Tobias Le Stard Midteen Corrections 2. a, c 1 6. Step 1. The predictor variable it the APR payments 1 mude on my entir cord. The susponse variable the APR at the true of my payments. We model their **3**6 E lelationship. Step 2: We leave me model by computing least 3 squerce errimates for the intexcept and 3 8 = po + p1 x + 6, where e. APR sate, & : payments on credit card 6: sendon error 3 Step 3: To assess the fir of the model, we compute residuely and investigate the sesiduel is. fixed-values flor as well as the Normal quantite flor (we look for normality) to 4. With the estimated model, we can determine enfidence interale for our stope, mean lesponse. For instance, we can test if there truly it a staturially significant boear lelationship between our variable (40: \$2 = 0 x, H2: \$2 +0) 7. rull print seceired 8. a) We breeze that indence for brearity of the variable & selation ship in the first sest graph. Many the second, we conclude that our error component has constant variance here the data points look unformly seartised darry, the soudervized residuals dor Stones a Normal destribution of residuals Ohe quantile

(h) full points received. Ja) The estimated standard deriation is 30.84 (Surved of sendual standard error in R). 6) Our sunner expects to hunn go. 80.82 more calonel for each albetronal mph increase in werage sunning speed c) We use a 2-sample t-test to determine if there a difference in me sample means. Define promo and Myane age at the set of hurning calories of our sunner in the same age group as our sunner, suspectively.

Thus, we test: Ho! Mounter "Manne-age against 4: Mounte + Mounte age Since [80.82 - 22.57, 80.82 + 22.517 = [58.31, 103.33] eventa contains value that full writing I standard dividion of the estimated average sate of morning enloying of no surround (This internal includes 100), we war say that othere is not a significant difference in means d) The Re value of this wodel it 0.4313. This within of determination measures how close me data is do one fitted values. In this case, there it not to Strong our model does not predict very well. Furthermore, However, the p-value of our slope it 0.00225, which is much smaller Than 0.05, indicating that Alege's a high probability that new model exceptly depict the linear Selectionship between our

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