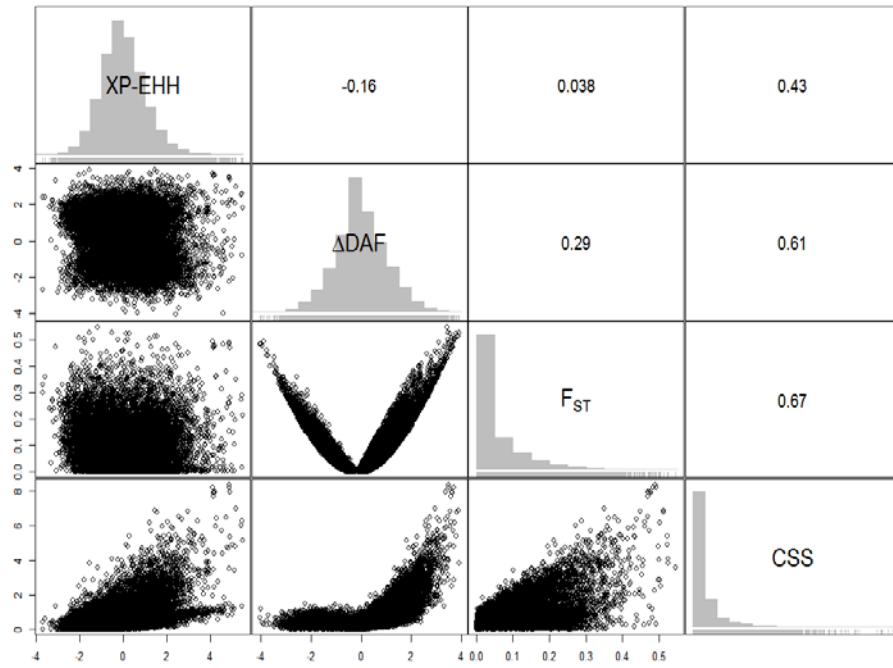
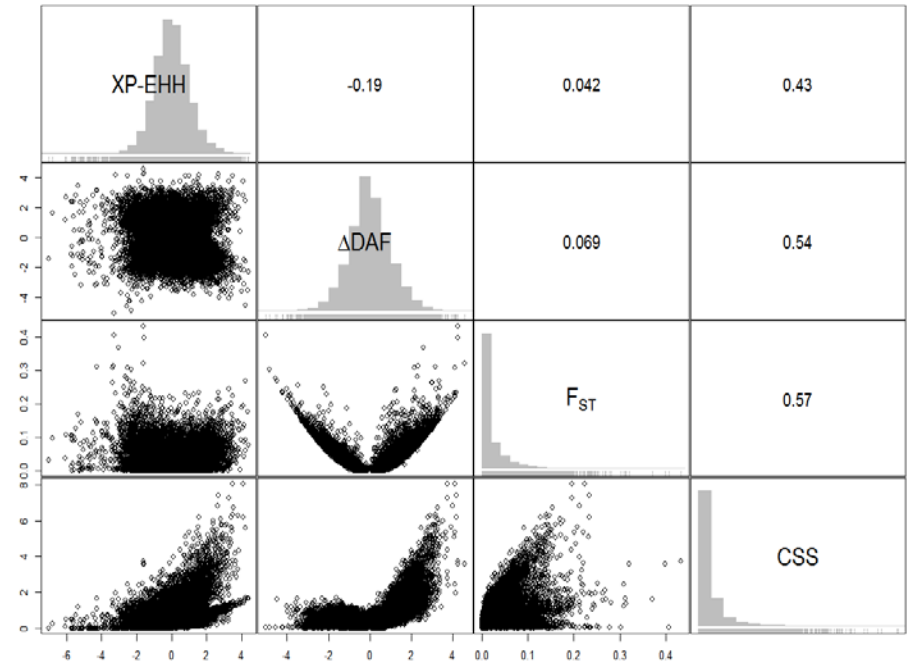


A Pairs Plot of Polled Cattle



B Pairs Plot of Double Muscle Cattle



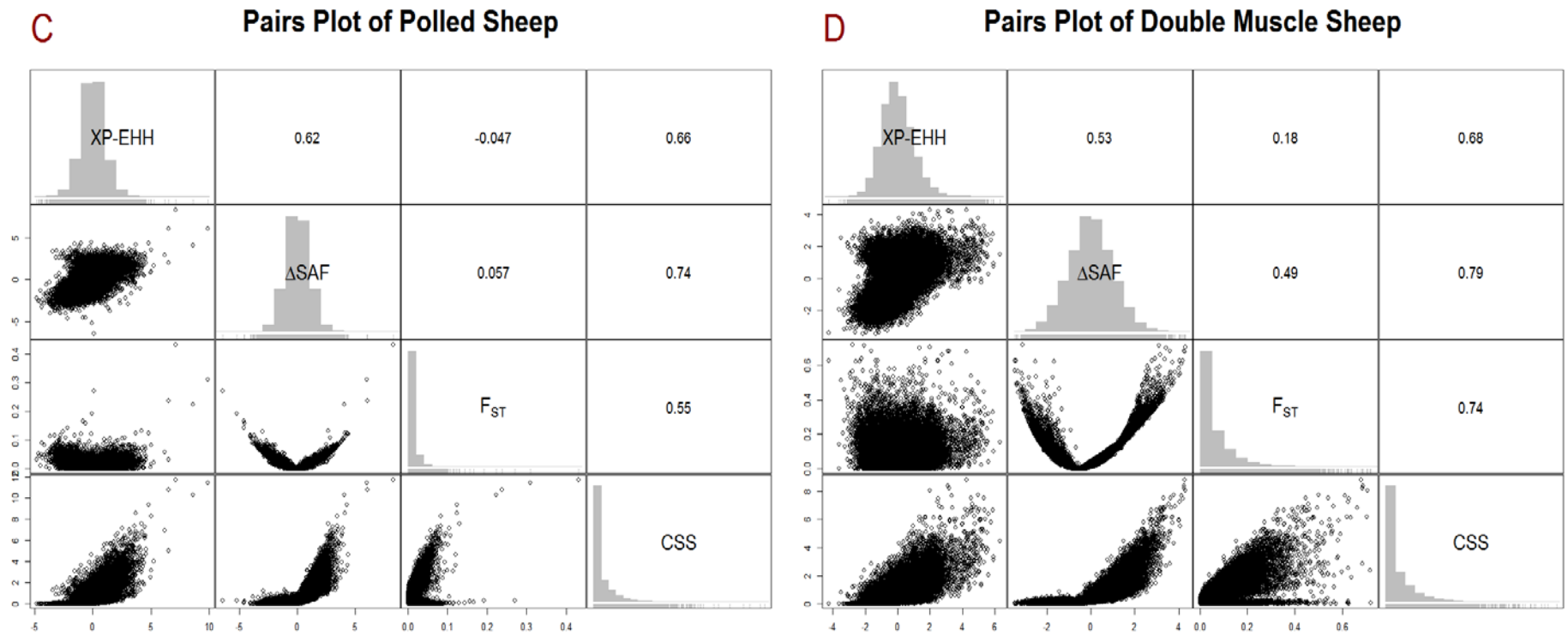


Figure S6. Genome-wide pairs plots (lower diagonals), histograms (diagonals) and correlations (upper diagonals) for constituent (XP-EHH, Δ SAF, F_{ST}) and composite selection signals (CSS) for polled (A), double muscle (B), cattle polledness (C) and double muscling (D) in sheep.