

| Disease site | Case | Delineation summaries | | | | | | | | | | | | | | | | | |
|--------------|------|-----------------------|----|-----------------------|-----------|-----------|------------|------|------|------|------|-------------------|------|-----|------|-----------|----|----|----|
| | | N _{obs} | | Volume [ml] (mean±SD) | | | | | | | | CI _{gen} | | | | RMSE [mm] | | | |
| | | | | GTV | | CTV | | GTV | | CTV | | PT | | MT | | PT | | MT | |
| | | PT | MT | PT | MT | PT | MT | PT | MT | PT | MT | PT | MT | PT | MT | PT | MT | PT | MT |
| HNSCC | 1 | 5 | 5 | 88 ± 23 | 69 ± 28 | 186 ± 23 | 156 ± 31 | 0.59 | 0.46 | 0.69 | 0.63 | 2.9 | 3.1 | 2.6 | 2.8 | | | | |
| | 2 | 3 | 4 | 39 ± 14 | 36 ± 17 | 109 ± 40 | 104 ± 32 | 0.49 | 0.40 | 0.50 | 0.50 | 7.0 | 3.8 | 8.6 | 4.5 | | | | |
| | 3 | 5 | 5 | 5 ± 3 | 5 ± 3 | 25 ± 8 | 24 ± 8 | 0.34 | 0.29 | 0.44 | 0.41 | 3.4 | 3.6 | 3.3 | 3.4 | | | | |
| | 4 | 5 | 5 | 15 ± 3 | 13 ± 5 | 54 ± 16 | 49 ± 18 | 0.61 | 0.40 | 0.51 | 0.47 | 1.4 | 3.2 | 2.8 | 3.0 | | | | |
| | 5 | 5 | 4 | 71 ± 19 | 51 ± 27 | 178 ± 48 | 133 ± 50 | 0.55 | 0.33 | 0.61 | 0.47 | 2.5 | 4.7 | 3.3 | 4.4 | | | | |
| (N)SCLC | 1 | 6 | 6 | 97 ± 70 | 66 ± 32 | 207 ± 155 | 179 ± 105 | 0.33 | 0.41 | 0.35 | 0.38 | 5.5 | 10.4 | 7.0 | 10.5 | | | | |
| | 2 | 6 | 6 | 296 ± 27 | 190 ± 29 | 469 ± 50 | 354 ± 76 | 0.78 | 0.66 | 0.77 | 0.64 | 2.1 | 3.0 | 2.7 | 4.2 | | | | |
| | 3 | 4 | 5 | 161 ± 39 | 90 ± 47 | 336 ± 119 | 226 ± 98 | 0.55 | 0.36 | 0.50 | 0.44 | 5.9 | 4.3 | 6.6 | 6.4 | | | | |
| | 4 | 6 | 6 | 643 ± 59 | 680 ± 209 | 942 ± 92 | 1035 ± 256 | 0.77 | 0.55 | 0.79 | 0.58 | 3.0 | 5.6 | 3.2 | 6.4 | | | | |
| | 5 | 6 | 6 | 120 ± 15 | 130 ± 79 | 243 ± 52 | 235 ± 122 | 0.73 | 0.30 | 0.69 | 0.35 | 2.0 | 14.0 | 2.9 | 14.0 | | | | |

Abbreviations/Symbols: (P/M)T, (pre/mid)-treatment; N_{obs}, Number of observers providing delineations; SD, standard deviation;

CI_{gen}, generalised conformity index; RMSE, root-mean-square error.