Recap

Stability Selection

random subsampling of half of the data:

$$I^{*1}, \dots, I^{*B} \text{ independent}$$

$$I^{*b} \underbrace{\text{random subsample}}_{\text{without replacement}} \subset \{1, \dots, n\}, |I^{*b}| = \lfloor n/2 \rfloor$$

feature selection algorithm $\hat{S}_{\lambda} \subseteq \{1, \dots, p\}$

stability of selected single features:

$$\hat{\Pi}_j(\lambda) = \mathbb{P}^*[j \in \hat{S}_\lambda(I^*)] pprox B^{-1} \sum_{b=1}^B I(j \in \hat{S}_\lambda(I^{*b}))$$

Why half-sampling? I.e., subsampling without replacement with $|I^{*b}| = |n/2|$?

Freedman (1977): sampling without replacement with subsample size $m = \lfloor n/2 \rfloor$ is closest to i.i.d. sampling *n* individuals with replacement (i.e., bootstrap resampling)

"closest" means w.r.t total variance distance

Connecting to false discoveries

$$\hat{S}_{\text{stable}} = \{j; \max_{\lambda \in \Lambda} \hat{\Pi}_j(\lambda) \ge \pi_{\text{thr}}\}$$

Choice of π_{thr} ?

as a measure for type I error control (against false positives):

$$V =$$
 number of false positives $= |\hat{S}_{\text{stable}} \cap S_0^c|$

where S_0 is the set of the true relevant features $\hat{S}_{\Lambda} = \bigcup_{\lambda \in \Lambda} \hat{S}(\lambda)$ $q_{\Lambda} = \mathbb{E}[|\hat{S}_{\Lambda}(\underbrace{I}_{random subsample})|]$