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miR-52	3'-UCGUGCCUUUGUAUACAUGCCCAC-5'
cdh-3 3'UTR WT	5`-UUUCAUCAUUUUCGAUUCAUUUUUACGGGUUCCUAAAUCU-3`
cdh-3 3'UTR MUT	5'-UUUCAUCAUUUUCGAUUCAUUUUUCGAAGUUCCUAAAUCU-3'
cdh-3 3'UTR DEL	5`-UUUCAUCAUUUUCGAUUCAUUUUUCCUAAAUCU-3`

FIGURE S3.—Mutations introduced into the full-length 3'UTR of *cdh-3*. 40 base pairs of the 3'UTR of *cdh-3* encompassing the predicted miR-52 target site is shown. Mutations introduced into the 3'UTR should abolish either some (MUT) or all (DEL) base pairing between the miRNA seed and the 3'UTR.