logi.S

Figure: Forest plot of coefficient β estimates with 99% confidence intervals for genes RP11−909M7.3, RP13−487P22.1, NLRP2, SNHG14, PEG10, hsa−mir−335, SNORD116−20, PWAR6, SNRPN, MEST, PWRN1, AL132709.5, WA, IGF2, KCNQ1OT1, WA.8, NAP1L5, MEG3, GRB10, INPP5F, ZNF331, PEG3, UBE3A, ZDBF2. Legend: estimate $\hat{\beta}$; 99 % conf. int. x-axis: coefficient β.