Additional file 4. A proof that the bounds are sharper than previous bounds.

For the lower bounds:

$$\operatorname{Risk}_{\operatorname{class}=i,j}^{\operatorname{LB}} = 1 - \min_{\substack{(i'\neq i)\in\{1,\dots,L_{1}\}\\(j'\neq j)\in\{1,\dots,L_{2}\}}} \left\{ \frac{1 - \operatorname{Risk}^{\operatorname{profile}=i,j}}{(1 - \operatorname{Risk}^{\operatorname{profile}=i,j}) \times (1 - \operatorname{Risk}^{\operatorname{profile}=i,j'})}, 1 \right\}$$
$$= \max_{\substack{(i'\neq i)\in\{1,\dots,L_{1}\}\\(j'\neq j)\in\{1,\dots,L_{2}\}}} \left\{ \frac{\operatorname{Risk}^{\operatorname{profile}=i,j} - \operatorname{Risk}^{\operatorname{profile}=i',j} - \operatorname{Risk}^{\operatorname{profile}=i,j'} + \operatorname{Risk}^{\operatorname{profile}=i',j} \times \operatorname{Risk}^{\operatorname{profile}=i,j'}}, 0 \right\}$$
$$\geq \max_{\substack{(i'\neq i)\in\{1,\dots,L_{2}\}\\(j'\neq j)\in\{1,\dots,L_{2}\}}} \left\{ \operatorname{Risk}^{\operatorname{profile}=i,j} - \operatorname{Risk}^{\operatorname{profile}=i',j} - \operatorname{Risk}^{\operatorname{profile}=i,j'}, 0 \right\}$$
$$= \operatorname{Sjölander} \text{ et al.'s lower bound.}$$

For the upper bounds:

Sjölander et al. (21) also derived an assumption-free upper bound for the specific interaction:

$$\min\left\{\operatorname{Risk}^{\operatorname{profile}=i,j}, \sum_{(i'\neq i)\in\{1,\dots,L_1\}} \left(1-\operatorname{Risk}^{\operatorname{profile}=i',j}\right), \sum_{(j'\neq j)\in\{1,\dots,L_2\}} \left(1-\operatorname{Risk}^{\operatorname{profile}=i,j'}\right)\right\}.$$
 However, this bound is

wrongly too sharp. In fact, the cumulative completion risk of the class = i, j sufficient-cause

interaction is bounded above, neither by $\sum_{(i'\neq i)\in\{1,\dots,L_1\}} (1-\text{Risk}^{\text{profile}=i',j})$ nor by

 $\sum_{\substack{(j'\neq j)\in\{1,\dots,L_2\}}} \left(1 - \text{Risk}^{\text{profile}=i,j'}\right), \text{ as dictated by Sjölander et al (21). To see why, let us assume that all }$

the diseased subjects in a population are caused by and only by sufficient-cause interactions, that is, all the profile = i, j diseased subjects had contracted the disease because of the completion of the class = i, j sufficient-cause interaction, for all $i \in \{1, ..., L_1\}$ and all $j \in \{1, ..., L_2\}$. Further, assume that the follow-up is very long (T is very large) so that $\operatorname{Risk}^{\operatorname{profile}=i,j} \approx 1$ for all $i \in \{1,...,L_1\}$ and all $j \in \{1, ..., L_2\}$. Sjölander et al.'s upper bound (21) then becomes nearly zero. Clearly, it should not be.