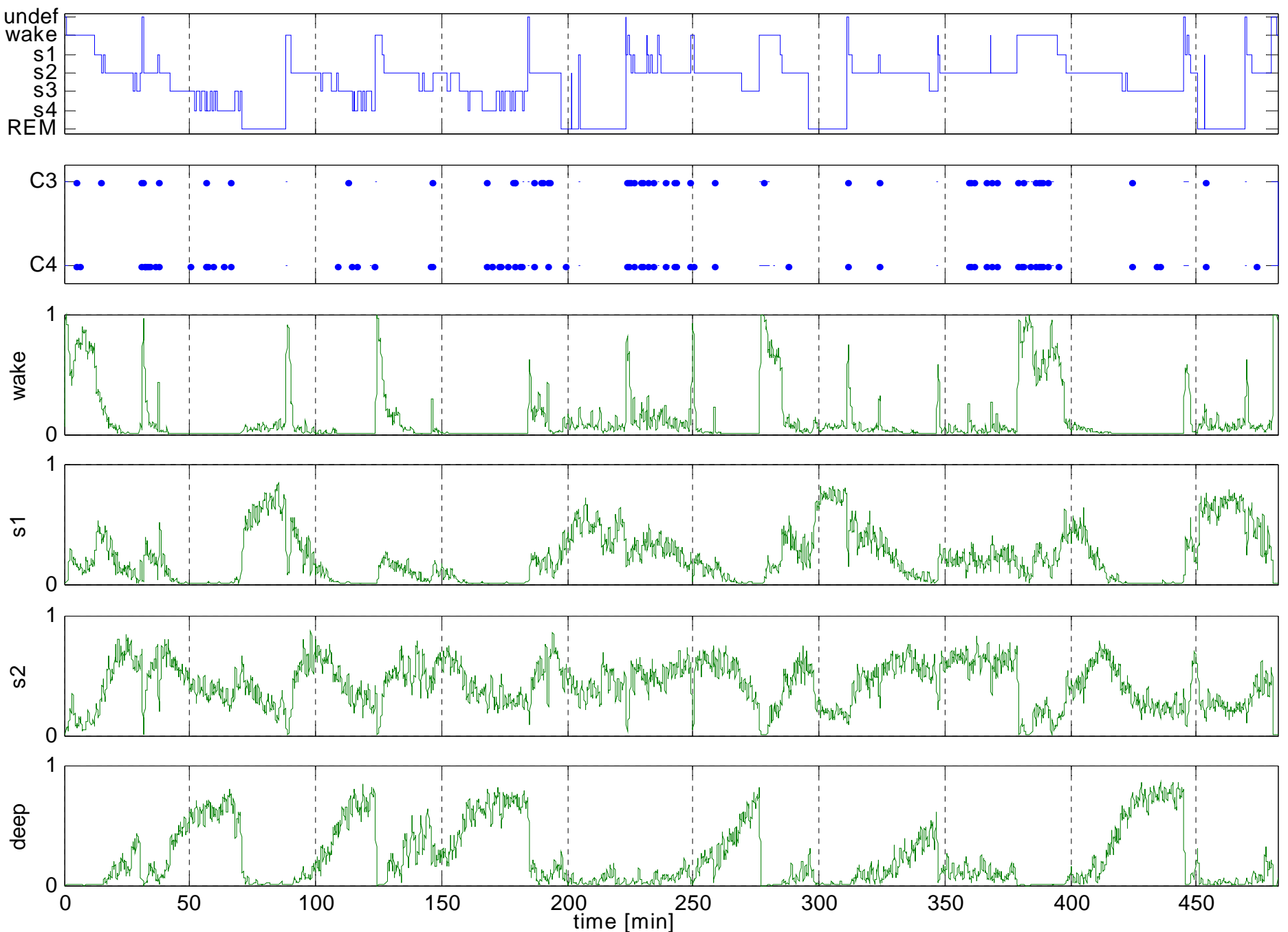
S1 vs S2



S2 vs deep (S3+S4)



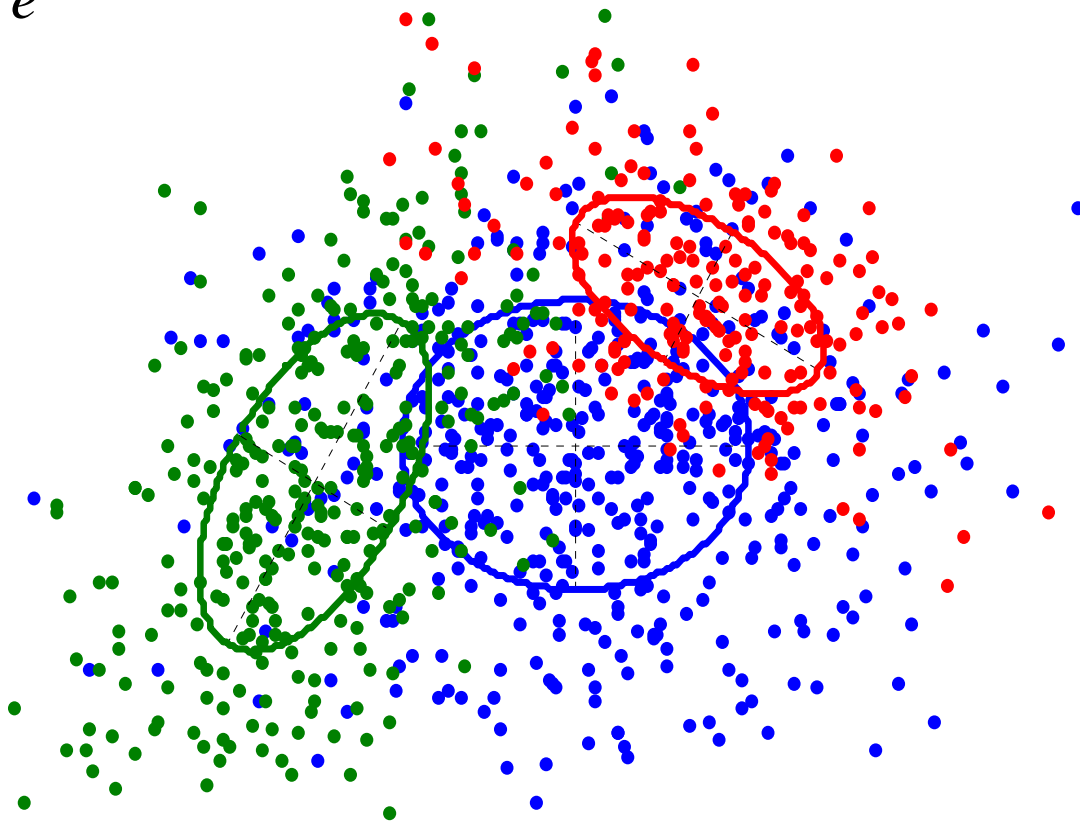
subject b0042, 2nd night, age 74, female, healthy control, filter 15 time points



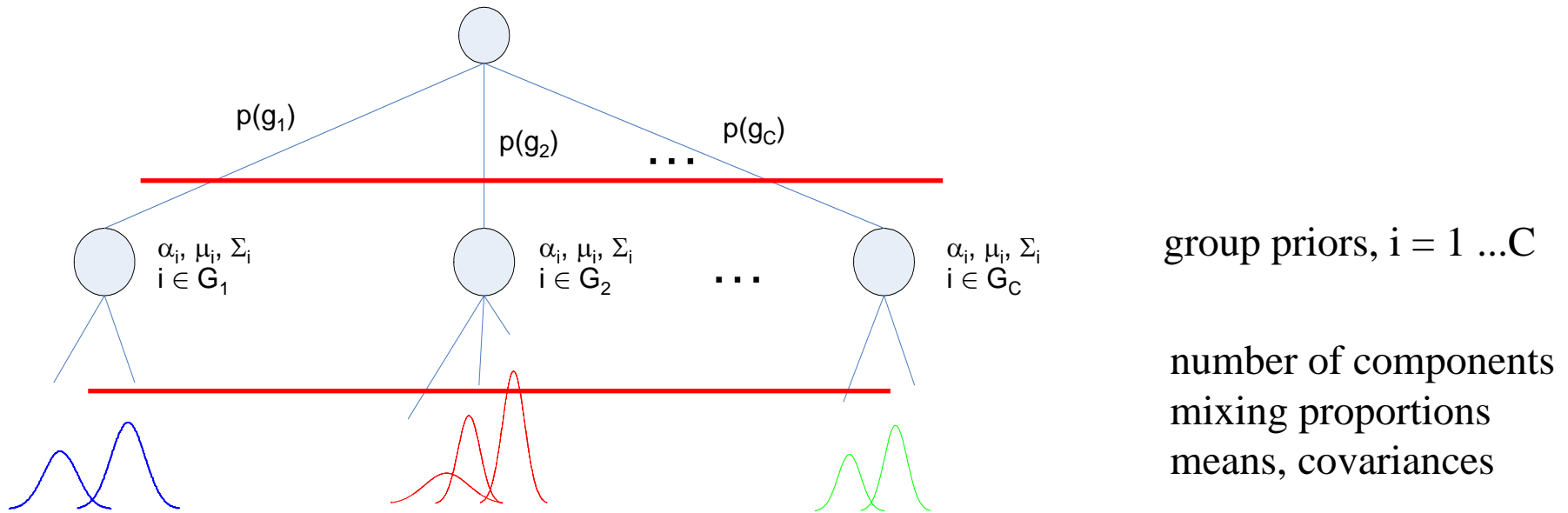
Gaussian Mixture Model (GMM)

$$p(x) = \sum_{k \in K} \alpha_k p(x | \theta_k) \quad \text{where} \quad \sum_{k \in K} \alpha_k = 1, \quad \alpha_k \geq 0$$

$$p(x | \theta_k) = (2\pi)^{-d/2} |\Sigma_k|^{-1/2} e^{-(x - \mu_k)^T \Sigma_k^{-1} (x - \mu_k) / 2}$$



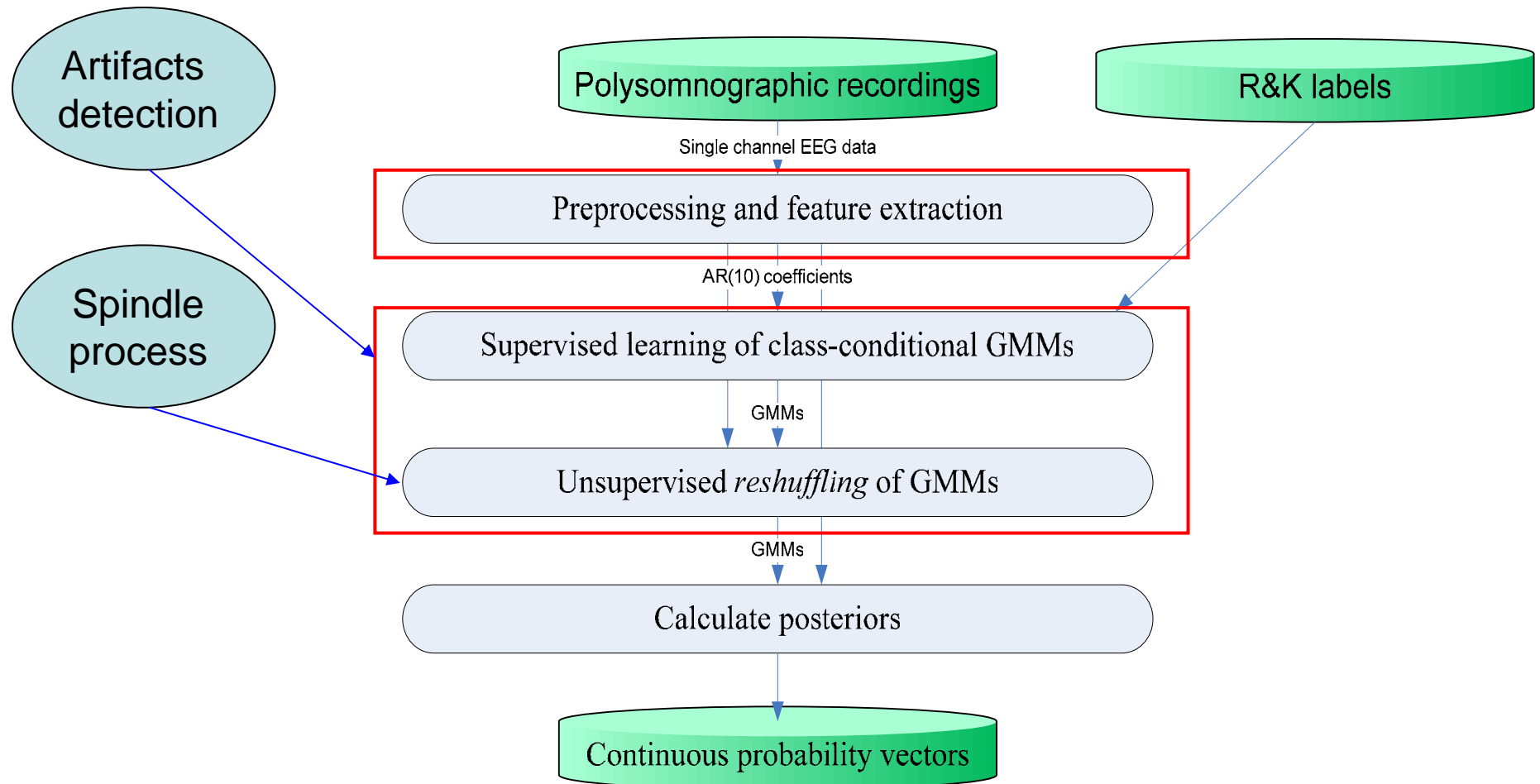
Structure of a Hierarchical GMM



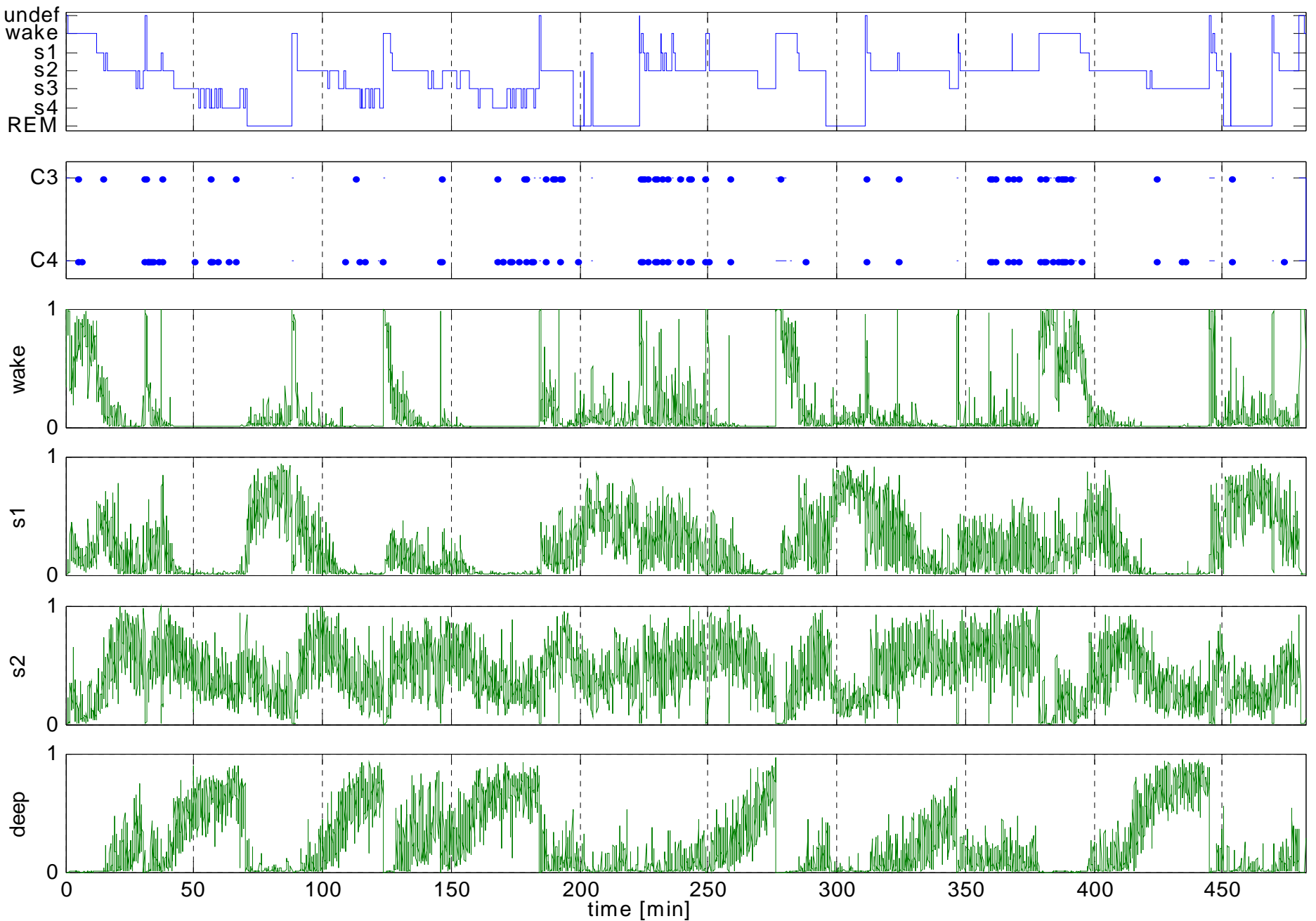
Step1: train mixtures of individual classes, set group priors

Step2: unsupervised (semi-supervised) *reshuffling*, keep structure

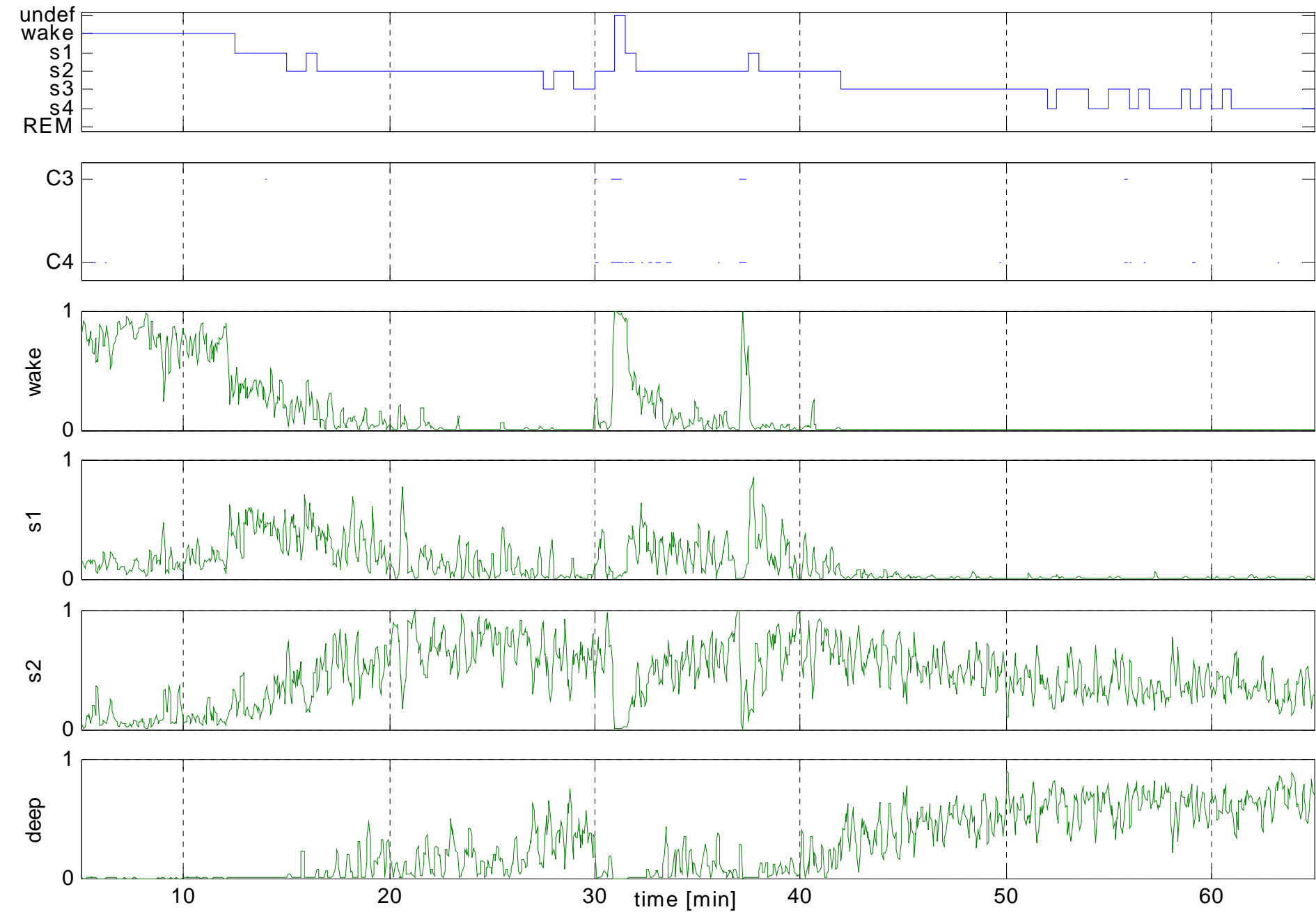
Data flow of the model building process



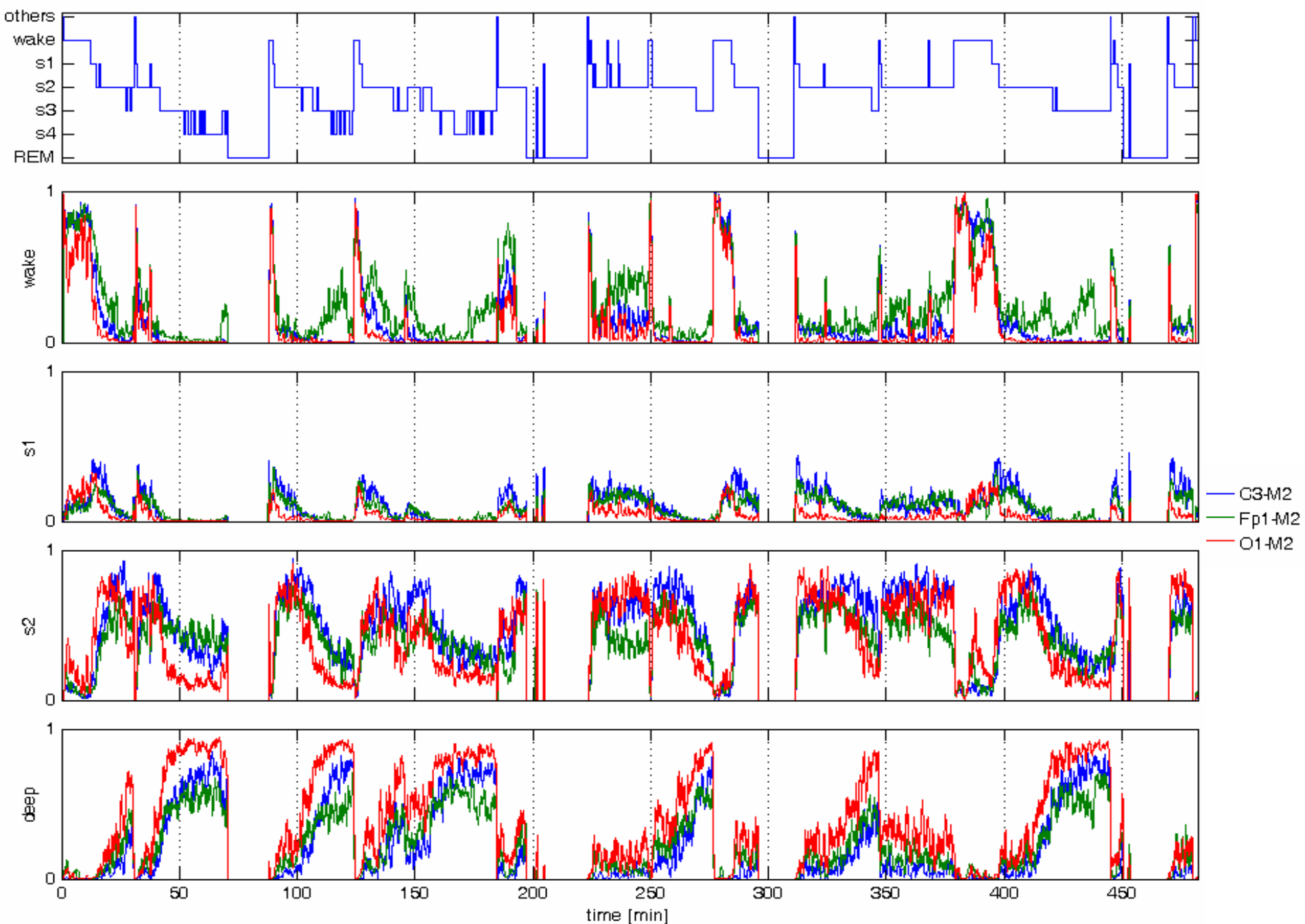
subject b0042, 2nd night, age 74, female, healthy control, filter 3 time points



subject b0042, 2nd night, age 74, female, healthy control, filter 3 time points



subject b0042, 2nd night, age 74, female, healthy control, filter 15 time points



Results

Normal Healthy Subjects (PSQI \leq 5)

- 176 healthy subjects (83 males and 93 females) aged between 20 and 95 years.

A) Night differences (p-values < 0.01)

Sleep Quality Index (SSA-1) (Saletu et al. 1987)

c_eff	-0.538	rk_eff	-0.542
c_wake_tsp	0.527	rk_wake_tsp	0.517
c_rAUC_wake	0.529		
c_s1_tst	0.328	rk_s1_tst	0.332
c_rAUC1_s1	0.336		
c_s2	-0.303	rk_s2	-0.325
c_rAUC_s2	-0.311		
		rk_s4	-0.245
c_rAUC2_deep	-0.401		
c_rAUC1_deep	-0.396		
c_rAUC_deep	-0.289		
c_fw	0.300	rk_fw	0.213

Mood (VAS score)

c_s2	0.244	rk_s2	0.258
c_s2_tst	0.201	rk_s2_tst	0.201
c_rAUC_s2	0.250		

Drowsiness (VAS score)

c_eff	-0.201	rk_eff	-0.229
c_wake_tsp	0.196	rk_wake_tsp	0.219
c_AUC_wake	0.205		
c_AUC1_deep	-0.197		

B) Individual Nights (p-values < 0.01)

Fine motor activity test
(Grunberger 1977)

Alphabetical cross-out test
(Grunberger 1977)

c_eff	0.291	rk_eff	0.223
c_wake_tsp	-0.387	rk_wake_tsp	-0.312
c_AUC1_tsp_wake	-0.458		
c_AUC2_tsp_wake	-0.458		
c_fw	-0.457	rk_fw	-0.307
c_fs	-0.423	rk_fs	-0.280
c_AUC1_s2	-0.374		
c_AUC_s1	-0.337		

c_wake_tsp	-0.307	rk_wake_tsp	-0.296
c_AUC1_tsp_wake	-0.381		
		rk_s1_tst	-0.211
c_AUC1_s1	-0.340		
c_AUC_s2	-0.250		
		rk_s4	0.299
c_fw	-0.354	rk_fw	-0.277
c_eff	0.297	rk_eff	0.281

Conclusions

- Continuous probabilistic sleep modeling shows **higher flexibility to model several characteristics of sleep** (sleep arousals, sleep stages transitions, delta process)
- The continuous sleep model has shown **the same or a higher level of information** about the sleep process in the investigated correlation and discrimination (not presented here) tasks
- The continuous sleep model can successfully **complement the R&K** standard for a more comprehensive description of sleep
- The probabilistic approach allows **principled sensor and different processes fusion** (spindle process, EOG or EMG modeling) and **a new sleep process interpretation**