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松井茂之の主な研究業績

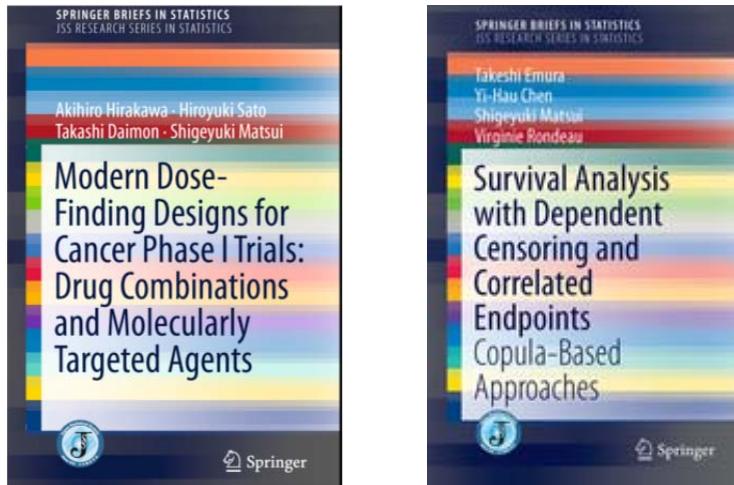
Book Editor:

1. Matsui S, Buyse M, Simon R. (editors). Design and Analysis of Clinical Trials for Predictive Medicine. CRC Press, 2015.
2. Matsui S, Crowley J (editors). Frontiers of Biostatistical Methods and Applications in Clinical Oncology, Springer, 2017.
3. 丹後俊郎, 松井茂之 (編). 新版医学統計学ハンドブック. 朝倉書店, 2018.



Book Author:

1. Hirakawa A, Sato H, Daimon T, Matsui S. Modern Dose-Finding Designs for Cancer Phase I Trials: Drug Combination and Molecularly Targeted Agents. Springer, 2018
2. Emura T, Chen YH, Matsui S, Rondeau V. Survival Analysis with Dependent Censoring and Correlated Endpoints. Copula-Based Approaches, Springer, 2018.



Book chapter:

1. Matsui S, Igeta M, Toyoizumi K. Biomarker-based phase II and III clinical trials in oncology. In Textbook of Clinical Trials in Oncology. (eds. S. Halabi and S. Michiels), CRC Press. (To appear).
2. Matsui S. (2017). Phase III clinical trial designs incorporating predictive biomarkers: an overview. In Frontiers of Biostatistical Methods and Applications in Clinical Oncology. (eds. S. Matsui, J. Crowley), Springer.
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9. Matsui S, Nonaka T, Choai Y. (2014). Design of phase III clinical trials with predictive biomarkers for personalized medicine. In *Developments in Statistical Evaluation of Clinical Trials* (eds K. van Montfort, J. Oud, W. Ghidey), Springer.
10. Hirakawa A, Matsui S. (2014). Dose-finding for two-agent combination phase I trials. In *Developments in Statistical Evaluation of Clinical Trials* (eds K. van Montfort, J. Oud, W. Ghidey), Springer.
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14. 松井茂之 (2010). 再発事象データの解析. 丹後俊郎, 小西貞則(編) 医学統計学の事典. 朝倉書店.
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論文業績

生物統計の方法論

1. Emura T, Matsui S, Chen HY. (2018). compound.Cox: univariate feature selection and compound covariate for predicting survival. *Computer Methods and Programs in Biomedicine* (In press).
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和文総説, コメント(計量生物学)

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