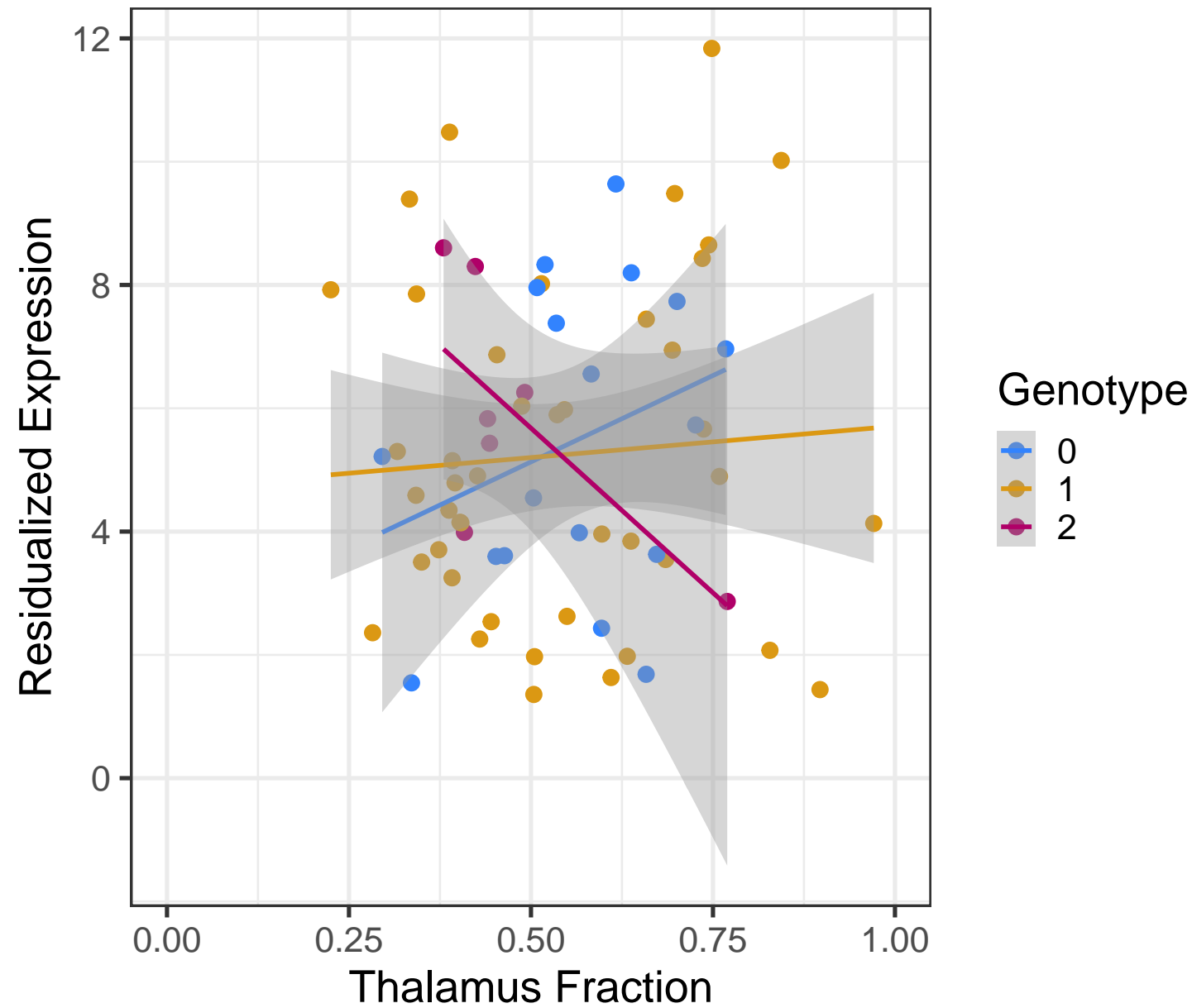
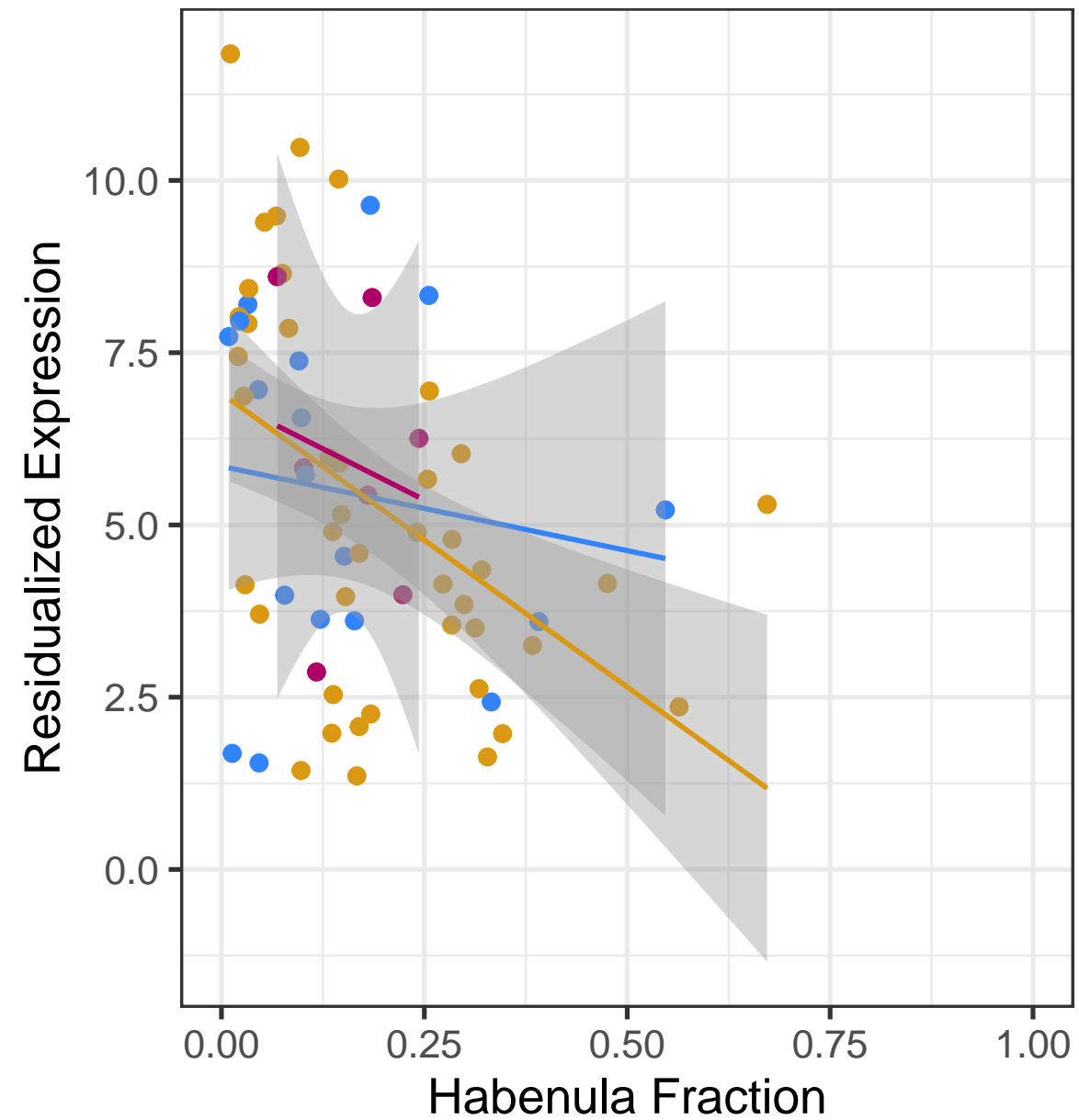
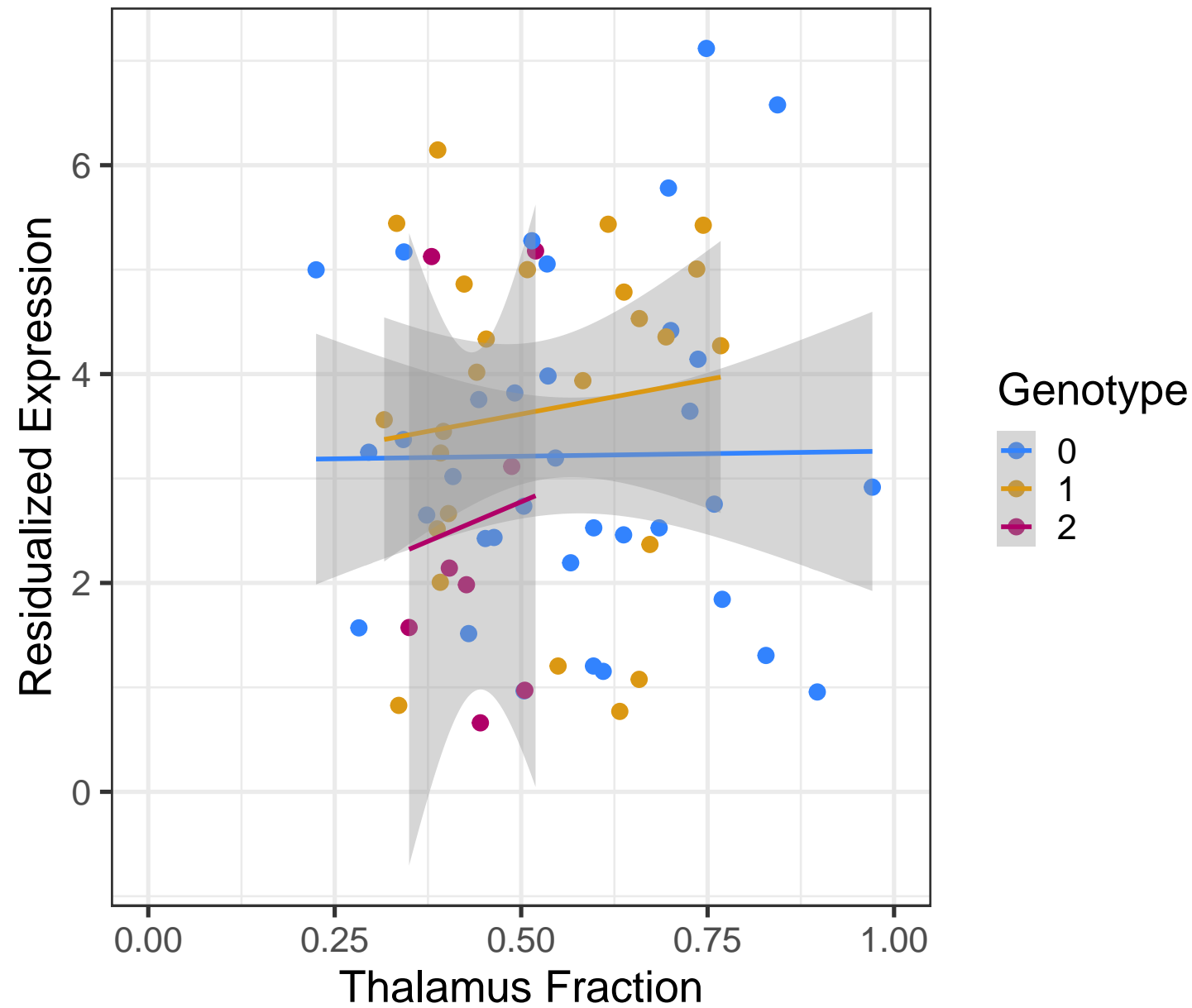
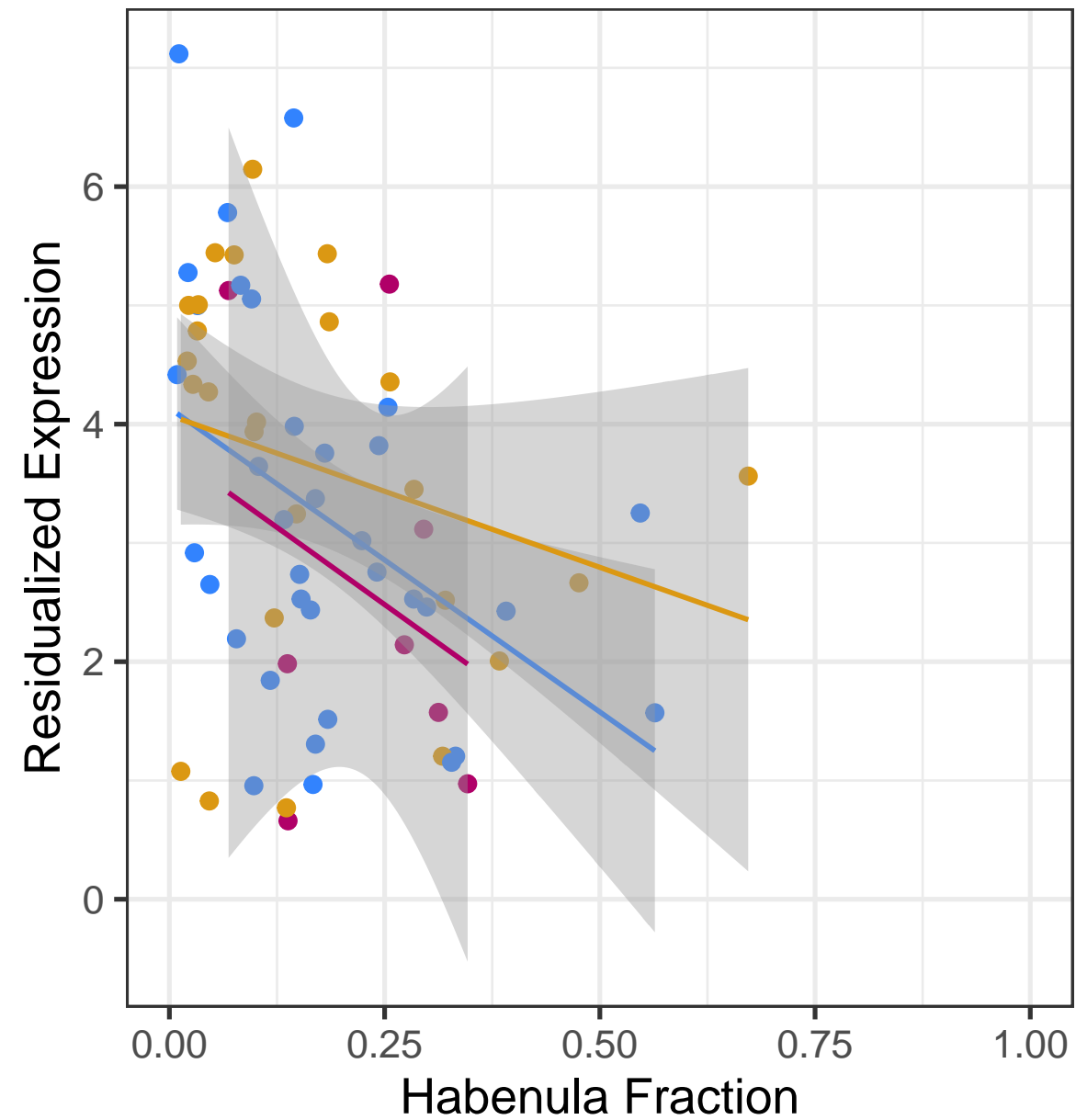


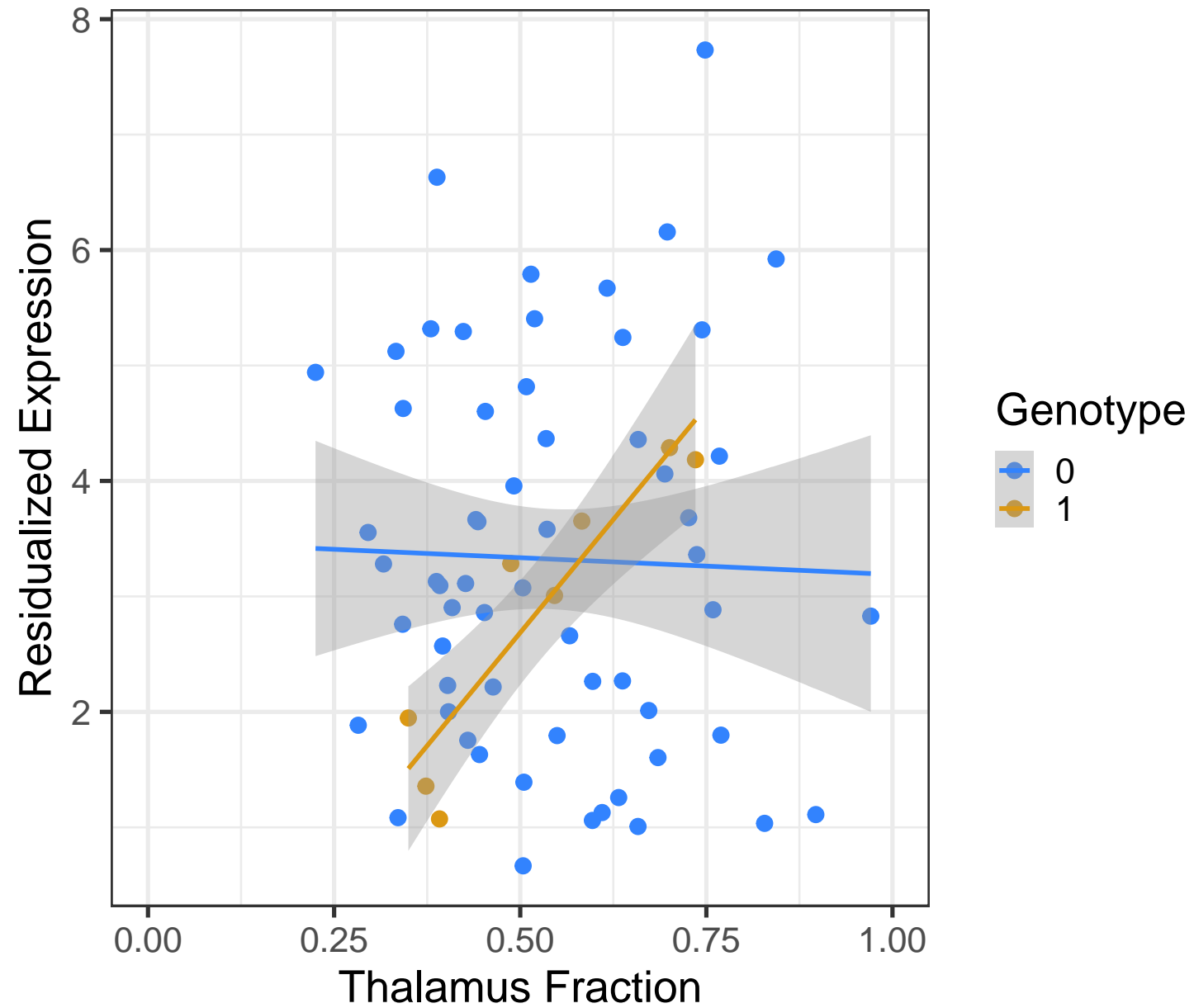
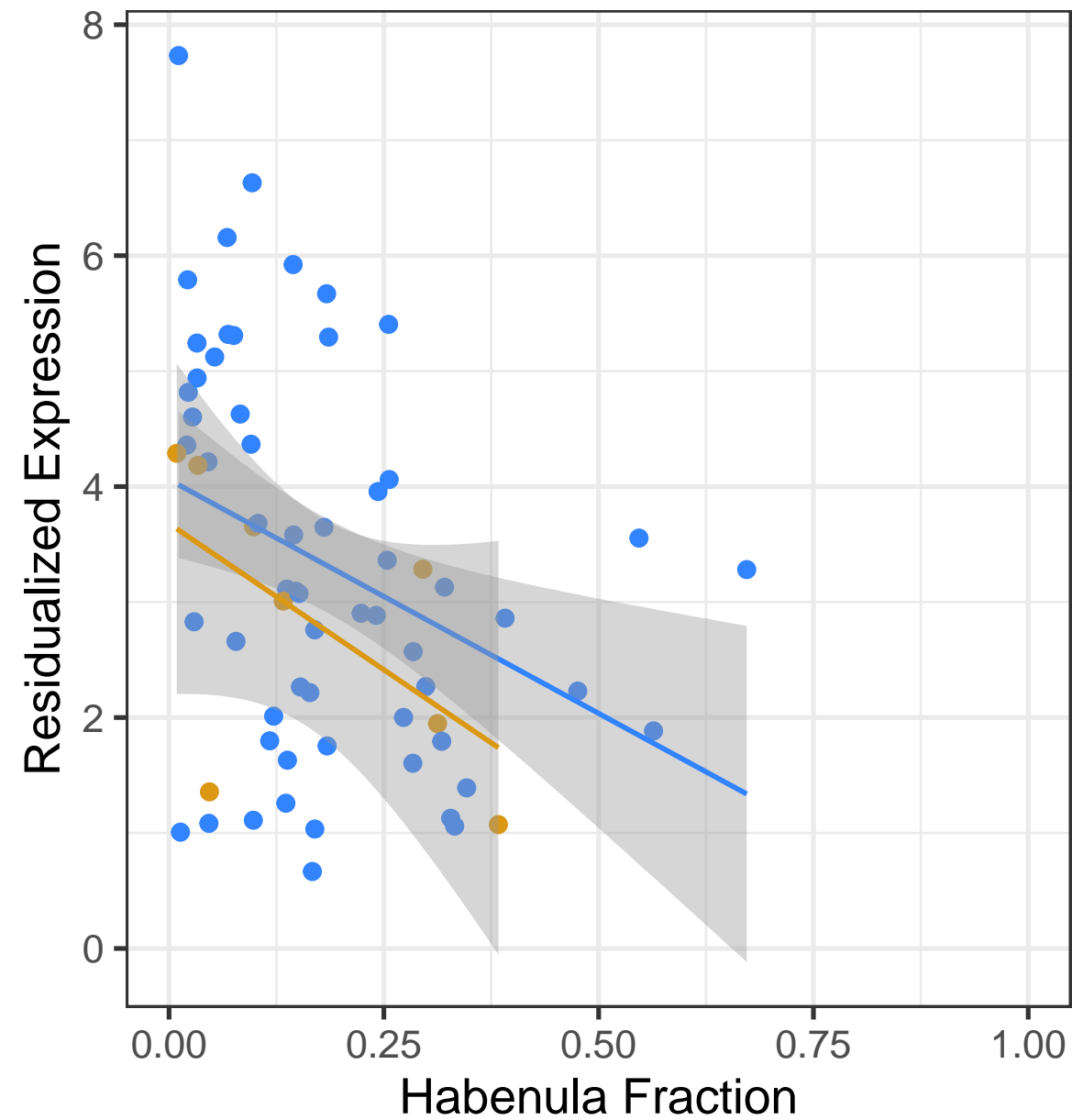
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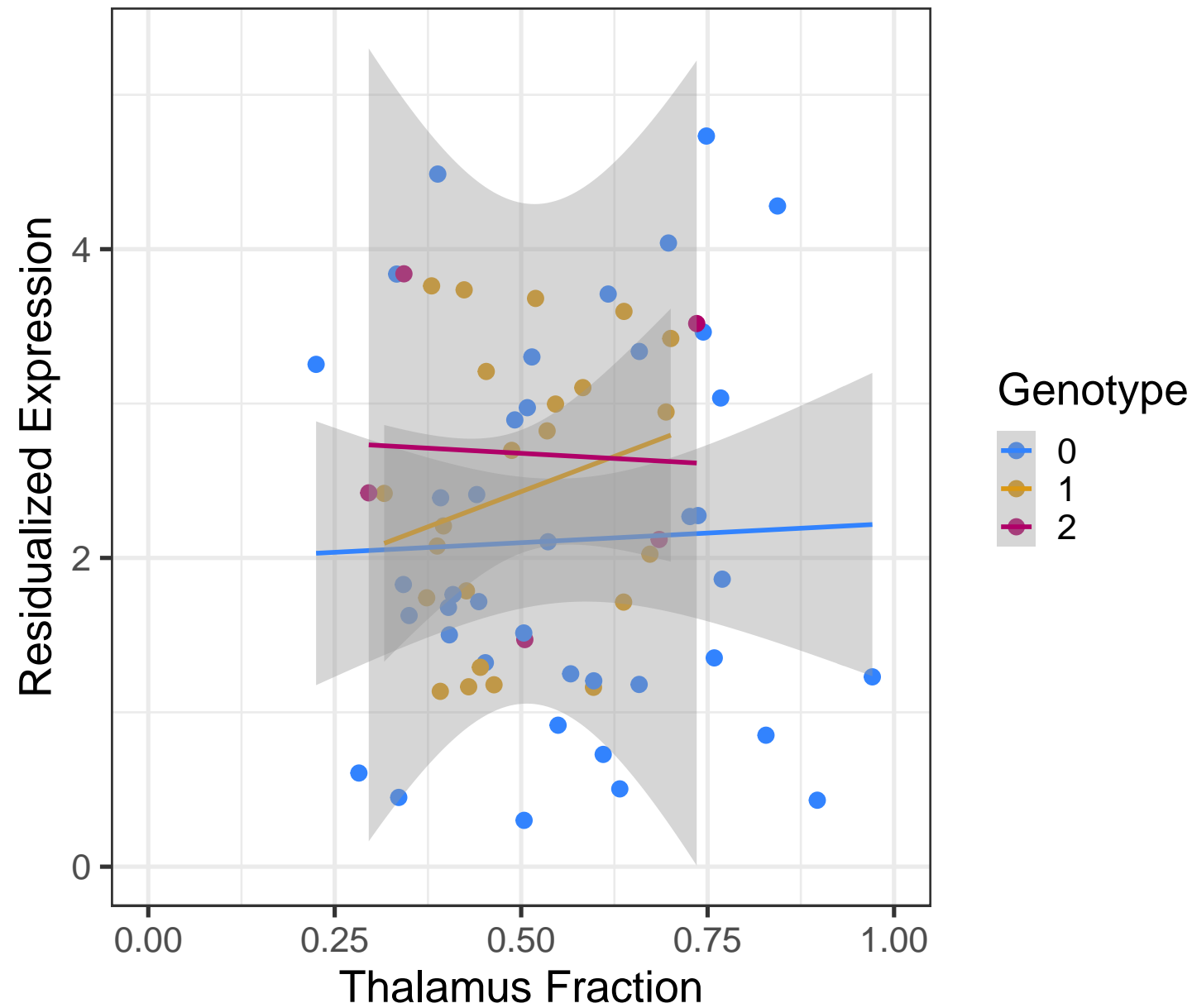
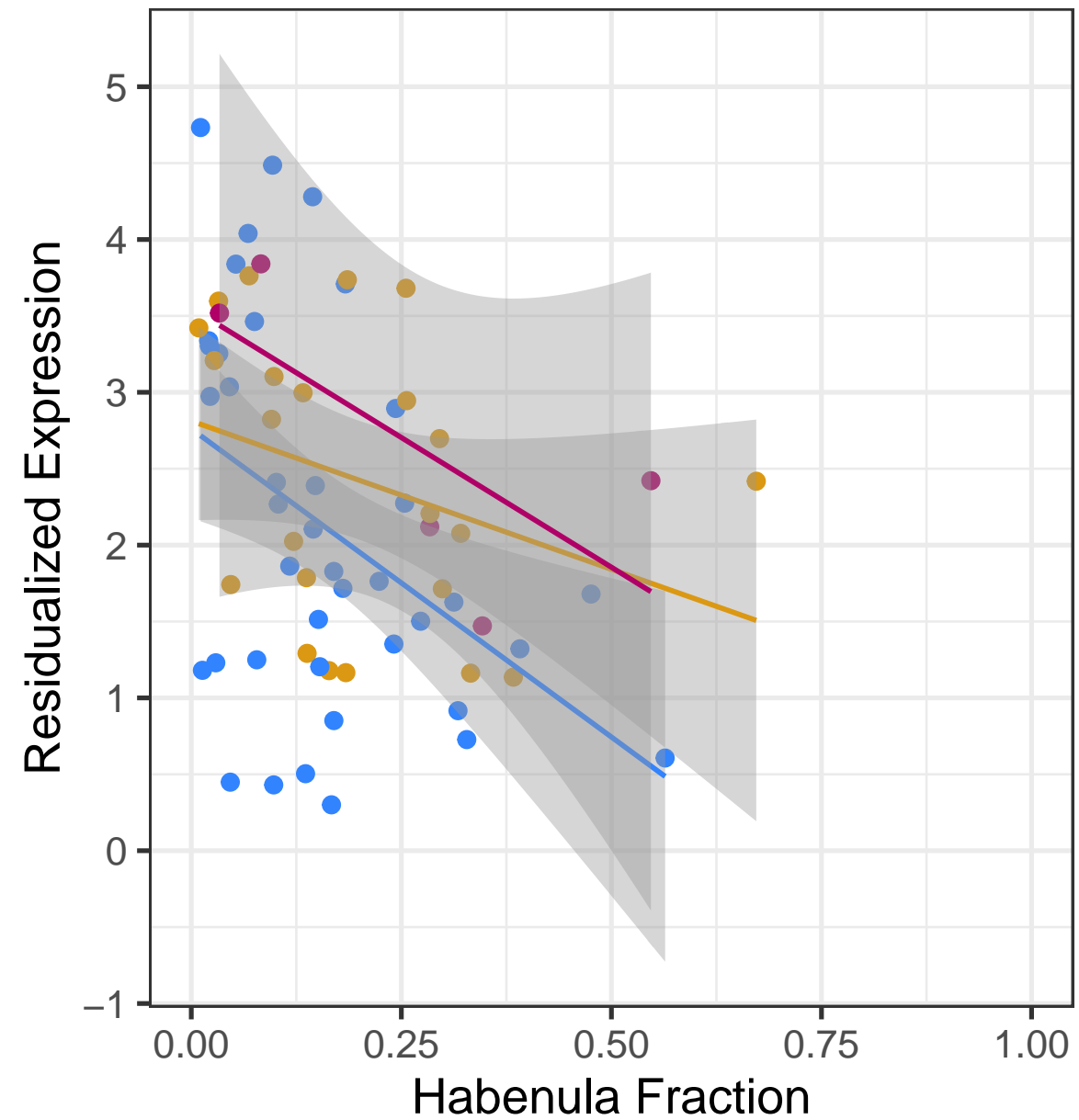
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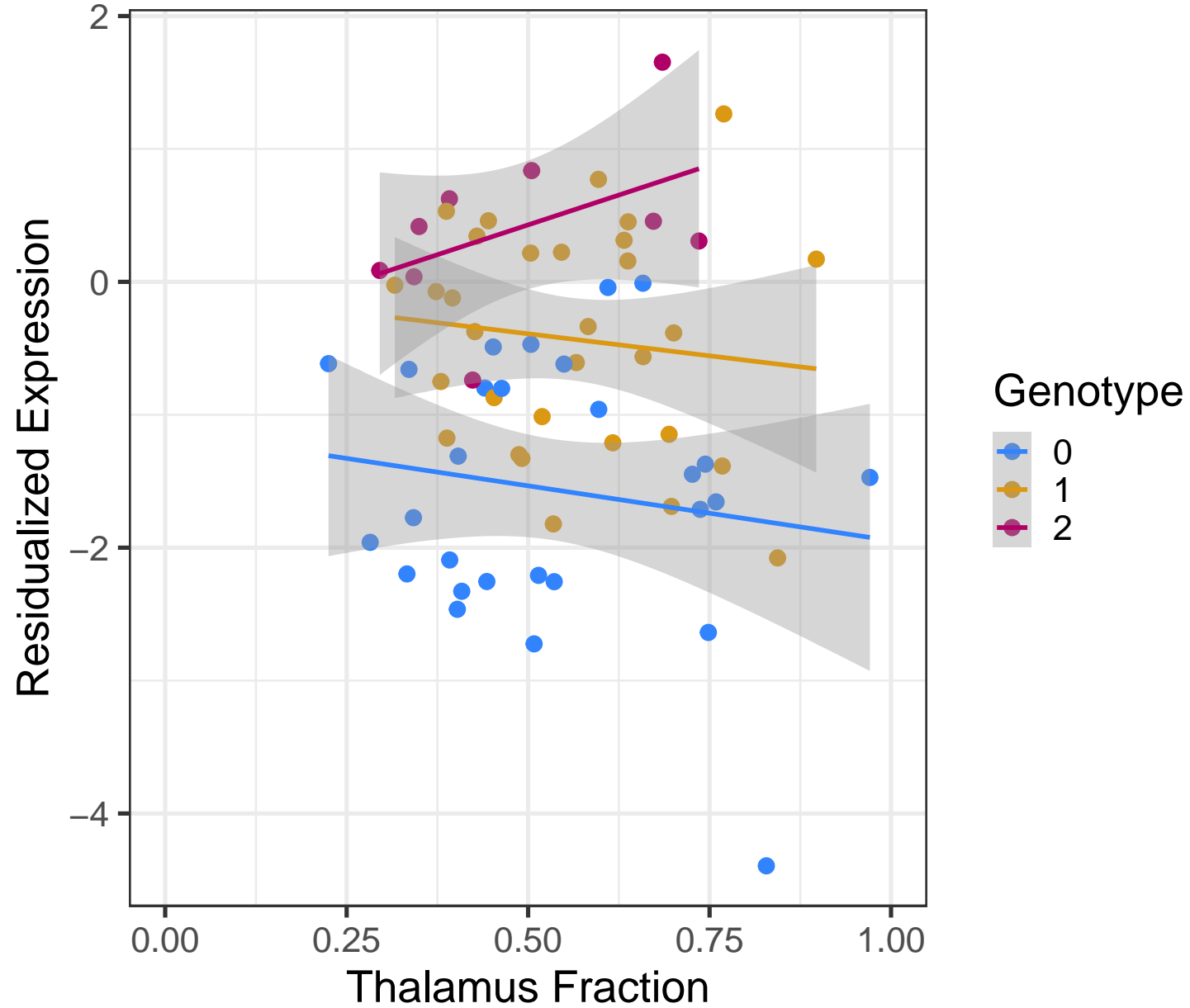
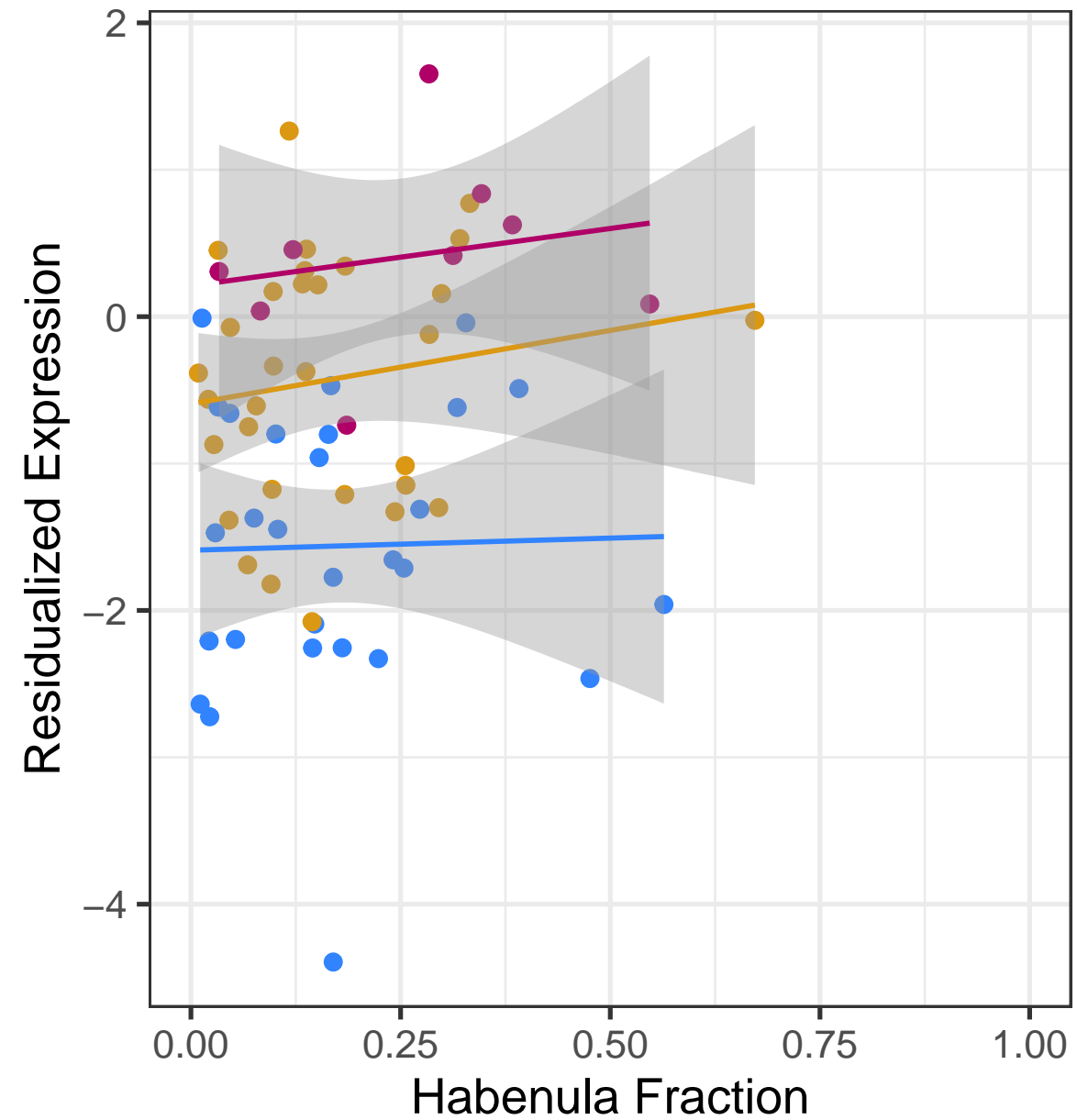
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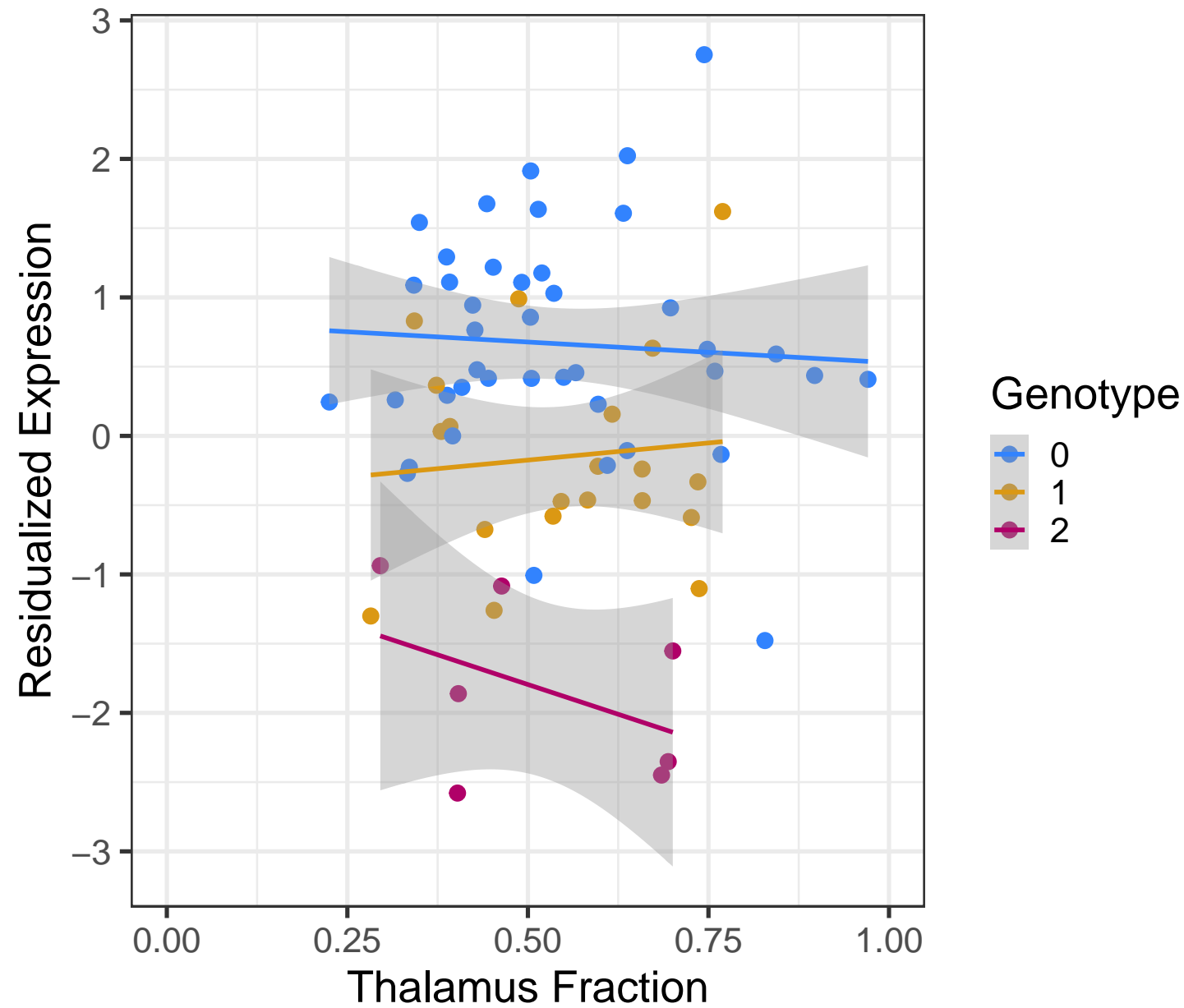
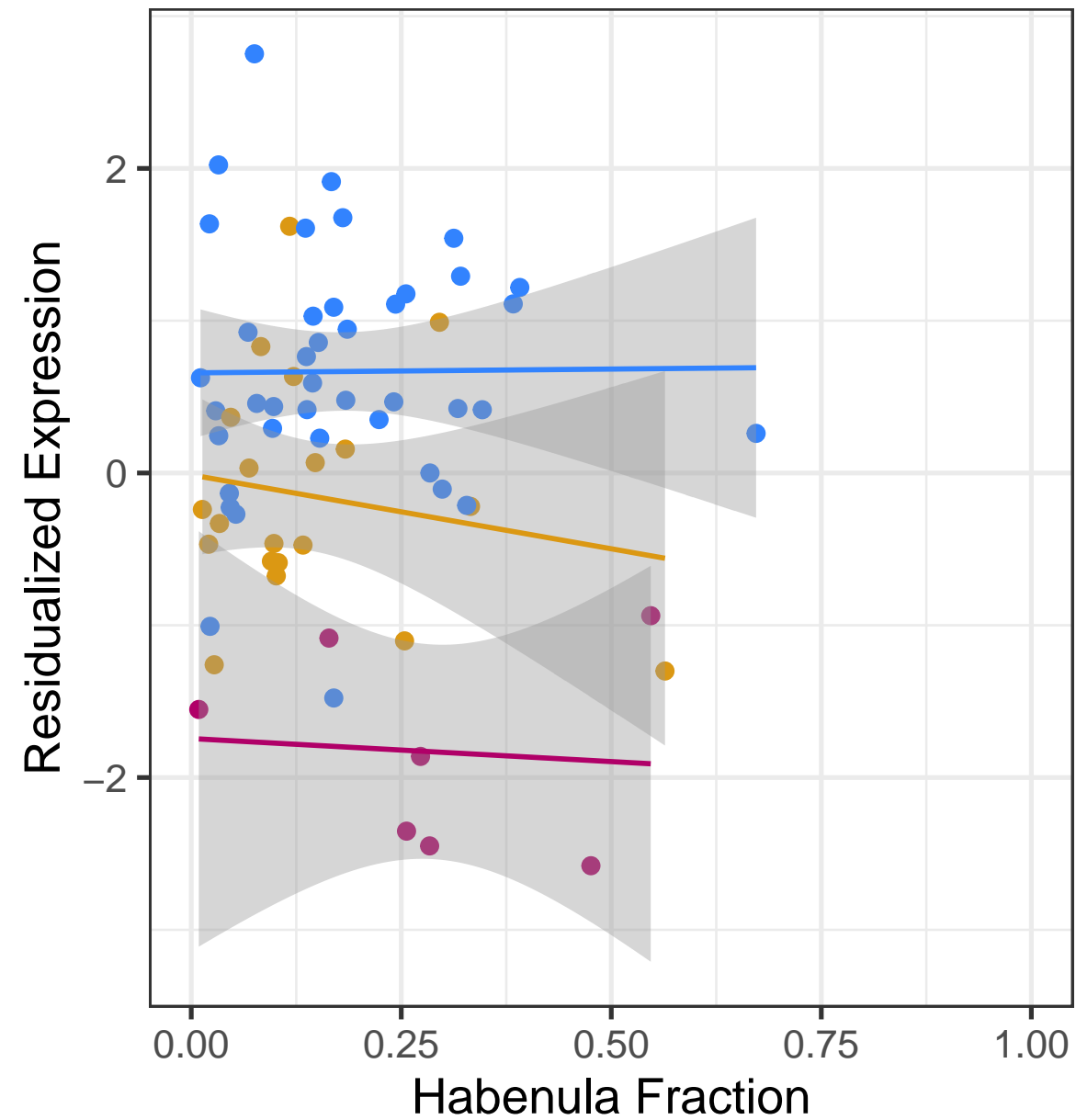
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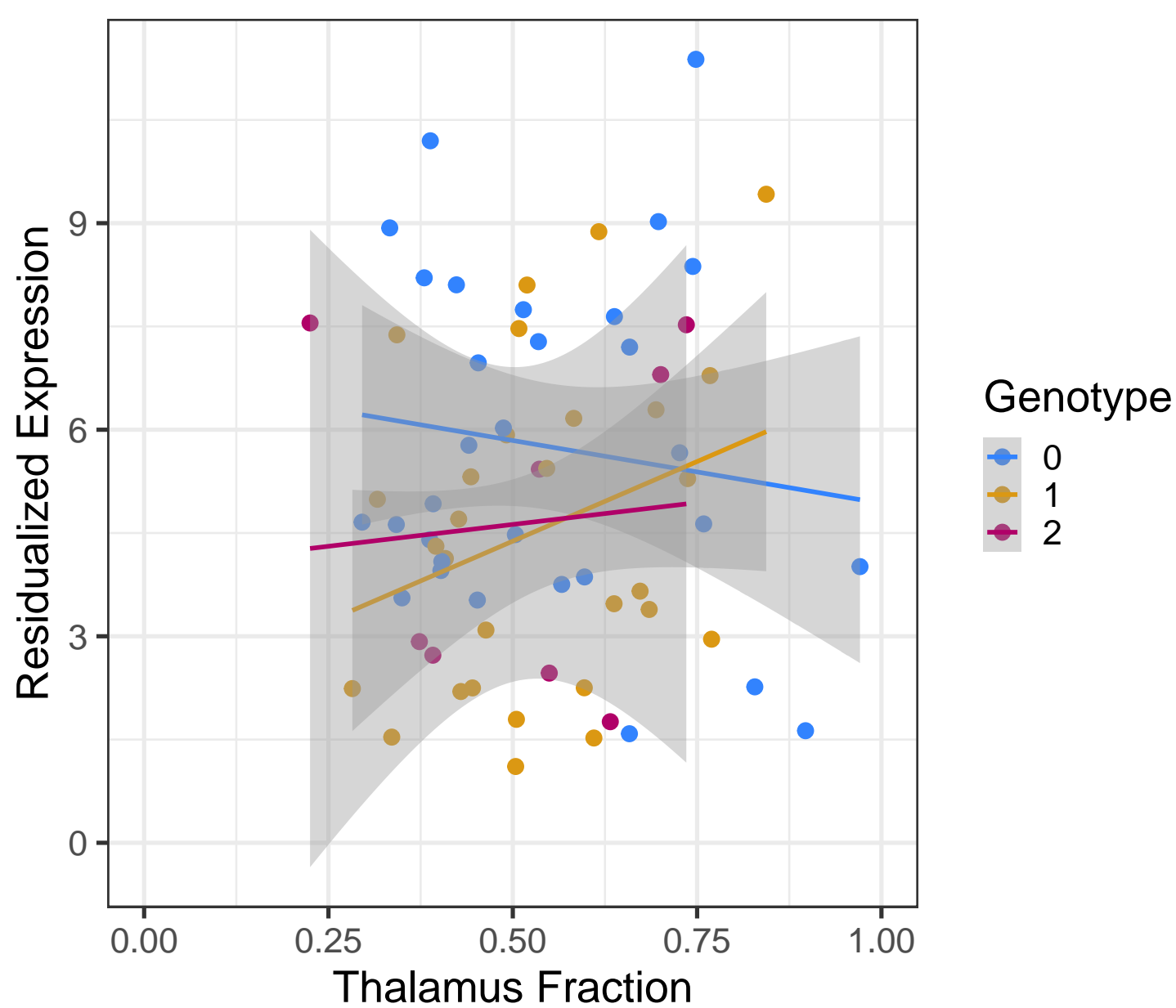
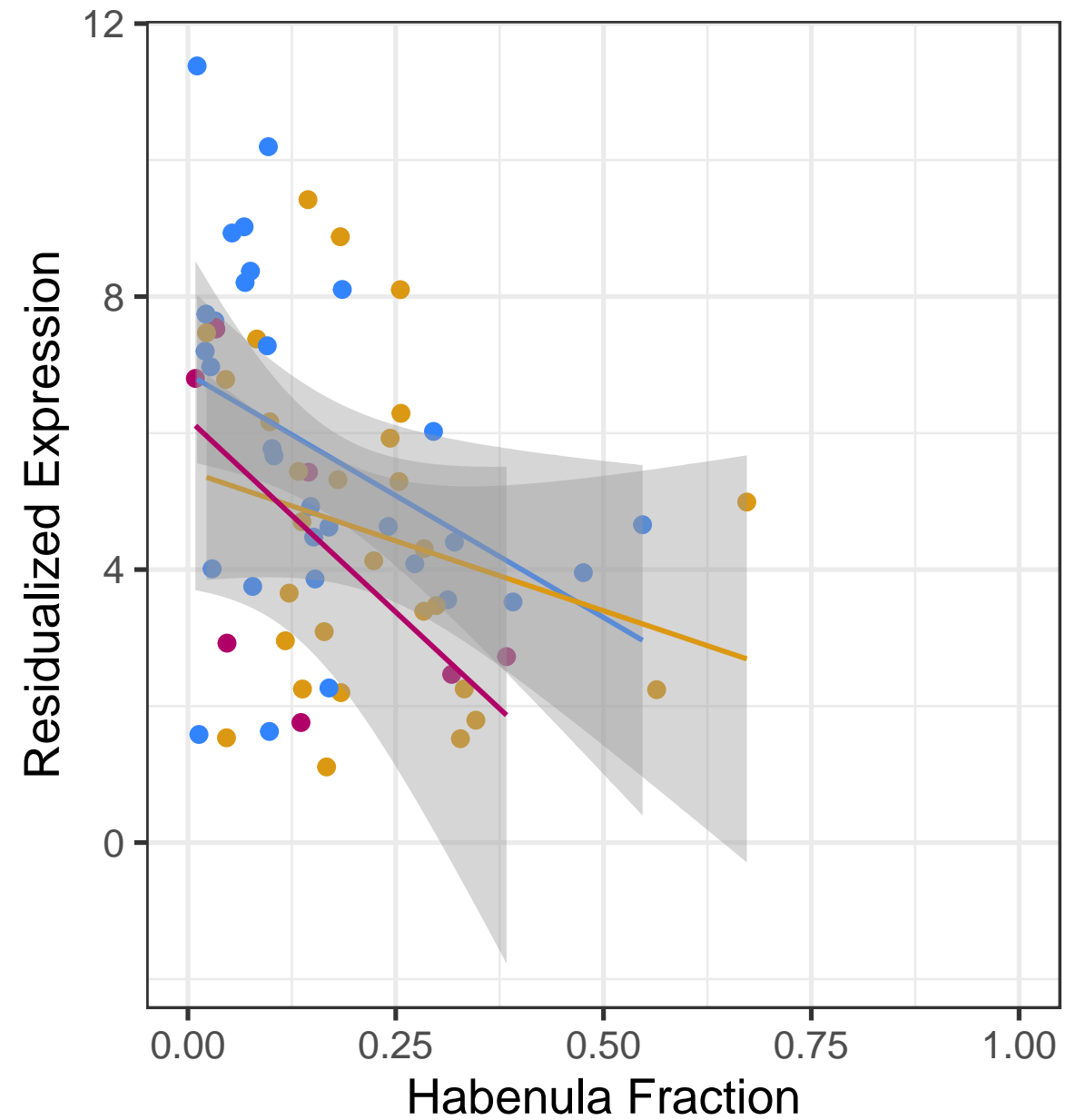
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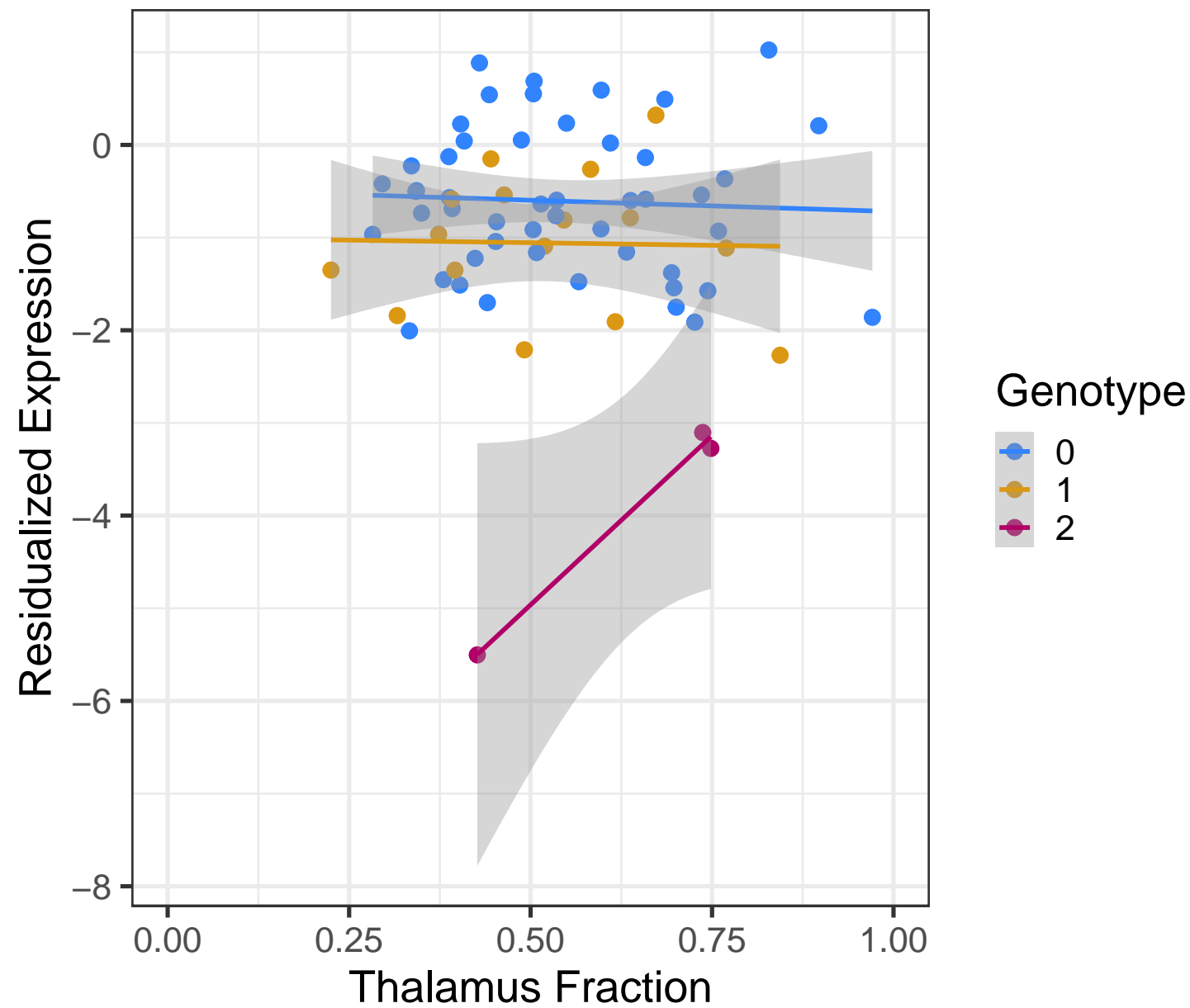
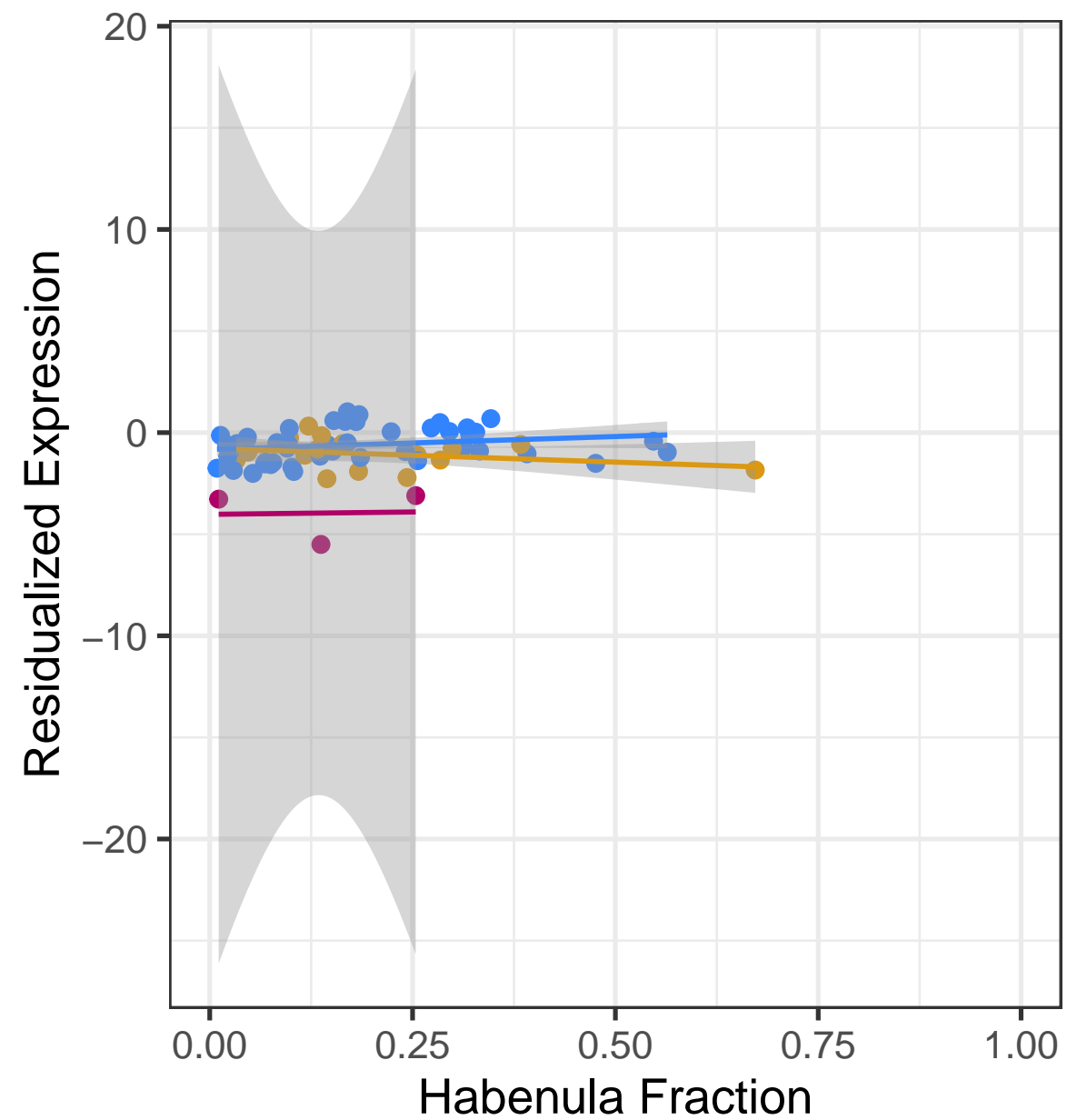
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VARs2: chr6:30934095:C:G

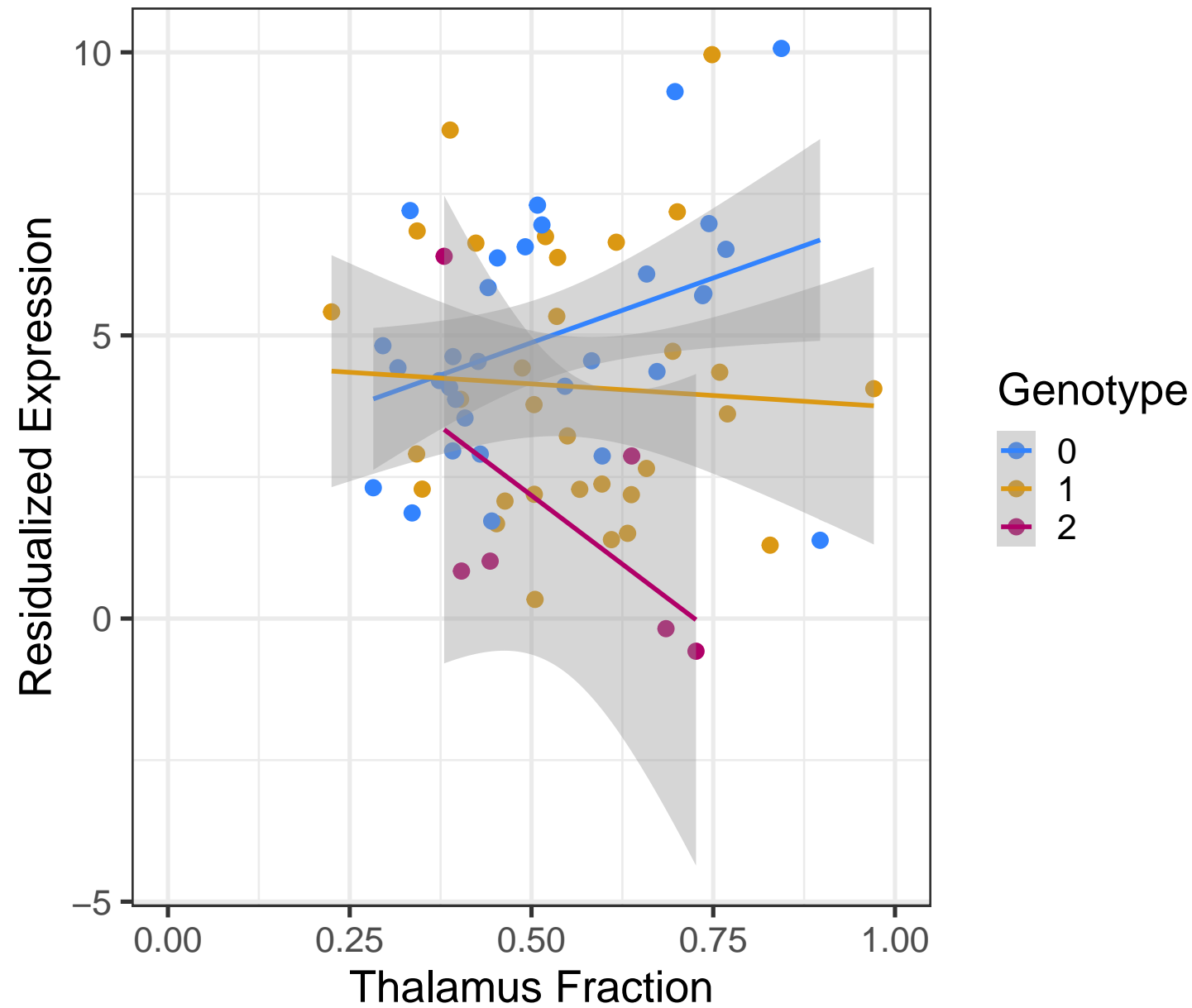
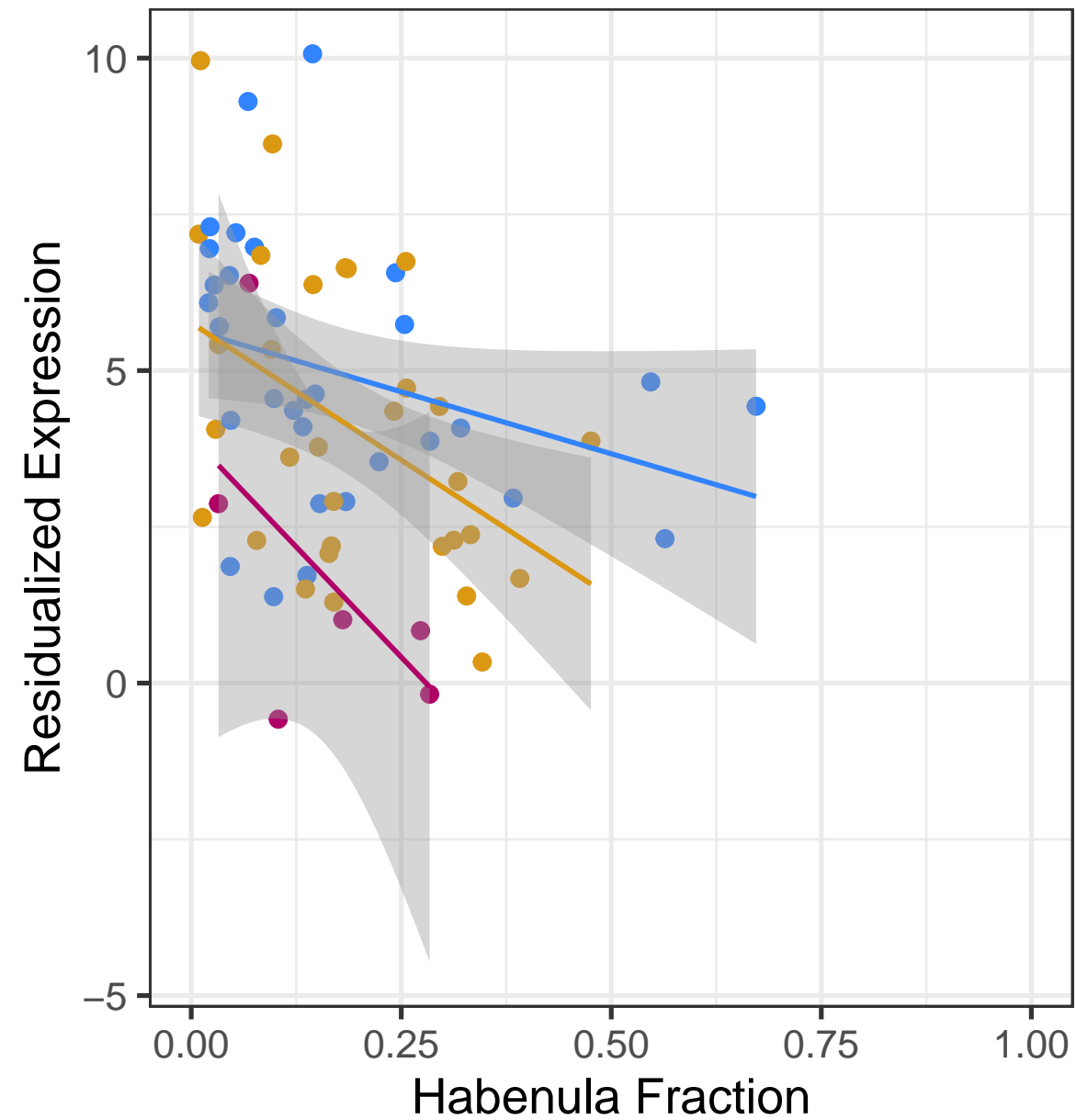


PSORS1C1: chr6:31025536:C:T

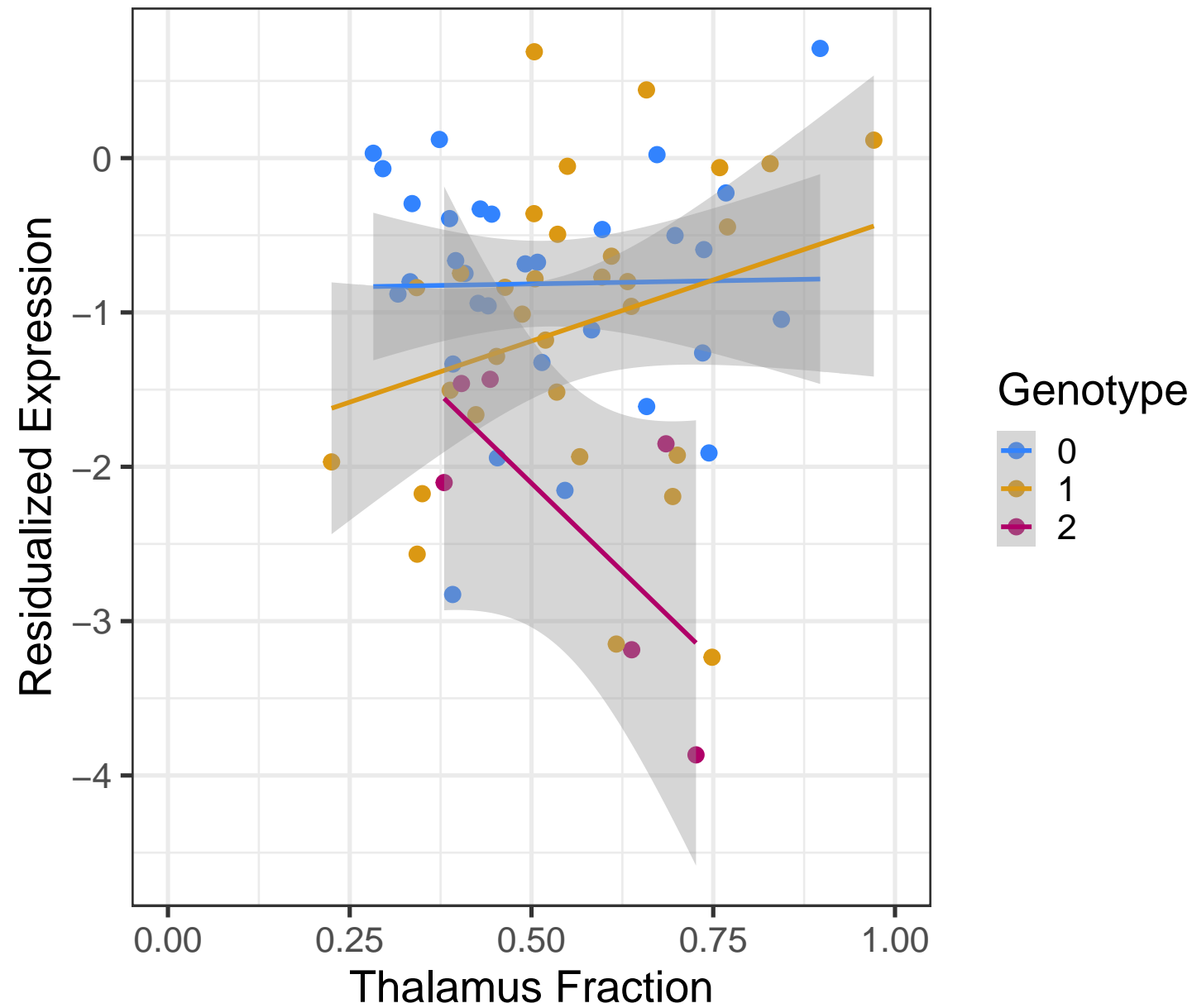
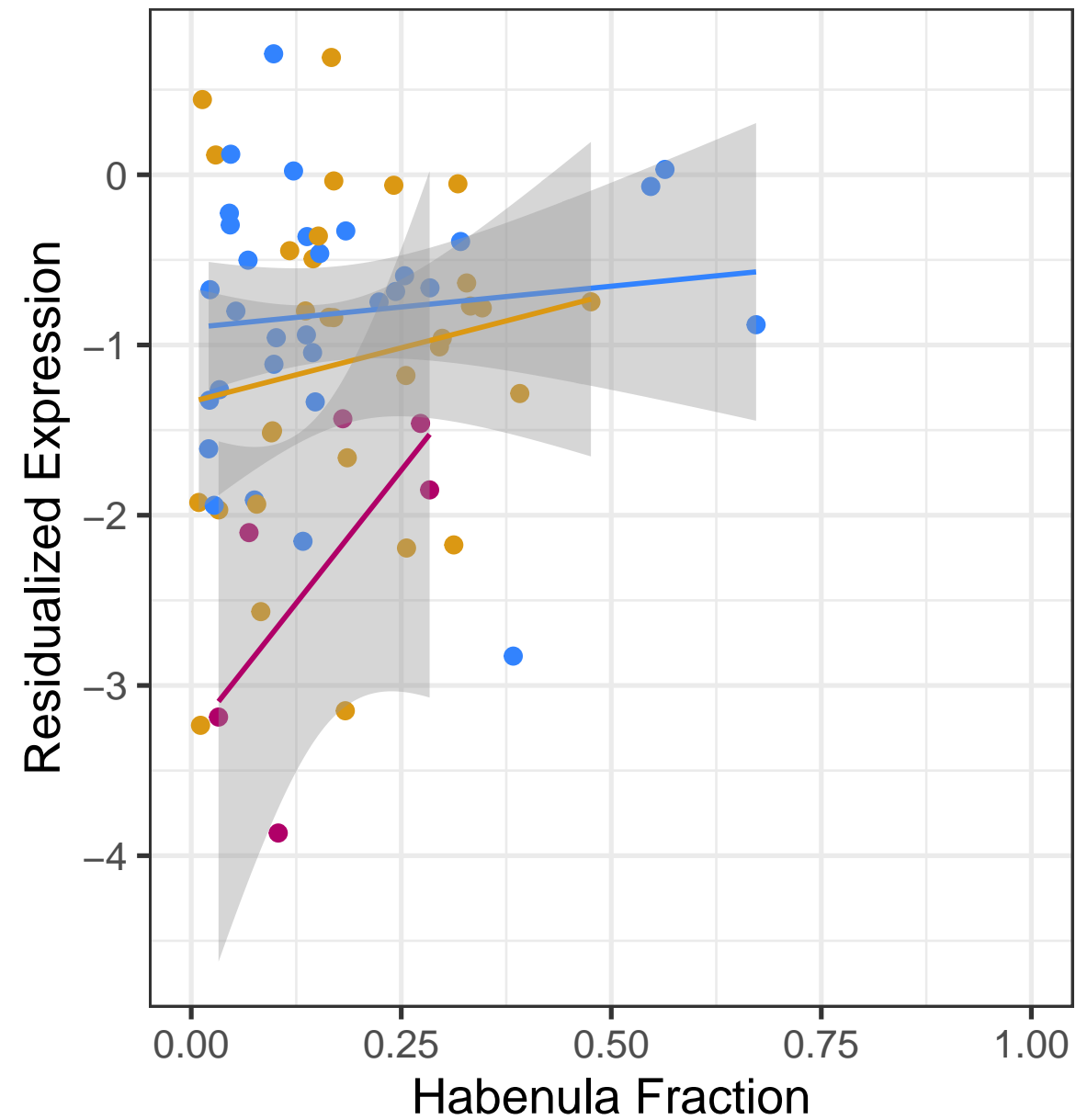




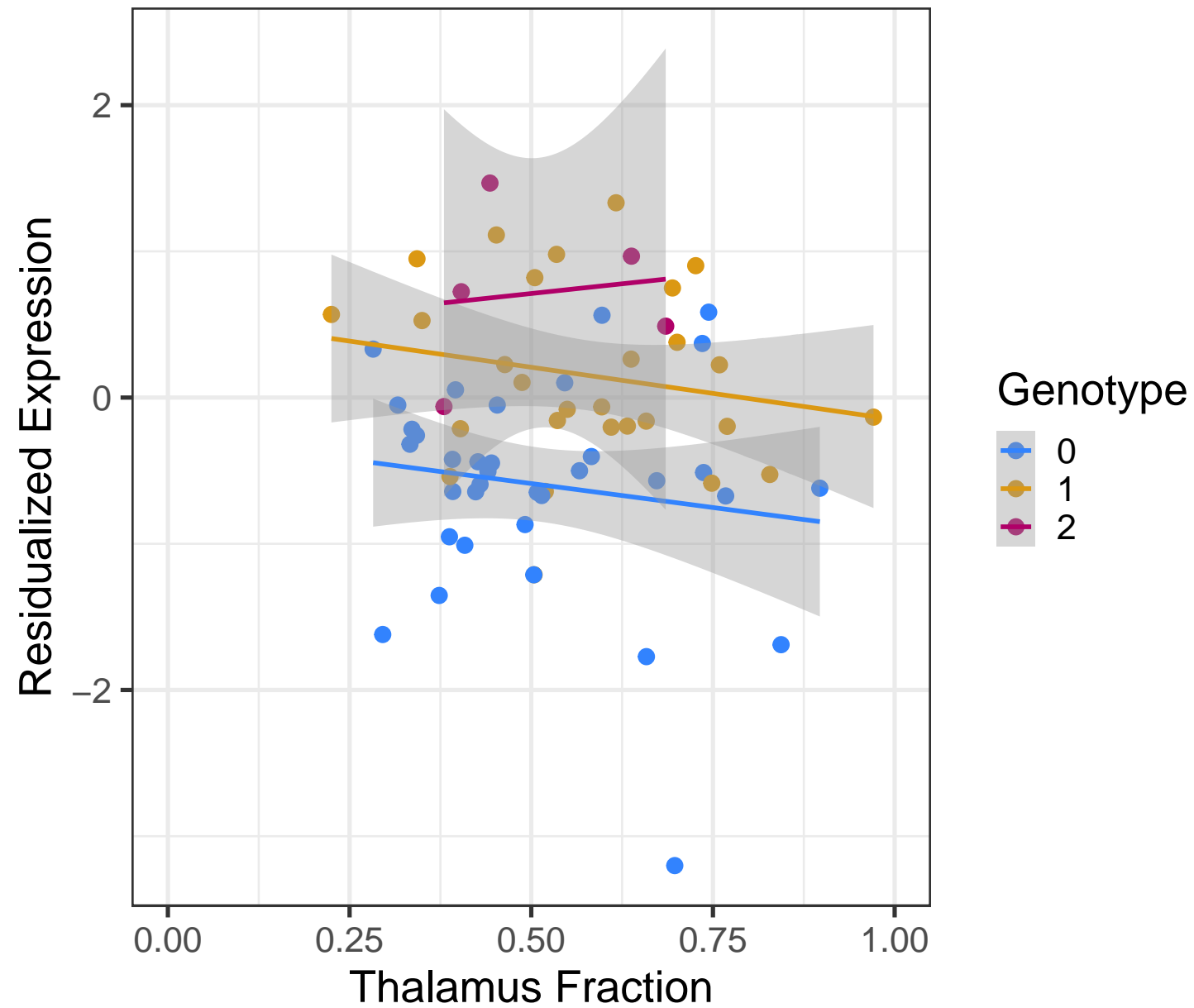
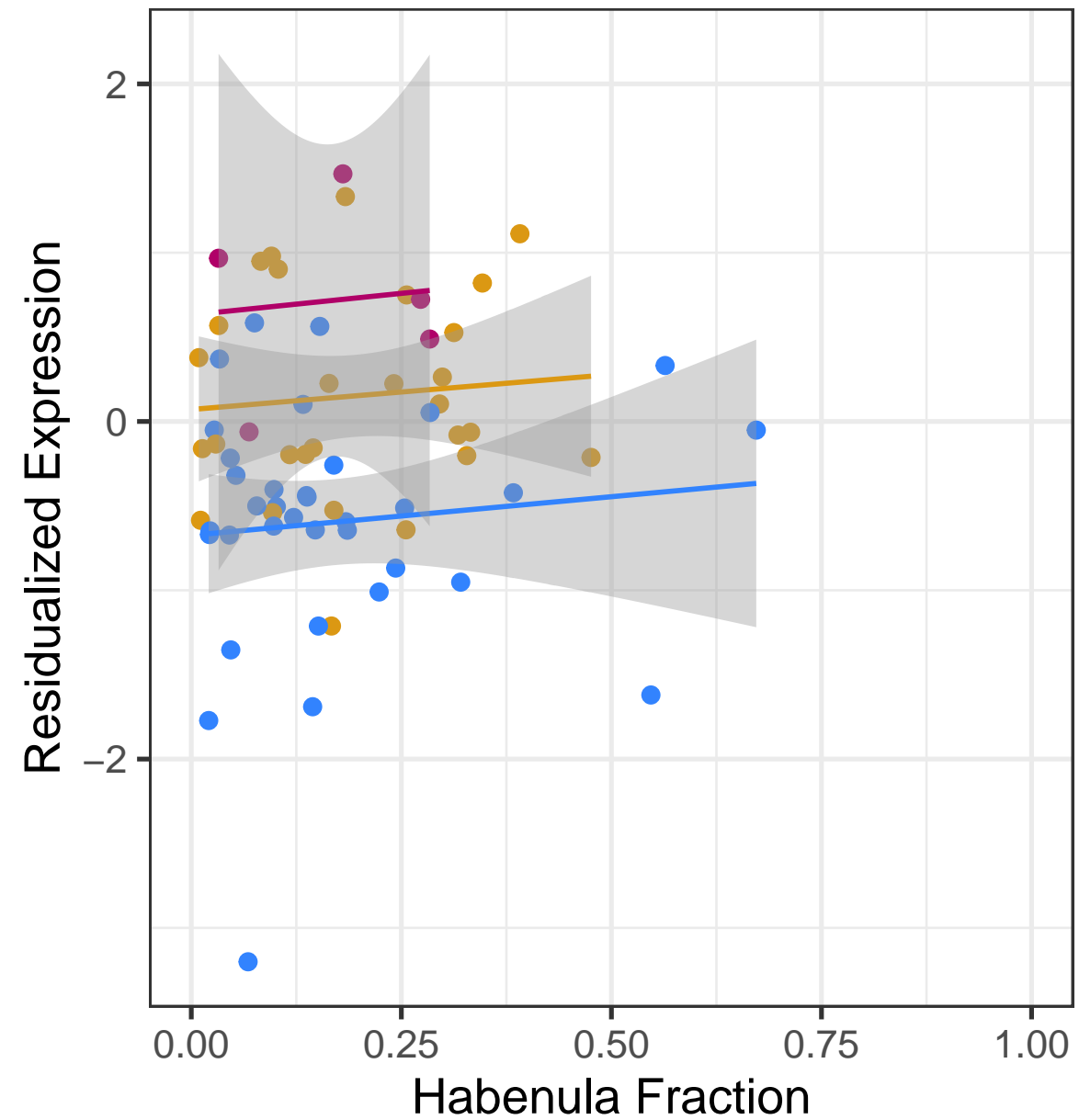
LRRC37A4P: chr17:45910839:A:G



DND1P1: chr17:45910839:A:G



KANSL1-AS1: chr17:46278268:C:A



LRRC37A: chr17:46002201:C:T

