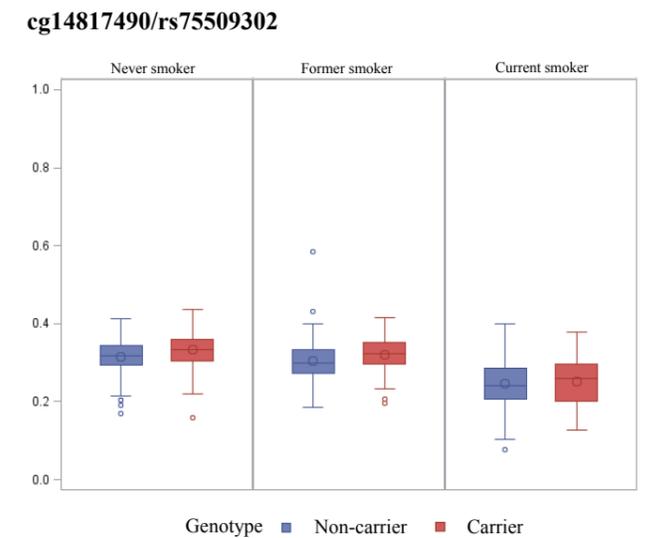
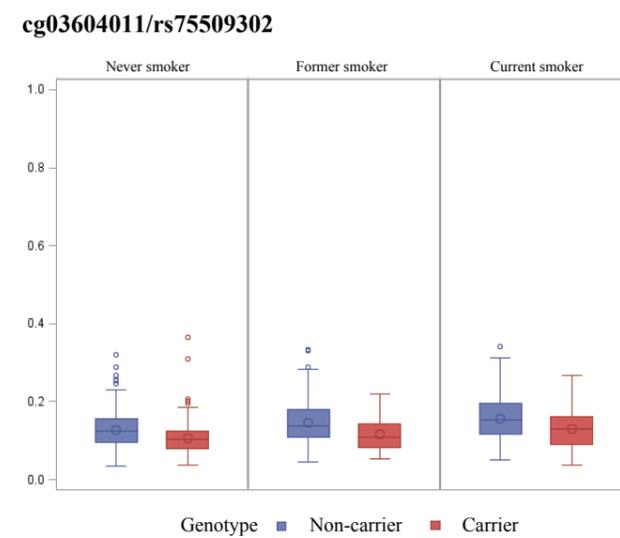
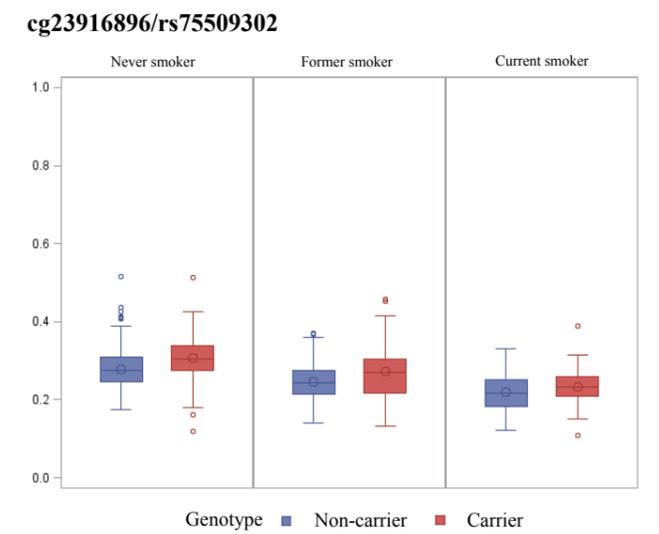
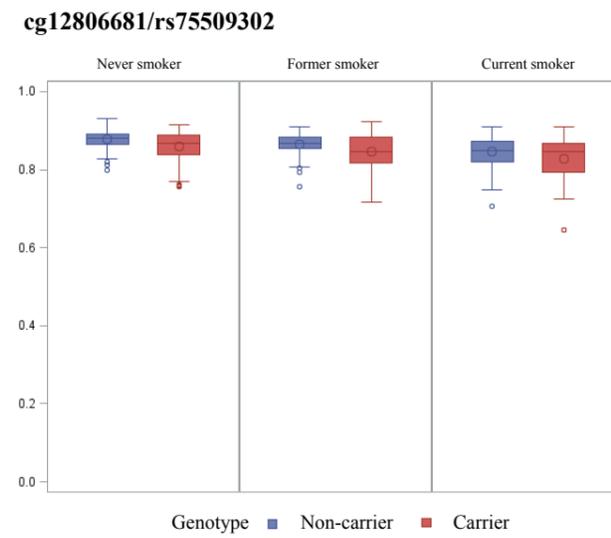
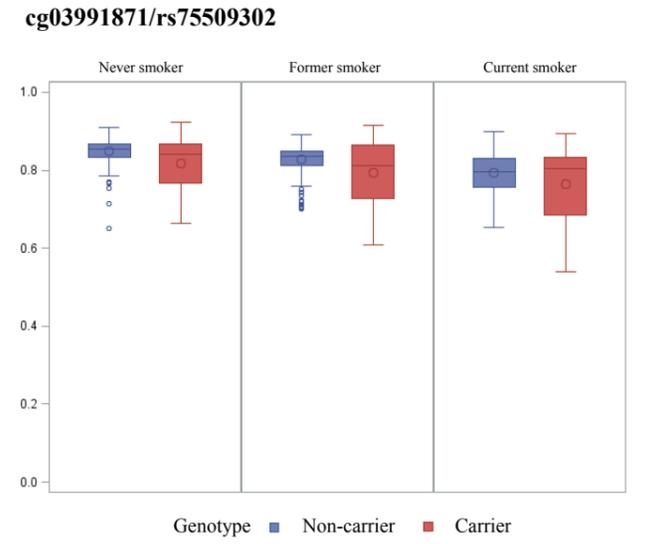
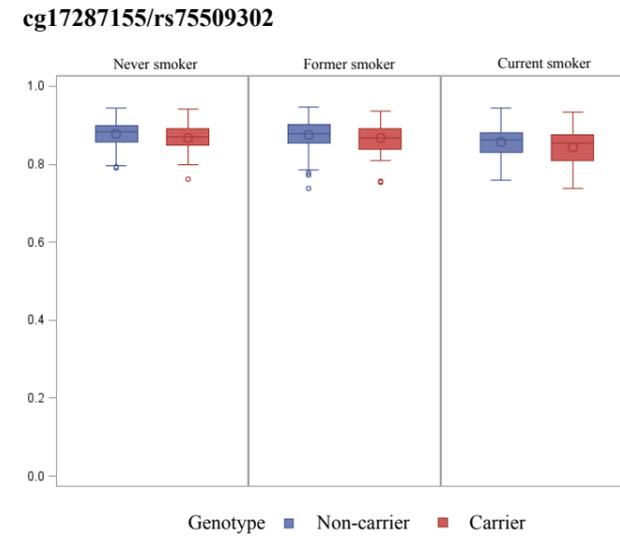
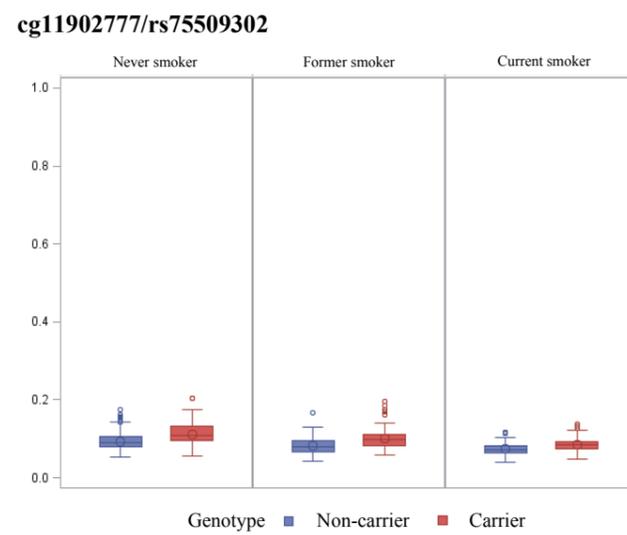
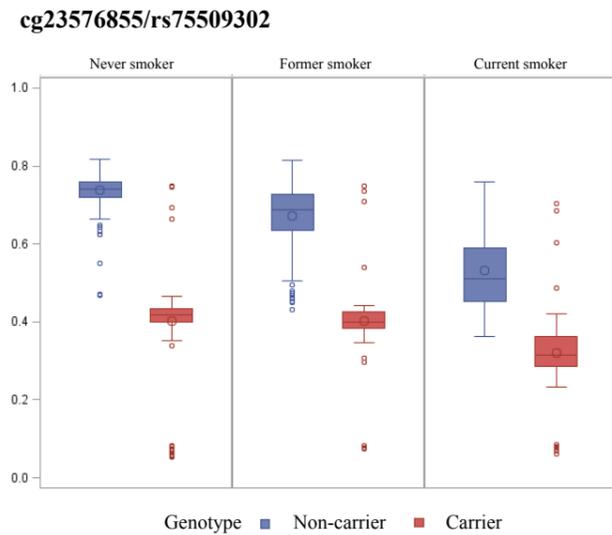
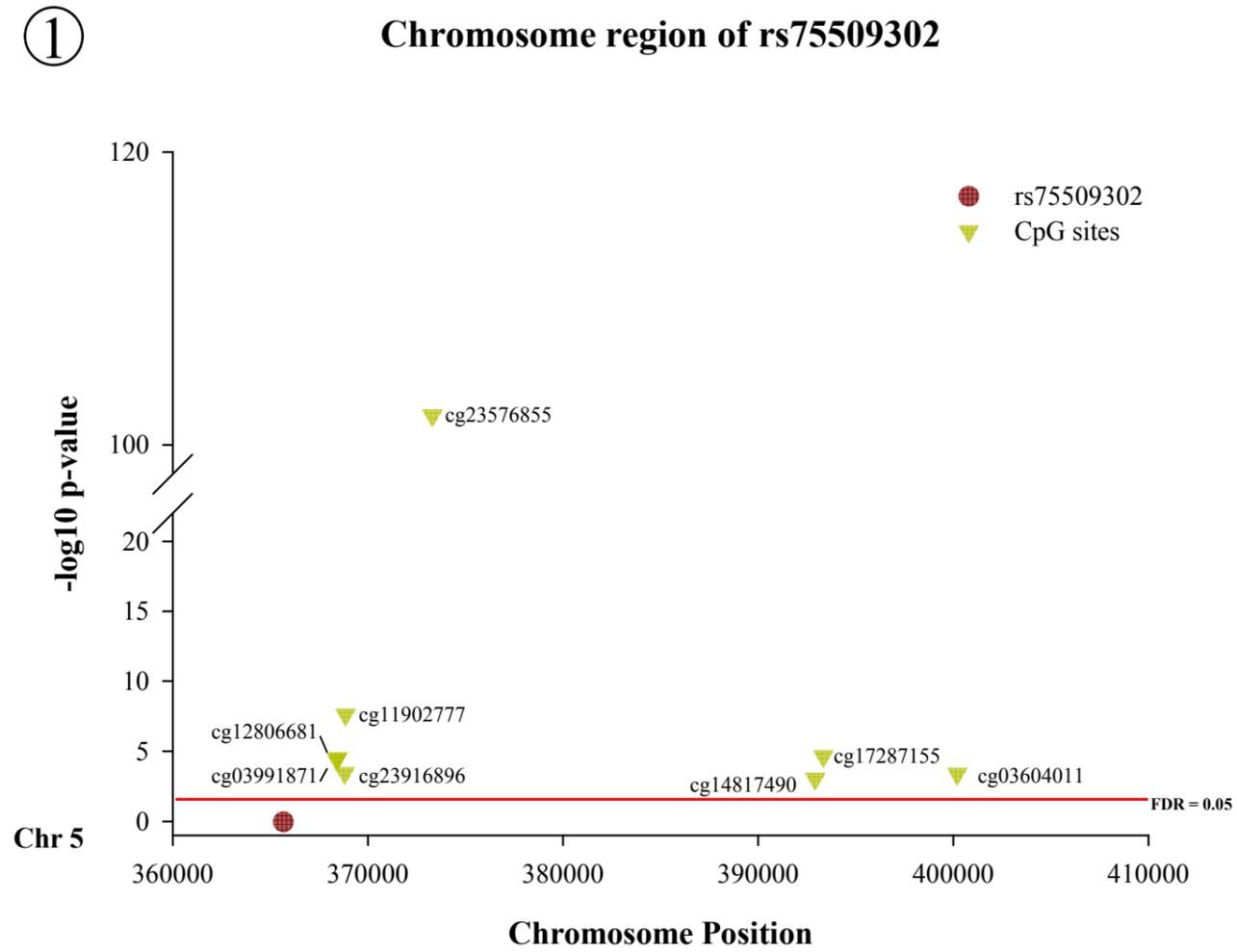
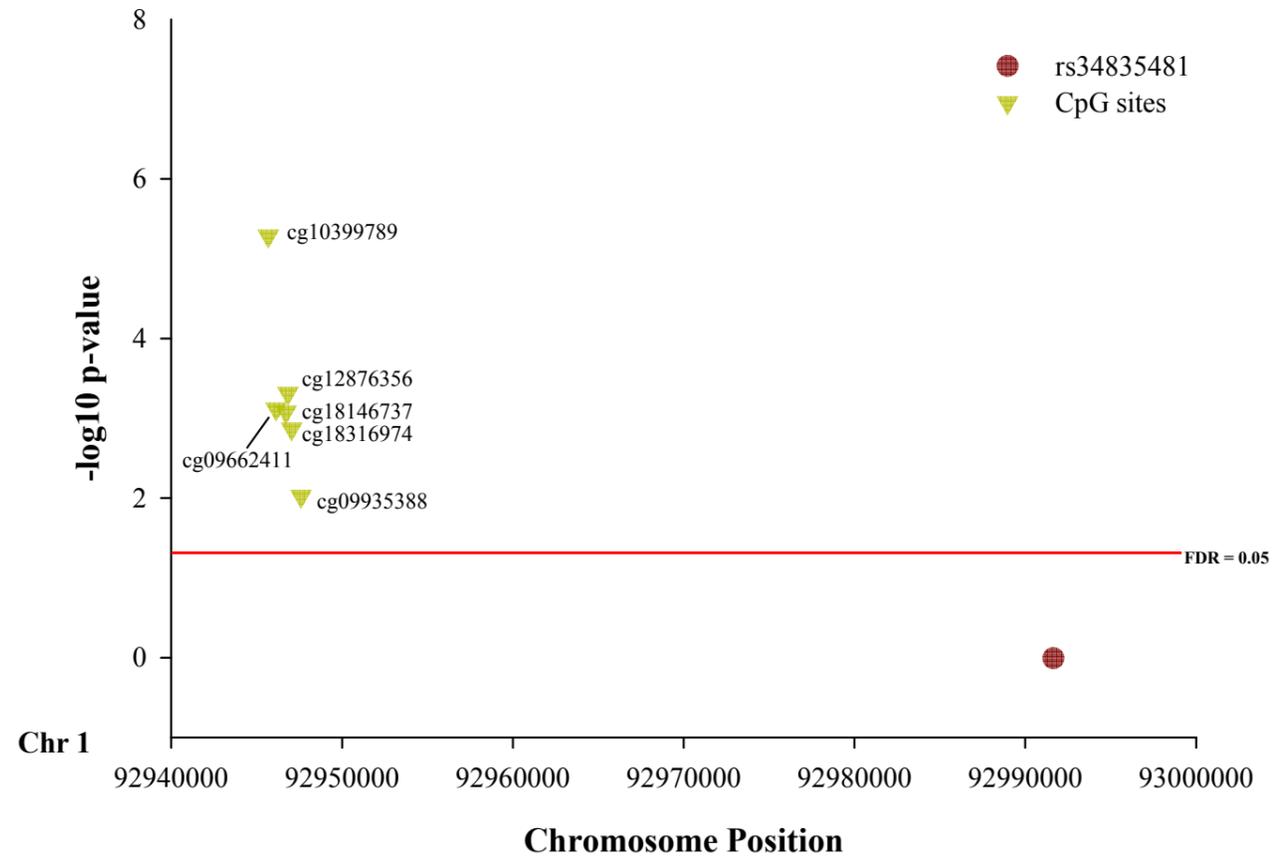


**Figure S2** Locations and distributions of methylation levels of 19 smoking-related CpG sites based on the three most frequently identified mQTLs (carrier/non-carrier) and smoking status in validation panel

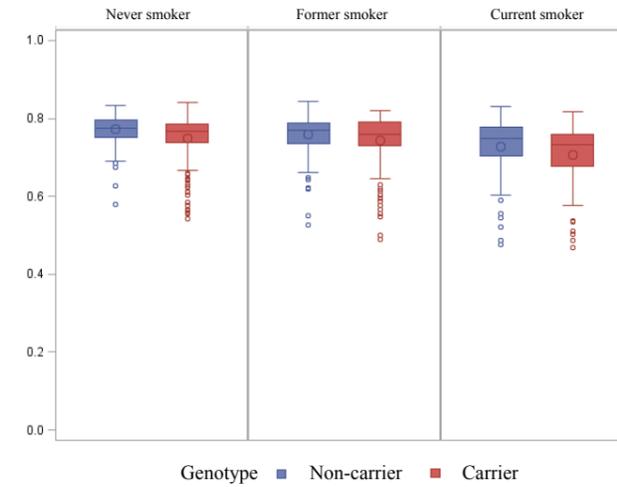


②

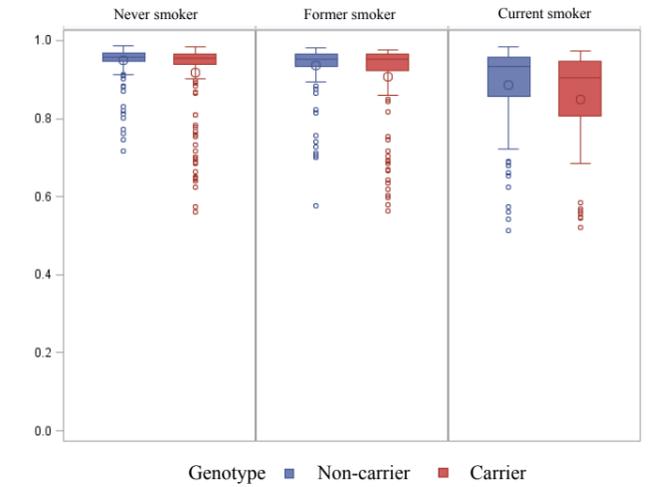
### Chromosome region of rs34835481



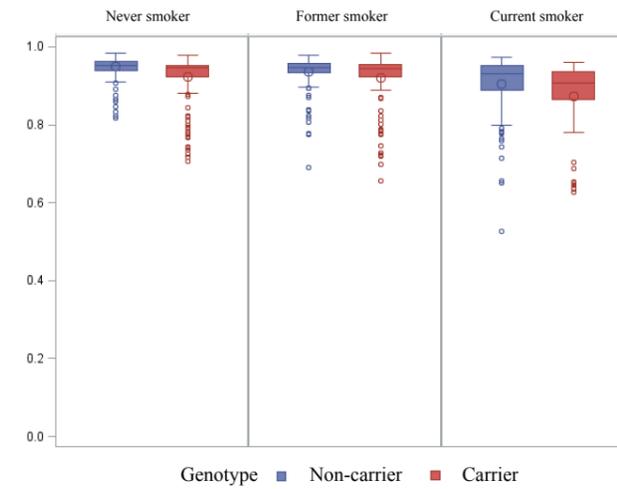
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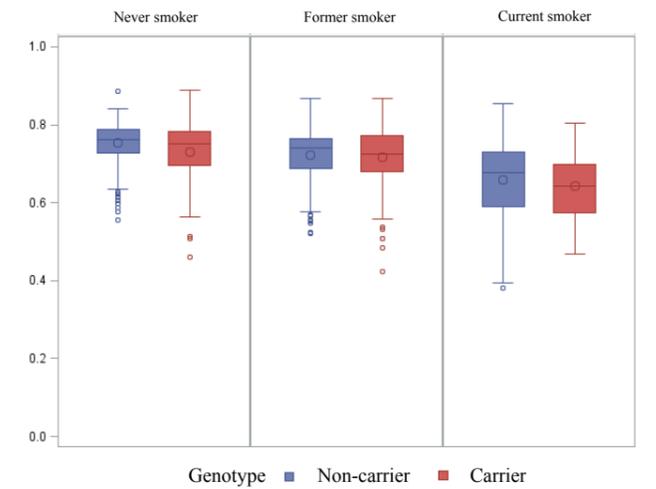
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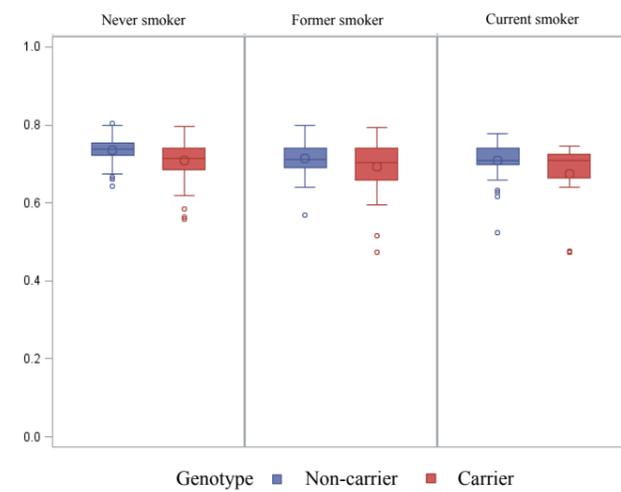
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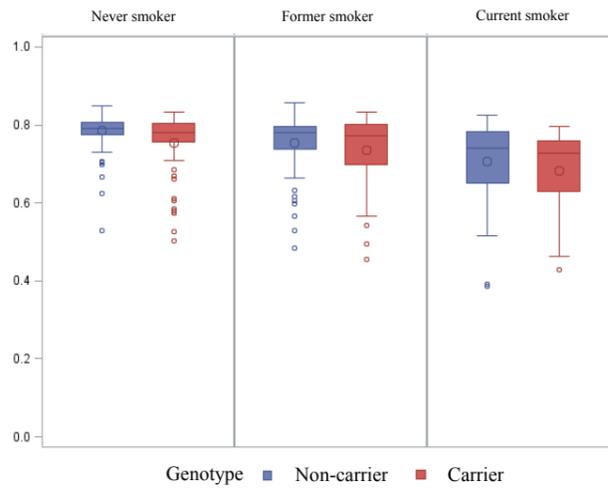
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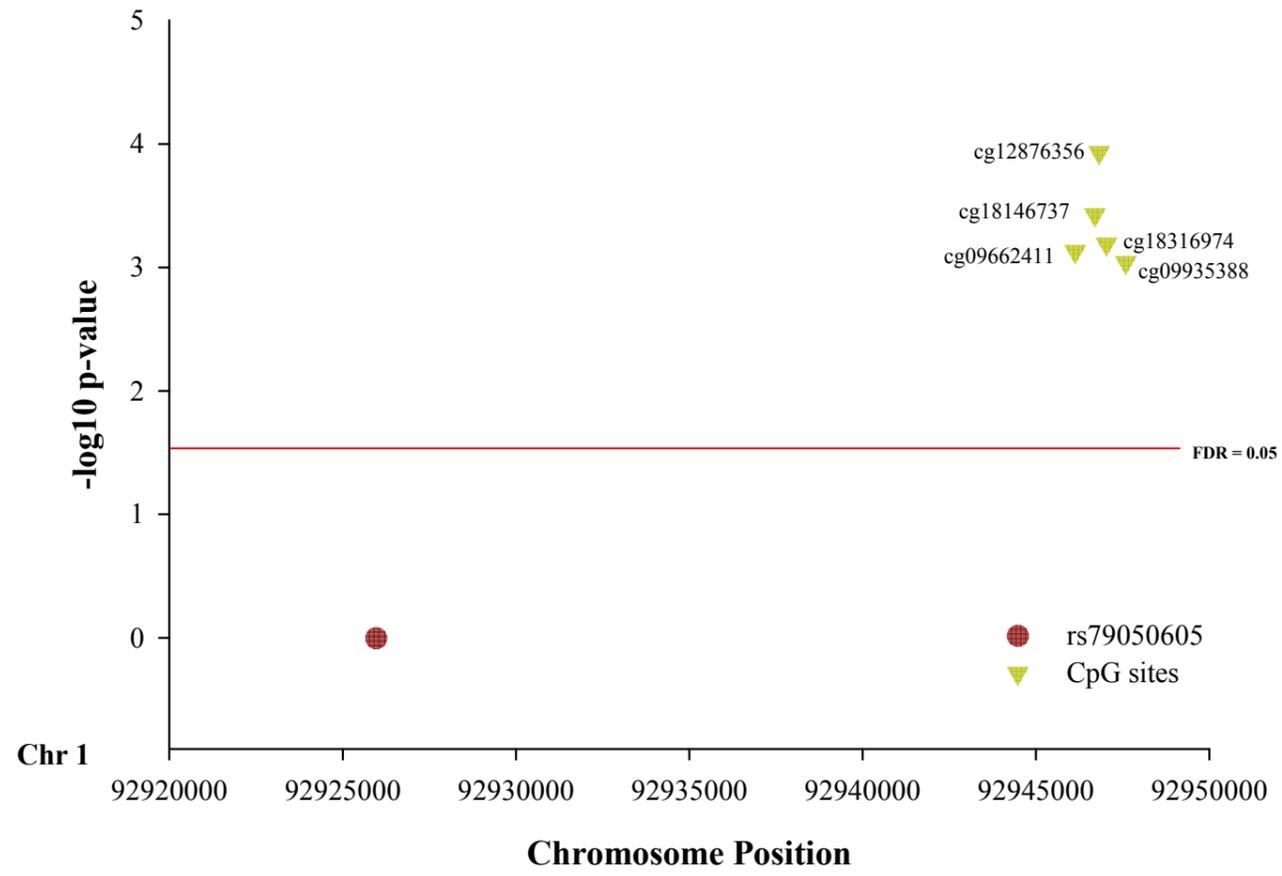


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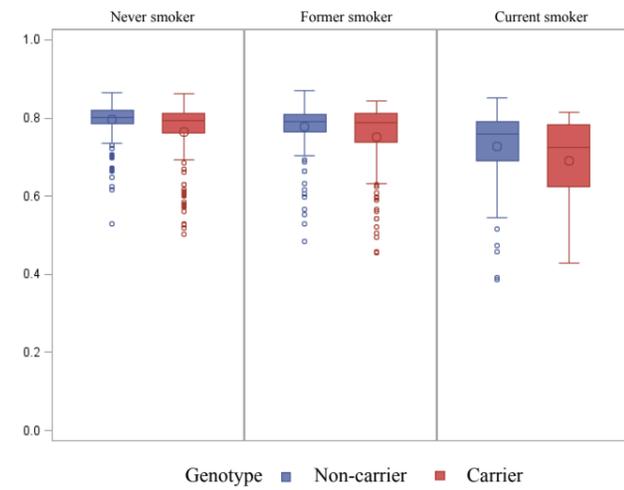


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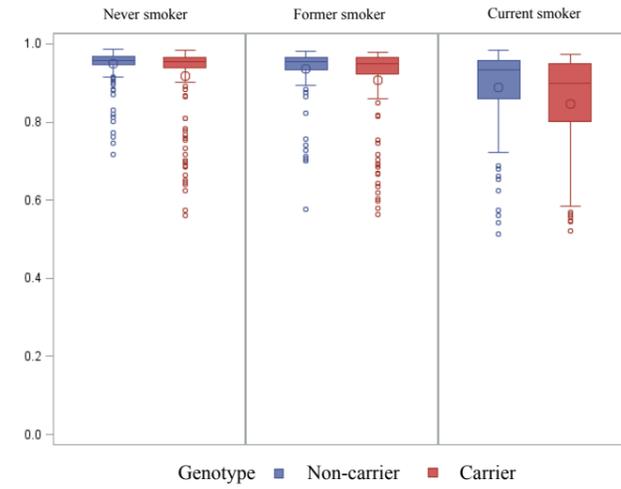
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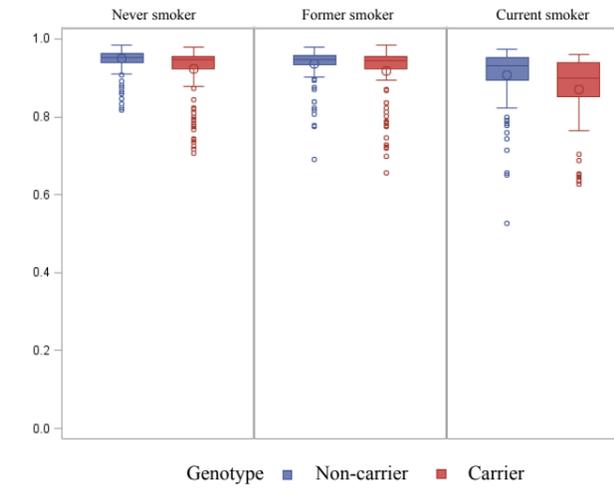
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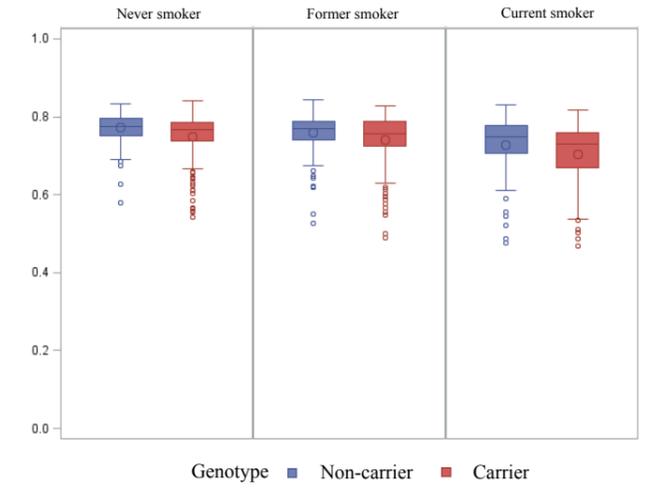
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cg18316974/rs79050605



cg09662411/rs79050605



cg09935388/rs79050605

