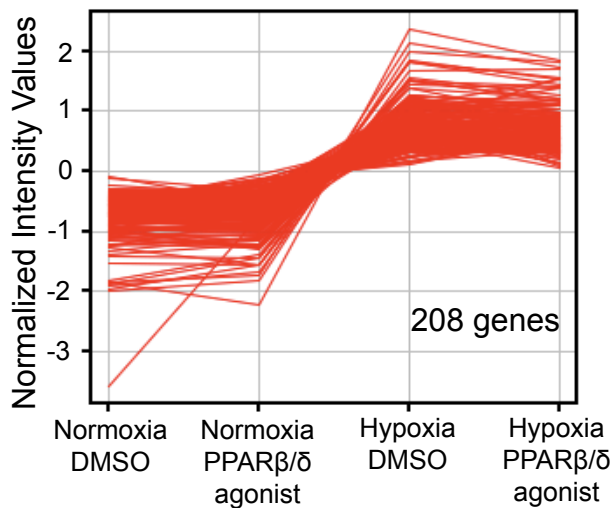
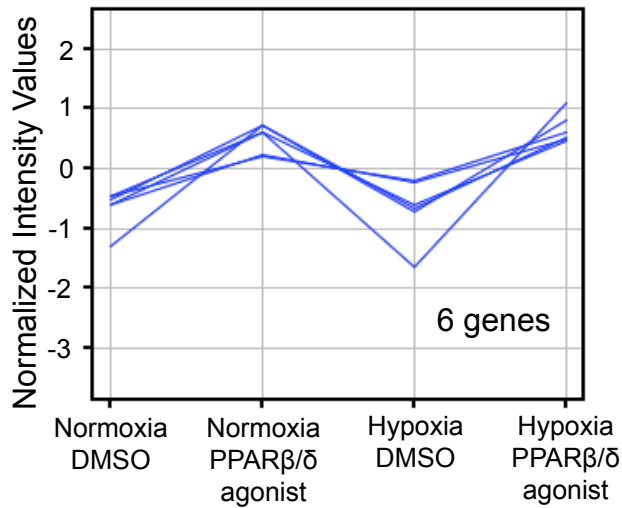


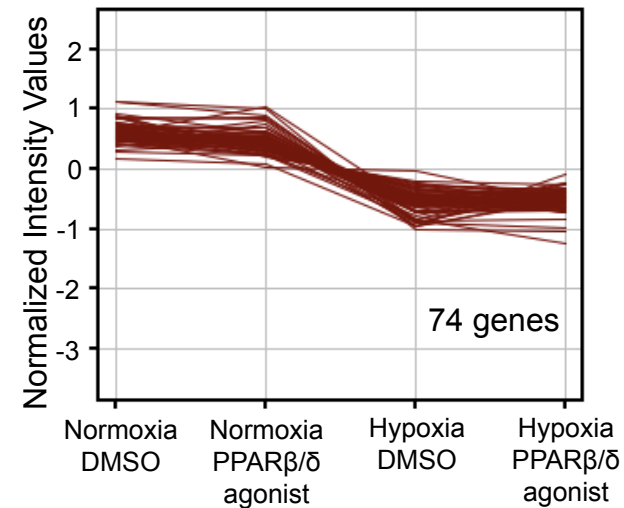
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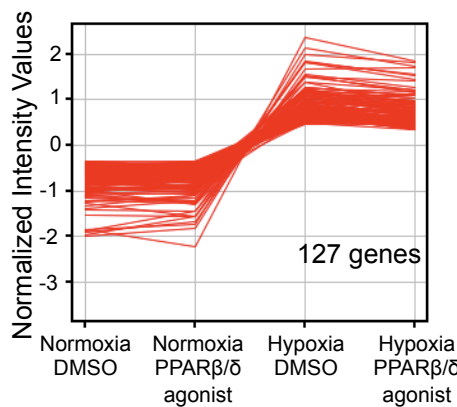
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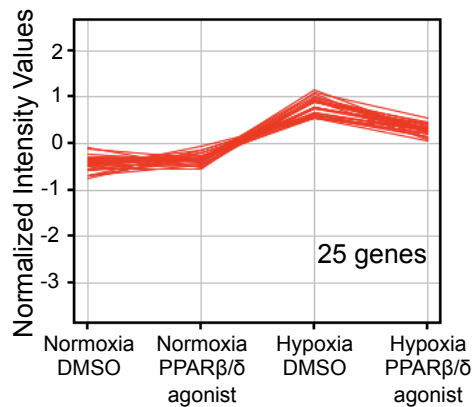
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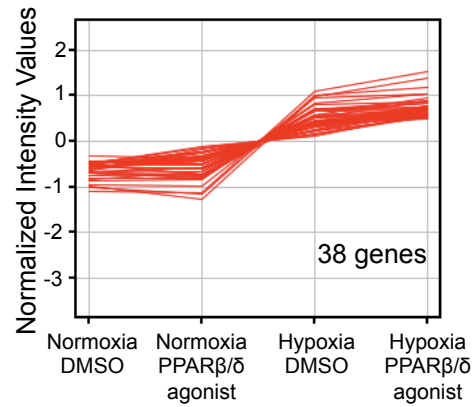
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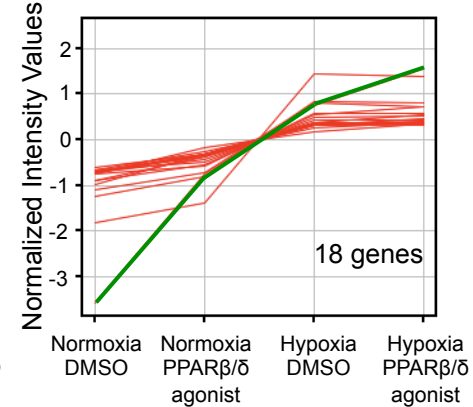
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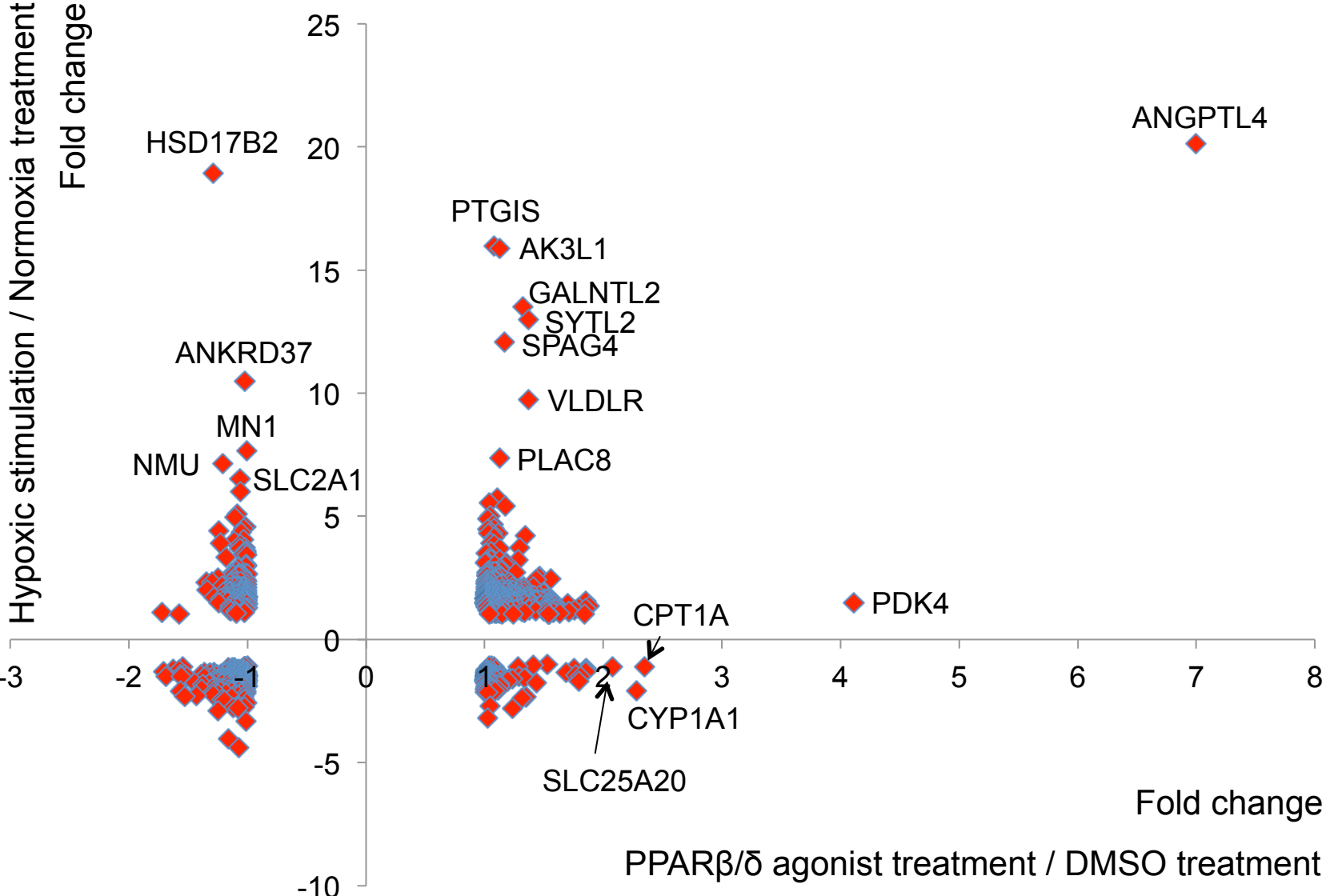


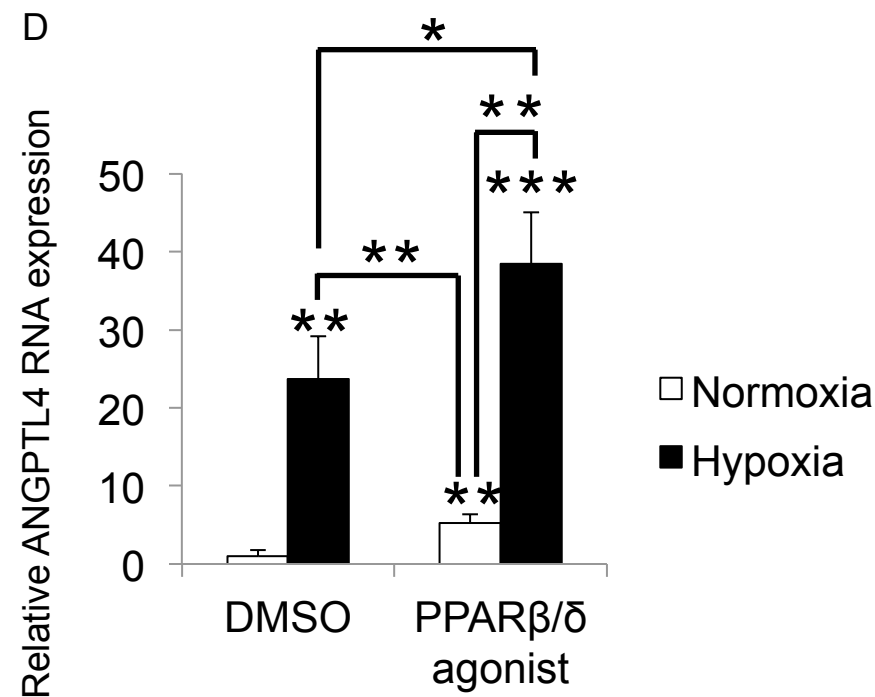
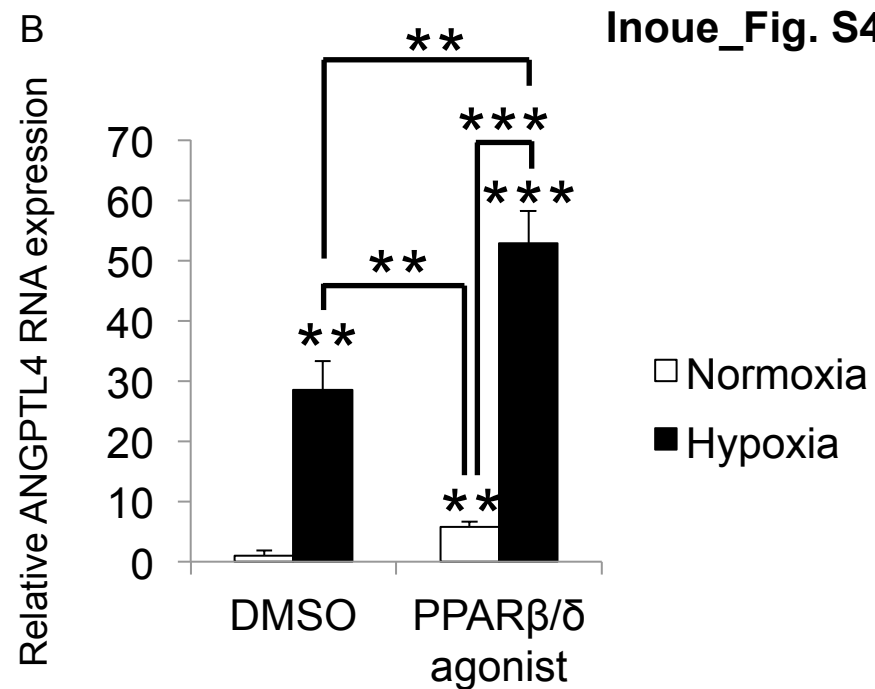
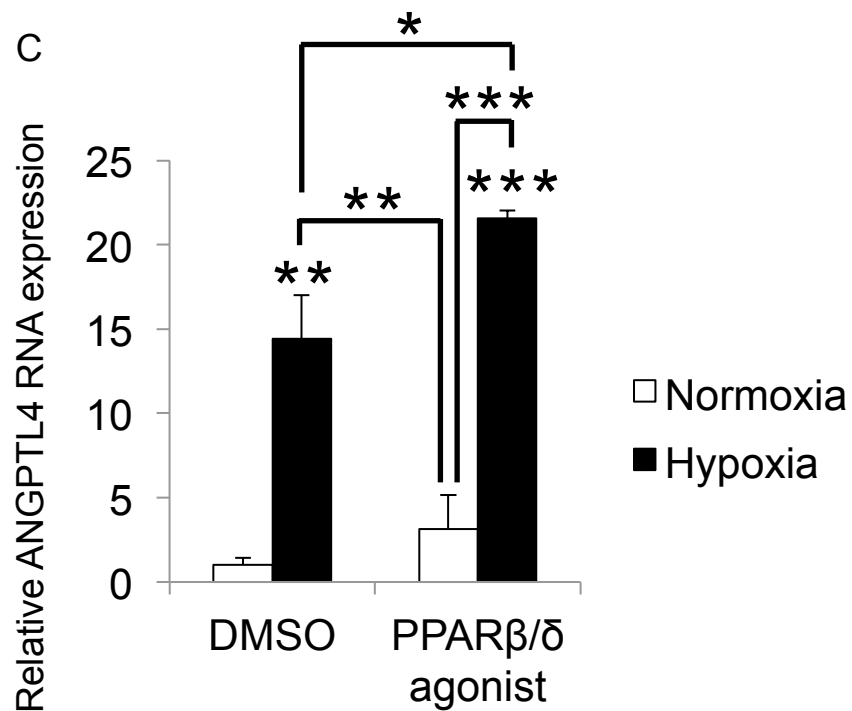
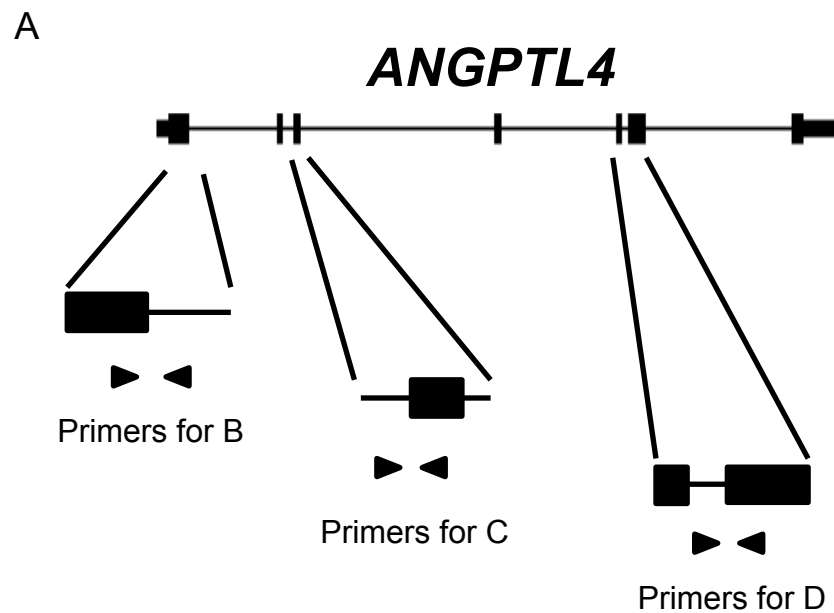
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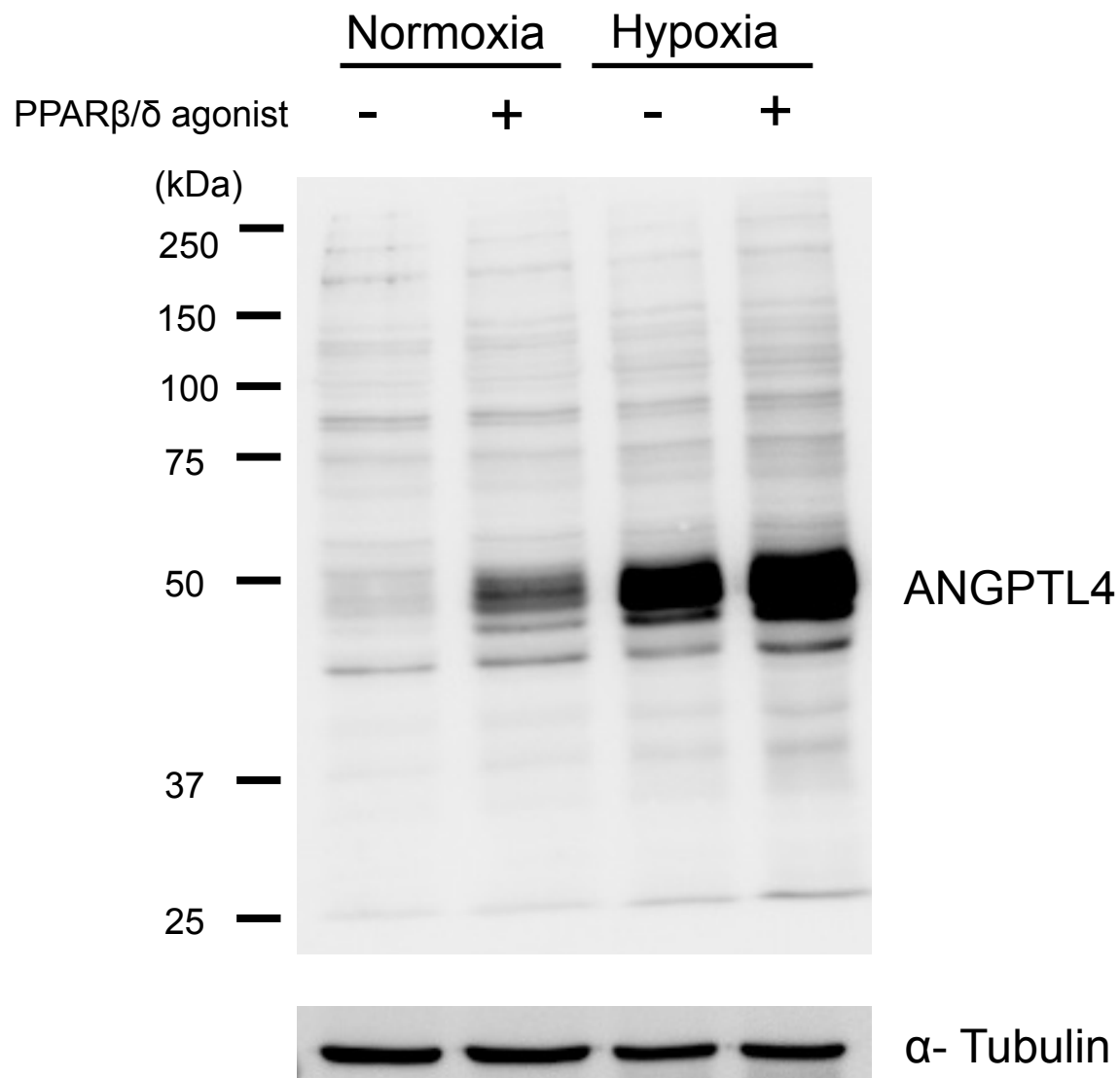


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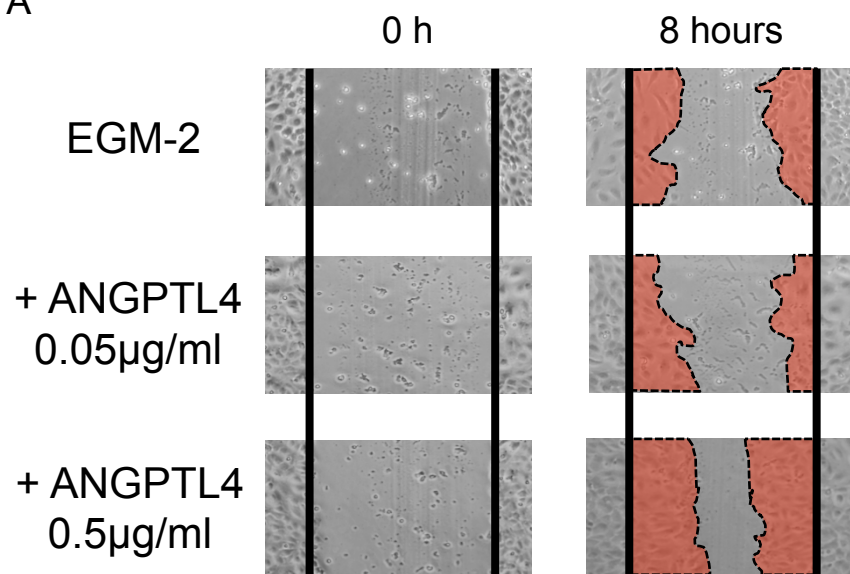




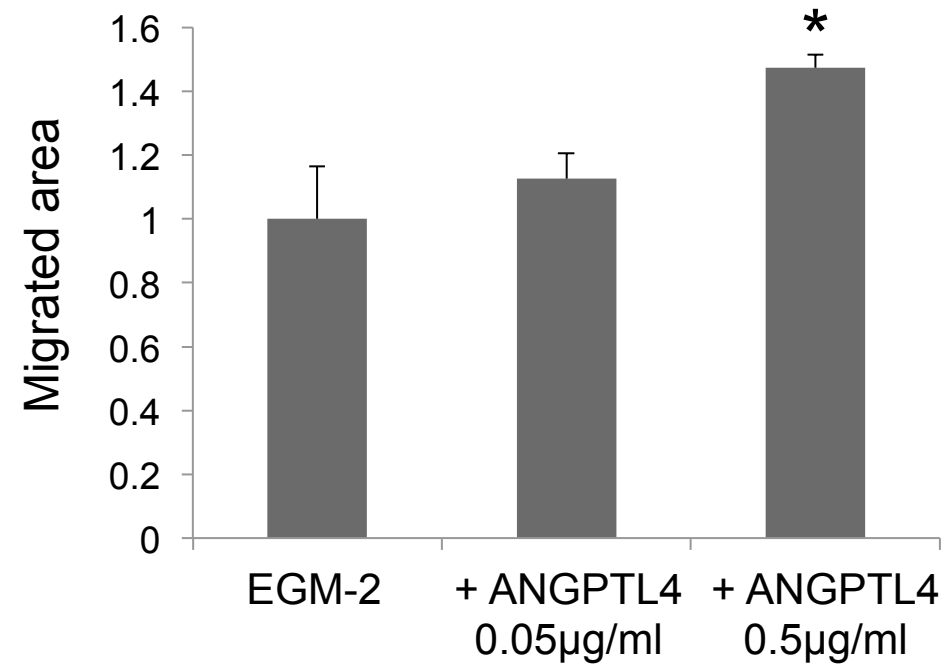




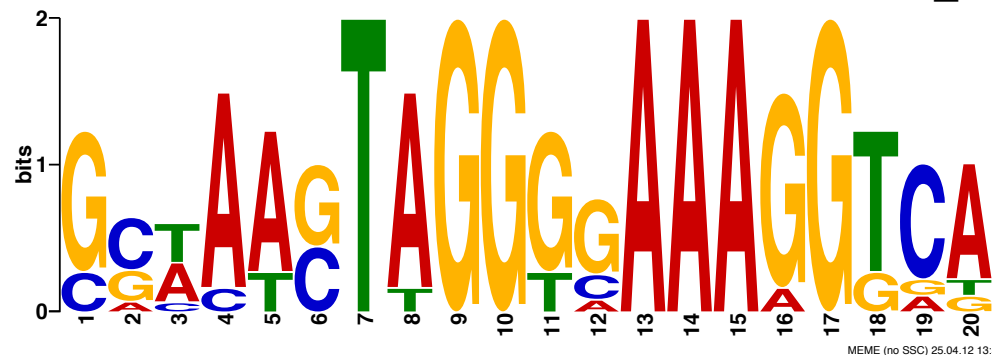
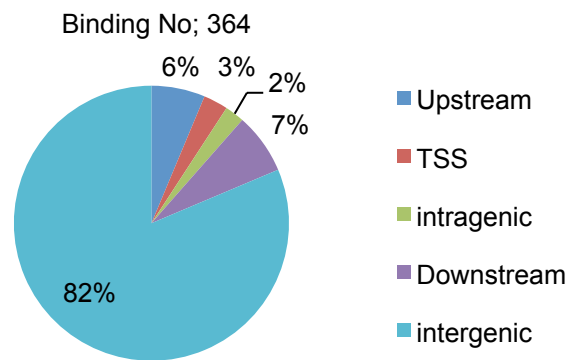
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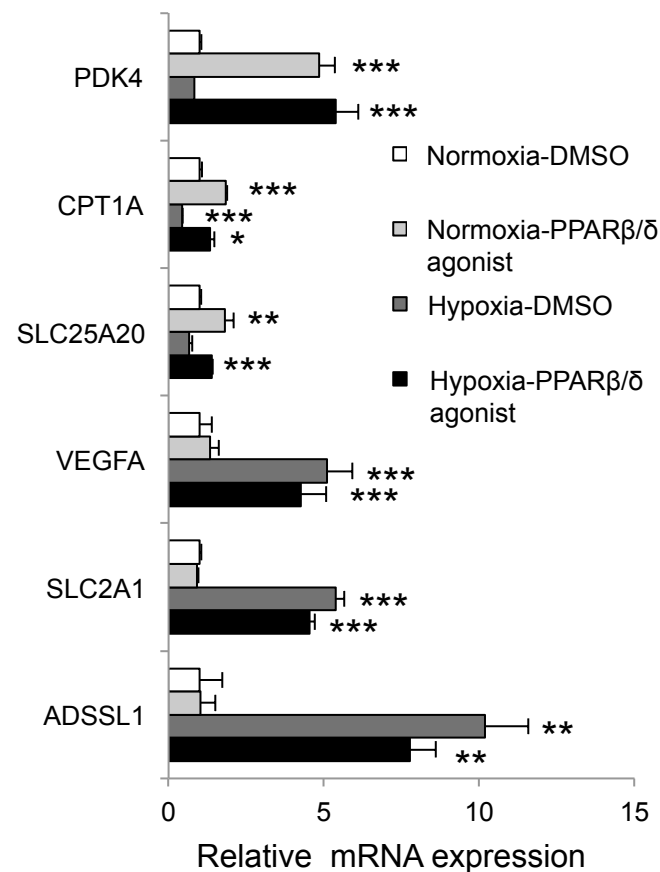
B



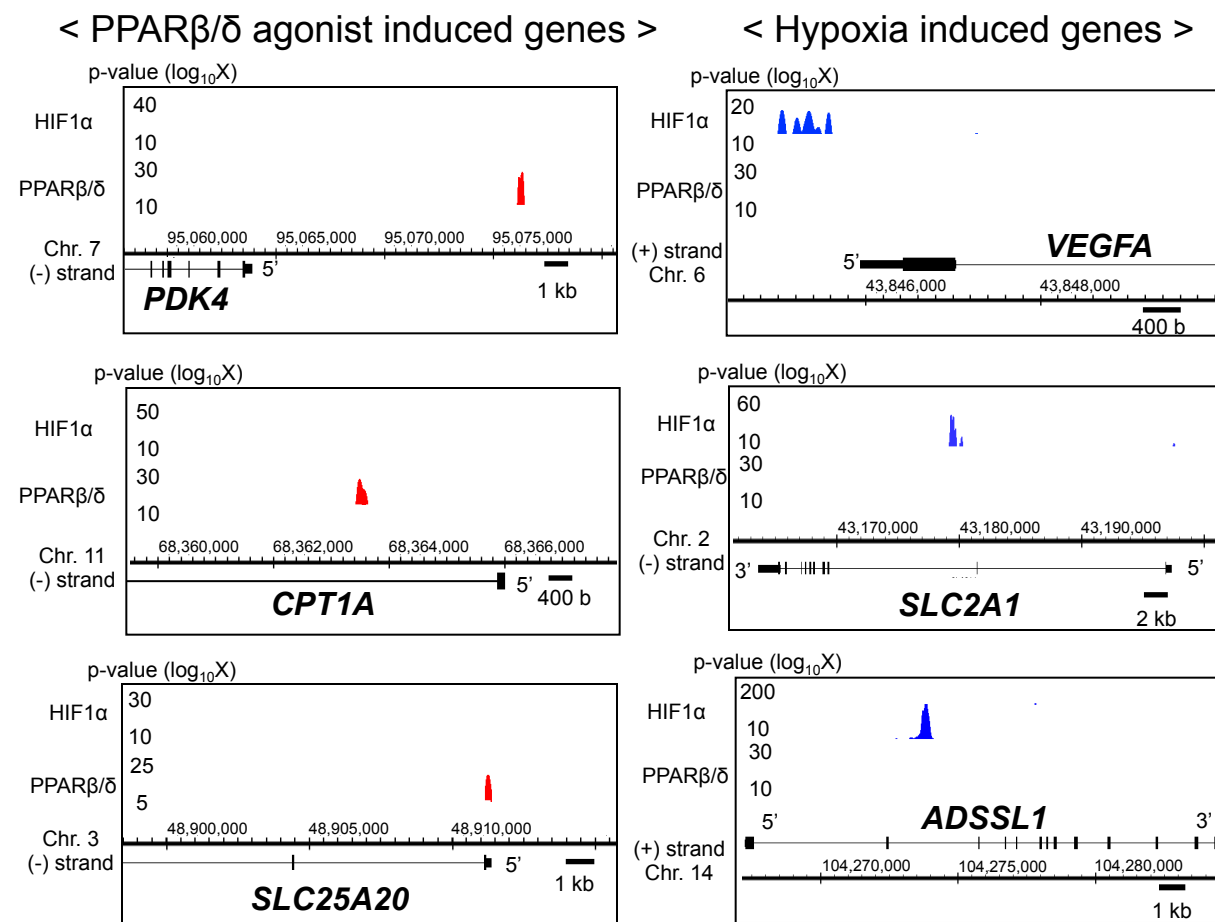
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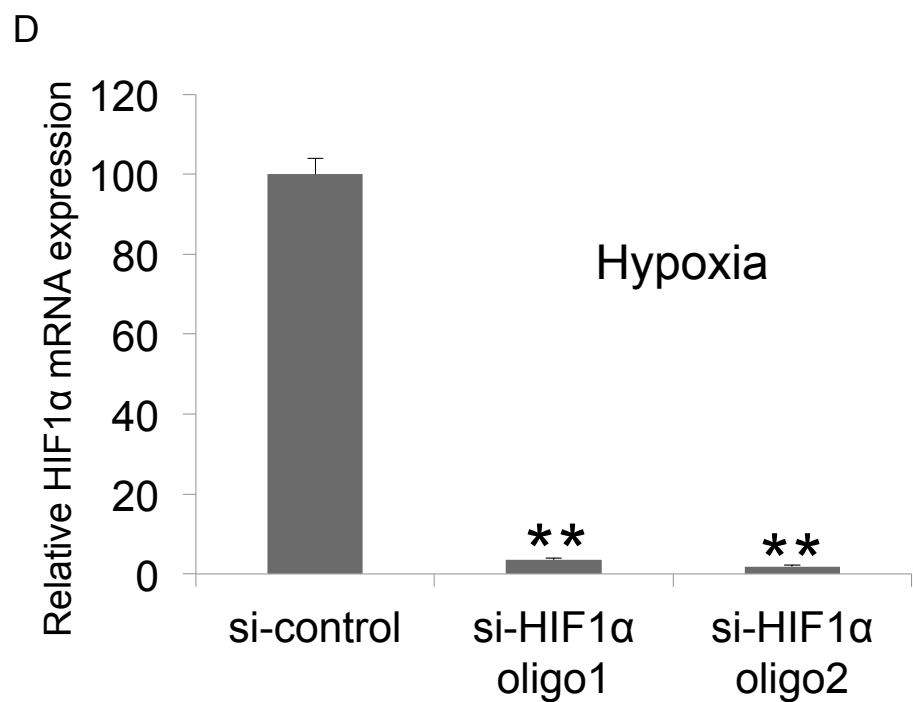
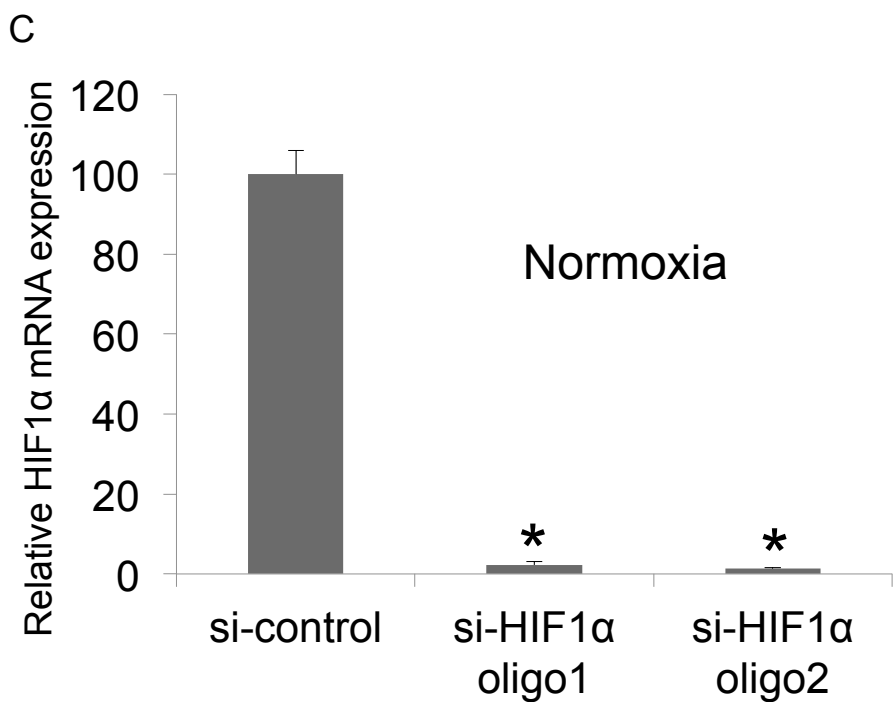
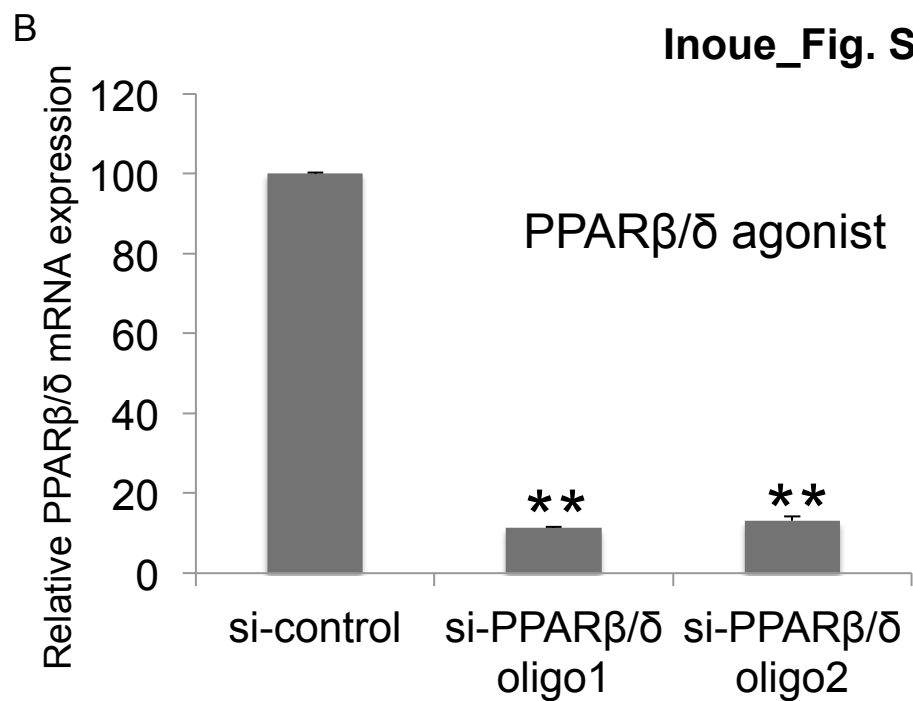
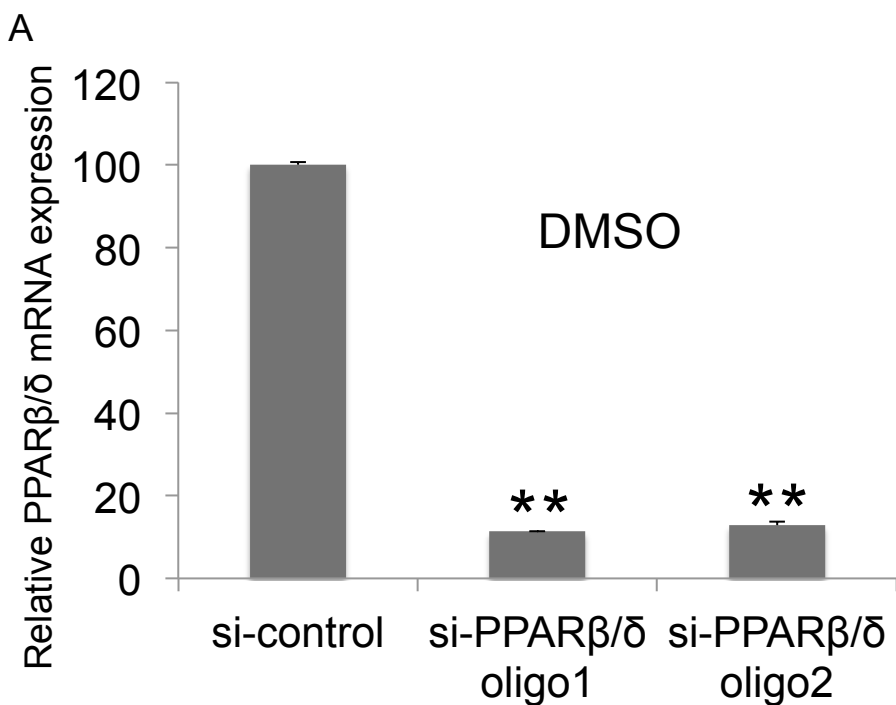


B

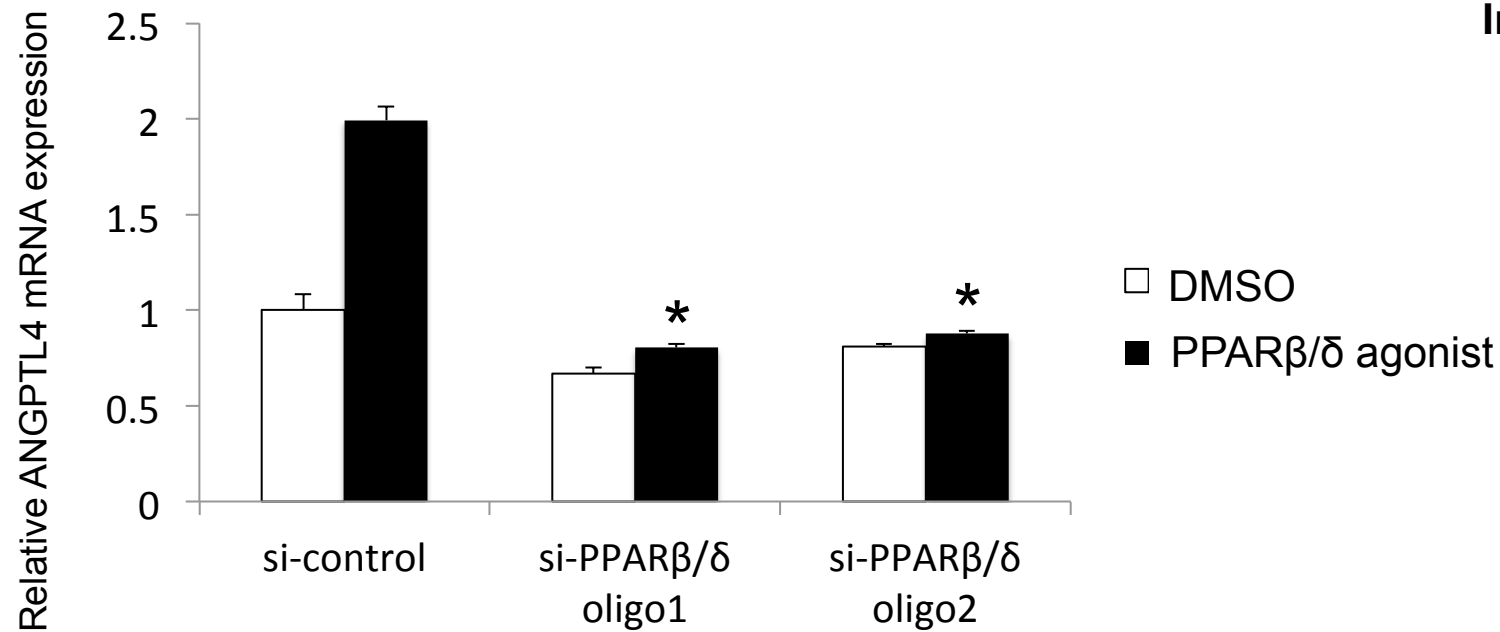


C

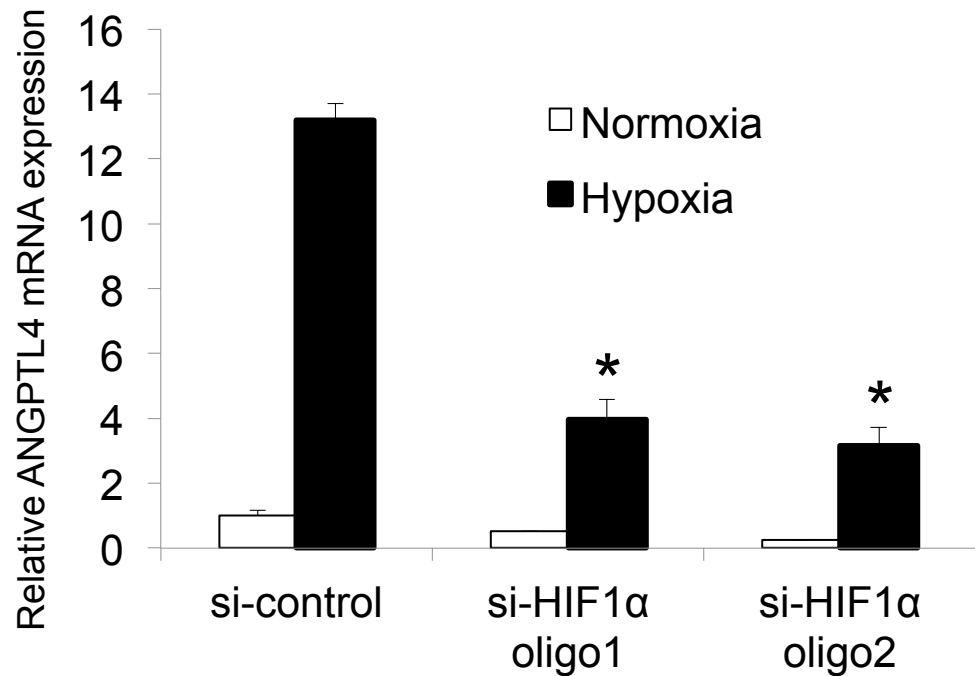




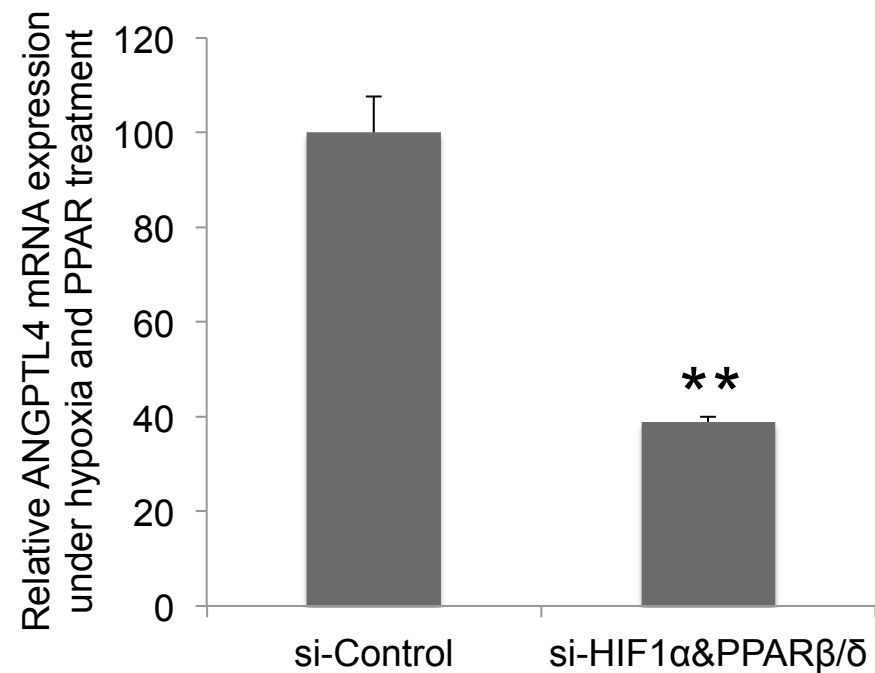
A



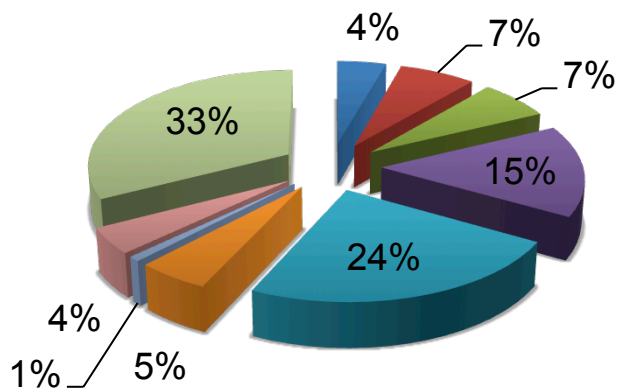
B



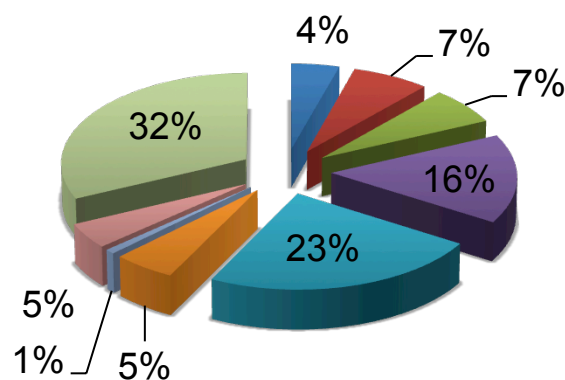
C



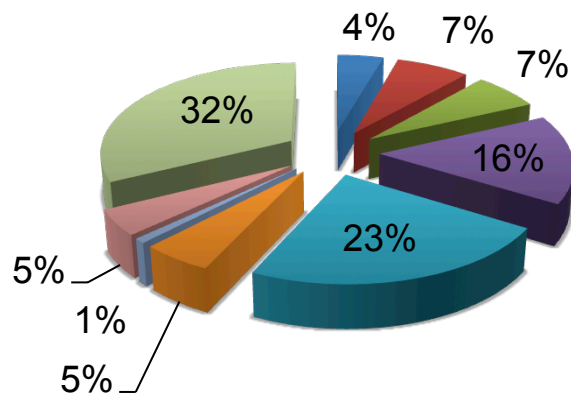
Normoxia + DMSO
Binding No: 33,221



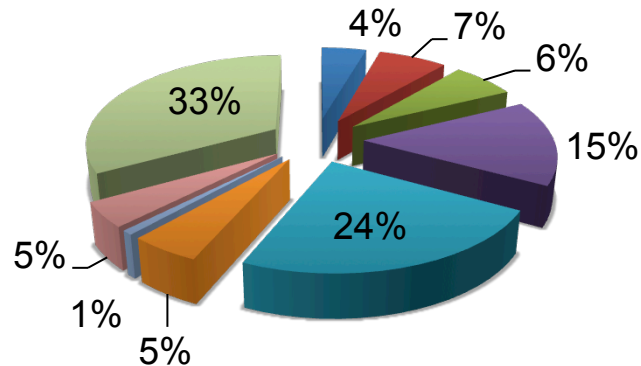
Normoxia + PPAR β/δ agonist
Binding No; 37,730



Hypoxia + DMSO
Binding No; 37,491

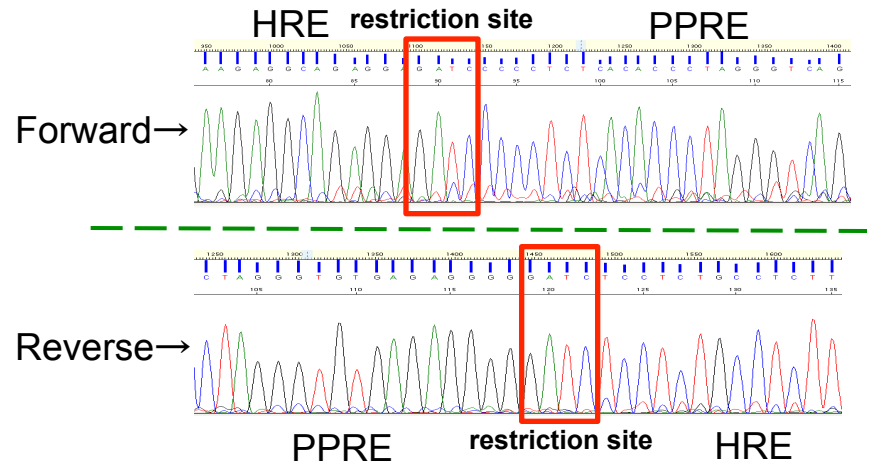


Hypoxia + PPAR β/δ agonist
Binding No; 37,050

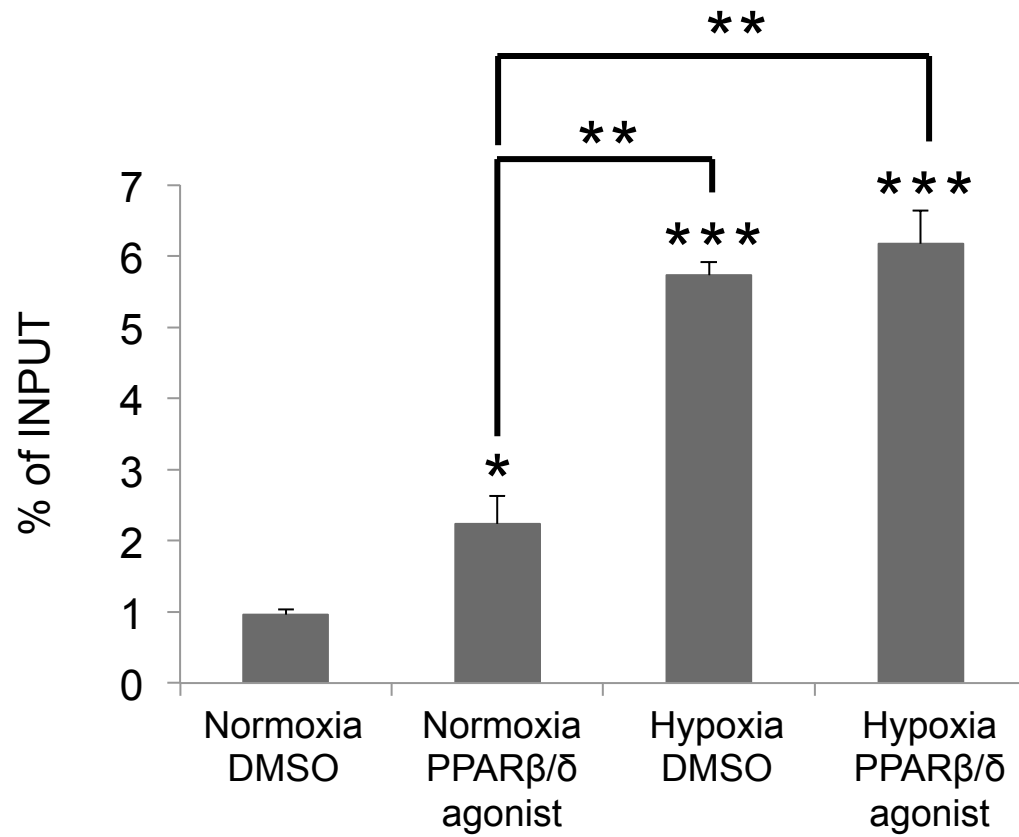


- Upstream
- TSS
- 5'UTR
- first intron
- other introns
- all exons
- 3'UTR
- Downstream
- intergenic

GGGAAGAGGCAGAGGAGATCCCCTCTCACACCCTAGGGTC



GACCCTAGGGTGTGAGTGGGGATCTCCTCTGCCTCTTCCC



p-value ($\log_{10}X$)