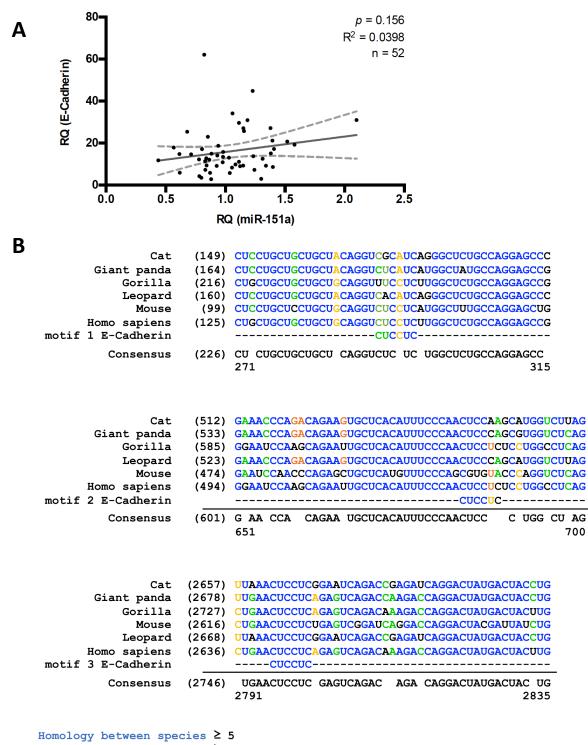
Supplemental Figure S9: E-cadherin and miR-151a correlation and miR-151a binding site conservation in E-cadherin mRNA. (A) The correlation between miR-151a and E-cadherin expression in NSCLC patient samples. The relative expression levels of miR-151a and E-cadherin were determined in 52 primary NSCLC samples by RT-qPCR. The potential correlation between miR-151a and E-cadherin expression was investigated by linear regression. The Line of Best Fit is indicated in black with grey dotted lines showing the 95% confidence interval (**p**=0.156, R2 = 0.0398). (**B**) mRNA sequence alignment of Homo sapiens (Z13009), Mouse (X06115), Leopard (XM_019455447), Gorilla (XM_004057872), Giant Panda (XM_002923207), and Cat (XM_003998189) as well as hsa-miR-151a (seed sequence, motif 1 to 3). Blue indicates homology between n≥5, green represents homology between n≥4 and orange represents homology between n≥ 3 species.



Homology between species ≥ 4 Homology between species ≥ 3