

## Appendix. Proofs.

### Residual association in situation 3

If  $M^* = M + \gamma_0 + \gamma_1 X + U$  where  $U$  is normally distributed with mean 0 and variance

$\sigma_u^2$ , the observed regression coefficient for  $X$  is approximately

$$\beta_1^* = \beta_1 - \beta_2^* \gamma_1 + \beta_2^* (1 - \lambda) \alpha_1 / \lambda, \text{ where } \lambda \text{ is the reliability ratio(5).}$$

Proof. First ignore the random error term  $U$  and write  $\tilde{M} = \gamma_0 + M + \gamma_1 X$ . When we consider the model  $\text{logit}(\Pr(Y=1 | \tilde{M}, X, C)) = \tilde{\beta}_0 + \tilde{\beta}_1 X + \tilde{\beta}_2 \tilde{M} + \tilde{\beta}_C^t C$ , it immediately follows that  $\tilde{\beta}_2 = \beta_2$ ,  $\tilde{\beta}_C^t = \beta_C^t$ ,  $\tilde{\beta}_0 = \beta_0 - \beta_2 \gamma_0$  and  $\tilde{\beta}_1 = \beta_1 - \beta_2 \gamma_1$ , with  $\beta_0$ ,  $\beta_1$ ,  $\beta_2$  and  $\beta_C$  the coefficients from the true logistic model (1). The measured intermediate then is  $M^* = \tilde{M} + U$ , with  $U$  normally distributed with mean 0, and variance  $\sigma_u^2$ . Using the formulas of Carroll et al<sup>3, p 52</sup> for bias in the regression coefficients for random measurement error, yields that the regression coefficients for exposure and intermediate in the model  $\text{logit}(\Pr(Y=1 | M^*, X, C)) = \beta_0^* + \beta_1^* X + \beta_2^* M^* + \beta_C^t C$ , are equal to  $\beta_2^* = \lambda \beta_2$  and

$$\beta_1^* = \beta_1 - \beta_2 \gamma_1 + \beta_2 (1 - \lambda) (E[M^* | X = 1, C] - E[M^* | X = 0, C]).$$

Using that  $E[M^* | X = 1, C] - E[M^* | X = 0, C] = \alpha_1 + \gamma_1$ , yields the required result.

### Residual association in situation 5

Suppose there is an interacting trigger  $T$ , which interacts with  $X$  such that  $M = c_0 + M^* + c_1 TX$ , with  $M^*$  the measured intermediate. It can be shown that in case of a rare disease

$$\Pr(Y=1 | M^*, X, C) \approx \exp(\beta_0 + \beta_1 X + \beta_2 (c_0 + M^*) + \beta_C^t C) \int \exp(\beta_2 c_1 t X) dF(t),$$

with  $F(t)$  the distribution function of  $T$ .

Proof: Because  $\text{logit}(\Pr(Y=1 | M, X, C)) = \beta_0 + \beta_1 X + \beta_2 M + \beta_C^t C$ , it follows that  $\text{logit}(\Pr(Y=1 | M^*, X, T, C)) = \beta_0 + \beta_1 X + \beta_2 (c_0 + M^* + c_1 TX) + \beta_C^t C$ .

Note that we do not observe  $T$ . If we perform a logistic regression analysis with  $M^*$  and  $X$  as covariates, we model  $\Pr(Y=1 | M^*, X, C)$ . This probability is equal to

$$\Pr(Y=1 | M^*, X, C) = \int \Pr(Y=1 | M^*, X, C, T=t) dF(t | M^*, X, C),$$

with  $F(t | M^*, X, C)$  the distribution function of  $T$  given  $M^*, X$  and  $C$ . If the trigger is not affected by the confounders, then, because  $T$  is unconditionally independent of  $M^*$  and  $X$ ,  $F(t | M^*, X, C) = F(t)$ . When the disease prevalence is low, odds ratios and relative risks are nearly equivalent and logistic models can be approximate by relative risk models.

Then

$$\Pr(Y=1 | M^*, X, C, T) \approx \exp(\beta_0 + \beta_1 X + \beta_2 (c_0 + M^* + c_1 TX) + \beta_C^t C).$$

It then follows that:

$$\Pr(Y=1 | M^*, X, C) \approx \exp(\beta_0 + \beta_1 X + \beta_2 (c_0 + M^*) + \beta_C^t C) \int \exp(\beta_2 c_1 tX) dF(t).$$

The integral in this expression can be simplified for several different distributions for  $T$ .

For example if the trigger is binary with  $p_T = \Pr(T=1)$ , then

$$\int \exp(\beta_2 c_1 tX) dF(t) = \exp(\beta_2 c_1 X) p_T + (1 - p_T).$$

$$\exp(\beta_1^*) = \Pr(Y=1 | M^*, X=1, C) / (\Pr(Y=1 | M^*, X=0, C))$$

$$\approx \exp(\beta_1) [\exp(\beta_2 c_1) p_T + (1 - p_T)], \text{ which leads to the result in (9).}$$

In case of additional random measurement error, assume that  $M^* = \tilde{M} + U$ , with

$$U \sim N(0, \sigma_u^2) \text{ and } M = c_0 + \tilde{M} + c_1 TX. \text{ Because } E[M | X, C] = \alpha_0 + \alpha_1 X + \alpha_2^t C, \text{ it}$$

follows that  $E[\tilde{M} | X, C] = \alpha_0 + \alpha_1 X + \alpha_2^t C - c_0 - p_T c_1 X$ . When using  $M^*$  instead of

$\tilde{M}$  in a logistic model, the formulas of Carroll et al<sup>3, p 52</sup> for bias in the regression

coefficients for classical measurement error, yield that  $\beta_1^* = \tilde{\beta}_1 + \beta_2^* (1 - \lambda)(\alpha_1 - p_T c_1) / \lambda$

and  $\beta_2^* = \lambda \tilde{\beta}_2$ . Combining this with (9) gives

$$\beta_1^* = \beta_1 + \log[\exp(\beta_2^* c_1 / \lambda) p_T + (1 - p_T)] + \beta_2^* (1 - \lambda)(\alpha_1 - p_T c_1) / \lambda$$

## Residual association in situation 6

If there is a post-hoc phenomenon, such that  $M^* = M + \gamma_0 + \gamma_1 Y + U$ , where

$U \sim N(0, \sigma_u^2)$ , then approximately:

$$\beta_2^* \approx \lambda \beta_2 + \frac{\gamma_1}{\sigma_M^2 + \sigma_U^2} \text{ and}$$

$$\beta_1^* = \beta_1 - \frac{\gamma_1 \alpha_1}{\sigma_M^2 + \sigma_U^2} + \left( \beta_2^* - \frac{\gamma_1}{\sigma_M^2 + \sigma_U^2} \right) (1 - \lambda) \alpha_1 / \lambda .$$

Proof: First ignore the random error term  $U$  and write  $M^* = \gamma_0 + M + \gamma_1 Y$ . Bayes' theorem gives that

$$\text{odds}(\Pr(Y = 1 | X, M^* = m^*, C)) = \frac{f_{X,C,M^*}(X, m^*, C | Y = 1) \Pr[Y = 1]}{f_{X,C,M^*}(X, m^*, C | Y = 0) \Pr[Y = 0]},$$

where  $f$  indicates the density function.

Note that  $f_{X,C,M^*}(X, m^*, C | Y = 1) = f_{X,C,M}(X, m^* - \gamma_0 - \gamma_1, C | Y = 1)$  and

$f_{X,C,M^*}(X, m^*, C | Y = 0) = f_{X,C,M}(X, m^* - \gamma_0, C | Y = 0)$ . Applying again Bayes' theorem yields:

$$\text{odds}(\Pr(Y = 1 | X, M^* = m^*, C)) = \frac{\Pr(Y = 1 | X, M = m^* - \gamma_0 - \gamma_1, C)}{\Pr(Y = 0 | X, M = m^* - \gamma_0, C)} \frac{f_{M|X,C}(m^* - \gamma_0 - \gamma_1 | X, C) f(X, C)}{f_{M|X,C}(m^* - \gamma_0 | X, C) f(X, C)}$$

Because  $M|X, C$  is normally distributed with constant variance, it is straightforward to show that

$$\frac{f_{M|X,C}(m^* - \gamma_0 - \gamma_1 | X, C)}{f_{M|X,C}(m^* - \gamma_0 | X, C)} = \exp \left( \frac{1}{\sigma_M^2} \left[ -0.5 \gamma_1^2 - \gamma_0 \gamma_1 - \gamma_1 \alpha_0 - \gamma_1 \alpha_1 X - \gamma_1 \alpha_2 C + \gamma_1 m^* \right] \right)$$

Using that, in case of a rare disease,  $\Pr(Y=0|X, M, C) \approx 1$ , and that in case of a rare disease  $\Pr(Y=1|X, M, C)$  can be approximate by a relative risk model, yields

odds( $\Pr(Y = 1 | X = 1, M^* = m^*, C) \approx$

$$\exp\left(\beta_0 - \frac{0.5\gamma_1^2 + \gamma_0\gamma_1 + \gamma_1\alpha_0}{\sigma_M^2} + \beta_2(-\gamma_0 - \gamma_1) + \left(\beta_1 - \frac{\gamma_1\alpha_1}{\sigma_M^2}\right)X + \left(\beta_c^t - \frac{\gamma_1\alpha_2^t}{\sigma_M^2}\right)C + \left(\beta_2 + \frac{\gamma_1}{\sigma_M^2}\right)m^*\right)$$

This gives  $\beta_1^* \approx \beta_1 - \frac{\gamma_1\alpha_1}{\sigma_M^2}$  and  $\beta_2^* \approx \beta_2 + \frac{\gamma_1}{\sigma_M^2}$

In case of random measurement error, with  $M^* = M + \gamma_0 + \gamma_1 Y + U$ , assume first that

$\tilde{M} = M + U$ . If  $\tilde{M}$  is used instead of  $M$  in the logistic model, the formulas of Carroll et al<sup>3</sup>,  
p<sup>52</sup> for bias in the regression coefficients for classical measurement error,

yield  $\tilde{\beta}_1 = \beta_1 + \beta_2(1 - \lambda)\alpha_1$  and  $\tilde{\beta}_2^* = \lambda\beta_2$ . Since  $M^* = \gamma_0 + \tilde{M} + \gamma_1 Y$ , we have

$\beta_1^* \approx \tilde{\beta}_1 - \frac{\gamma_1\alpha_1}{\sigma_{\tilde{M}}^2}$  and  $\beta_2^* \approx \tilde{\beta}_2 + \frac{\gamma_1}{\sigma_{\tilde{M}}^2}$ . This leads to the final results:

$$\beta_2^* \approx \lambda\beta_2 + \frac{\gamma_1}{\sigma_{\tilde{M}}^2}$$

and  $\beta_1^* \approx \beta_1 - \gamma_1\alpha_1 / \sigma_{\tilde{M}}^2 + \beta_2(1 - \lambda)\alpha_1 = \beta_1 - \gamma_1\alpha_1 / \sigma_{\tilde{M}}^2 + \left(\beta_2^* - \frac{\gamma_1}{\sigma_{\tilde{M}}^2}\right)(1 - \lambda)\alpha_1 / \lambda$ .