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)

$$E_{y_0x} [MSE (f (x))] = E_{y_0x} [(y_0 - f (x))^2]$$

$$= E_x [E_{y_0|x} [(y_0 - f (x))^2 | x]]$$

$$= E_x [Var (y_0|x) + (E (y_0|x) - f (x))^2], \qquad (6.1)$$