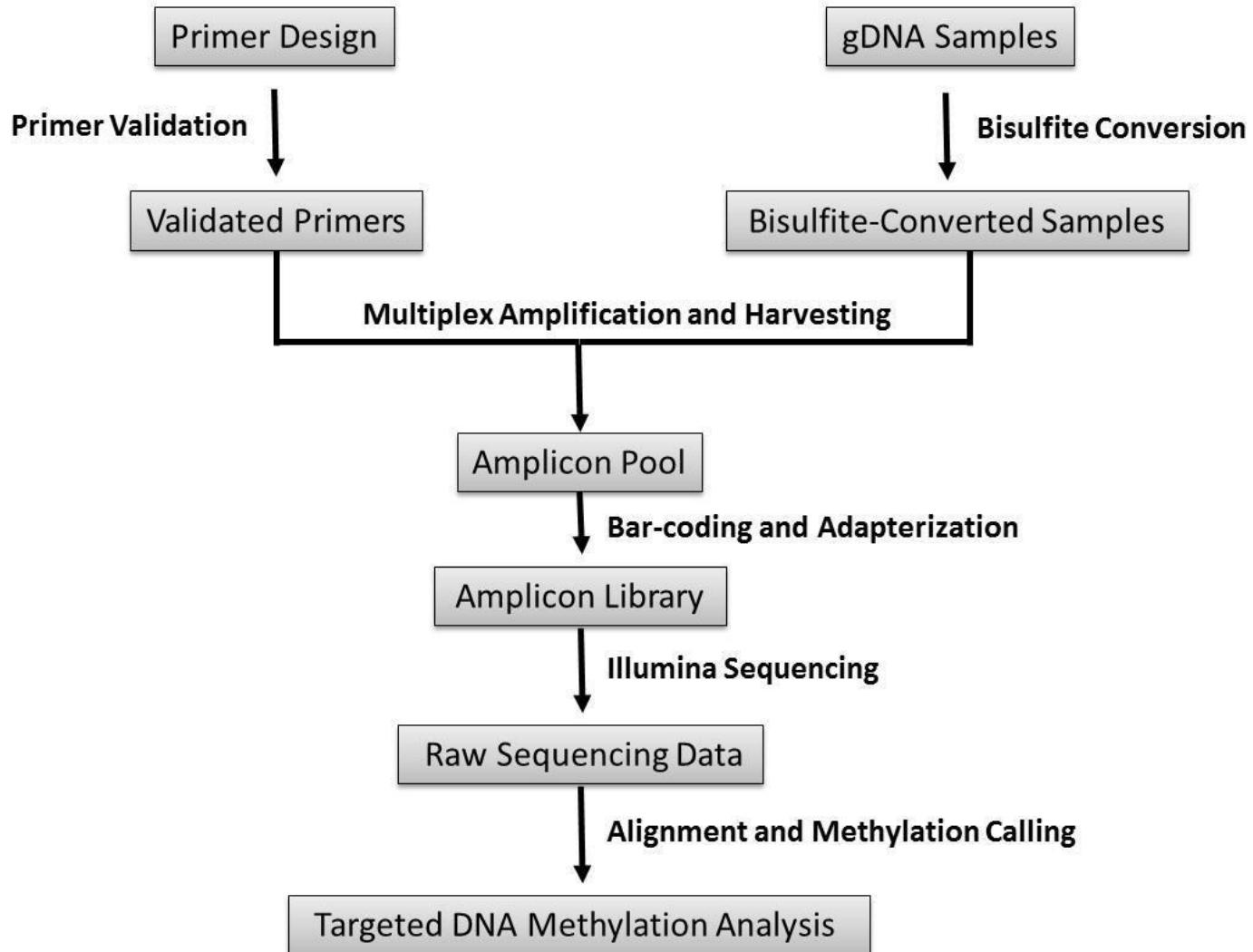


# Supplement # 1

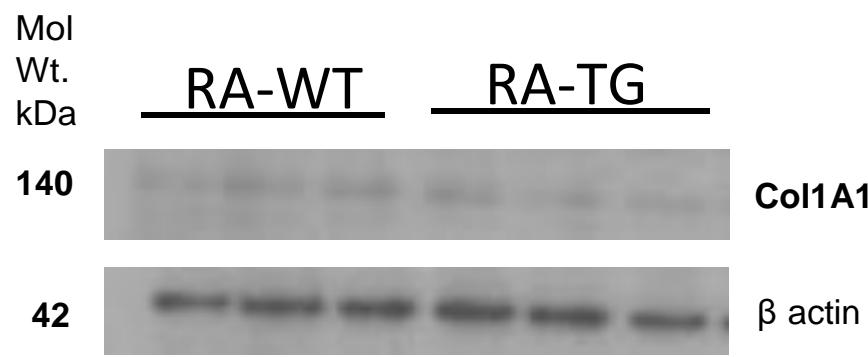


# Supplement # 2

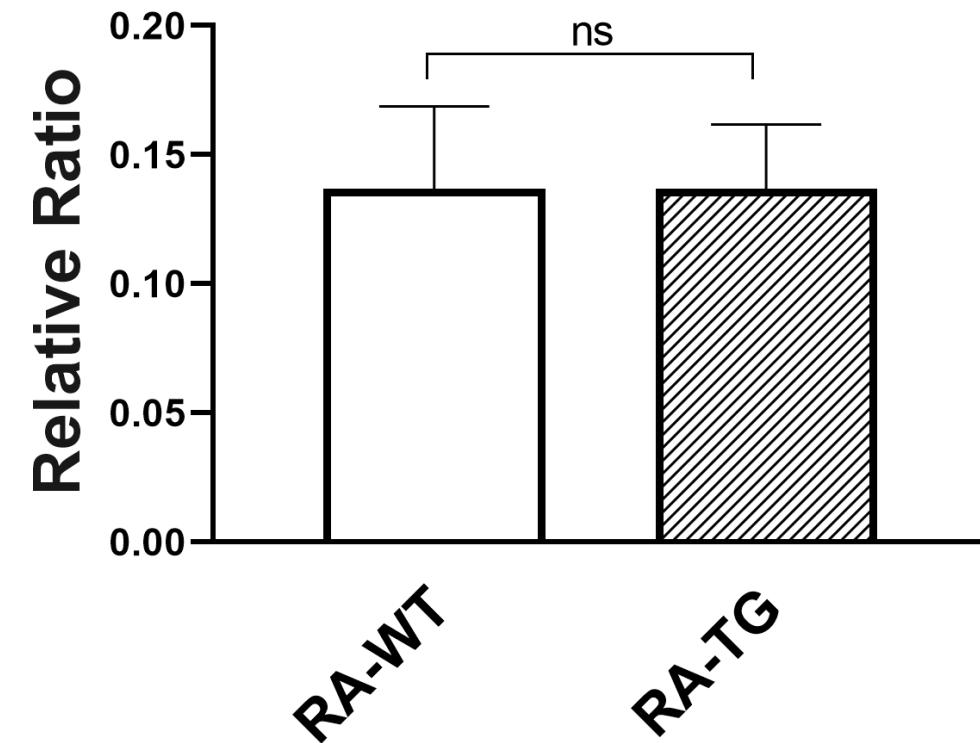
CGGCATCCGGTGGGCTCCTCTCCCTCCCCAGGCAAGCTGCAGCATTTGAGAGTGGCGATGGCAGACGGTGGC  
CCCCTCAGAGCAGCAAAGCTGAATAAACTGGATGGCAAGTATGGATCGCCCTGTACAATCTACTGCTCAGCCC  
TGAGGCCCGAGCCCGTTACTGCCTTACAAGCTTGCCAAGGGACAGTGCTTAAGGTGGAGAACTCAATGACCCC  
GCCAGCCCCACTGCCCTGCCAGACCCACCCCCAGGGCATCCTGGCCTGTCCCTGTCCCCTTCCACAAC  
TTGGGCTTCAGGCATCGACAGGGGAGCTGGTAAGATGCAAACCCCTCCAAGGACGCTCAGCCAATTTCAGGACTT  
CCTGTCTCTGGCCCTCTGGCTAGGAACAGTTAGTTCTCAATGCCCTCCCTGGATCTCTTGAGCTTCAGGCC  
TTCCTCACTGACACACTACTCGACCAGTTGCCAATCTTGCGATCTGAAGGGTTCTGGCCACCTGTCCCTG  
GCTGAAACCCAGCCCCCTAACAGAACGCTAGTGTAGAACAGGTAAGGCCCTAGAACATGCTGGAGTGCTTGG  
GACGGTGGTCCACTTTCCGGCTCCTCCCTGTCTCCATTACACTTTTAGATCCCAGAAATCTGGGATCG  
CCTGGAGAGAGAGAACAAAGGAAATGGCAGGCTATCGCCAAGCACCAGCTCAGCACGTATTAGCCTCTCGGA  
GAAGGATCTTGTCAACAAAGCACAGAGGTAAGTCCACTGAGTGGCTGAAGTCACGTGCATAGGACCAAGGAGGG  
AAGGTACACTAGTGCTCTGTGCACCTACCTGCCCGGTCCATTAGCTCTCAGCTCTCTCGACTCCTCCCCAGGTG  
GGCTGAAACCTACAGGCTGGATGTCCTAGAGGCAGTAGCTCCGGAGGCCCTGGCTACTGCAAACGAGA  
GGCCTCCAAGCGCTGCTCCAGATGCCAGAATGTGTGGTATTGCTGCAGGTGAGGCAGATCCTGGACCTTAGTGC  
CCCCCTAACGCTCAACCCCTGGGTCTCCTACAAACCCCTGTCTTCAGCTGCAGGCCACCCCGCTGCAGGGAGTGT  
CAAGTCAGCACTGGGAGAACGACCGAAAGACATGTGTTCTAGCAGCCAAGGTGACAGAGCCAAGTGAAGGC  
TAGCTGCCAGGGCCGACAGAACAGAGCGCACCCAGACTGTGACTGAGCTTCAGGGACTCGTGGCTCGCCCCGTCA  
GACCTCAATTCCCTGGTGAGCACAGCTGAGTCAGTAGAGCTGCAAACCACACACTCTTCACTCCTGCCCA  
CCCAGCGAATAACGAGAGGACCAGGCCACAGCATCAGCAAACCTGGCAGACAGGGTTGGGGGTGAAGCAAGGGC  
CGGATGTGGAGACCCCTTCCTCGCACAATAAACGAGACTCATGAAACGCAACAGTGTGGCGCTGTACAGTC  
TCCTGGGGTGGGGAGAACGACTGGACGGAAAGATTGGGAAGGGCTCTTGAAAGGGCCACCCCTGTGCCCGT  
GCCCGCCCACAAGTCCACACGGTCGGCTCGCCTAGGGTCCAGCGCTCCTCGGCTCTTCCCAGGCCCT  
GCCTGGATCTGGGGCGGTGCGGAAGTCGCGGCCCTGCCCTCGCCGCTCGTACTCACTAGCTGCTA  
GTCCCGTACAGCACCGGCTGACCGGGCATGTGGCGGAGCCAGAACTCATGAACTACGCGAGCTGGCACCGT  
CCGGCGCATGGTCCAGGCCGTACCCGGCTGGAGCGTGCCAAACGCTCTGCATGCCCGGGTACAACACGCA  
ATCCGTACAGCAGCACGTCCCAGGTGCTGGCCACCGTTCCAGCCTGCAGGGCCACACGACACCGTGGTGC  
ACCTCTGTGGAGACTTCATCTGGGGCGTCGTGCGAAGGGCTACAGTC

Small Letters = Repeat Sequences; Capital Letters = Unique Sequence; Dark Gray= Regions not covered by Targeted Sequencing Amplicons, Light Gray= Regions of Interest; CG= CG of interest >mm10. DNA range=chr9:107549843-107551842 5'pad=0 3'pad=0 Estimated No. of amplicons: 8.

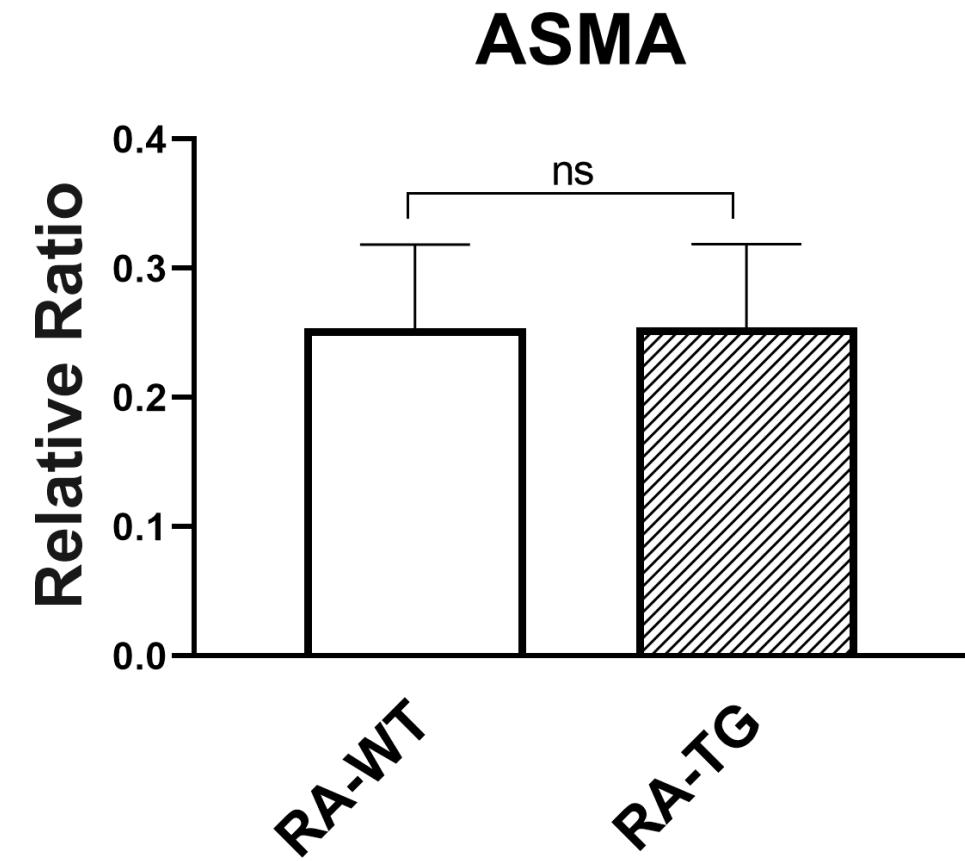
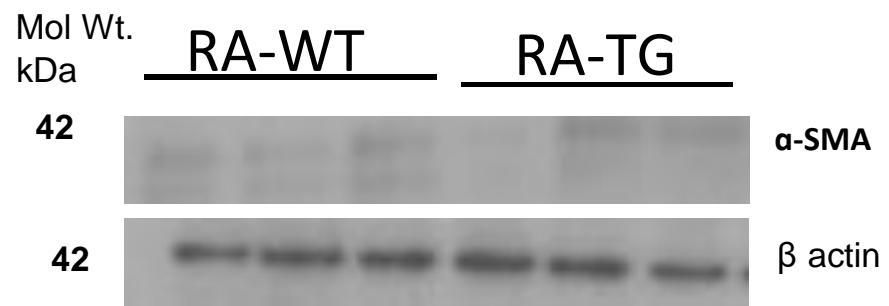
# Supplement # 3A



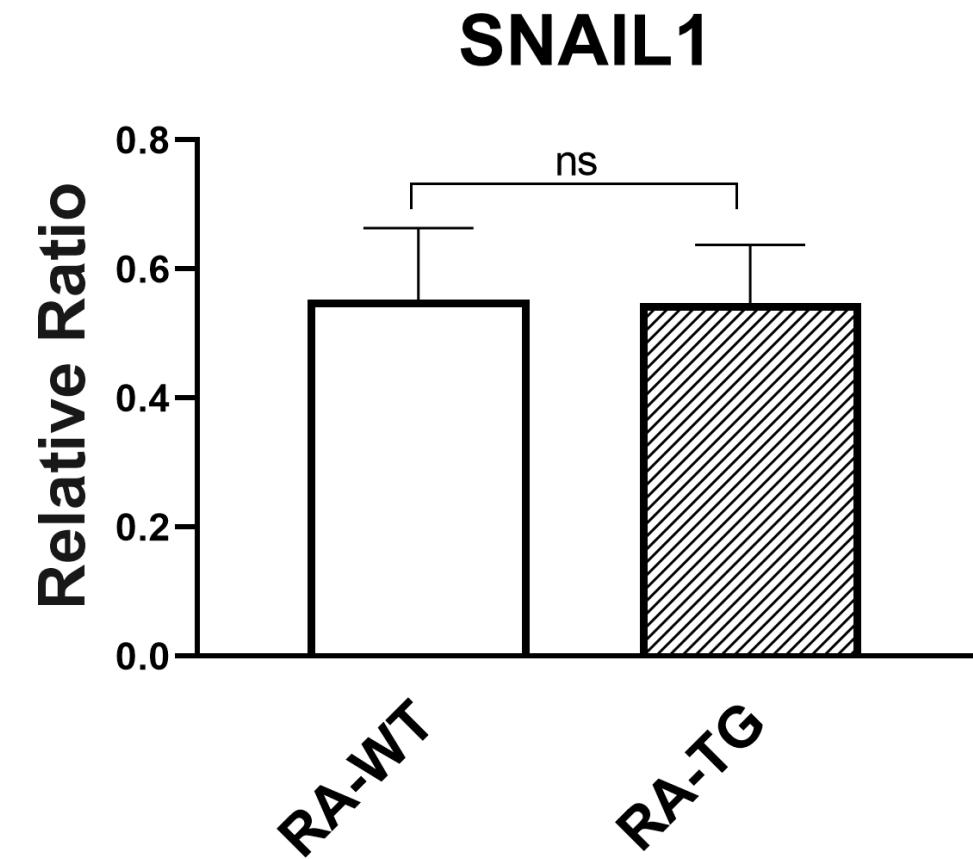
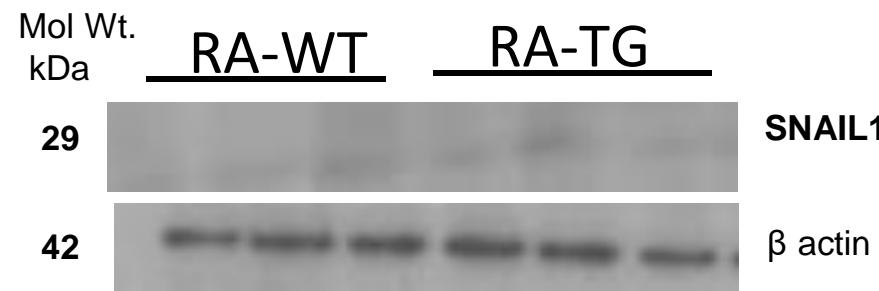
**Collagen 1 (Col1A1)**



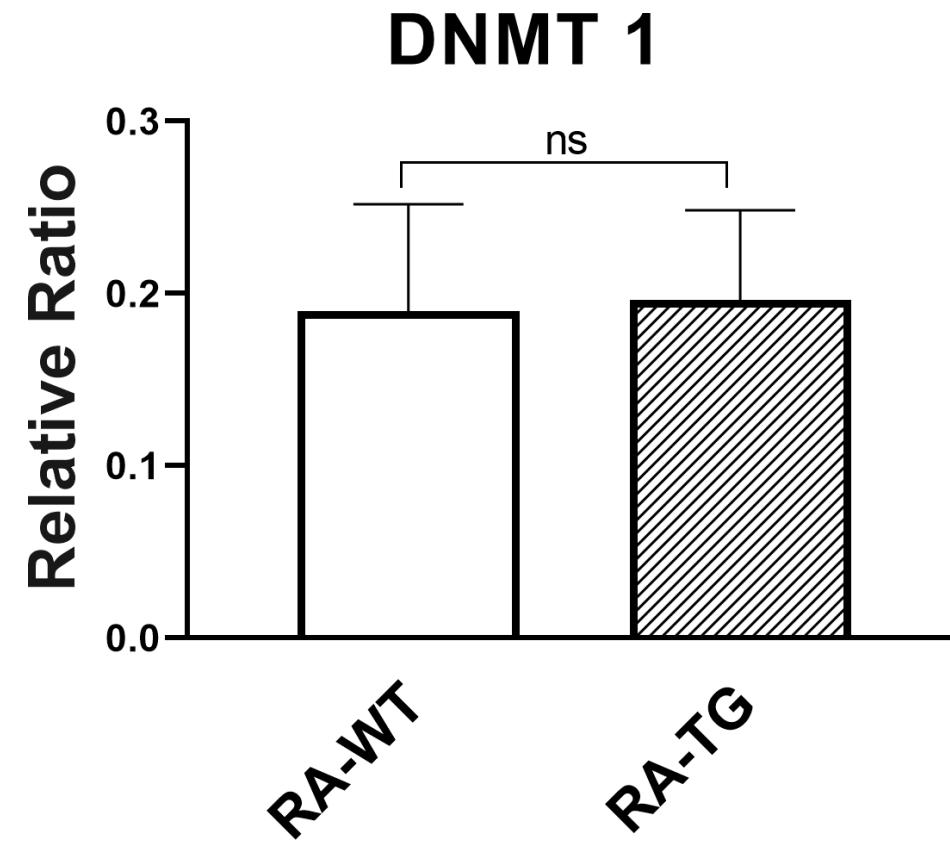
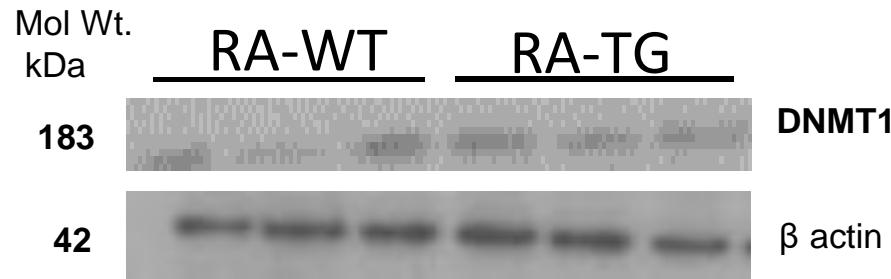
# Supplement # 3B



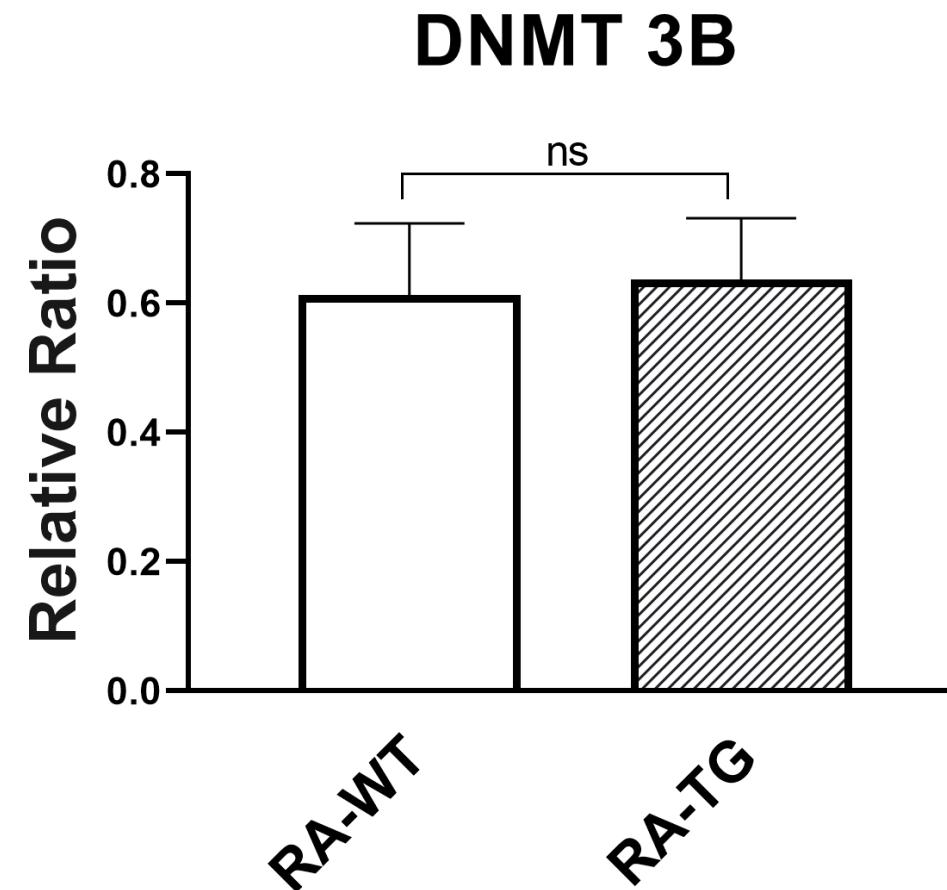
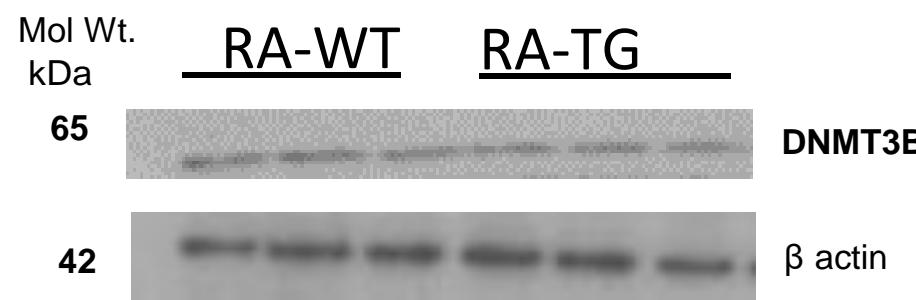
# Supplement # 3C



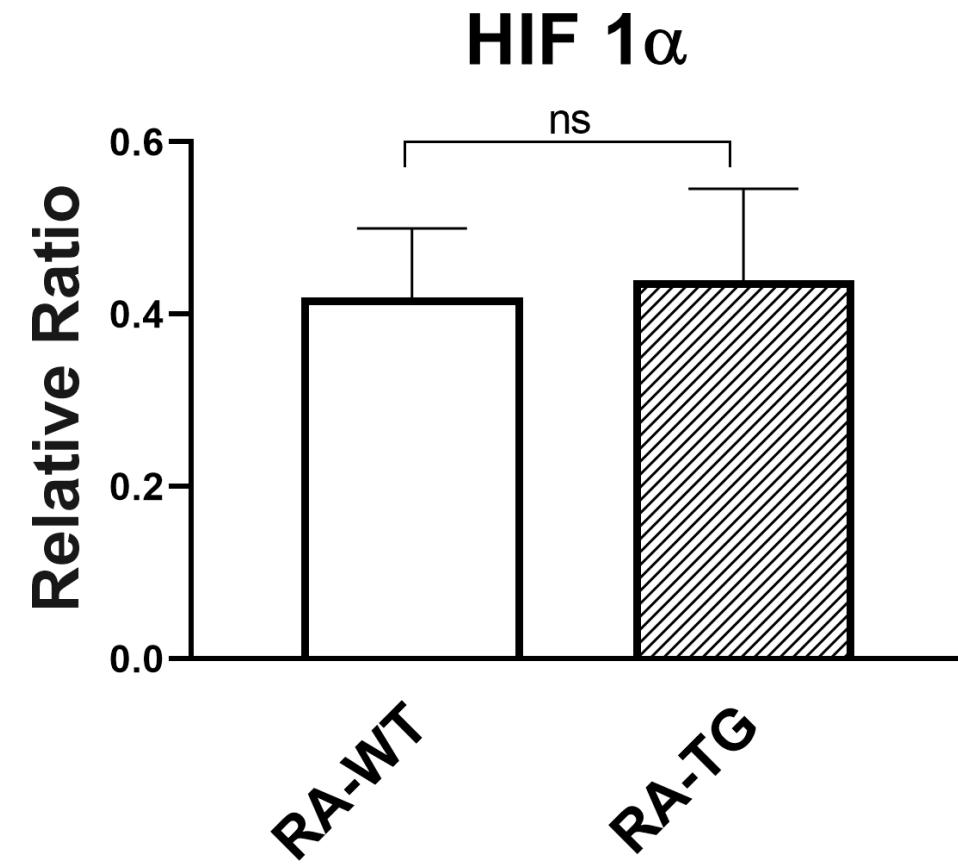
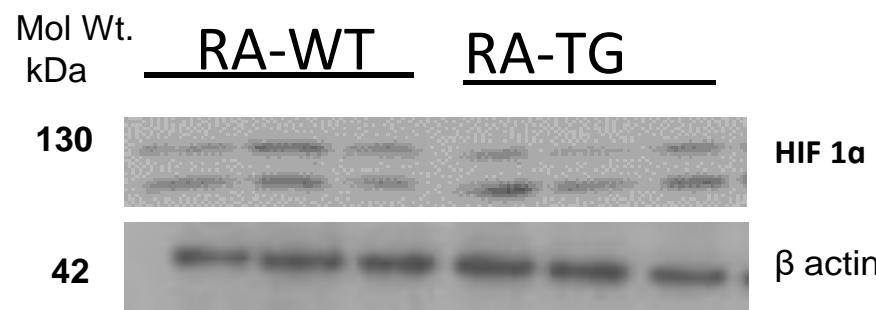
# Supplement # 3D



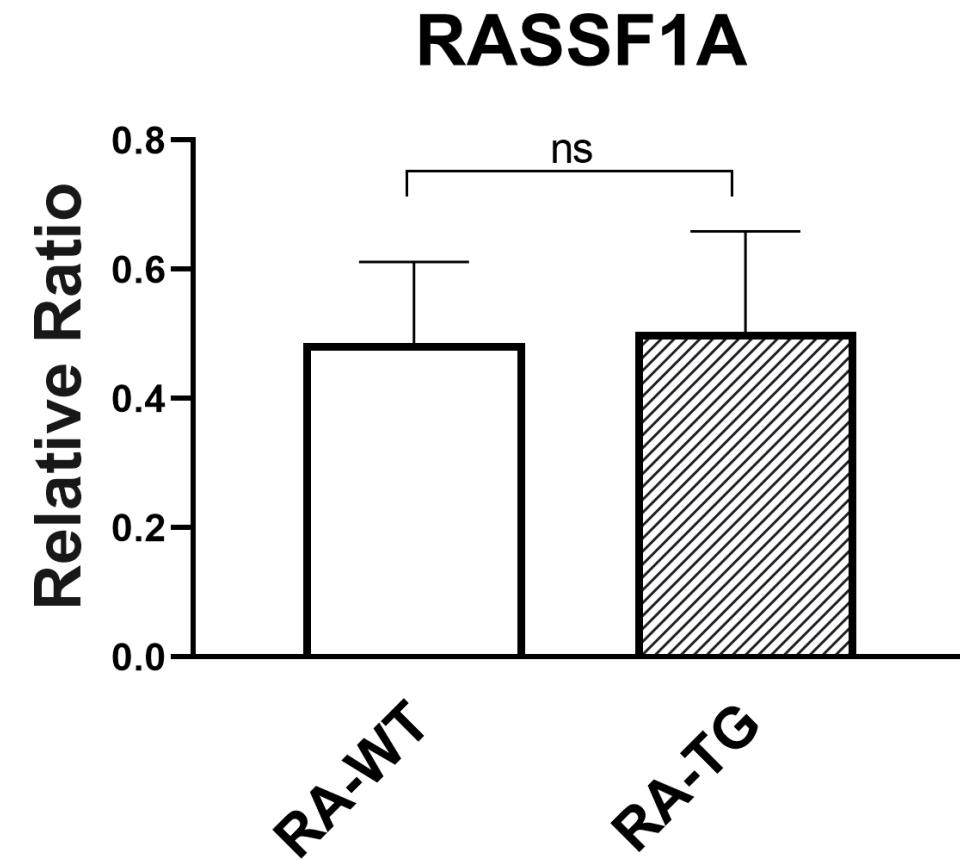
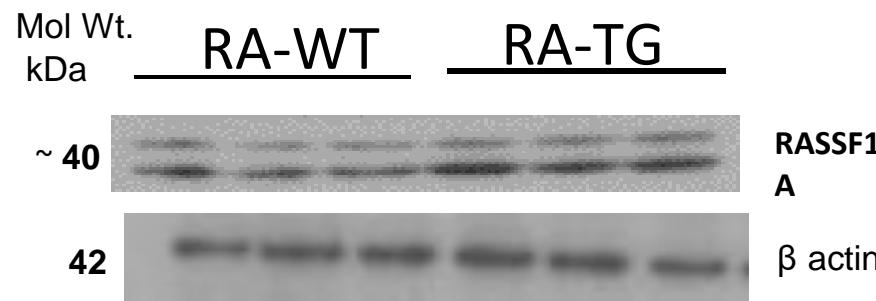
# Supplement # 3E



# Supplement # 3F



# Supplement # 3G



# Supplement # 3H

