## **GAS** Power Calculator Relevant Equations

GAS Power Calculator Copyright 2017 Jennifer Li Johnson University of Michigan School of Public Health Department of Biostatistics Center for Statistical Genetics <a href="http://csg.sph.umich.edu/abecasis/gas\_power\_calculator">http://csg.sph.umich.edu/abecasis/gas\_power\_calculator</a>

$$AAfreq = (DAF)^2 \tag{1}$$

$$ABfreq = 2 * (DAF) * (1 - DAF)$$
(2)

$$BBfreq = (1 - DAF)^2 \tag{3}$$

Multiplicative Disease Model

$$x = [(GRR)^2, GRR, 1.0] (4)$$

Additive Disease Model

$$x = [2.0 * GRR - 1.0, GRR, 1.0]$$
 (5)

Dominant Disease Model

$$x = [GRR, GRR, 1.0] \tag{6}$$

Recessive Disease Model

$$x = [GRR, 1.0, 1.0] (7)$$

$$AAprob = \frac{x[0] * prev}{x[0] * AAfreq + x[1] * ABfreq + x[2] * BBfreq}$$
 (8)

$$ABprob = \frac{x[1]*prev}{x[0]*AAfreq + x[1]*ABfreq + x[2]*BBfreq} \quad (9)$$

$$BBprob = \frac{x[2] * prev}{x[0] * AAfreq + x[1] * ABfreq + x[2] * BBfreq} \quad (10)$$

$$casesDAF = \frac{AAprob*AAfreq + ABprob*ABfreq*0.5}{prev} \quad (11)$$

$$controlsDAF = \frac{(1 - AAprob) * AAfreq + (1 - ABprob) * ABfreq * 0.5}{1 - prev}$$
(12)

$$V cases = casesDAF * (1 - casesDAF)$$
 (13)

$$V controls = controls DAF * (1 - controls DAF)$$
 (14)

$$ncp = \frac{casesDAF - controlsDAF}{\sqrt{\left(\frac{Vcases}{cases} + \frac{Vcontrols}{controls}\right) * 0.5}}$$
(15)

$$C = -ninv(\alpha * 0.5) \tag{16}$$

$$P = ndist(-C - ncp, false) + ndist(C - ncp, true)$$
 (17)

## Notes:

- Invalid model if AAprob > 1
- Vcases: Cases Variance
- Vcontrols: Controls Variance
- ncp: noncentrality parameter
- C: significance threshold
- ndist: Standard normal distribution adapted from ID Hill, "The Normal Integral" Applied Statistics, Vol 22, pp. 424-427 available here.
- ninv: Inverse normal distribution adapted from Wichura's PPND16, Algorithm AS241, Applied Statistics Vol 37 1988 pp 477 484 available here.

For definitions of the other variables click here.