

1. Page 2,

CHANGE: "This requires equating the expected proportion of genotypes shared identical by descent, given a pedigree, with the observed phenotypic correlation between relatives."

TO: "This requires equating the expected proportion of the genome shared identical by descent, given a pedigree, with the observed phenotypic correlation between relatives."

(see note0101.pdf at <https://github.com/SorensenD/SLGDS>)

2. Page 258, eqn. (6.1),

CHANGE second line to (brackets missing):

$$\begin{aligned}
 E_{y_0x} [\text{MSE} (f(x))] &= E_{y_0x} [(y_0 - f(x))^2] \\
 &= E_x [E_{y_0|x} [(y_0 - f(x))^2 | x]] \\
 &= E_x [\text{Var} (y_0|x) + (E (y_0|x) - f(x))^2], \quad (6.1)
 \end{aligned}$$