

## Tissue Extraction Troubleshooting Guide

Problem	Solution
Poorly homogenized tissue	<ul style="list-style-type: none"> <li>Repeat homogenization</li> </ul>
Low or no DNA yield	<ul style="list-style-type: none"> <li>Keep the sample in the magnetic separation device when removing supernatant during the binding or washing steps.</li> <li>Remove the supernatant without disturbing the pellet of beads by angling the pipette tip away from the pellet.</li> <li>Ensure Binding Buffer, Wash Buffer 1 and Wash Buffer 2 have had Isopropanol and Ethanol added correctly.</li> </ul>
Remains of MAGneat beads in elution	<ul style="list-style-type: none"> <li>Leave on the magnetic stand until elution buffer is clear.</li> <li>Centrifuge eluate for 2-5 minutes at maximum speed. Remove supernatant free of beads.</li> </ul>
Poor 260/230 Ratio	<ul style="list-style-type: none"> <li>Centrifuge eluate for 2-5 minutes at maximum speed. Remove supernatant and dispense into new tube ready to measure.</li> </ul>
If MAGneat beads are being lost throughout the extraction	<ul style="list-style-type: none"> <li>Increase magnetizing time throughout the protocol.</li> <li>Remove the supernatant without disturbing the pellet of beads by angling the pipette tip away from the pellet.</li> </ul>
Issues with downstream applications due to ethanol carryover	<ul style="list-style-type: none"> <li>Increase air drying time to ensure MAGneat beads have dried before adding elution.</li> </ul>