## SIMULTANEOUSLY OPTIMIZING DOSE AND SCHEDULE OF A NEW CYTOTOXIC AGENT

<u>T. Braun</u><sup>†1</sup>, P. Thall<sup>2</sup>, H. Nguyen<sup>2</sup>, M. de Lima<sup>2</sup>

<sup>1</sup> University of Michigan, Ann Arbor, MI, USA;

† E-mail: tombraun@umich.edu

Recently, a new type of phase I design was proposed that determines a maximum tolerated schedule (MTS), rather than a conventional maximum tolerated dose (MTD), in which the MTS is defined as the maximum number of courses of a cytotoxic agent that can be given without causing unacceptable cumulative toxicity (Braun et al., *Biometrics*, 2005, 61: 335-343). We expand this previous work by providing methodology to simultaneously optimize both dose and schedule in a multi-course phase I setting.

Specifically, we evaluate J doses,  $j=1,2,\ldots J$ , and K nested schedules, each of which is a sequence of administration times. The hazard of toxicity for a single administration of dose j is parameterized as a triangle having base of length  $b_j+c_j$  and area equal to  $a_j$ , with the height of the triangle,  $2a_j/(b_j+c_j)$ , occurring at  $u=b_j$ . The total cumulative hazard for any enrolled subject is simply the sum of the areas of the triangular hazards for each administration received before an interim analysis.

Prior to a new subject enrolling, we combine the interim information of enrolled subjects in a standard censored data likelihood and use Bayesian methods to adaptively update the 3J hazard parameter estimates, and thereby, the MTS. We examine our approach via simulations motivated by an actual clinical trial and show our algorithm has excellent operating characteristics in a trial of N=60 patients seeking to find the best combination of J=3 doses and K=4 schedules.

<sup>&</sup>lt;sup>2</sup>M.D. Anderson Cancer Center, Houston, TX