

Diagnostic Performance of Regional Systematic Biopsy for Prostate Cancer

Stratified by PI-RADS and Histologic Zones

ELECTRONIC SUPPLEMENTARY MATERIAL

Supplementary Table S1. Diagnostic performances of different prostate sampling schemes in TZ

| PI-RADS | | n | PCa | p^* | Kappa* | csPCa | p^* | Kappa* |
|---------|--------|-----|------------|---------|--------|------------|---------|--------|
| 3 | TB+SB | 167 | 61 (36.5) | - | - | 38 (22.8) | - | - |
| | TB | 167 | 46 (27.5) | < 0.001 | 0.796 | 33 (19.8) | 0.063 | 0.911 |
| | SB | 167 | 52 (31.1) | 0.004 | 0.880 | 34 (20.4) | 0.125 | 0.929 |
| | TB+RSB | 167 | 58 (34.7) | 0.25 | 0.961 | 38 (22.8) | 1 | 1 |
| 4 | TB+SB | 141 | 112 (79.4) | - | - | 95 (67.4) | - | - |
| | TB | 141 | 99 (70.2) | < 0.001 | 0.758 | 87 (61.7) | 0.008 | 0.876 |
| | SB | 141 | 95 (67.4) | < 0.001 | 0.697 | 80 (56.7) | < 0.001 | 0.777 |
| | TB+RSB | 141 | 110 (78.0) | 1 | 0.979 | 95 (67.4) | 1 | 1 |
| 5 | TB+SB | 419 | 404 (96.4) | - | - | 393 (93.8) | - | - |
| | TB | 419 | 397 (94.7) | 0.016 | 0.802 | 388 (92.6) | 0.063 | 0.906 |
| | SB | 419 | 393 (93.8) | 0.001 | 0.719 | 384 (91.6) | 0.004 | 0.841 |
| | TB+RSB | 419 | 404 (96.4) | 1 | 1 | 393 (93.8) | 1 | 1 |

*Compared with TB+SB

TB = targeted biopsy; SB = systematic biopsy; RSB = regional systematic biopsy;

mpMRI = multiparametric magnetic resonance imaging; PI-RADS = Prostate Imaging

Reporting and Data System; PCa = prostate cancer; csPCa = clinically significant prostate cancer; TZ = transitional zone

Supplementary Table S2. Diagnostic performances of different prostate sampling schemes compared with TB+RSB in PZ

| PI-RADS | | n | PCa | <i>p</i> * | Kappa* | csPCa | <i>p</i> * | Kappa* |
|---------|--------|-----|------------|------------|--------|------------|------------|--------|
| | TB+RSB | 158 | 63 (39.9) | - | - | 51 (32.3) | - | - |
| 3 | TB | 158 | 51 (32.3) | < 0.001 | 0.836 | 43 (27.2) | 0.008 | 0.879 |
| | SB | 158 | 61 (38.6) | 0.625 | 0.947 | 49 (31.0) | 0.625 | 0.941 |
| | TB+RSB | 325 | 269 (82.8) | - | - | 243 (74.8) | - | - |
| 4 | TB | 325 | 234 (72.0) | < 0.001 | 0.697 | 218 (67.0) | < 0.001 | 0.815 |
| | SB | 325 | 255 (78.5) | < 0.001 | 0.823 | 228 (70.1) | < 0.001 | 0.885 |
| | TB+RSB | 505 | 486 (96.2) | - | - | 483 (95.6) | - | - |
| 5 | TB | 505 | 482 (95.4) | 0.125 | 0.901 | 479 (94.9) | 0.125 | 0.913 |
| | SB | 505 | 484 (95.8) | 0.5 | 0.948 | 481 (95.2) | 0.5 | 0.954 |

*Compared with TB+RSB

TB = targeted biopsy; SB = systematic biopsy; RSB = regional systematic biopsy; PI-RADS = Prostate Imaging Reporting and Data System; PCa = prostate cancer; csPCa = clinically significant prostate cancer; PZ = peripheral zone

Supplementary Table S3. Diagnostic performances of different prostate sampling schemes compared with TB+RSB in TZ

| PI-RADS | | n | PCa | p^* | Kappa* | csPCa | p^* | Kappa* |
|---------|--------|-----|------------|---------|--------|------------|---------|--------|
| | TB+RSB | 167 | 58 (34.7) | - | - | 38 (22.8) | - | - |
| 3 | TB | 167 | 46 (27.5) | < 0.001 | 0.833 | 33 (19.8) | 0.063 | 0.911 |
| | SB | 167 | 52 (31.1) | 0.146 | 0.838 | 34 (20.4) | 0.125 | 0.929 |
| | TB+RSB | 141 | 110 (78.0) | - | - | 95 (67.4) | - | - |
| 4 | TB | 141 | 99 (70.2) | < 0.001 | 0.778 | 87 (61.7) | 0.008 | 0.876 |
| | SB | 141 | 95 (67.4) | < 0.001 | 0.681 | 80 (56.7) | < 0.001 | 0.777 |
| | TB+RSB | 419 | 404 (96.4) | - | - | 393 (93.8) | - | - |
| 5 | TB | 419 | 397 (94.7) | 0.016 | 0.802 | 388 (92.6) | 0.063 | 0.906 |
| | SB | 419 | 393 (93.8) | < 0.001 | 0.719 | 384 (91.6) | 0.004 | 0.841 |

*Compared with TB+RSB

TB = targeted biopsy; SB = systematic biopsy; RSB = regional systematic biopsy; PI-RADS = Prostate Imaging Reporting and Data System; PCa = prostate cancer; csPCa = clinically significant prostate cancer; TZ = transitional zone