



# Webinar: Pushing the limits of sensitivity for ultra-low and single-cell NGS RNA applications



Takara's **Switching Mechanism at 5' End of RNA Template (SMART)** technology delivers industry leading sensitivity for your challenging and low-input samples. SMART-Seq provides full-length transcript information; enabling in-depth analysis of transcript isoforms, gene fusions, and point mutations in your NGS libraries.

Please join us to see how SMART-Seq® ultra-low and single cell technologies can help you get more information out of every sequencing experiment.

**Speaker: Christina Ulrich, PhD**  
Territory Manager II, Takara Bio USA, Inc

This webinar is hosted by the  
**CPRIT UHealth Cancer  
Genomics Center (CGC)**

DATE: November 3, 2022  
TIME: 12:00 -1:00 PM

**Have Questions?**  
Contact your Territory Manager,  
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Please register for this webinar with the CGC today :  
<https://forms.gle/67DmxdpgBx81VkvUz5>



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