



Fig. S6. Genetic variations contribute to domestication in Asian and African rice. a-d Divergent regions identified by F_{ST} using SVs: Manhattan plot of F_{ST} values between *Or* and *Osi* (a), F_{ST} values between *Or* and *Osj* (b), F_{ST} values between *Ob* and *Og* (c), and F_{ST} values between Asian and African rice (d). e-h Divergent regions identified by F_{ST} using SNPs: Manhattan plot of F_{ST} values between *Or* and *Osi* (e), between *Or* and *Osj* (f), between *Ob* and *Og* (g), and between Asian rice and African rice (h). i-k Divergent regions identified by D_{xy} using SVs: Manhattan plot of D_{xy} values between *Or* and *Osi* (i), between *Or* and *Osj* (j), and between *Ob* and *Og* (k). l-n Divergent regions identified by D_{xy} using SNPs: Manhattan plot of D_{xy} values between *Or* and *Osi* (l), between *Or* and *Osj* (m), and between *Ob* and *Og* (n). Manhattan plots are at 20 kb resolution; dashed lines indicate the top 5% of D_{xy} or F_{ST} values and black lines indicate known genes. *Osi*, *Osj*, *Or*, *Og*, and *Ob* refer to *O. sativa indica*, *O. sativa japonica*, *O. rufipogon*, *O. glaberrima*, and *O. barthii*, respectively.