

### **ABRAXIS® Affinity Capture & Extraction (ACE) Kit -Microcystins PN 520100**

### **Prepare Samples**

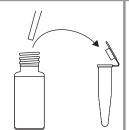
## **Urine Samples**

## **Blood Serum Samples**

#### **\*Centrifuge Samples**

Thaw frozen sample immediately before

testing.
Add 1.4 mL of thawed sample to a Protein LoBind tube. Centrifuge samples at 2,000 x g for 5 minutes to pellet any precipitates.



#### **\*Centrifuge Samples**

Thaw frozen sample immediately before testing. Add 1.4 mL of thawed sample to a Protein LoBind tube. Centrifuge samples at 10,000 x g for 5 minutes to separate any precipitates or flocculants.



#### 2. Transfer Samples

Transfer the sample supernatant to a new Protein LoBind tube and discard tube with pellet.

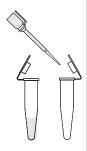


#### 2. Transfer & Dilute Samples

Transfer 0.5 mL of the sample supernatant (clear liquid portion) to a new Protein LoBind tube and discard tube with précipitate.

Add 1.0 mL of Seri-Standard Sample Diluent/Zero Standard, vortex well (1:3 dilution).

NOTE: Beads must be able to move through serum easily and be visible against the magnet. If diluted serum is still too thick to easily pipet, additional Seri-Standard Sample Diluent/Zero Standard may be added. Account for any additional dilutions in final calculation.



#### 3. Run Affinity Capture & Extraction (ACE) Kit

The samples are now ready to be run along with the standards in the ACE Kit.

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**Please Note:** Prior to running the Affinity Capture & Extraction (ACE) Kit, please prepare the standards, using the flowchart provided in the Standard Set and samples using the flowchart on the backside of this flowchart.

#### 1. Re-suspend Beads

Re-suspend beads with gentle shaking or rotating.



#### 8. Separate Beads from Solution

Pulse spin tubes in microcentrifuge then place tubes in magnetic separator for 2-3 minutes. Discard the supernatant.



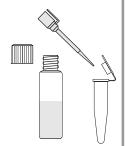
#### 2. Addition of Beads

Add 25  $\mu$ L of the bead solution for each standard and sample to a Protein LoBind tube. Pool according to user quide instructions.



#### 9. Wash Beads

Add 500 μL of the Sample Wash Buffer to the bead solution and vortex.



#### 3. Separate Beads from Solution

Place tubes into magnetic separator for 1 minute then discard supernatant



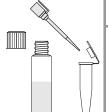
#### **10. Separate Beads from Solution**

Pulse spin tubes in microcentrifuge then place tubes in magnetic separator for 2-3 minutes. Discard the supernatant.



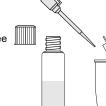
#### **4. Wash Beads**

Add 75  $\mu$ L of the Bead Wash Buffer to each standard and sample and vortex.



## 11. Add Elution Buffer/Sample Wash Ruffer

Add either elution buffer or sample wash buffer to the tubes and vortex gently. (For which, please see user guide)



Place tubes into a mixer or thermomixer at 1400 rpm for 5 minutes at the correct temperature.

#### **5. Separate Beads from Solution**

Place tubes into magnetic separator for 1 minute then discard supernatant



#### 12. Separate Beads from Solution

Pulse spin tubes in microcentrifuge tube then place tubes in magnetic separator for 2-3 minutes.



#### **6. Re-suspend Beads**

Add 25  $\mu L$  of the Bead Wash Buffer to each standard and sample and vortex.

Separate pool into 25 µL per Protein LoBind Tube.



#### **13. Transfer Elution Supernantant**

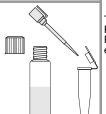
Transfer the elution supernatant to new Protein LoBind tube and discard beads. Please see user guide for additional postelution modifications.



## 7. Addition of Standards and Samples

Add 1 mL of the standards and samples (previously prepared) to the bead solution and vortex.

Place tubes into mixer at 1400 rpm for 5 minutes at room temperature



#### **14. Run Down Stream Assay**

Use the elution supernatant to run the downstream assay.

To run the downstream assay, please refer to the user guide/flow chart provided with the assay. DISCARD the standards that are provided within the downstream assay and replace them with the extracted standards from either the ABRAXIS® Uri-Standards Set Microcystins or the ABRAXIS® Seri-Standards Set Microcystins.

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