

**Application setting for CREATININE ENZYMATIC, Diagnosticum Zrt, ADVIA 1650, 7/2008, Trinec**

<p>Analytical conditions</p> <p>R1 volume: 72</p> <p>R2 volume: 0</p> <p>R3 volume: 24</p> <p>R4 volume: 0</p> <p>R1 diluent volume: 0</p> <p>R2 diluent volume: 0</p> <p>R3 diluent volume: 0</p> <p>R4 diluent volume: 0</p> <p>Serum reac.svol: 16</p> <p>Serum dil. method: Stand</p> <p>Serum dil.s.vol: 30</p> <p>Serum dil. volume: 120</p> <p>Serum dil. Posit: 0</p> <p>Urine set</p> <p>Reaction time: 10 min</p> <p>Reagent 1 stir: Weak</p> <p>Reagent 2 stir: Weak</p> <p>Reagent 3 stir: Weak</p> <p>Reagent 4 stir: Weak</p>	<p>Sub Param# 39: 1 Up Down</p> <p>Sub-anly.conditions</p> <p>Name: ORE Digits: 2</p> <p>M-wave. L: 545</p> <p>S-wave. L: EPA</p> <p>Analy.mthd: EPA</p> <p>Calc.mthd: STD</p> <p>Qualit.judg: Not do Qualit.set</p> <p>Rea-time correct. form.</p> <p>Reanalysis conditions</p> <p>Serum reac. smp. vol (u): 16</p> <p>Serum dilut.method (u): Special</p> <p>Serum dil. smp. vol (u): 10</p> <p>Serum diluent vol (u): 90</p> <p>Serum diluent posi (u): 0</p> <p>Serum reac. smp. vol (d): 3</p> <p>Serum dilut.method (d): None</p> <p>Serum dil. smp. vol (d): 0</p> <p>Serum diluent vol (d): 0</p> <p>Serum diluent posi (d): 0</p> <p>Urine set</p> <p>Rerun.cond. H</p>	<p>Standards setting</p> <p>BLK H: 9.99999 Multi-STD</p> <p>BLK L: -9.9999</p> <p>STD H: 9.99999</p> <p>STD L: -9.9999</p> <p>FV: 0.00000</p> <p>Abnm(serum)H: 1770</p> <p>Abnm(serum)L: -99999</p> <p>Abnm(urine)H: 999999</p> <p>Abnm(urine)L: -99999</p> <p>Normal value set</p> <p>Calculation method setting</p> <p>M-DET.P.i: 0 S-DET.P.p: 44</p> <p>M-DET.P.m: 97 S-DET.P.r: 45</p> <p>M-DET.P.n: 96</p> <p>Check D.P.I: 0</p> <p>Limit value: 0.003</p> <p>Variance: 10</p> <p>* Reaction rate method</p> <p>Cycle: 3</p> <p>Factor: 3</p> <p>React.typ: Dec.</p> <p>E2 corre: Not do</p> <p>Blank (u): 9.9999</p> <p>Blank (d): -9.999</p> <p>Sample (u): 9.9999</p> <p>Sample (d): -9.999</p> <p>* Endpoint method</p> <p>Reabsorb (u): 9.9999</p> <p>Reabsorb (d): -9.999</p> <p>Prozone form: None</p> <p>Prozone limit: 9.999</p> <p>Prozone judge: Upper limit</p> <p>Judge limit: 9.999</p> <p>M-DET.P.m: 0 S-DET.P.p: 0</p> <p>M-DET.P.n: 0 S-DET.P.r: 0</p> <p>IMA setting</p>
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Calib. Curve info.	<p>calc. Mthd: 1-point</p> <p>Axis conv: No conv.</p> <p>Formula: Linear</p> <p>Points: 2</p>
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<p>Item Analysis Conditions (Urine)</p> <p>Urine reac. S. vol: 16</p> <p>Urine dil. Method: Special</p> <p>Urine dil. Smp. Vol: 2</p> <p>Urine diluent vol: 148</p> <p>Urine diluent Posi: 0</p>	<p>Multi-Standards setting</p> <p>Formula: Logit Log 3</p> <p>Axis conv: No convert</p> <p>Points: 6</p> <table border="1"> <thead> <tr> <th></th> <th>FV</th> <th>Dil. Meth</th> <th>Dil. Smp. vo</th> <th>Diluent vol</th> <th>Diluent posi</th> <th>STD-L</th> <th>STD-L</th> </tr> </thead> <tbody> <tr><td>BLK</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></tr> <tr><td>1</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></tr> <tr><td>2</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></tr> <tr><td>3</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></tr> <tr><td>4</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></tr> <tr><td>5</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></tr> </tbody> </table>		FV	Dil. Meth	Dil. Smp. vo	Diluent vol	Diluent posi	STD-L	STD-L	BLK								1								2								3								4								5							
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<p>Reanalysis conditions setting</p> <p>Variance (*)</p> <p>Absorbance (U)</p> <p>Absorbance (D)</p> <p>Absorbance limit (u)</p> <p>Absorbance limit (d)</p> <p>Cell blank (N)</p> <p>Abnormal v.limit (H)</p> <p>Abnormal v.limit (L)</p> <p>Normal val. Limit (h)</p> <p>Normal val. Limit (l)</p> <p>Reagent shortage ®</p> <p>Overflow (/)</p> <p>Safety (s)</p> <p>Prozone (P)</p> <p>Effect. Nbr.o.pnts (n)</p> <p>Calibration c</p> <p>Reanalysis ®</p>	<p>Reanalysis Conditions (Urine)</p> <p>Urine reac. Smp. Vol (u): 3.00</p> <p>Urine dilut.method (u): None</p> <p>Urine dil. Smp. Vol (u): 0</p> <p>Urine diluent posi (u): 0</p> <p>Urine reac. Smp. Vol (d): 3.00</p> <p>Urine dilut. Method (d): None</p> <p>Urine dil. Smp. Vol. (d):</p> <p>Urine diluent vol (d):</p> <p>Urine diluent posi (d):</p> <p>Notes:</p> <p>* Data entered by the user</p> <p>█ Dilution of the method</p>
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