

SM04755, a Potential Disease-Modifying Treatment for Tendinopathy, Modulates the Wnt Pathway via Inhibition of CLKs and DYRK1A

V. Deshmukh, PhD, A. L. O'Green, MS, M. Ibanez, MS, T. Seo, MS, and Y. Yazici, MD
 Samumed, LLC, San Diego, CA

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Background

- Tendinopathy is associated with pain, inflammation, tendon degeneration, and failed healing. Despite the high prevalence of tendinopathy, its underlying pathogenesis is not fully understood¹
- Wnt signaling plays an important role in tendinopathy² by modulating inflammation, tenocyte lineage specification, protease production, and tendon homeostasis^{3,4}
- SM04755, a novel, topical, small-molecule Wnt pathway inhibitor, has previously been shown to inhibit inflammation, protect tenocytes, and increase tenocyte differentiation in nonclinical models⁵
- The mechanism of action of SM04755 leading to Wnt pathway inhibition, tenocyte differentiation and protection, and anti-inflammatory activity is described

Results

Figure 1. SM04755 was a potent inhibitor of CDC-like kinases (CLKs) and dual-specificity tyrosine phosphorylation-regulated kinase 1A (DYRK1A). Inhibition of CLKs and DYRK1A inhibited the Wnt pathway in rTDSCs

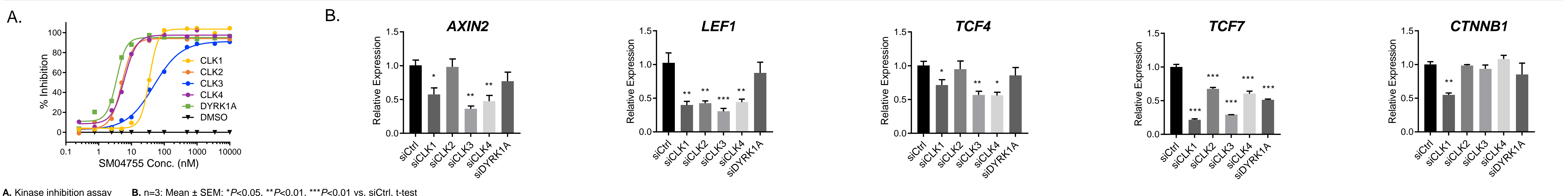


Figure 2. Inhibition of CLKs and DYRK1A induced tenocyte differentiation in vitro

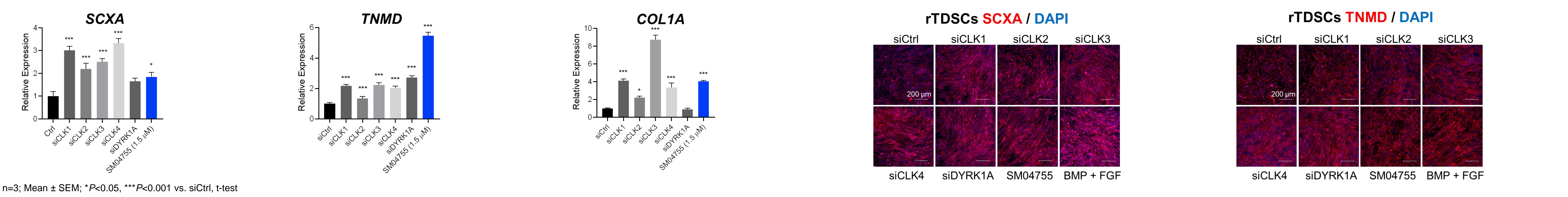


Figure 3. Inhibition of CLK1, 2, 4, and DYRK1A reduced catabolic protease expression in vitro

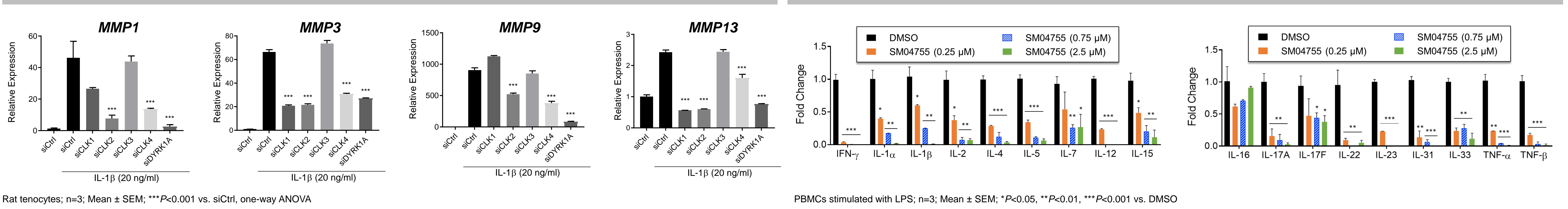
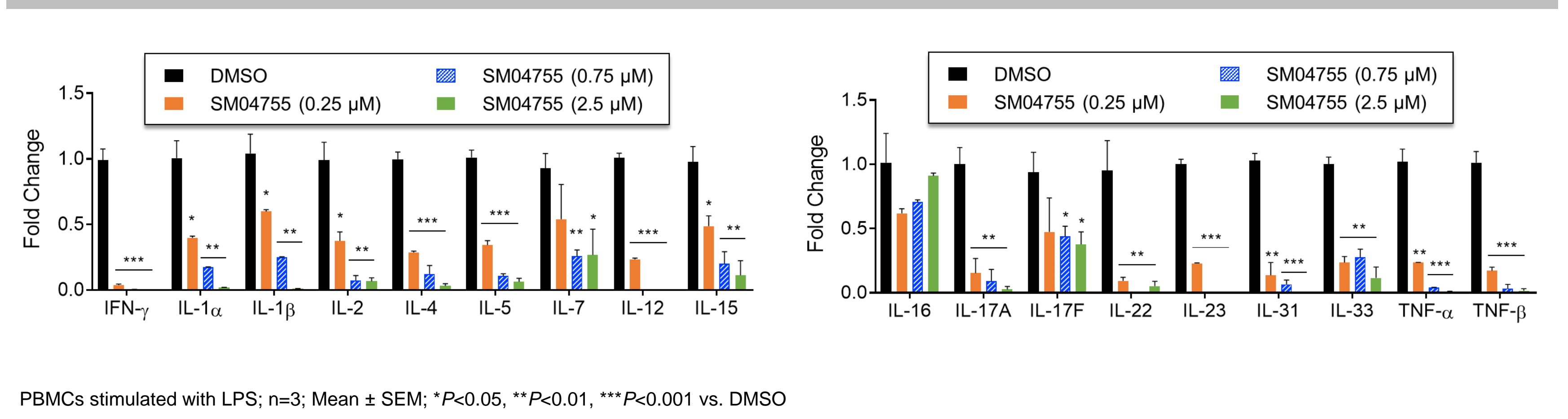


Figure 4. SM04755 demonstrated anti-inflammatory effects in vitro



Conclusions

- SM04755 inhibited intranuclear CLKs and DYRK1A, leading to Wnt pathway inhibition
- CLK and DYRK1A inhibition induced tenocyte differentiation and reduced tendon-destroying proteases in tenocytes
- SM04755 inhibited inflammatory signaling mediators and cytokine production
- SM04755, as a single agent, may potentially benefit symptoms and provide disease modification in tendinopathy
- Human tendinopathy trials are planned

Methods

- A kinome screen (318 kinases) was performed. Kinase inhibition was assessed by Thermo Fisher Z'-LYTE™ and LanthaScreen kinase assays (Fig. 1A)
- SM04755 and siRNA knockdown effects on gene expression in rat tendon-derived stem cells (rTDSCs) (Fig. 1B) and rat tenocytes (Fig. 3) were measured by qRT-PCR using TaqMan® primers. Gene expression was normalized to GAPDH
- SM04755 and siRNA knockdown effects on tenocyte marker expression in rTDSCs were assessed by immunostaining (Fig. 2)
- SM04755 effects on cytokine production in PBMCs stimulated with LPS were measured by MSD-based ELISA (Fig. 4)

References

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