

Learning Individualized Treatment Rules with Many Treatments: A Supervised Clustering Approach Using Adaptive Fusion

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Individualized Decision Making

- Example: Personalized Medicine

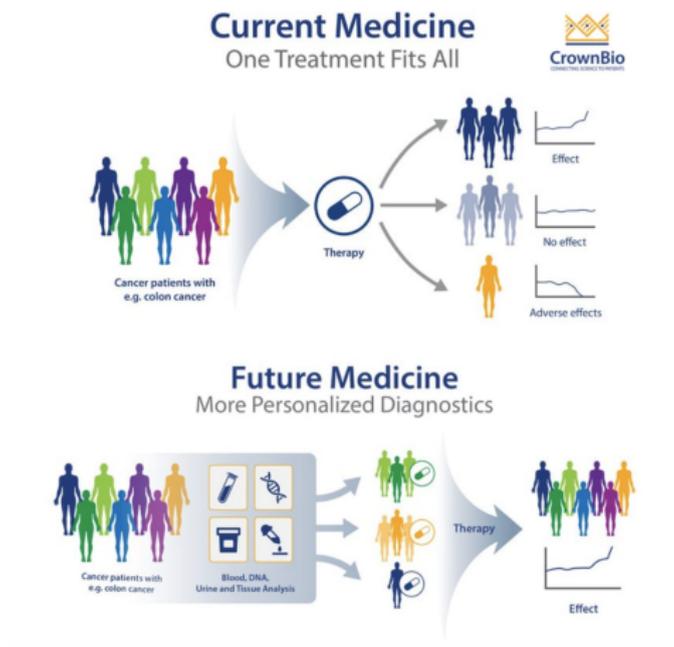


Figure 1: Transition from “one size fits all” to personalized medicine.¹

¹<https://blog.crownbio.com/pdx-personalized-medicine>



- Individualized decision making problems:
 - Making **personalized decision** based on **individualized information**
 - **Goal**: find the **best decision** that optimizes a specified **criterion**
- Focus on precision medicine:
 - *Individualized cancer treatment*: **tailoring therapies** based on **patients' genomic biomarkers** to **optimize** future **health status**
- Data $(Z, A, Y) \in \mathcal{Z} \times \mathcal{A} \times \mathbb{R}$
 - ① **Features** $Z \in \mathcal{Z} \subseteq \mathbb{R}^p$
 - ② **Assigned treatment** $A \in \mathcal{A} = \{1, 2, \dots, M\}$
 - ③ **Reward** $Y \in \mathbb{R}$ (larger the better)
- Individualized Treatment Rule (ITR) $D: \mathcal{Z} \rightarrow \mathcal{A}$
- Goal: Learn optimal ITR $D^* \in \mathcal{D}$ that maximizes the value function $\mathcal{V}(D)$

$$D^* \in \arg \max_{D \in \mathcal{D}} \left\{ \mathcal{V}(D) = \mathbb{E}[Y | A = D(X)] \right\}$$



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- 1 *Many treatments* available but *limited* observations for some specific treatments:
 - Large treatment space:
 - In *Patient-Derived Xenograft*: $|\mathcal{A}| > 20$ [Rashid et al., 2020]
 - Unbalanced structure of treatment assignment:
 - In *Sequenced Treatment Alternatives to Relieve Depression (STAR*D)*: number of patients who received the cognitive therapy v.s. venlafaxine is only around 1:3 [Rush et al., 2004]
 - In *Type 2 Diabetes*: observations of baseline treatments such as Metformin and Insulin would **dominate** others in electronic health record database [Montvida et al., 2018]
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- ② Treatments in large treatment space may work *similarly* for patients:
- *STAR*D* study: treatment options are combined into two class (one involves Selective Serotonin Reuptake Inhibitors (SSRI) + another one not) because treatments within same class have similar treatment effects [Liu et al., 2018, Pan and Zhao, 2021]
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- Supervised clustering

- Cluster the relationship $Y \sim Z \times A$ with *fusion penalty*:

$$\min_{\zeta} \left\{ \underbrace{\mathbb{E}_n \left[\mathcal{L} \left(Y, \sum_{a \in \mathcal{A}} \mathbb{I}[A = a] T(Z, \zeta_a) \right) \right]}_{\text{Loss}} + \underbrace{\sum_{1 \leq l < t \leq M} p \lambda_n (\|\zeta_l - \zeta_t\|_1)}_{\text{Fusion penalty}} \right\},$$

★ Heterogeneous treatment effects

where ζ_a 's are treatment-specific coefficients

- Convex minimization problem with *loss + fusion penalty* balanced by λ_n
- Interpretation:** maximize goodness of fit, while minimize heterogeneity among treatments simultaneously



- Clustering process can be visualized by a *dendrogram plot*:

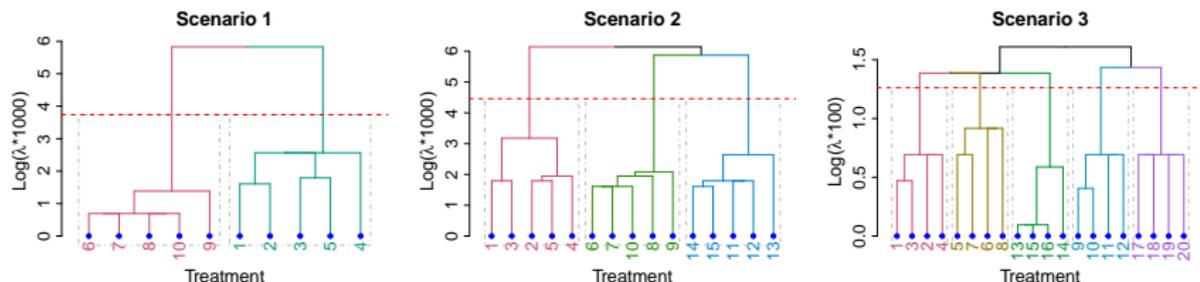


Figure 2: **Solution path** of estimated treatment group structure as λ_n increases. The true treatment group memberships are demonstrated with different colors. The red dotted horizontal lines show the best tuned λ_n using cross-validation.

- $\lambda_n = 0$: no penalty is imposed, hence no clustering pattern
- $\lambda_n \uparrow$: fusion penalty encourages similar treatments to merge together
- λ_n large enough: all treatments will be merged together



We also

- Solved fusion problem with adaptive proximal gradient algorithm effectively
- Proposed a novel *group-lasso* based method to select important variables
- Provided theoretical guarantee for estimating treatment group structure
- Conducted both simulation studies and real data analysis on cancer treatment to illustrate the superior performance of our method

😊 **Thanks for your listening!**

🌞 **Welcome to join our poster session if you have more questions.**



-  Liu, Y., Wang, Y., Kosorok, M. R., Zhao, Y., and Zeng, D. (2018).
Augmented outcome-weighted learning for estimating optimal dynamic treatment regimens.
Statistics in medicine, 37(26):3776–3788.
-  Montvida, O., Shaw, J., Atherton, J. J., Stringer, F., and Paul, S. K. (2018).
Long-term trends in antidiabetes drug usage in the us: real-world evidence in patients newly diagnosed with type 2 diabetes.
Diabetes care, 41(1):69–78.
-  Pan, Y. and Zhao, Y.-Q. (2021).
Improved doubly robust estimation in learning optimal individualized treatment rules.
Journal of the American Statistical Association, 116(533):283–294.
-  Rashid, N. U., Lockett, D. J., Chen, J., Lawson, M. T., Wang, L., Zhang, Y., Laber, E. B., Liu, Y., Yeh, J. J., Zeng, D., et al. (2020).
High-dimensional precision medicine from patient-derived xenografts.
Journal of the American Statistical Association, pages 1–15.





Rush, A. J., Fava, M., Wisniewski, S. R., Lavori, P. W., Trivedi, M. H., Sackeim, H. A., Thase, M. E., Nierenberg, A. A., Quitkin, F. M., Kashner, T. M., et al. (2004).

Sequenced treatment alternatives to relieve depression (star* d): rationale and design.

Controlled clinical trials, 25(1):119–142.

