

A. Additional information on TCP calculation

GTV was considered to be the histopathology proven and defined cancer volume in the prostate. TCP was calculated according to the Poisson distribution as [17-21]:

$$TCP = P_B = P_{GTV}(\{D\}, V) = \prod_{i=1}^M e^{-\rho_{cell} \Delta V_i P(D_i)}$$

With $P(D_i)$ the survival probability for the dose D_i [18,19]:

$$P(D_i) = e^{-\left(\frac{EQD2_i}{D50}\right)(e\gamma - \ln \ln 2)} \quad \text{or equivalently} \quad P(D_i) = e^{-a EQD2_i \left(1 + \frac{2Gy}{\alpha/\beta}\right)}$$

M was the total number of voxels within GTV or of bins of the differential dose volume histogram for total dose in the GTV. ρ_{cell} represented the cancer cell density in GTV and ΔV_i was the voxel volume or the total volume corresponding to the i^{th} dose bin. D50 was the equieffective dose for 2Gy per fraction resulting to 50% tumor control probability and γ was the maximum normalized dose-response gradient [19,26]. EQD2_i was the equieffective dose for 2Gy per fraction for the total dose D_i for the i^{th} -voxel or i^{th} -dose bin [9,21]:

$$EQD2_i = D_i \left(1 + \frac{D_i/N}{\alpha/\beta}\right) / \left(1 + \frac{2Gy}{\alpha/\beta}\right)$$

With N the total number of fractions for delivering the total dose D_i . α/β values described the curvature of the cell survival curve described by the linear-quadratic model, where α described the slope of the initial linear part of the curve.

B. Additional information on NTCP calculation

NTCP from non-uniform dose distributions were calculated using the relative seriality model [19,26-28]:

$$P_1 = 1 - \prod_{j=1}^{N_{organs}} (1 - P_1^j) = 1 - \prod_{j=1}^{N_{organs}} \left(1 - \left[1 - \prod_{i=1}^{M_j} (1 - P^j(D_i)^{s_j})^{\Delta v_i} \right]^{1/s_j} \right)$$

With $P(D_i)$ the complication probability rate for the dose D_i and was given by the same formula as for TCP, where here D50 is the equieffective dose for 2Gy per fraction resulting to 50% complication probability. Δv_i was the volume fraction being irradiated to dose D_i and s was the parameter which expresses the degree of seriality (the value varies from s close to zero for nearly parallel organs and upwards for increasing seriality).

The maximum normalized dose-response gradient γ in the survival probability function was be calculated from the parameter k of the corresponding logit probability distribution according to the following:

$$k = 4 \gamma_{50} \quad \text{and} \quad \gamma_{50} = \frac{\ln 2}{2}(e\gamma - \ln \ln 2), \quad \text{thus} \quad \gamma = \frac{\frac{k}{2\ln 2} + \ln \ln 2}{e}$$

with γ_{50} the slope of the dose-response curve at the 50% response level.