

Medical Technology Assessment Committee (MTAC)
 Clinical Knowledge Development Support, Kaiser Permanente Washington

Evidence table

Authors, Design, Aim	Patient's characteristics	Outcomes	Conclusion, RoB (QUADAS-2)																																
Waterhouse et al., 2019 Design: Retrospective study Aim: To assess the performance of ConfirmMDx in African American (AA) men.	N=211 AA Patients were men undergoing repeat biopsy within 30 months from a negative index biopsy. Age: 43-86 yrs; PSA: 0.6-61.6 ng/mL; 60% had normal DRE. GS: 66% had GS of 6; 34% had GS ≥7 Clinical stage: 69% T1c On repeat bx results: 62% had no PCa, 38% had PCa (66% GS ≤6; 33% GS ≥7). Index test consisted of Multiple qMSP assay (MDxHealth). The reference standard was histopathology on repeat prostate bx.	<p>The clinical performance (detection of any PCa) included: Sensitivity: 74.1% (63.1%-83.1%) Specificity: 60.0% (51.1%-68.5%) PPV: 53.6% (47.4%-59.6%) NPV: 78.8% (71.5%-84.6%)</p> <p>Sensitivity & specificity in this study were comparable to studies with predominantly Caucasian population.</p> <p>Detection of high-grade PCa (GS ≥7): Sensitivity: 77.8% (57.7%-91.4%) Specificity: 52.7% (45.2%-60.1%) PPV: 19.4% (15.8%-23.7%) NPV: 94.2% (88.7%-97.1%)</p> <p>No significant differences in sensitivity or specificity were observed between age groups (<55; 55-69; ≥70)</p> <p>The high NPV demonstrates that repeat biopsy can be avoided. A test with high NPV may be clinically useful.</p> <p>2*2 tables: Detection of any cancer at biopsy</p> <table border="1" data-bbox="821 987 1472 1084"> <thead> <tr> <th></th> <th>PCa pos</th> <th>PCa neg</th> <th></th> </tr> </thead> <tbody> <tr> <td>CMDx pos</td> <td>60</td> <td>52</td> <td>112</td> </tr> <tr> <td>CMDx neg</td> <td>21</td> <td>78</td> <td>99</td> </tr> <tr> <td></td> <td>81</td> <td>130</td> <td></td> </tr> </tbody> </table> <p>Detection of high-grade cancer at biopsy</p> <table border="1" data-bbox="821 1143 1331 1317"> <thead> <tr> <th></th> <th>HG PCa pos</th> <th>HG PCa neg</th> <th></th> </tr> </thead> <tbody> <tr> <td>CMDx pos</td> <td>21</td> <td>87</td> <td>108</td> </tr> <tr> <td>CMDx neg</td> <td>6</td> <td>97</td> <td>103</td> </tr> <tr> <td></td> <td>27</td> <td>184</td> <td></td> </tr> </tbody> </table>		PCa pos	PCa neg		CMDx pos	60	52	112	CMDx neg	21	78	99		81	130			HG PCa pos	HG PCa neg		CMDx pos	21	87	108	CMDx neg	6	97	103		27	184		<p>Conclusion: the study validates the use of ConfirmMDx to help guide repeat biopsy decision in AA. ConfirmMDx has a good performance in this AA population. Its high NPV shows that it may be clinically useful for classification of AA men who had an initial negative biopsy.</p> <p>RoB: High Patient selection: low Index test: Unclear (Were the index test results interpreted without knowledge of the results of the reference standard? Unclear) Reference standard: Unclear (Were the reference standard results interpreted without knowledge of the results of the index test? Unclear) Flow & timing: Low</p>
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<p>Stewart et al., 2013</p> <p>Design: retrospective study</p> <p>(Methylation Analysis to Locate Occult Cancer (MATLOC study))</p> <p>Aim: to demonstrate high NPV and clinical utility (by averting unnecessary biopsy)</p>	<p>The authors included 483 patients with two consecutive biopsies where the first biopsy was negative, and the repeat biopsy was positive (cases) or negative (controls); Mean age 63 years (45-84); PSA mean 7.61 (range 0.4 to 50 ng/ml); Percentage of patients with normal DRE 73%;</p> <p>Most patients had GS=6</p> <p>Histopathology biopsy: benign 73%.</p> <p>Patients were classified in two groups: Group 1, cases, with positive repeat biopsy result (n=87), and group 2 which have negative repeat bx referenced as controls (n=396).</p> <p>The index test was the qMSP assay. The reference test was histopathology result from repeat prostate bx. PSA > 4 ng/ml was more frequent in controls than in cases.</p>	<table border="1" data-bbox="915 354 1713 513"> <thead> <tr> <th></th> <th>Sensitivity (CI)</th> <th>Specificity (CI)</th> <th>NPV (CI)</th> <th>PPV(CI)</th> <th>Accuracy</th> </tr> </thead> <tbody> <tr> <td>3-gene model</td> <td>68% (57%-77%)</td> <td>64% (59%-69%)</td> <td>90% (87%-93%)</td> <td>29% (25%-33%)</td> <td>65%</td> </tr> </tbody> </table> <p>DNA methylation was a significant independent predictor of cancer on repeat biopsy (OR 3.17, 95% CI 1.81–5.53, p <0.0001). Atypical cells but not HGPIN were also predictor of cancer (OR 3.17 (95% CI 1.31–7.70)).</p>		Sensitivity (CI)	Specificity (CI)	NPV (CI)	PPV(CI)	Accuracy	3-gene model	68% (57%-77%)	64% (59%-69%)	90% (87%-93%)	29% (25%-33%)	65%	<p>Conclusion: The Findings show that the test has moderate sensitivity & specificity with high NPV and can predict repeat biopsy outcome. The test could decrease unnecessary repeat biopsy by 64%. The high NPV indicates a low false negative. The DNA methylation is an independent risk factor for detecting prostate cancer before repeat biopsy. The increased NPV could help avert unnecessary repeat biopsies.</p> <p>Rob: High</p> <p>Patient selection: High (Case control was not avoided)</p> <p>Index test: unclear (Were the index test results interpreted without knowledge of the results of the reference standard? Unclear)</p> <p>Reference standard: high (Is the reference standard likely to correctly classify the target condition? High risk)</p> <p>Flow & timing: low</p> <p>Applicability concerns: Low for Patient selection, Index Test, & reference standard.</p>
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<p>Partin et al., 2014: Detection of Cancer Using Methylated Events in Negative Tissue (DOCUMENT trial)</p> <p>Design: retrospective study</p> <p>Aim: To validate the performance of an epigenetic test.</p>	<p>N=350 patients Pt characteristics (n=320), mean age: 62 yrs, PSA (mean ng/mL): 6.1, % of normal DRE: 60%, Histopathology bx 1: No abnormality: 58%.</p> <p>All patients received repeat biopsy within 2 years; those with negative results were controls (n=228) and patients with positive results were cases (n=92). Index test was the multiplex methylation (MDxHealth, Irvine, California). Inclusion criteria was defined as a minimum of 8 cores per biopsy. The reference test was histopathology result on repeat prostate bx.</p>	<p>Sensitivity 62% (51%-72%), Specificity 64% (57%-70%), NPV 88% (85%-91%)</p> <p>The test was found to be the most significant independent predictor of patient outcome (OR 2.69, 95% CI 1.60-4.51) on repeat biopsy after controlling for age, PSA, DRE, first biopsy histopathological characteristics and race.</p> <p>Atypical cells were also found to be a significant predictor of patient outcome (OR, 2.37; P=0.0465).</p> <p>In a subgroup of African Americans (small sample size): Sensitivity 77% (95% CI 46–95) Specificity 66% (95% CI 46–82) NPV 93% (85% CI 82–97)</p>	<p>The findings have shown that the epigenetic test is a significant and independent predictor of prostate cancer detection in a repeat biopsy collected within 24 months. Its NPV is high and as result the test can help avoid unnecessary repeat biopsies if it is added to other known risk factors.</p> <p>Rob: High Patient selection: high (case control was not avoided) Index test: Low Reference standard: High (Is the reference standard likely to correctly classify the target condition? High risk) Flow & timing: Low Applicability concerns: Low for Patient selection, Index Test, & reference standard.</p>
<p>Van Neste et al., 2016</p> <p>Design: retrospective study</p> <p>Aim: To assess the performance of a DNA-methylation assay, to predict men at risk of harboring high-grade cancer.</p>	<p>Cohorts from the MATLOC & DOCUMENT trials, appraised above, were combined (n=803 patients). Therefore, patients' characteristics are similar to those studies. All men had a negative index biopsy followed by either a positive (179 men) or negative (624 men) repeat biopsy. The index test & reference standard are identical to that of MATLOC & DOCUMENT trials. Patients were classified according to the histopathological outcome of the repeat biopsy: High risk (GS ≥7, n=67), indolent PCa (GS ≤6, n=106), Control (no PCa, n=)</p>	<p>Overall NPV (for finding low levels of DNA methylation) 89.2% for all cancers. PPV 28.2% for all cancers. None of these cancers were detected at the time of index biopsy. The assay had a significantly increased PPV (P < 0.001) compared to current clinical practice.</p> <p>Regarding high-grade PCa detection (GS ≥7), NPV was 95.9%.</p> <p>DNA methylation was the strongest predictor of high-grade cancer (AUC 0.661 (0.572–0.751) p=0.001) upon repeat biopsy.</p> <p>Risk score (DNA-methylation intensity and traditional clinical risk factors) AUC: 0.762 (0.679 – 0.844)</p>	<p>The risk score may help to detect patients with high-grade cancer in comparison to PSA and PCPTRC.</p> <p>The risk of bias of the study is high. This is due to the exclusion of patients with GS 6, long duration between biopsies (>2 years), small sample size in high-grade cancer, conflict of interest in the form of financial interest between some authors and the laboratory.</p>

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	No significant differences in PSA, DRE, time between biopsies were reported. Histopathology significantly differ between groups. HGPIN predominated in the groups Age was significantly different in the groups (controls, GS≤6, GS≥7): 62.5/62.0, 63.3/64.0, 65.6/66.0 p<0.001.	Decision curve analysis: In comparison with PSA and PCPTRC (Prostate Cancer Prevention Trial Risk Calculator), there is a higher benefit that high-grade PCa was reported on repeat bx for risk score.	
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References

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