

ZF-TF mediated epigenetic repression of SCN9A gene as a therapeutic approach for painful peripheral neuropathies

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Neuropathic pain is one of the most difficult pain syndromes to manage



Given the high unmet need and lack of effective treatments, there is an urgent need to develop novel therapeutics for the treatment of chronic neuropathic pain



Mutations in the SCN9A gene (Nav1.7) are linked to inherited pain disorders



- NavI.7 is a voltage gated sodium channel expressed in the Dorsal Root Ganglion (DRG)
- Alterations in NavI.7 activity directly regulate pain levels in several genetic disorders, validating NavI.7 as a therapeutic target for pain
- Lowering Nav1.7 is expected to reduce pain without adversely affecting other sensory functions
- High structural similarities among Nav channels has made it challenging to develop Nav1.7 selective inhibitors



Zinc Finger based epigenetic regulation



Require binding just two DNA targets per cell to regulate RNA and protein levels



Highly potent, specific and optimizable



Derived from naturally occurring human genes



No modifications or edits to the genome





Blocking pain transmission to the brain has the potential to treat multiple pain indications



- Blocking NavI.7 in the DRG is expected to prevent the transmission of nociceptive pain signals to the brain
- This allows us to target multiple neuropathic pain indications, regardless of the cause of the pain
- Reducing pain by inhibiting Nav1.7 is not predicted to be associated with any CNS adverse effects



Zinc finger-mediated repression of Nav1.7 as a potent and specific therapeutic avenue for neuropathic pain







Developmental path to identify mouse and human ZF-repressors targeting the Nav1.7 gene





The efficacy of ZF-Rs was evaluated in the Spared Nerve Injury (SNI) neuropathic pain model

- SNI is the most validated mouse neuropathic pain model ("Gold standard")
- Surgically induced hypersensitivity to pain





- Mechanical and cold induced pain were measured before (baseline) and 4 weeks after ZF-R treatment
- Scn9a repression in DRG was evaluated at the bulk and single-cell (nociceptor) level
- Gabapentin was used as a positive control and administered one hour before the pain measurement on day 28



SNI model

Mouse specific ZF-Rs induced up to 70% bulk repression of Scn9a in DRGs







Potent repression of Scn9a mRNA in nociceptors of mouse lumbar DRG





In vivo repression of mouse *Scn9A* reverses pain hypersensitivity in a mouse model of neuropathic pain





SNI model

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ZF-Rs repressed SCN9A by more than 90% over a wide dose range with no detectable off-target activity



- SCN9A was the only significantly regulated transcript out of >20,000 genes analyzed
- No repression of any other Nav channel was observed



- Genes down-regulated
- Genes up-regulated
- SCN9A



13

3E5

1E5

MOI

1E4

3E4

3E3

The potency and safety of the human ZF-R candidates were evaluated in nonhuman primates





ZF-Rs repressed the expression of SCN9A in a dose-dependent manner in multiple DRG levels





- Comparable data were obtained in sacral DRG
- Multiple DRGs were evaluated for each level per animal

ZF-Rs were well tolerated in nonhuman primates



These results support the initiation of IND-enabling GLP Toxicology study



Conclusion



- Mouse ZF-Rs potently repressed Scn9a at the bulk and single-cell level in mouse DRG
- IT-L administered ZF-Rs reversed pain hypersensitivity in the SNI model of neuropathic pain



- Human ZF-Rs potently repressed SCN9A >90% in human iPSC-derived neurons
- ZF-Rs were highly specific with no off-target activity detected, including no repression of any other Nav channels



- Human ZF-Rs repressed SCN9A by up to 40-60% at all DRG levels in NHP
- ZF-Rs were well tolerated at all doses tested with no adverse findings related to treatment
- These results support the continued progression to IND-enabling nonhuman primate study





Thank you

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Sangamo Therapeutics Inc.