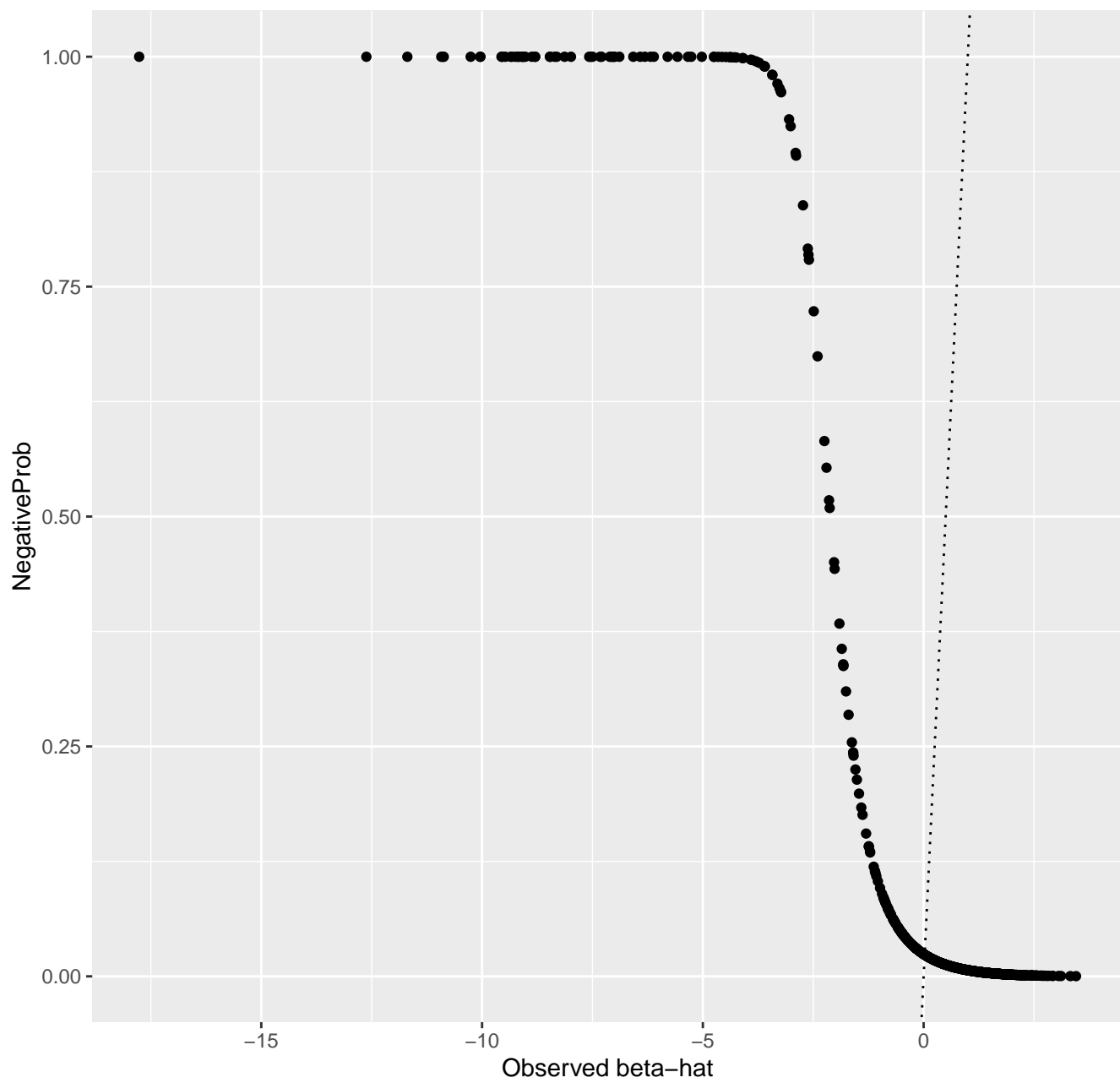
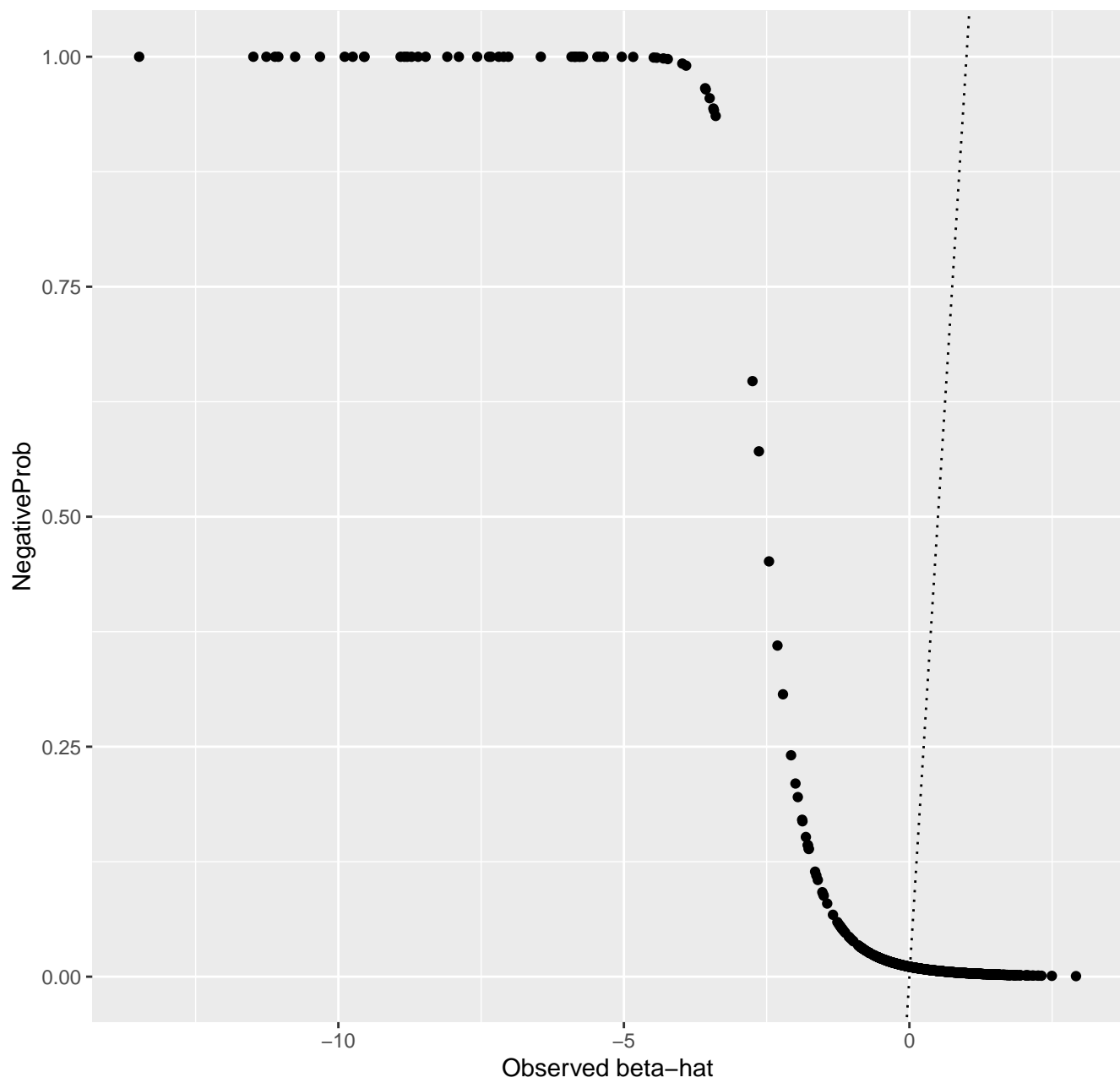


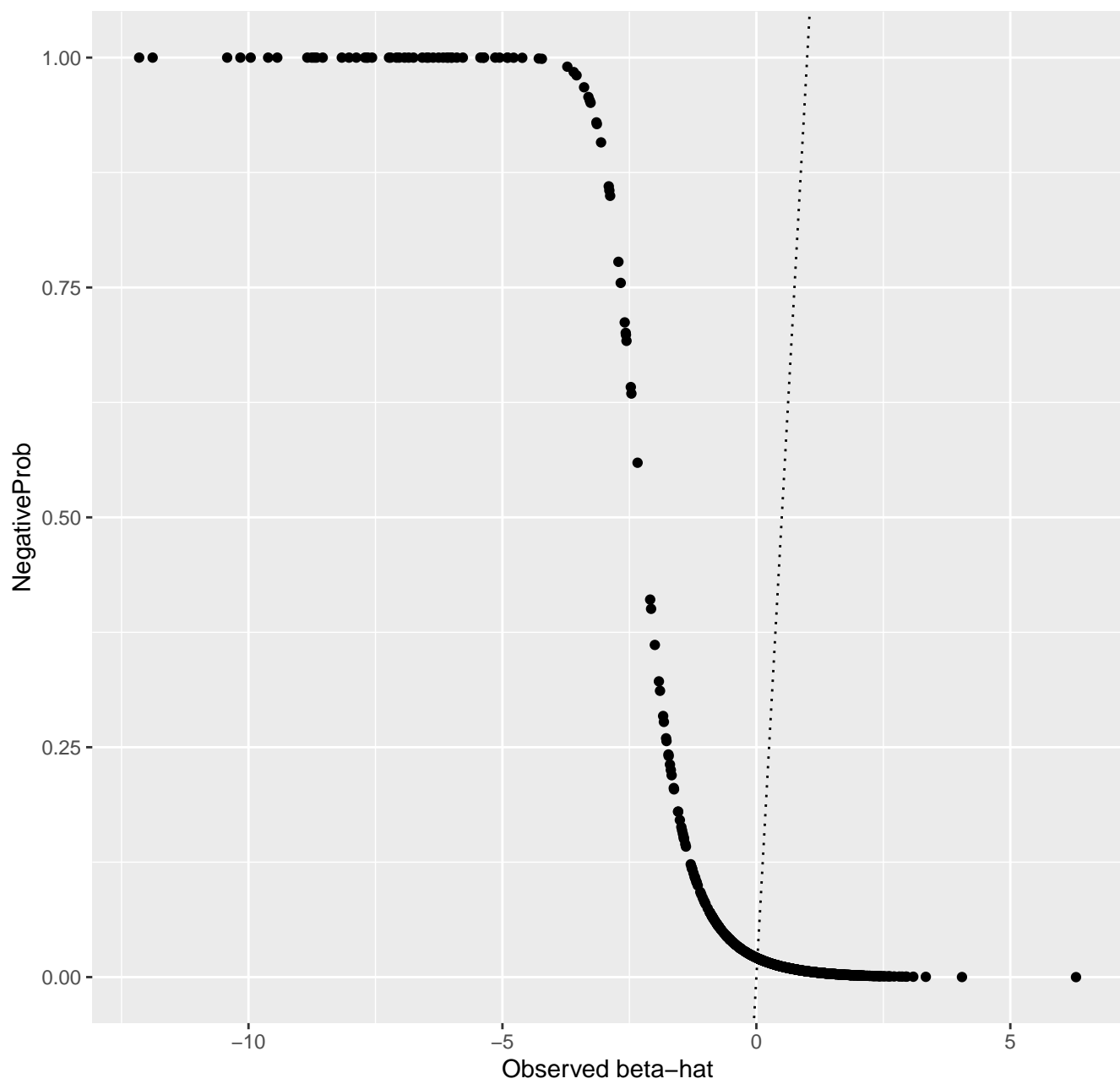
HNF1B_(6928) negative



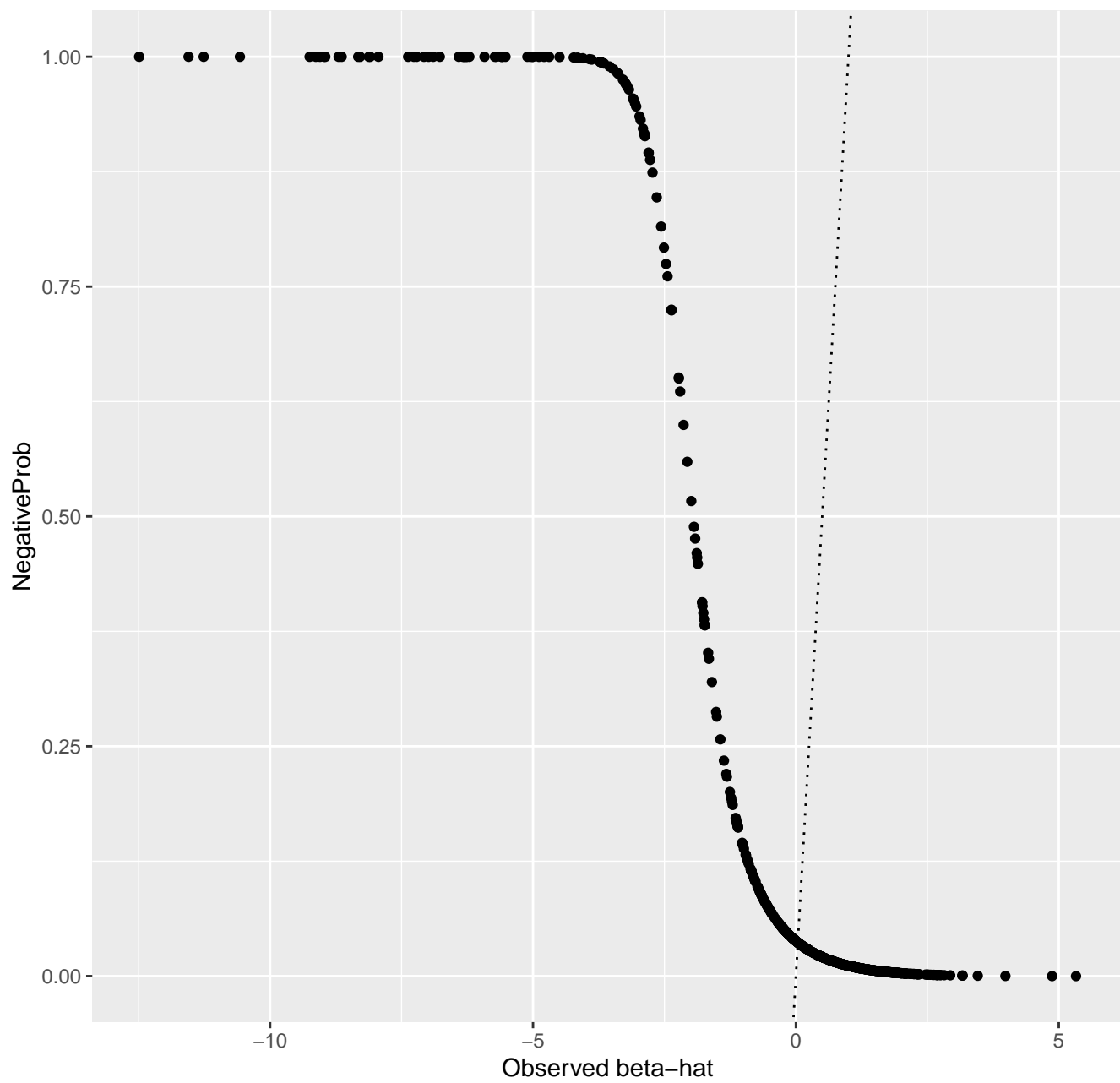
PAX8_(7849) negative



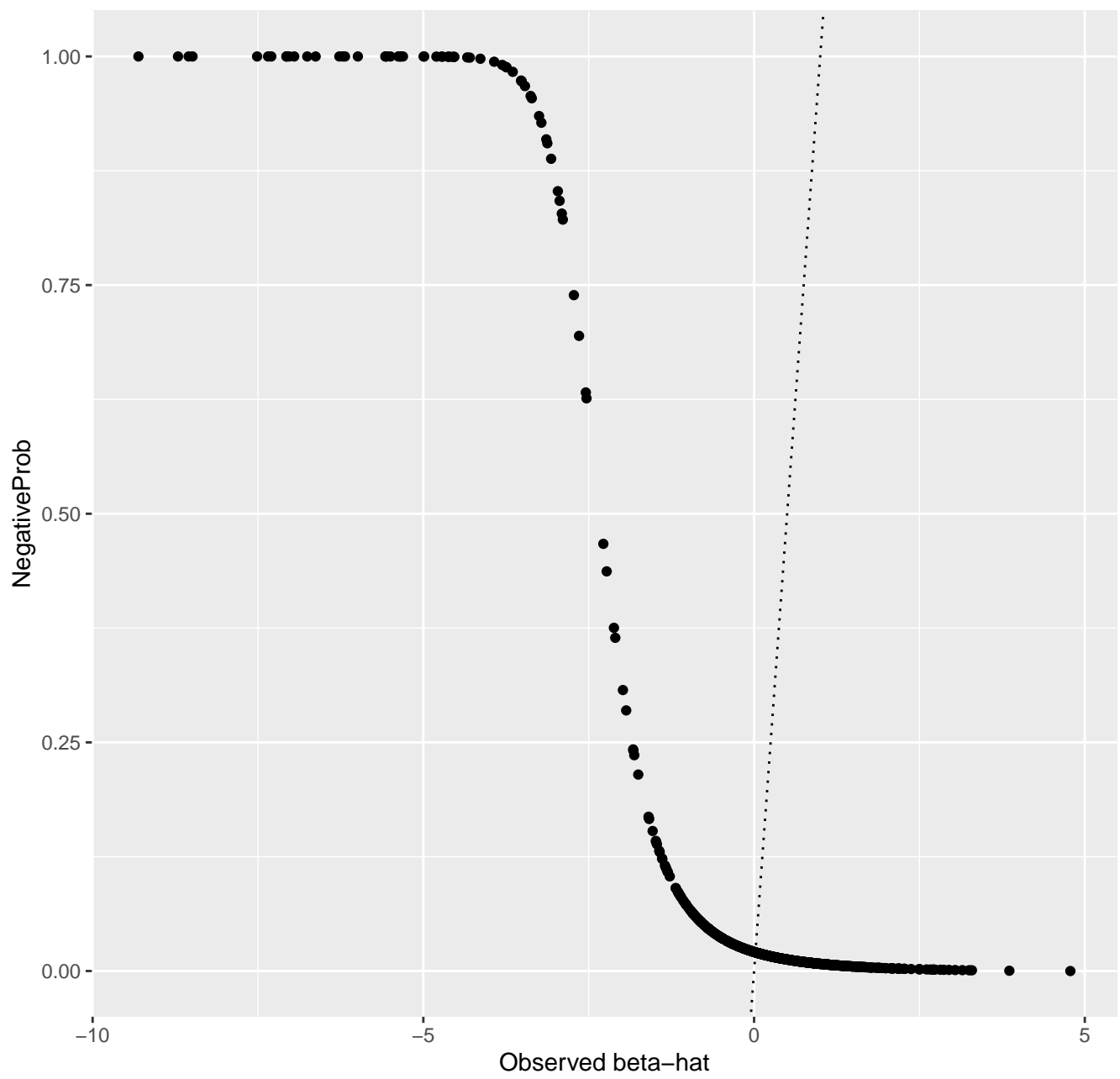
BRAF_(673) negative



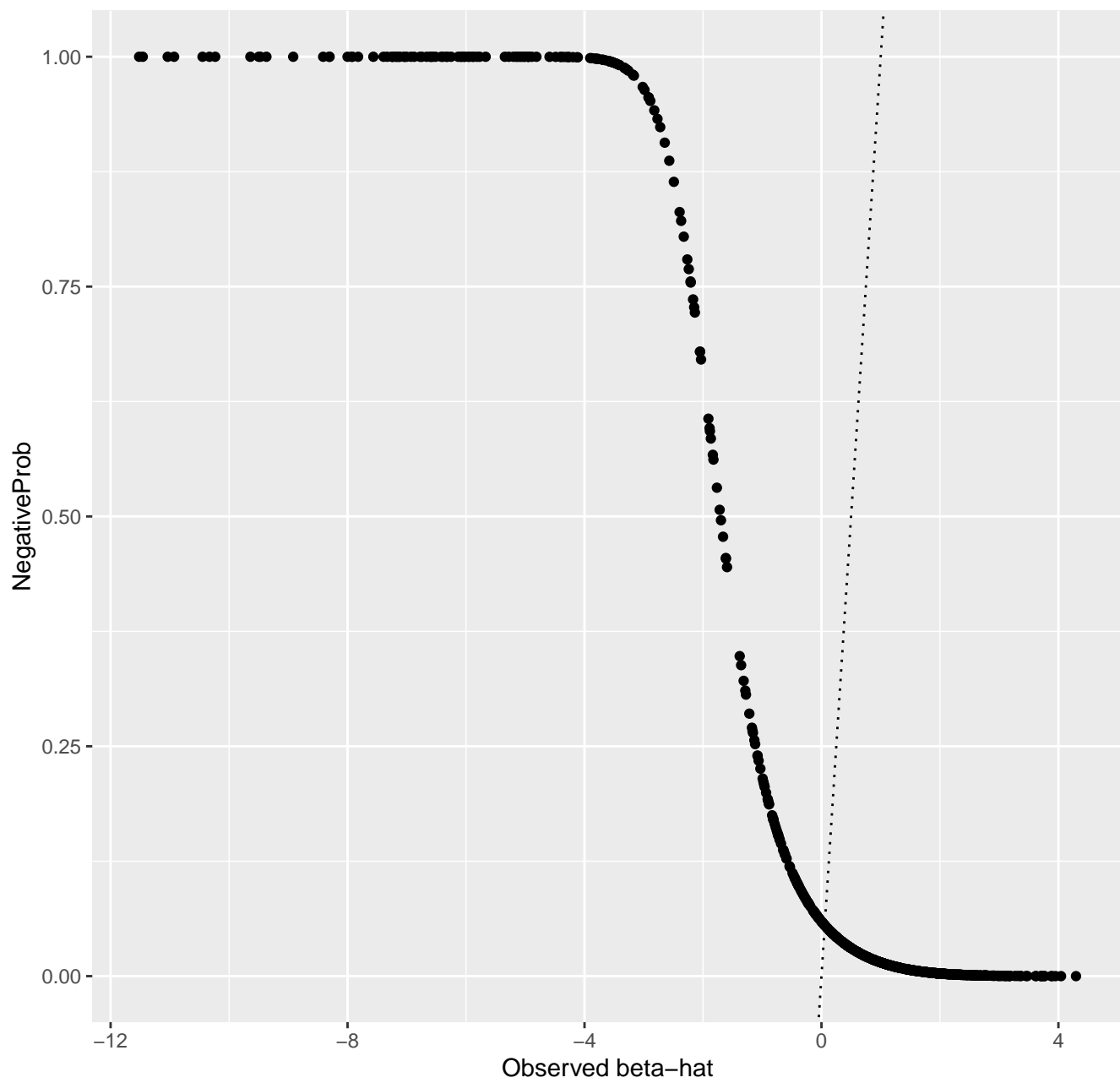
CTNNB1_(1499) negative



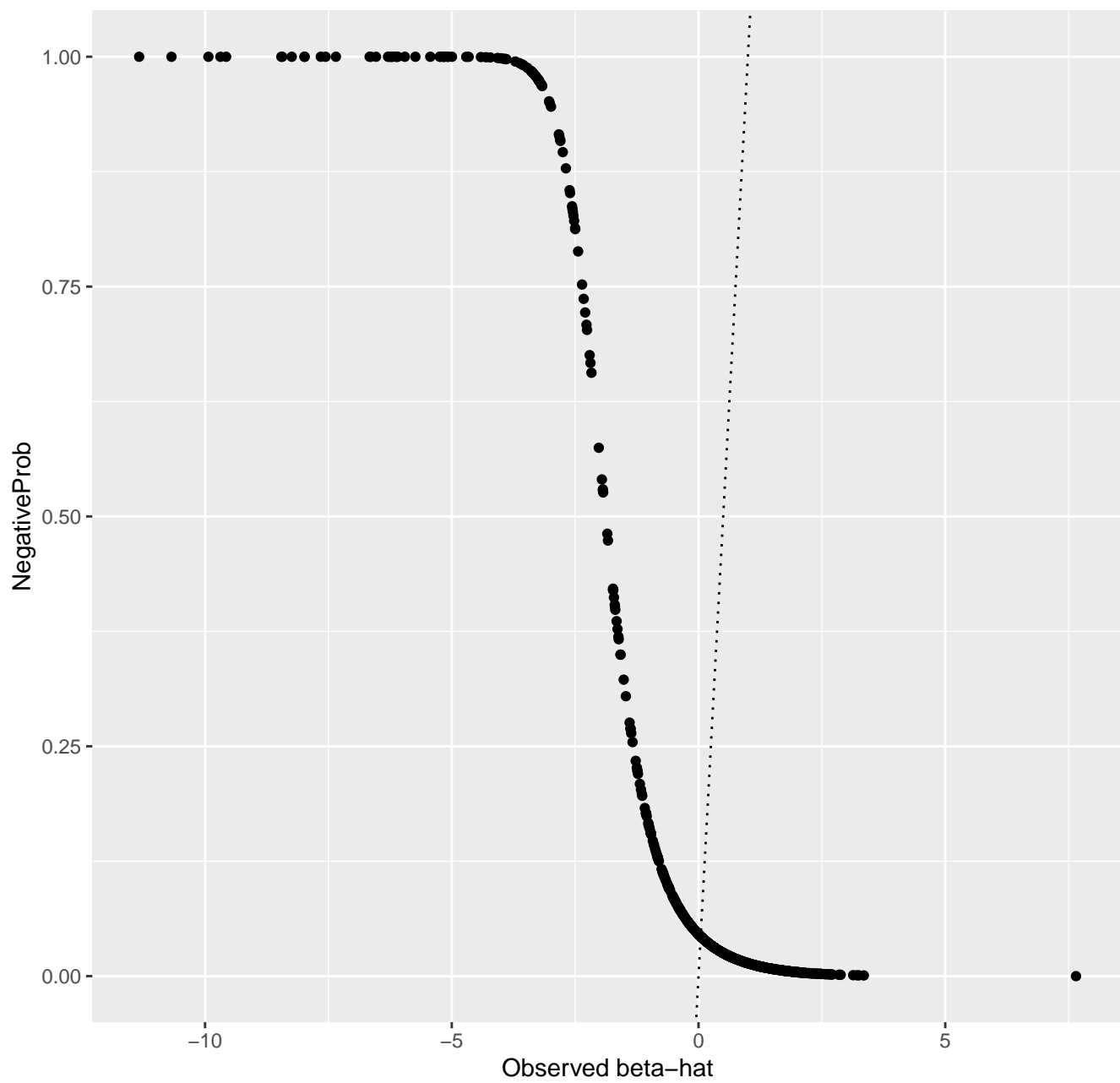
TCF7L2_(6934) negative



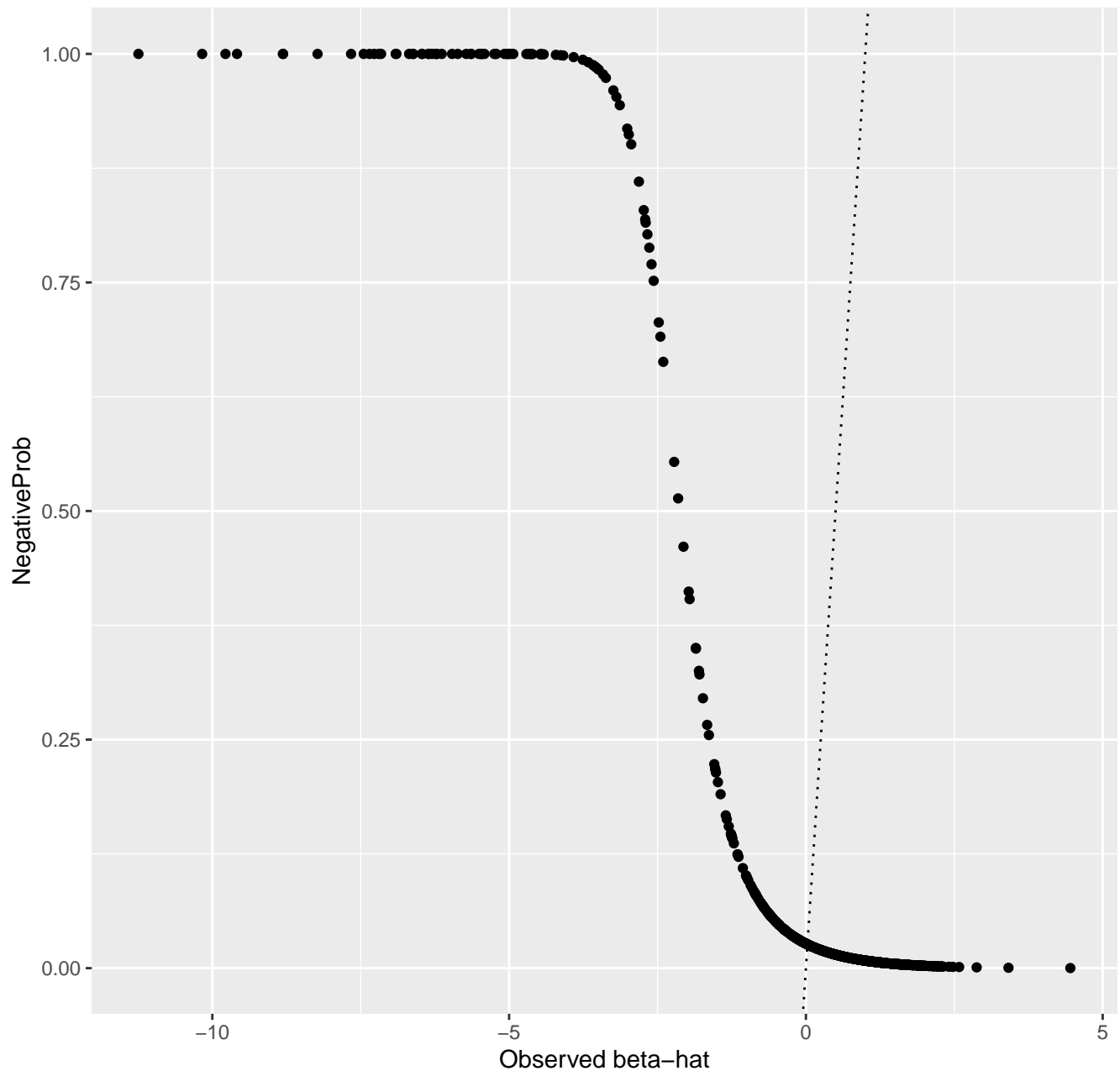
KRAS_(3845) negative



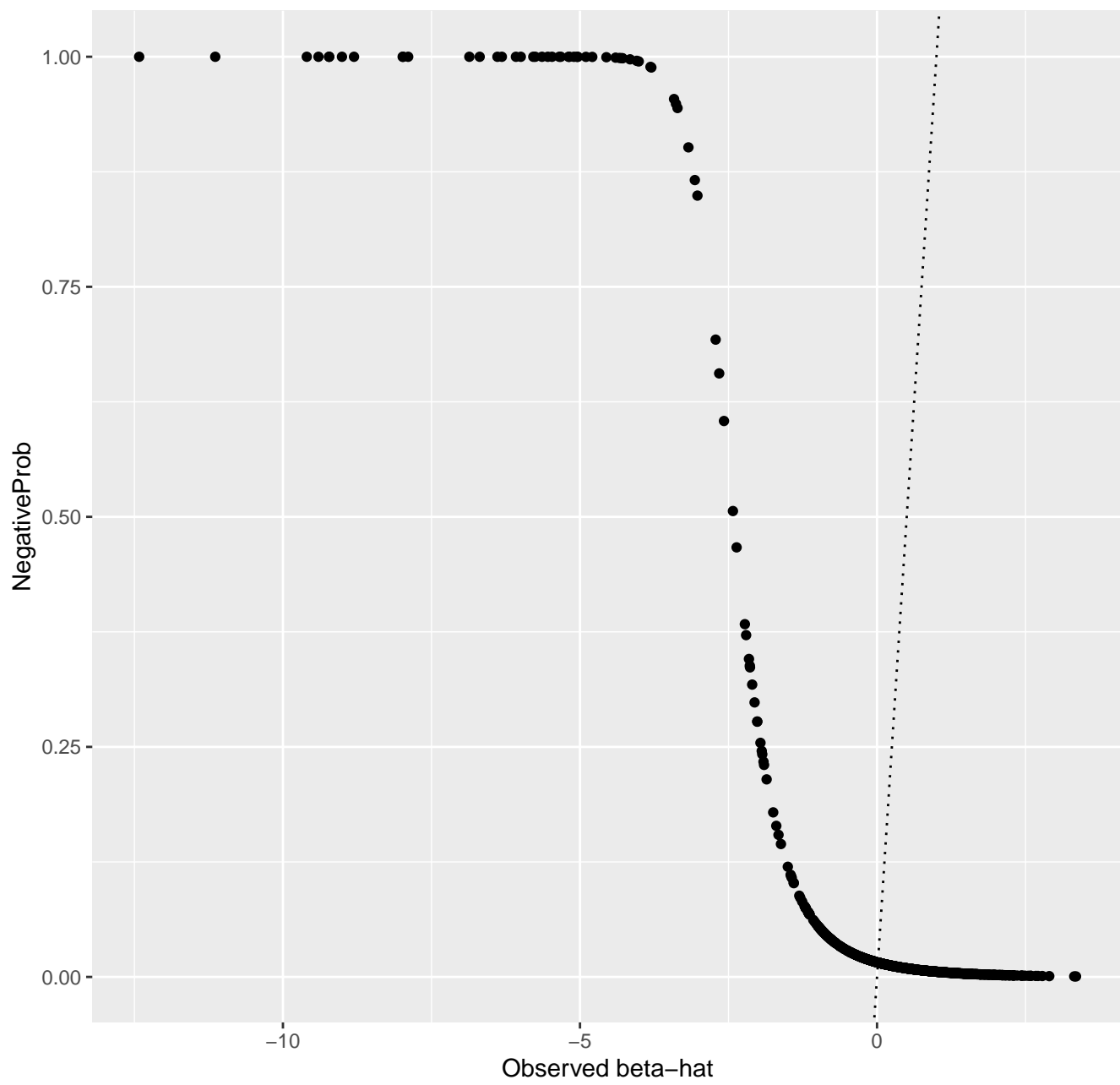
FGFR1_(2260) negative



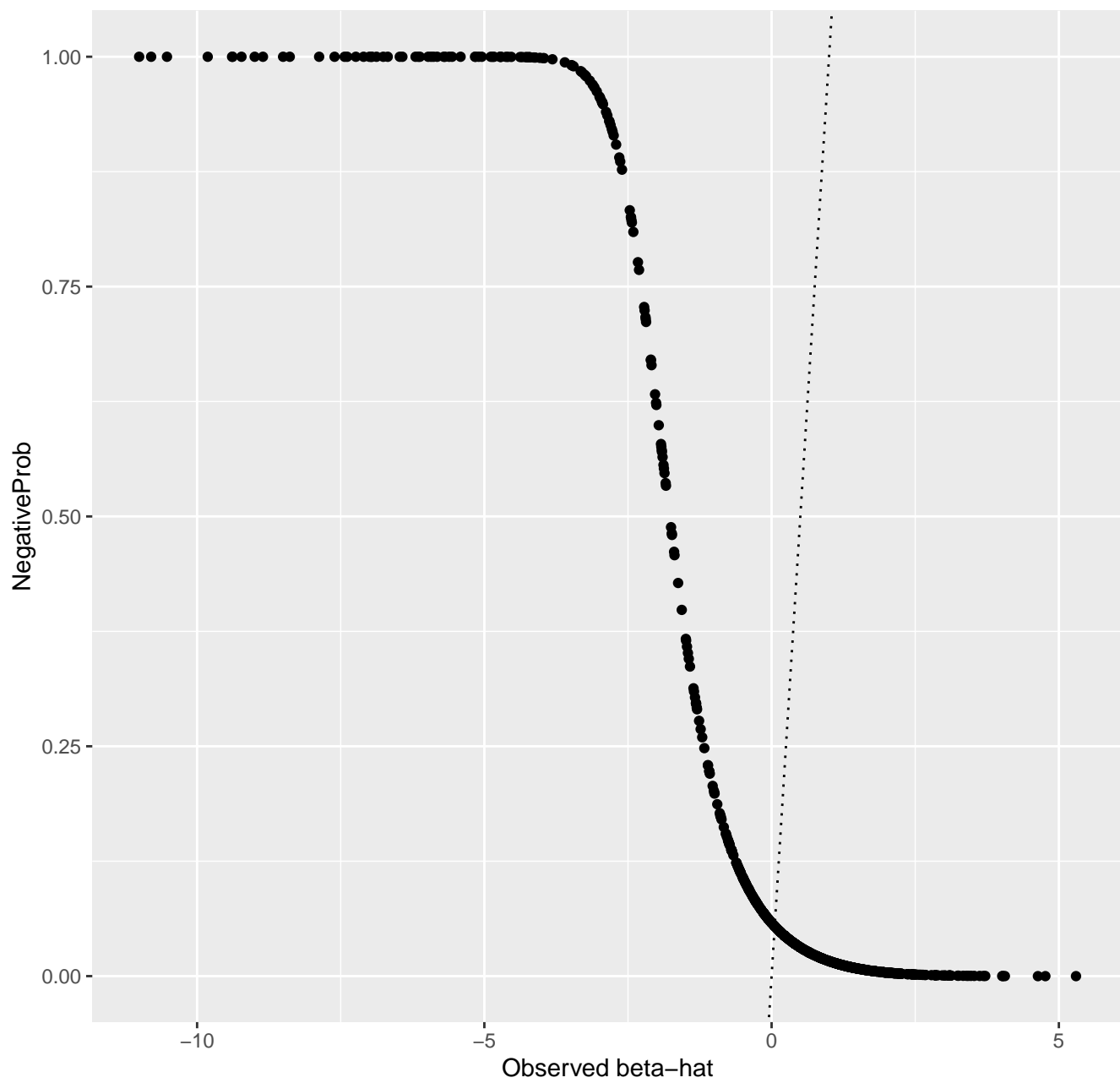
TP63_(8626) negative



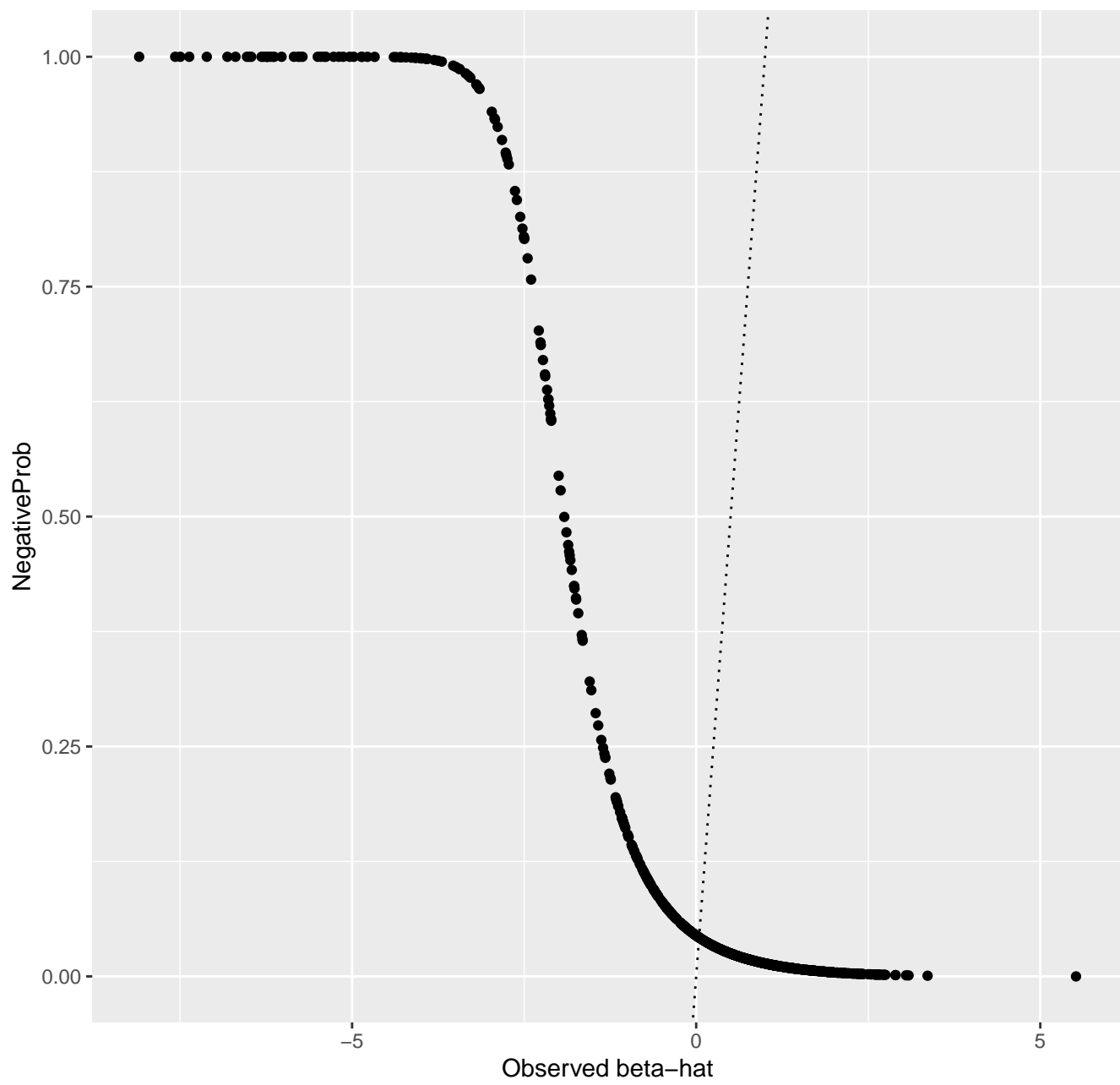
CCND2_(894) negative



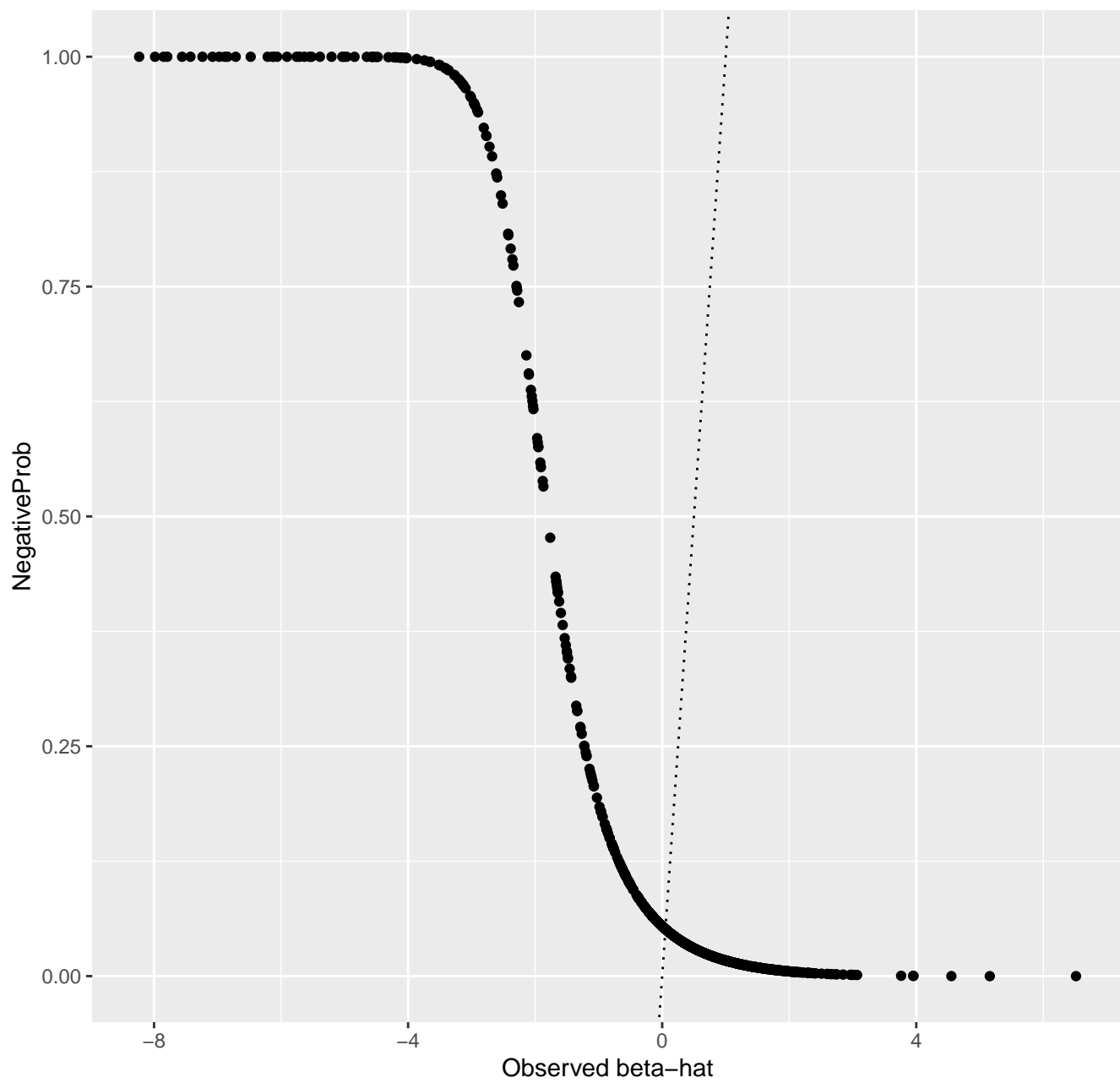
EIF2AK4_(440275) negative



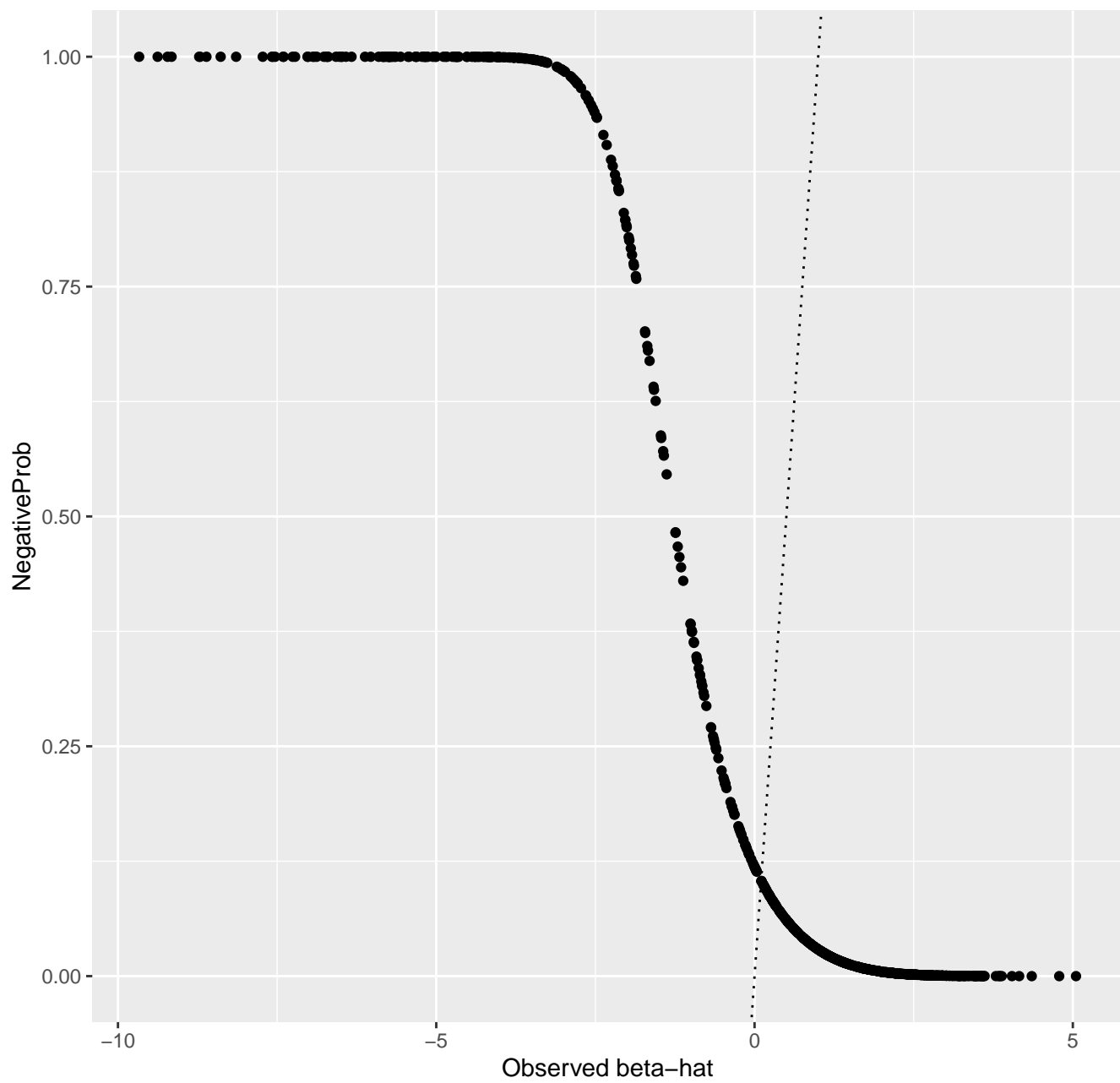
PARD6B_(84612) negative



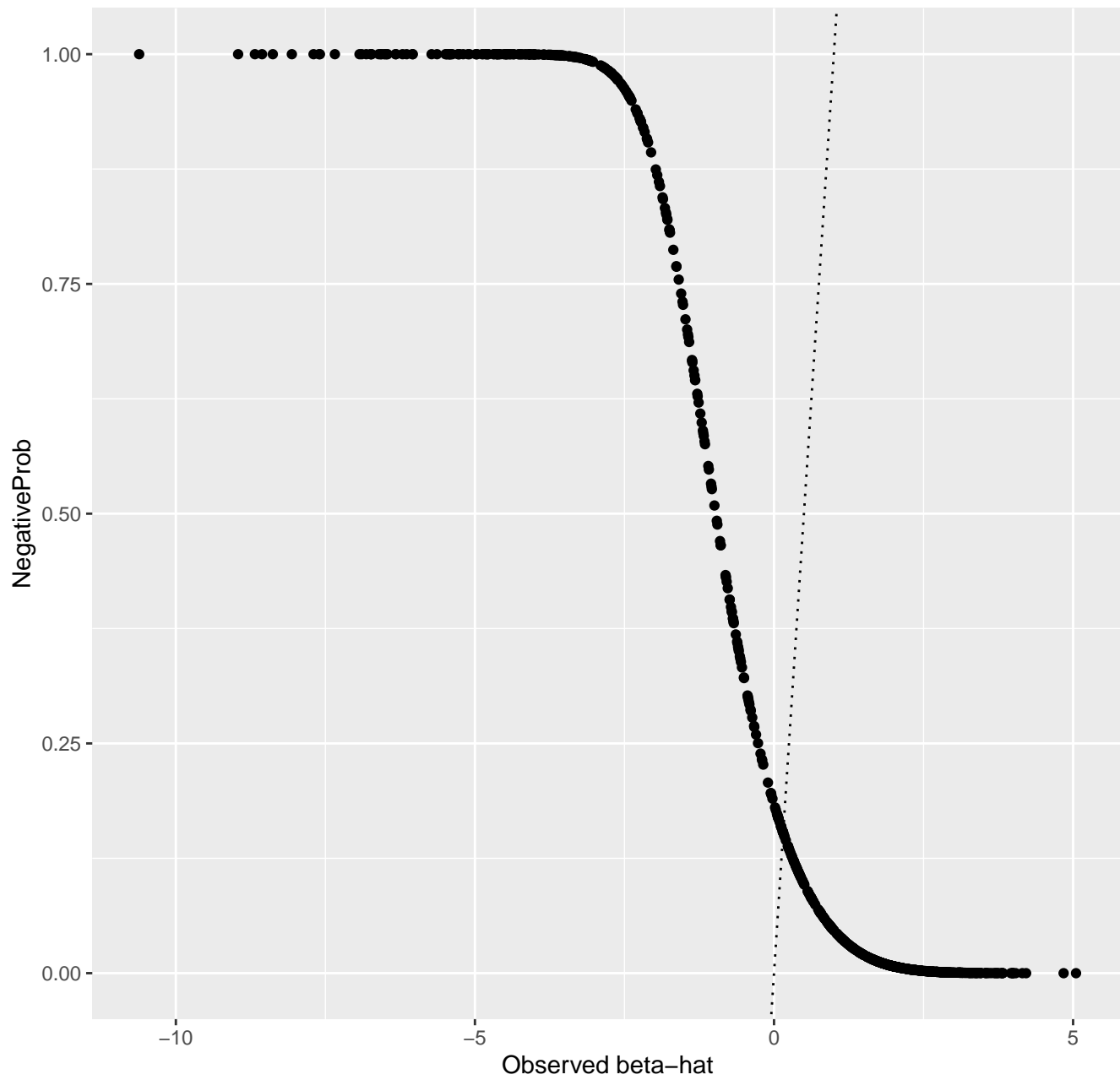
ZEB2_(9839) negative



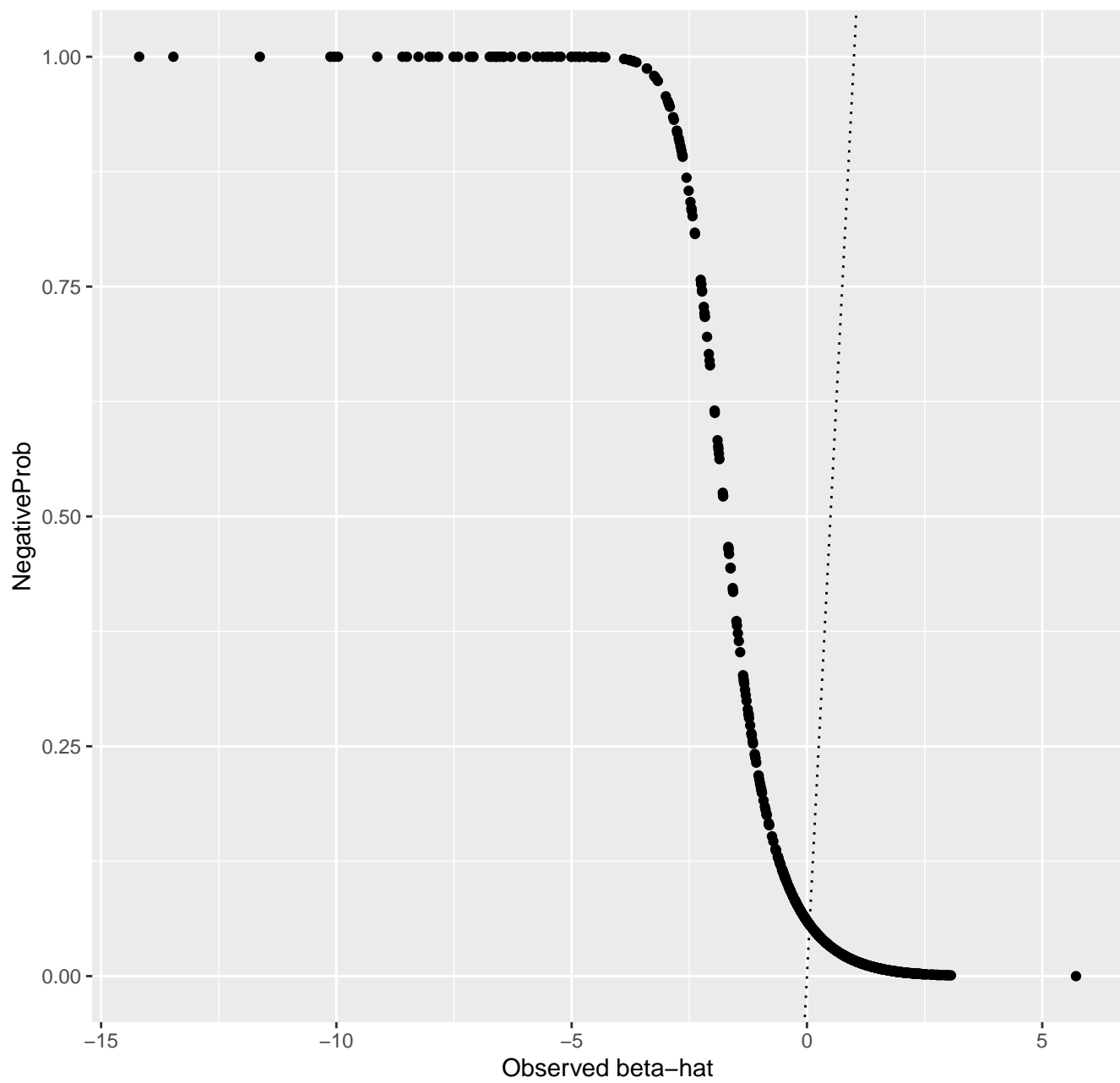
RPP25L_(138716) negative



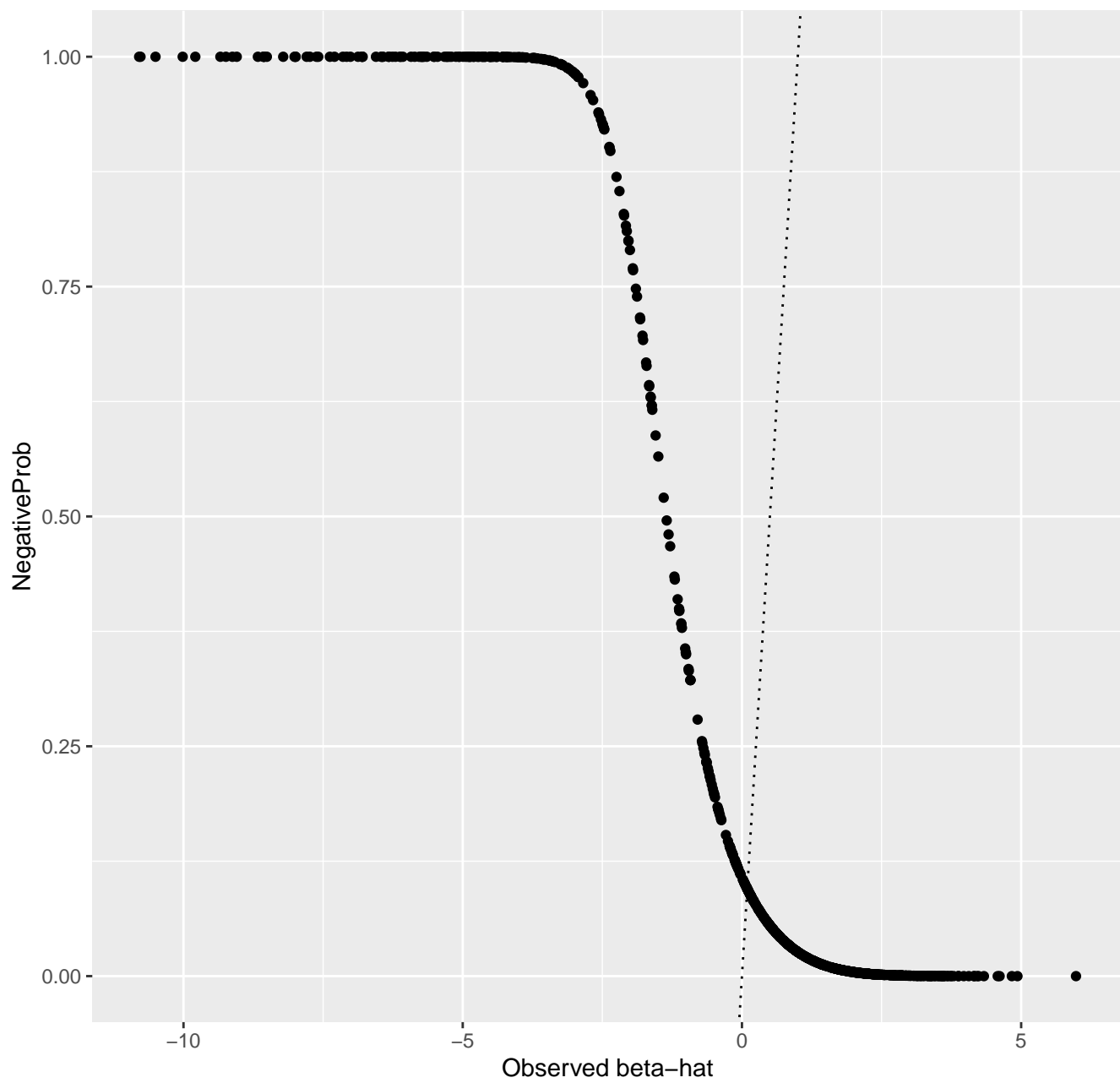
STX4_(6810) negative



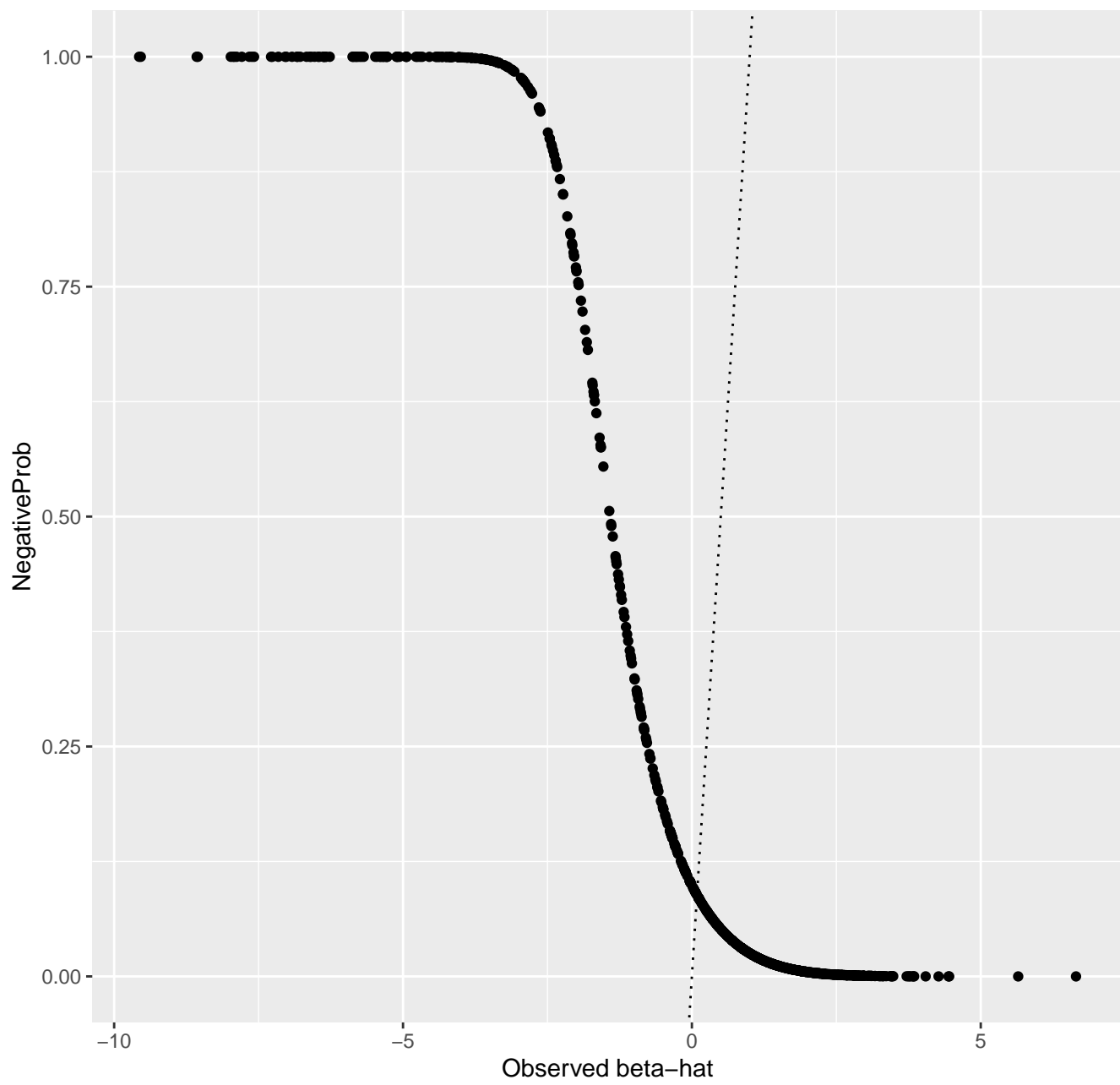
VPS4B_(9525) negative



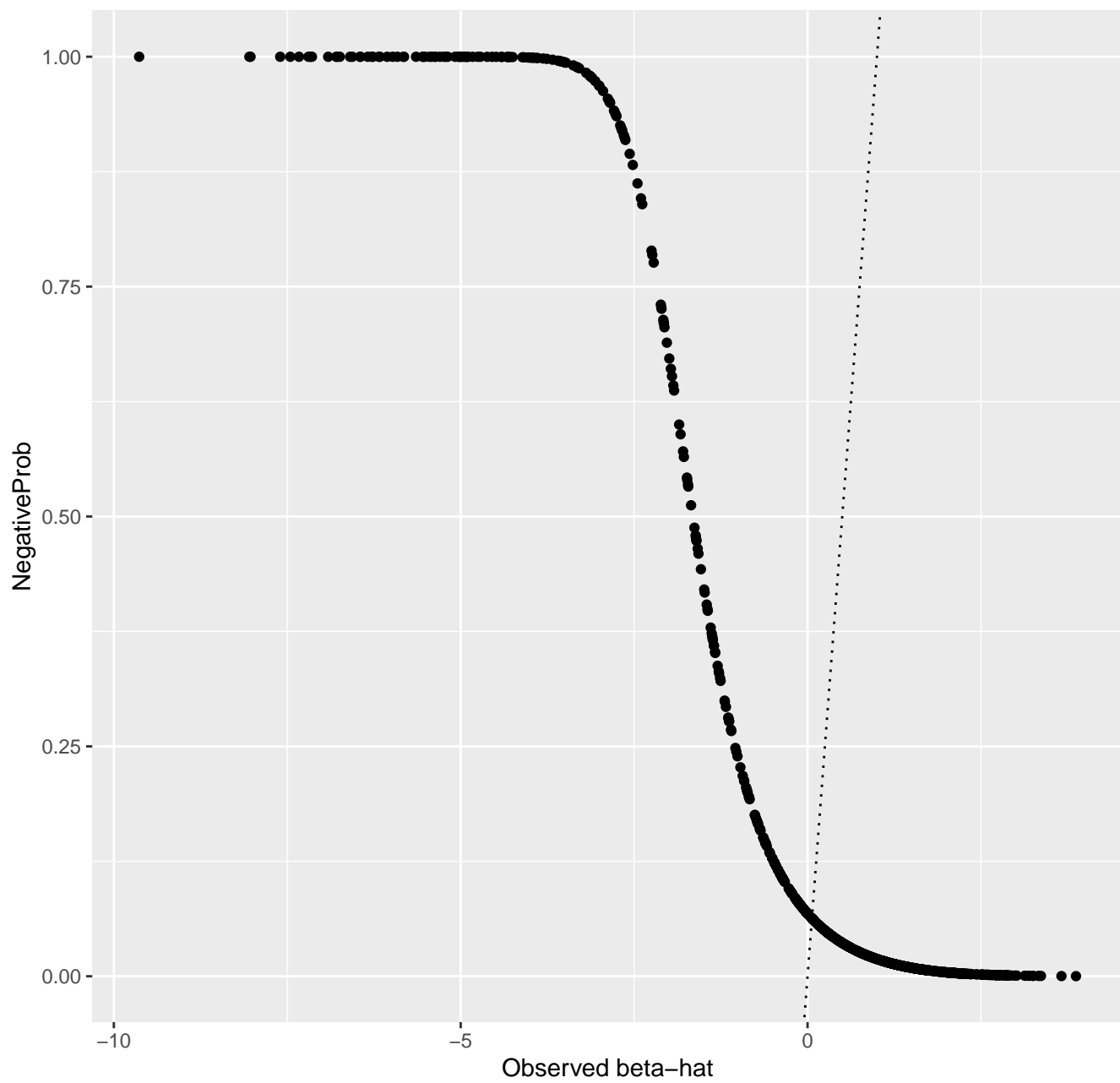
MDM2_(4193) negative



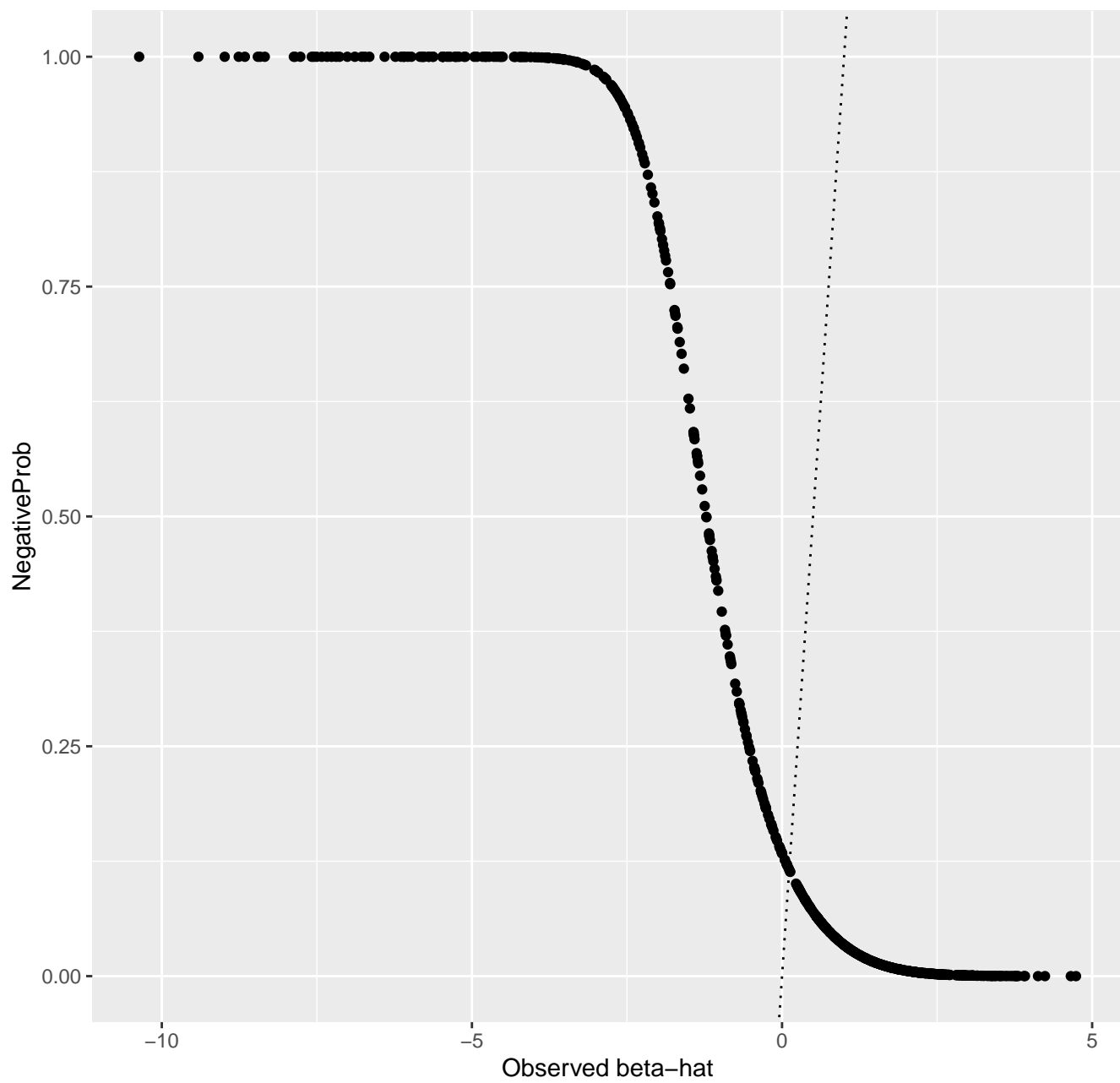
KLF5_(688) negative



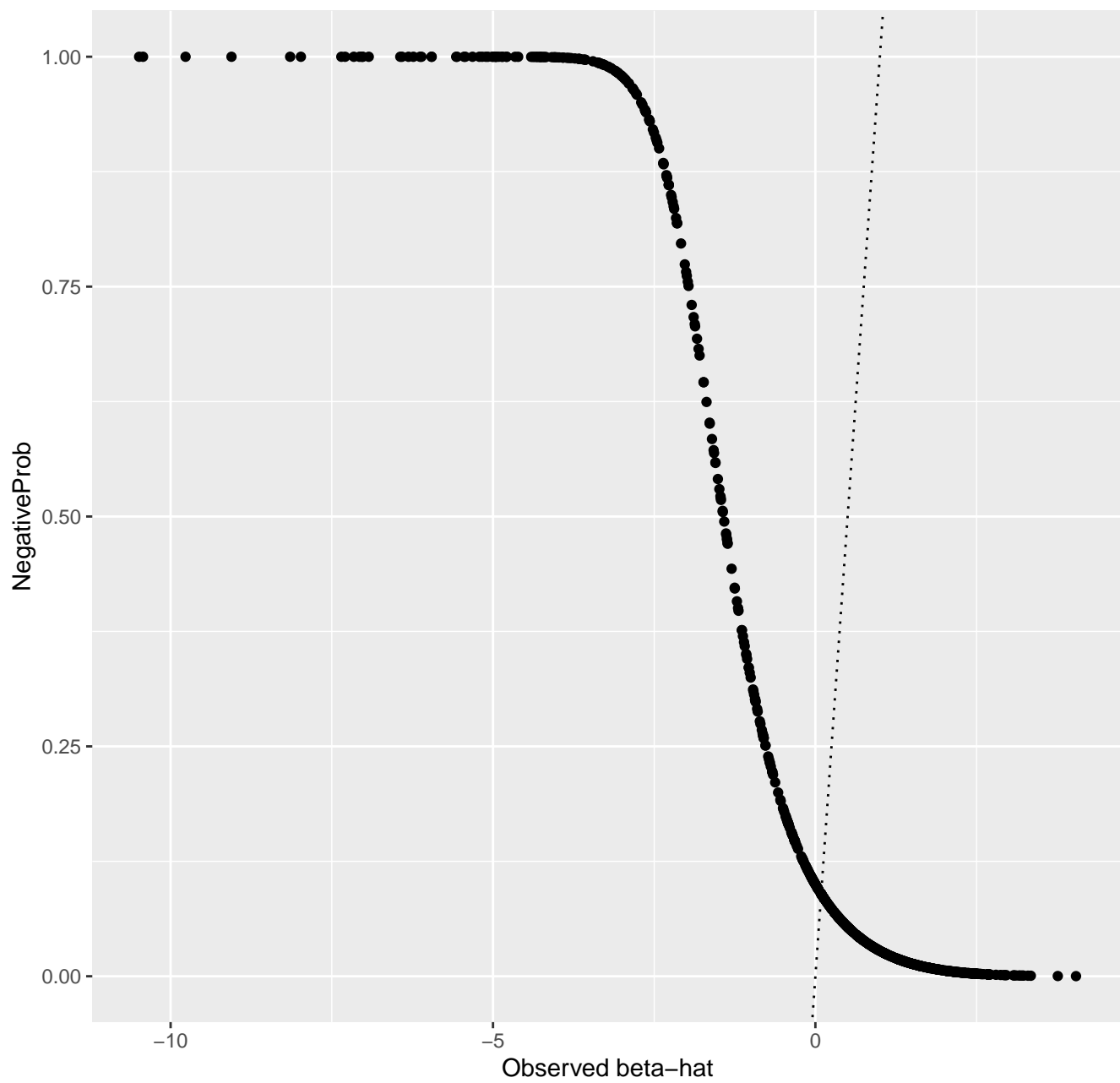
STAMBP_(10617) negative



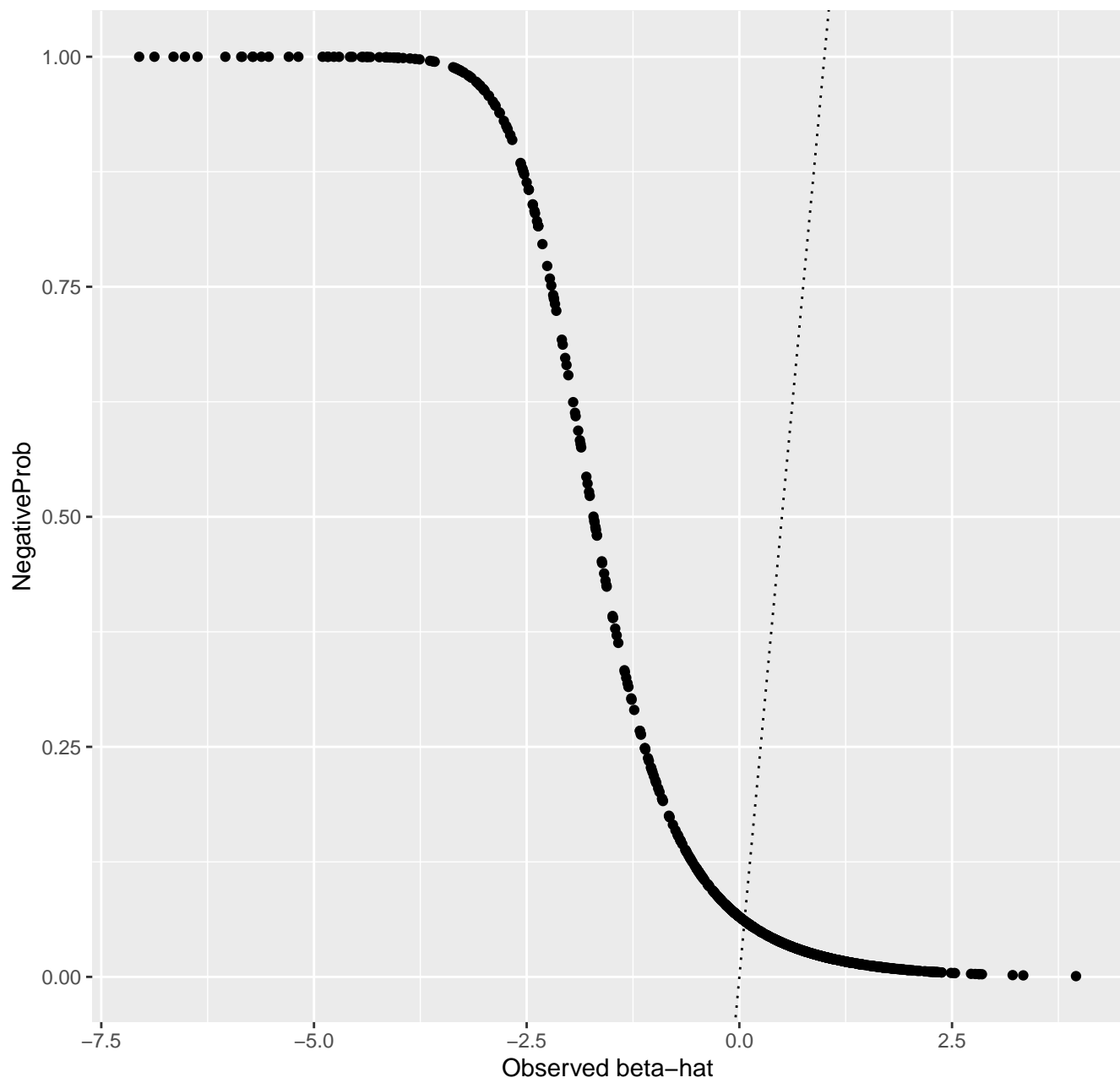
DHR SX_(207063) negative



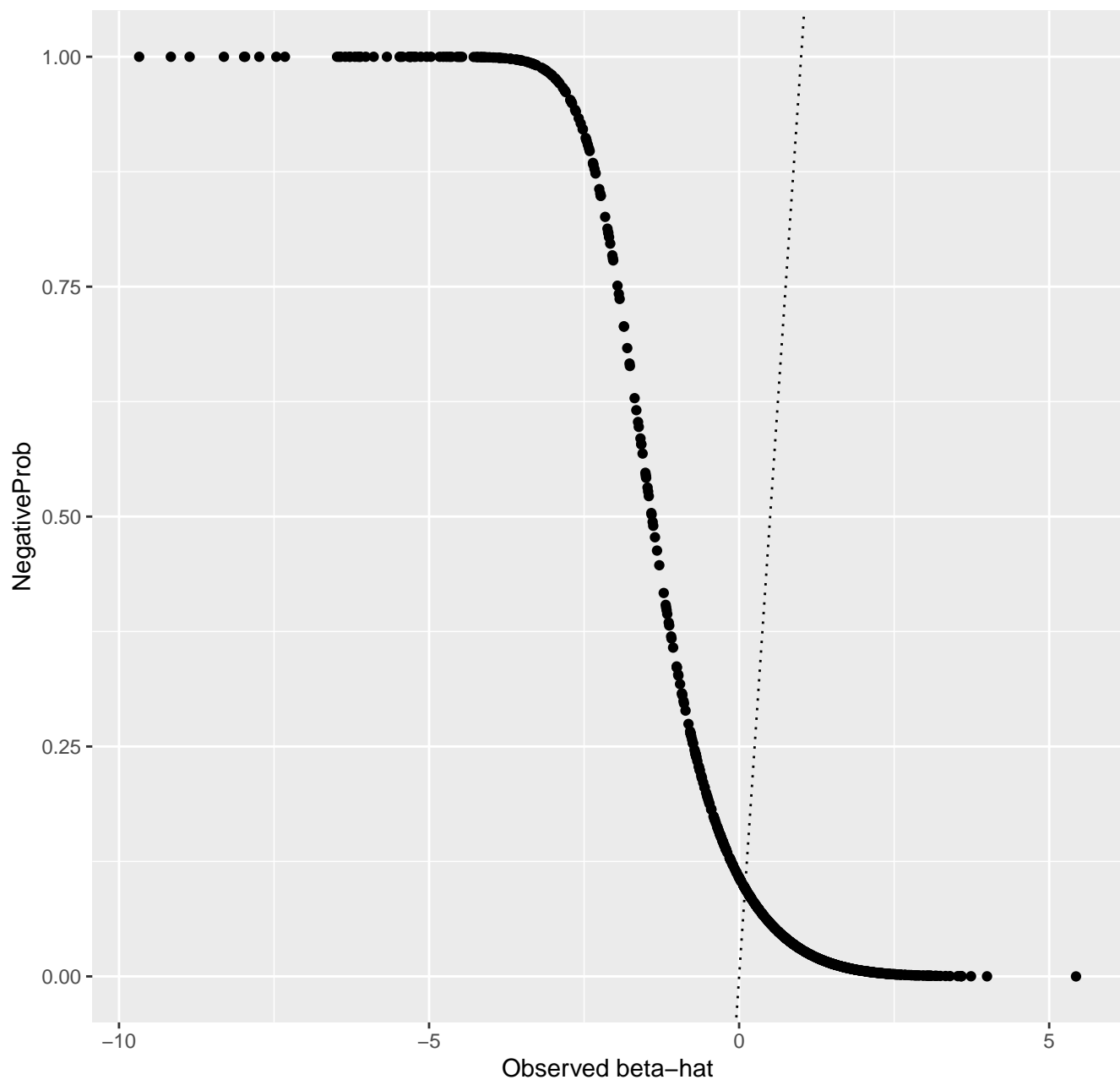
EGFR_(1956) negative



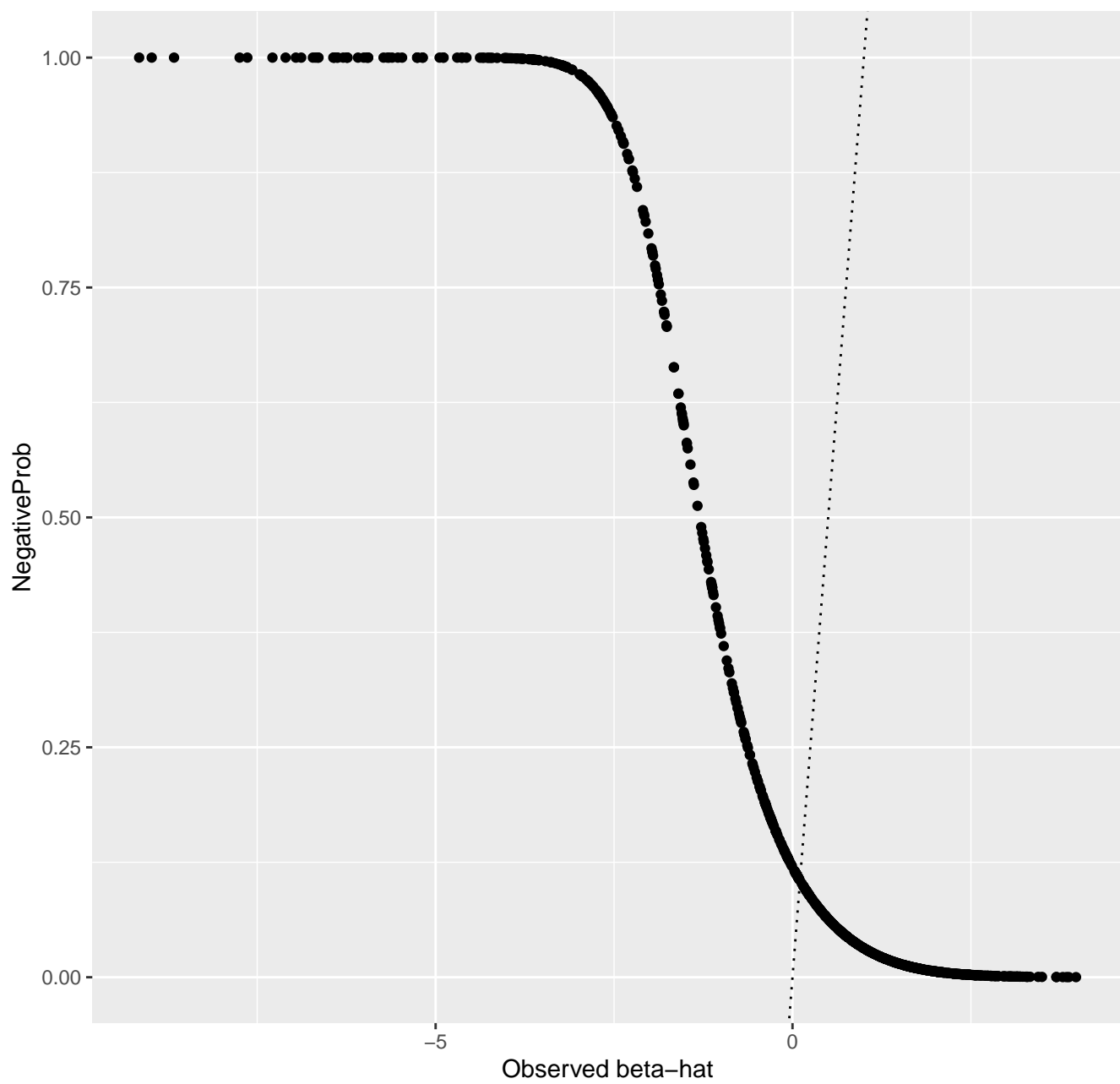
GRHL2_(79977) negative



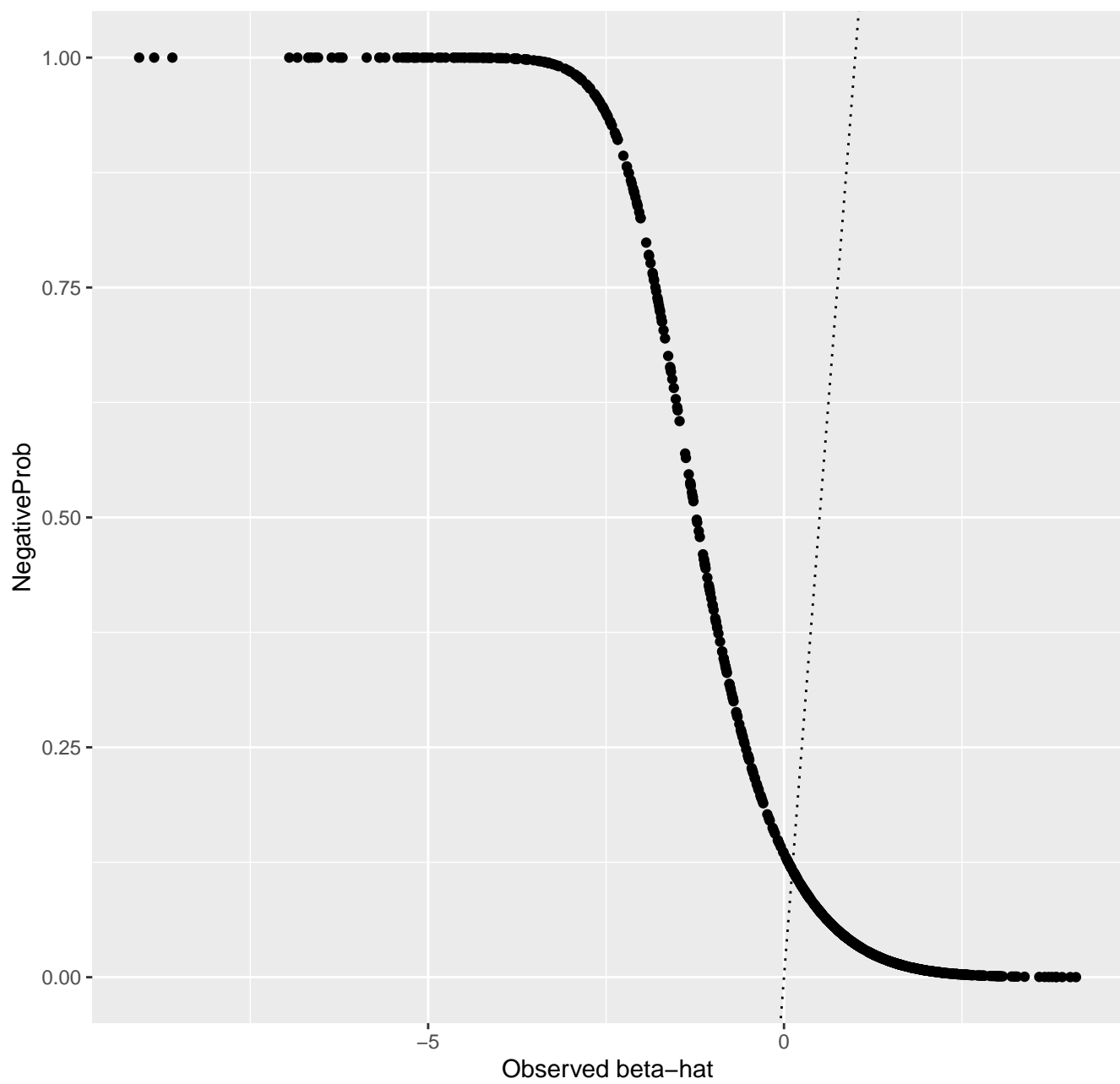
SLC7A1_(6541) negative



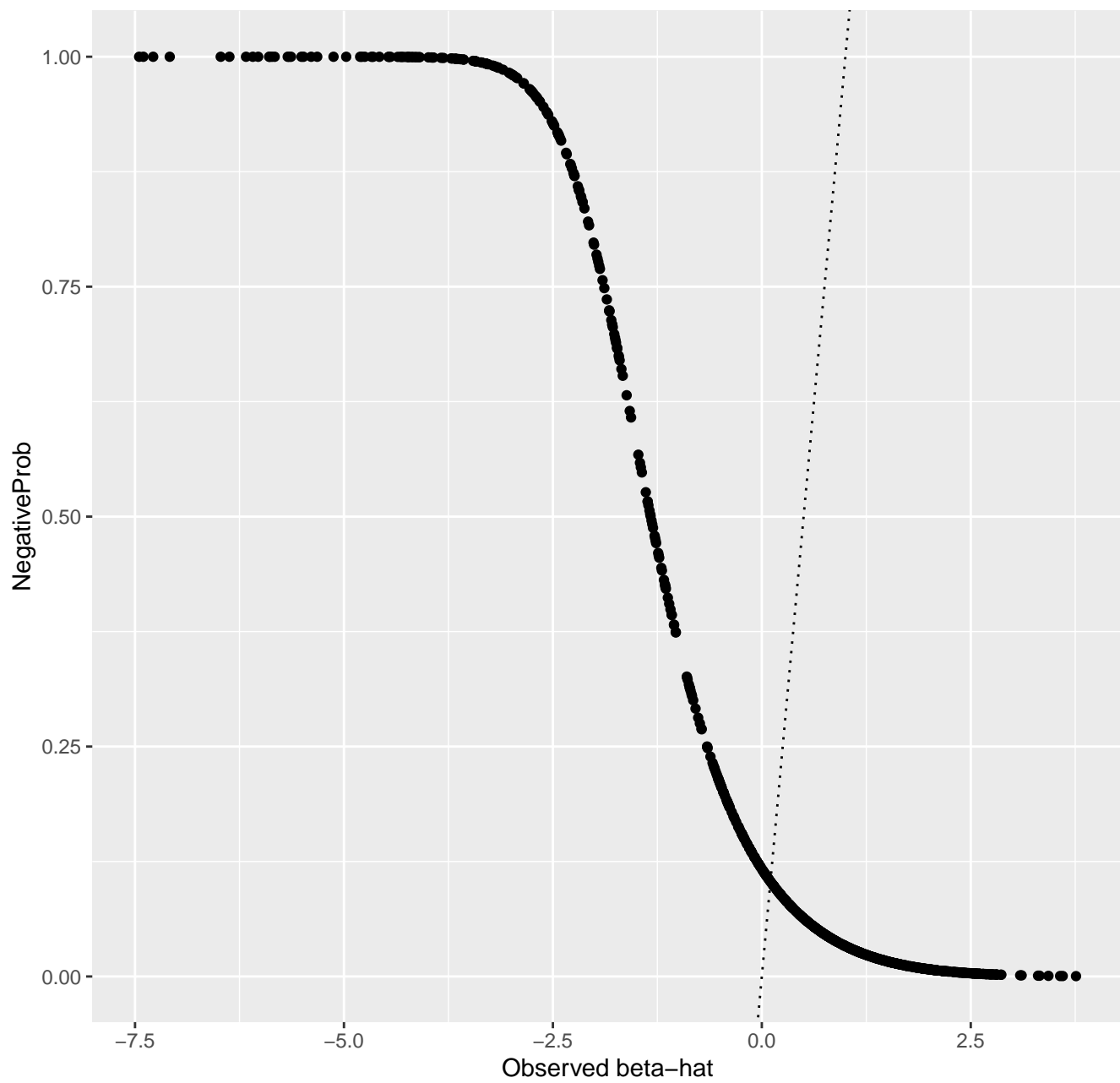
CYB5B_(80777) negative



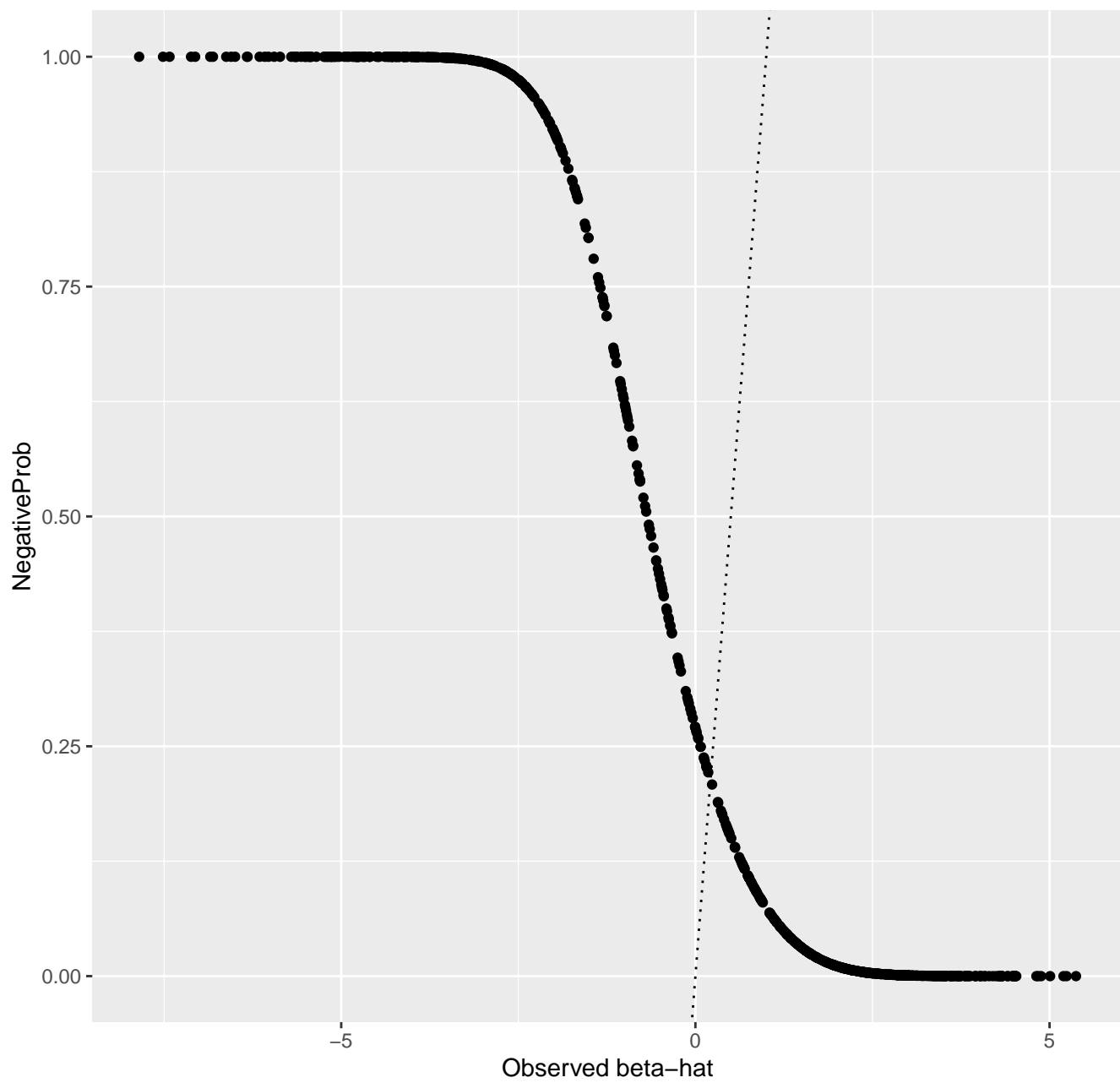
IGF1R_(3480) negative



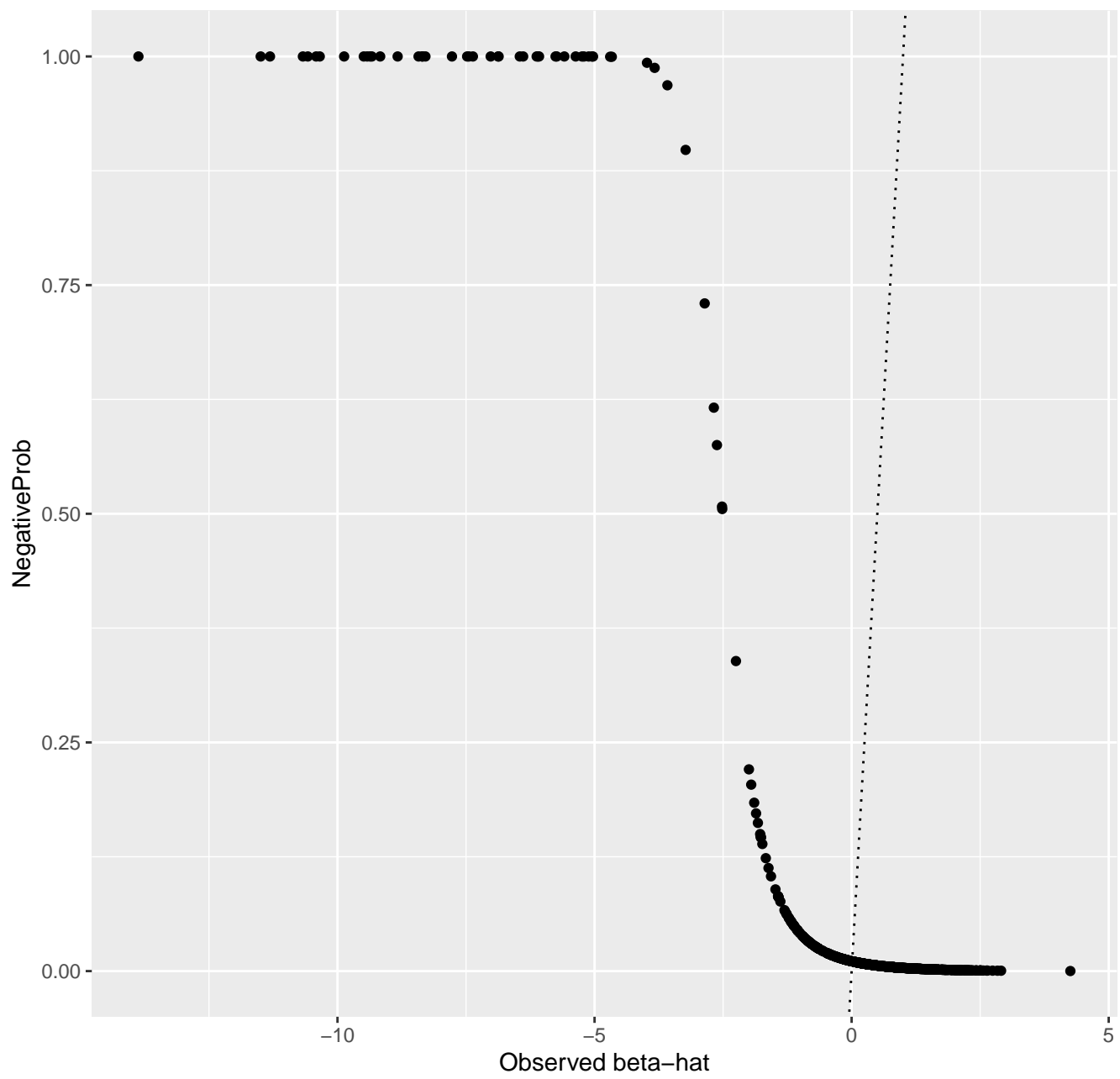
STXBP3_(6814) negative



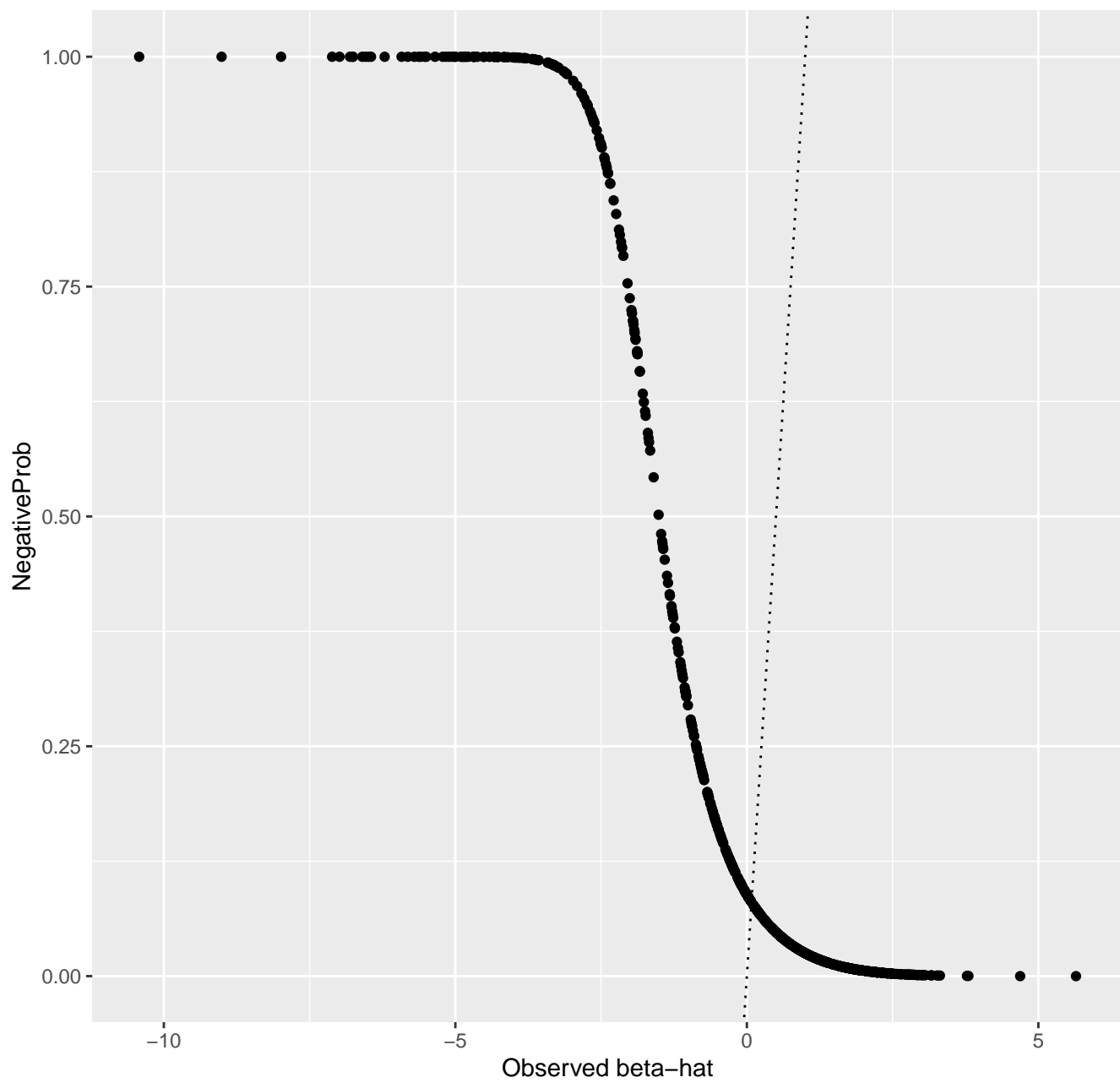
FAM50A_(9130) negative



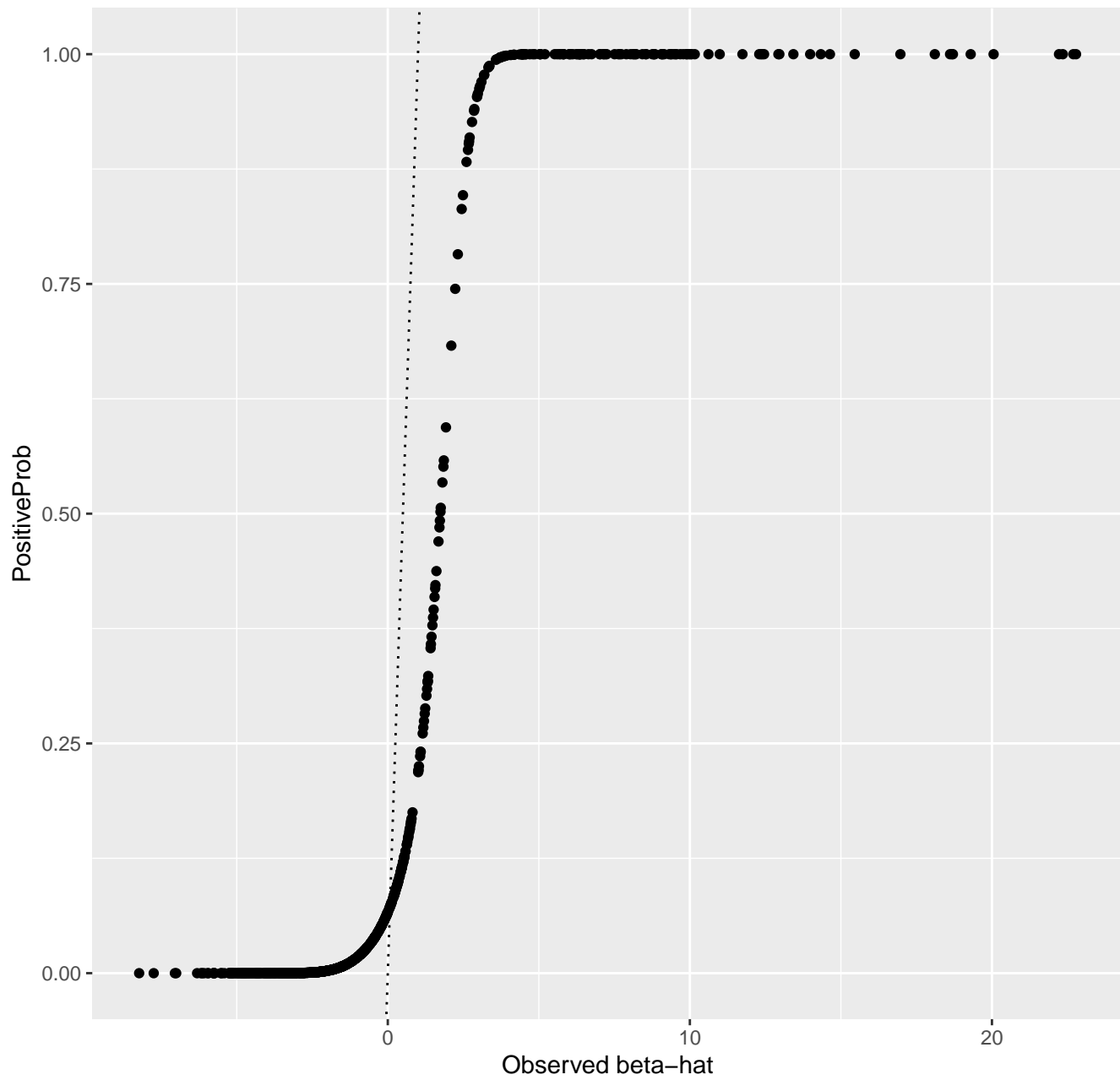
IRF4_(3662) negative



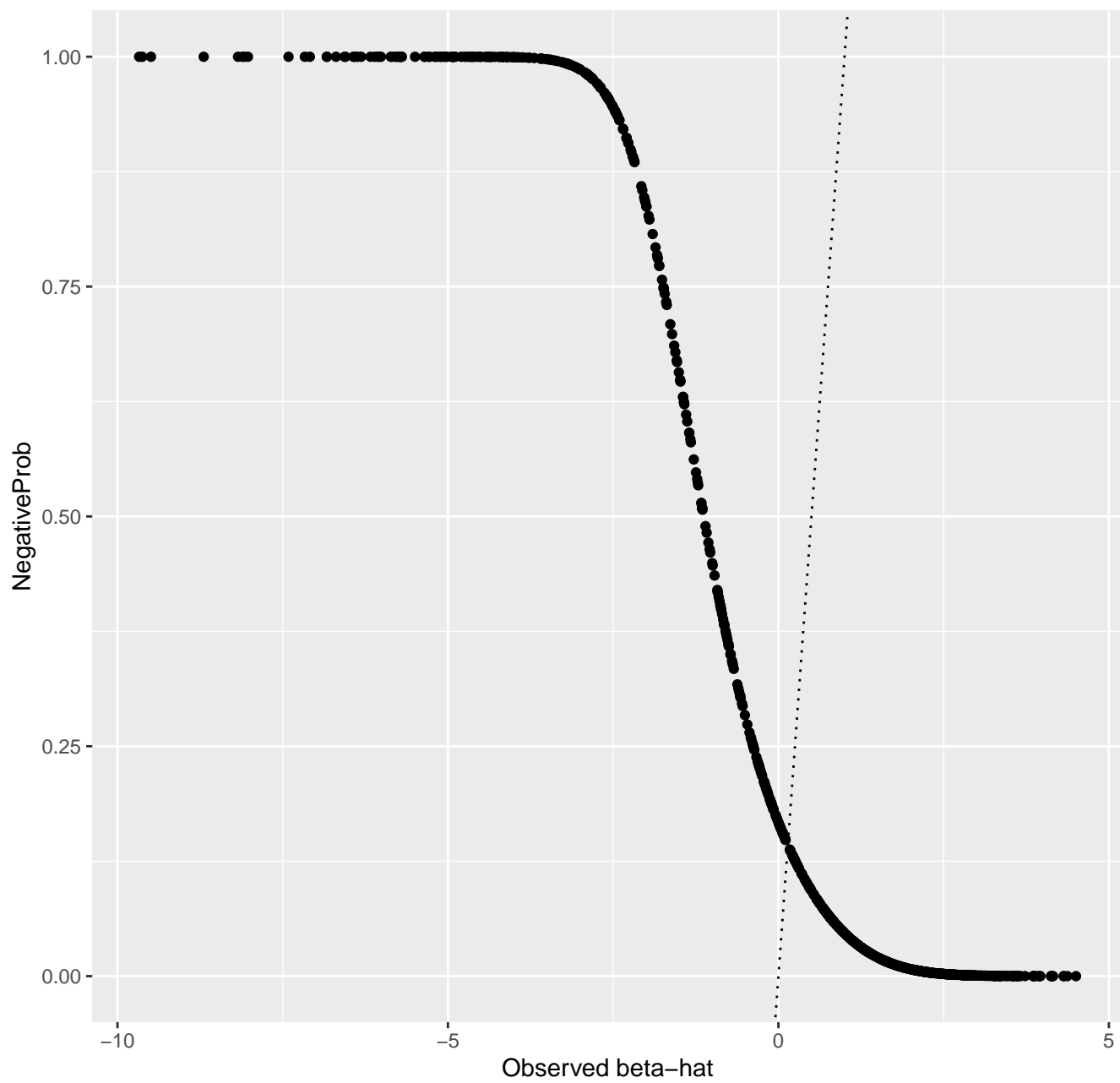
KIF18B_(146909) negative



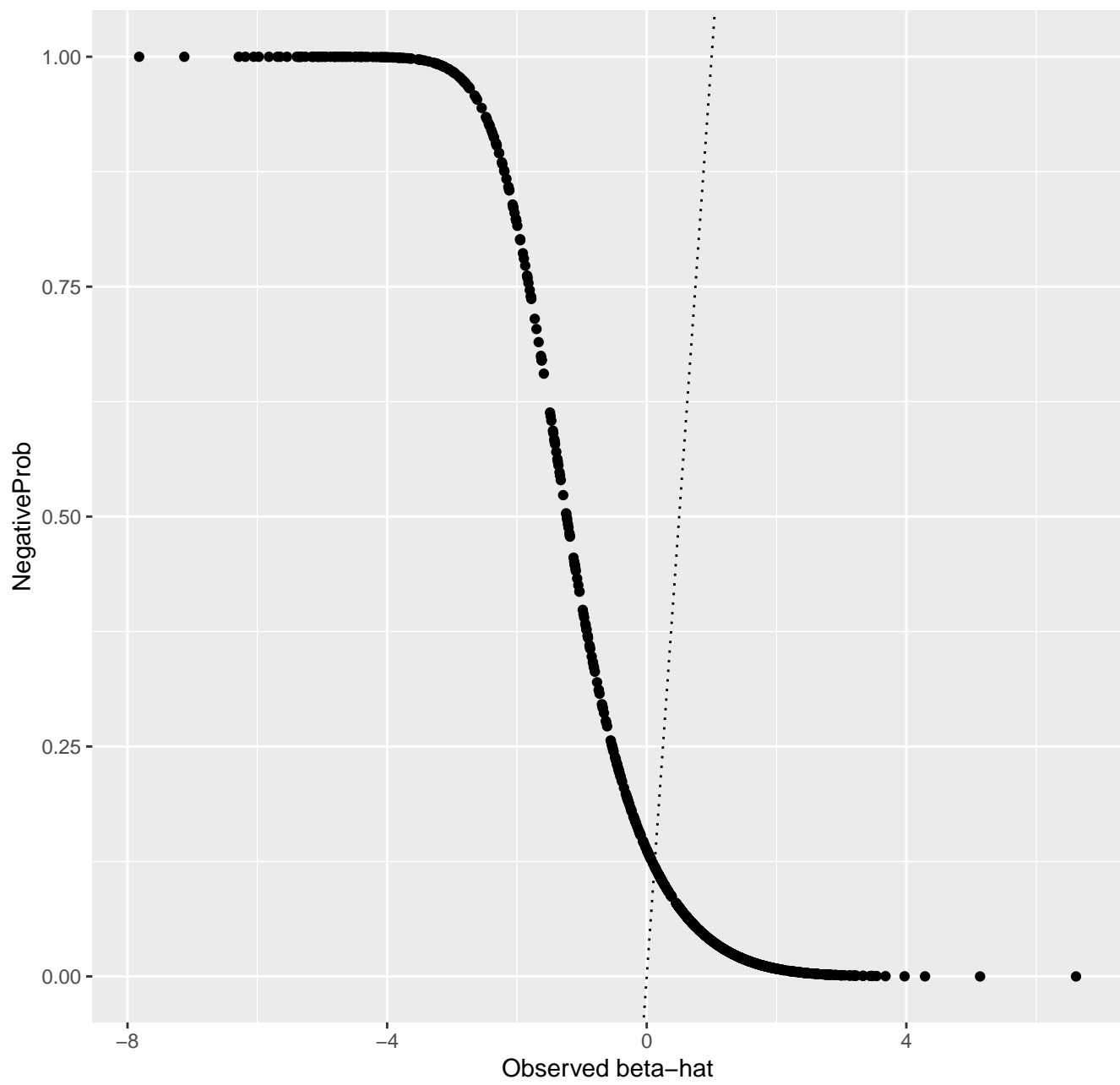
TP53_(7157) positive



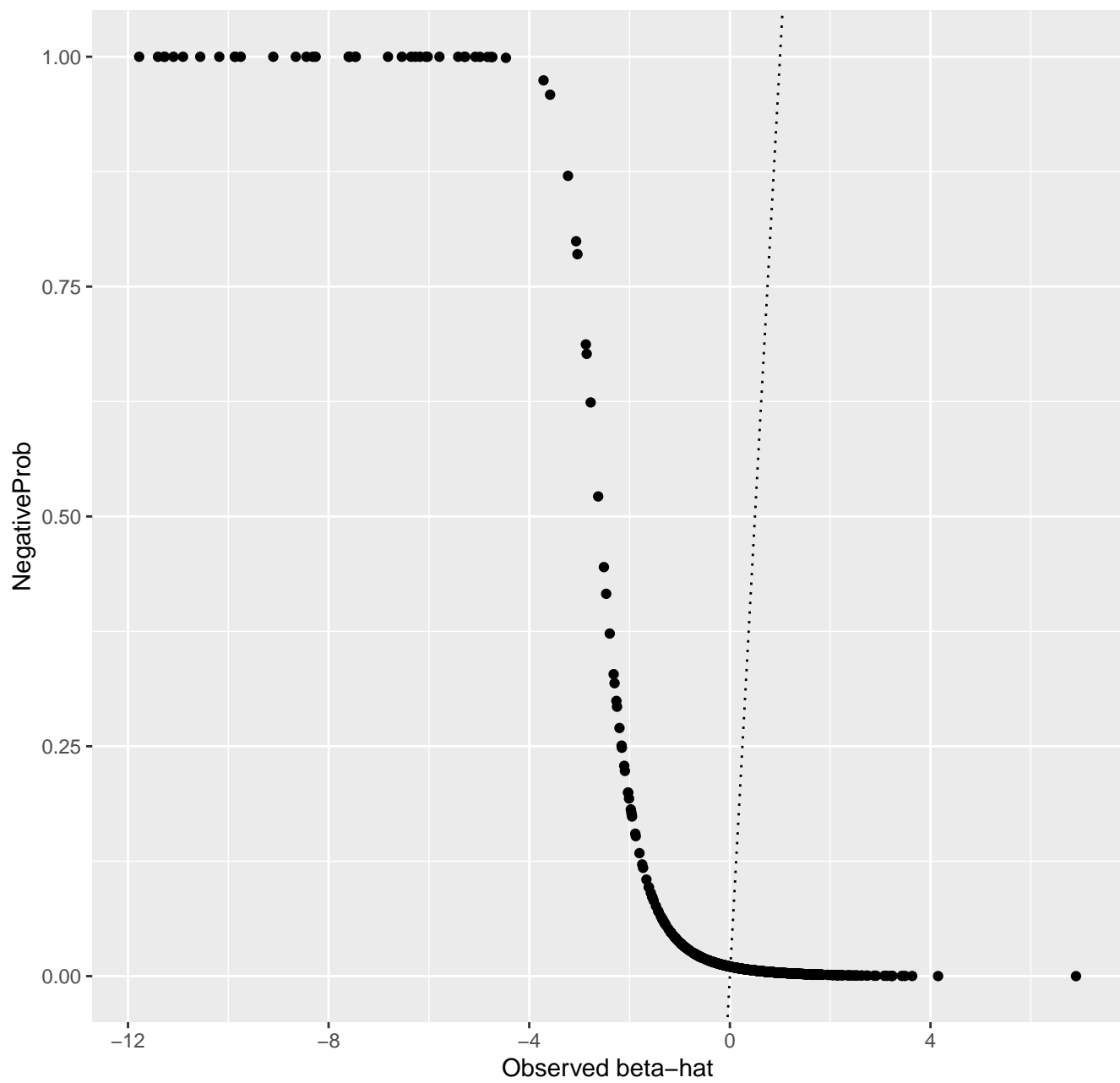
UBIAD1_(29914) negative



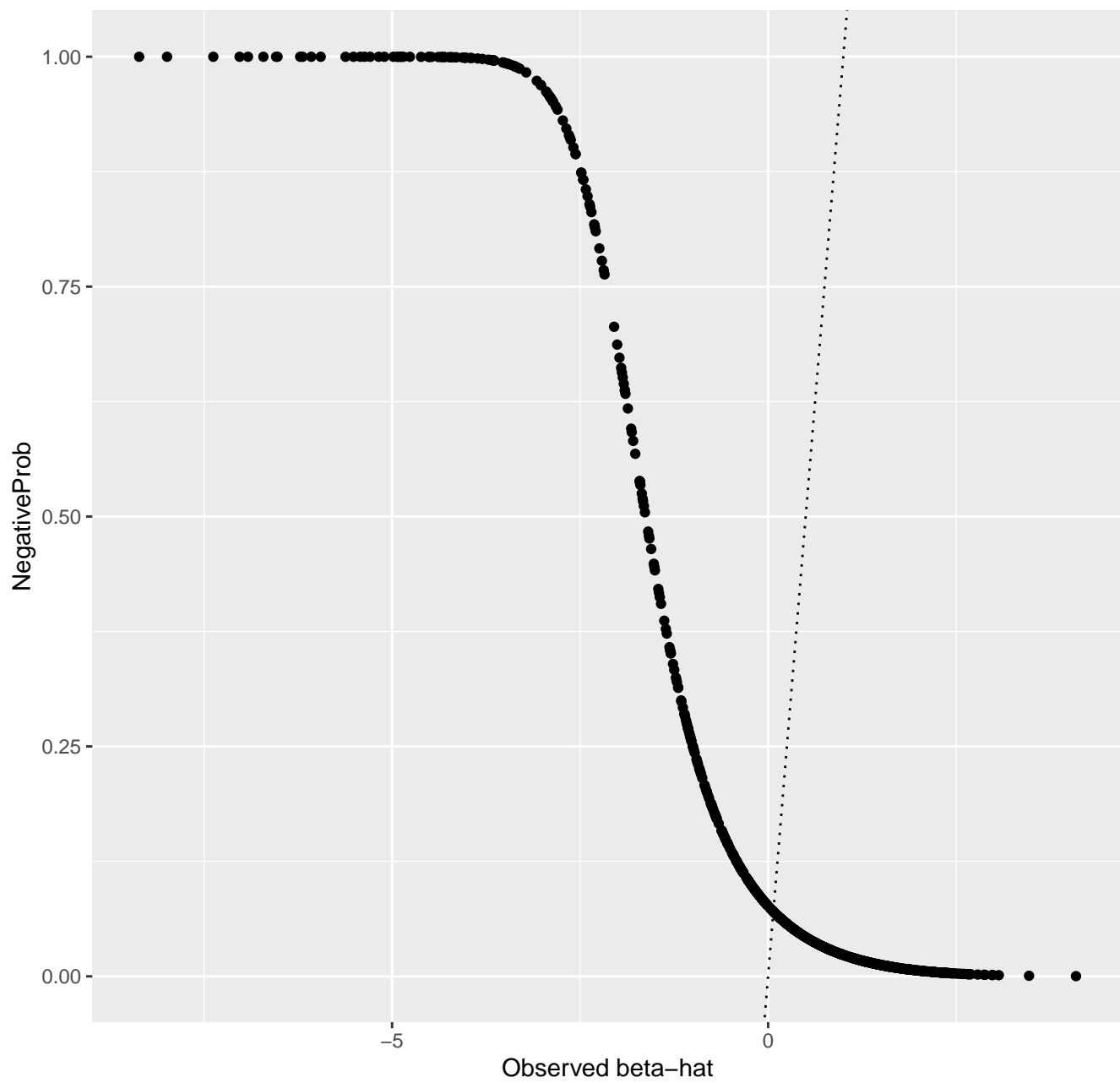
PTDSS1_(9791) negative



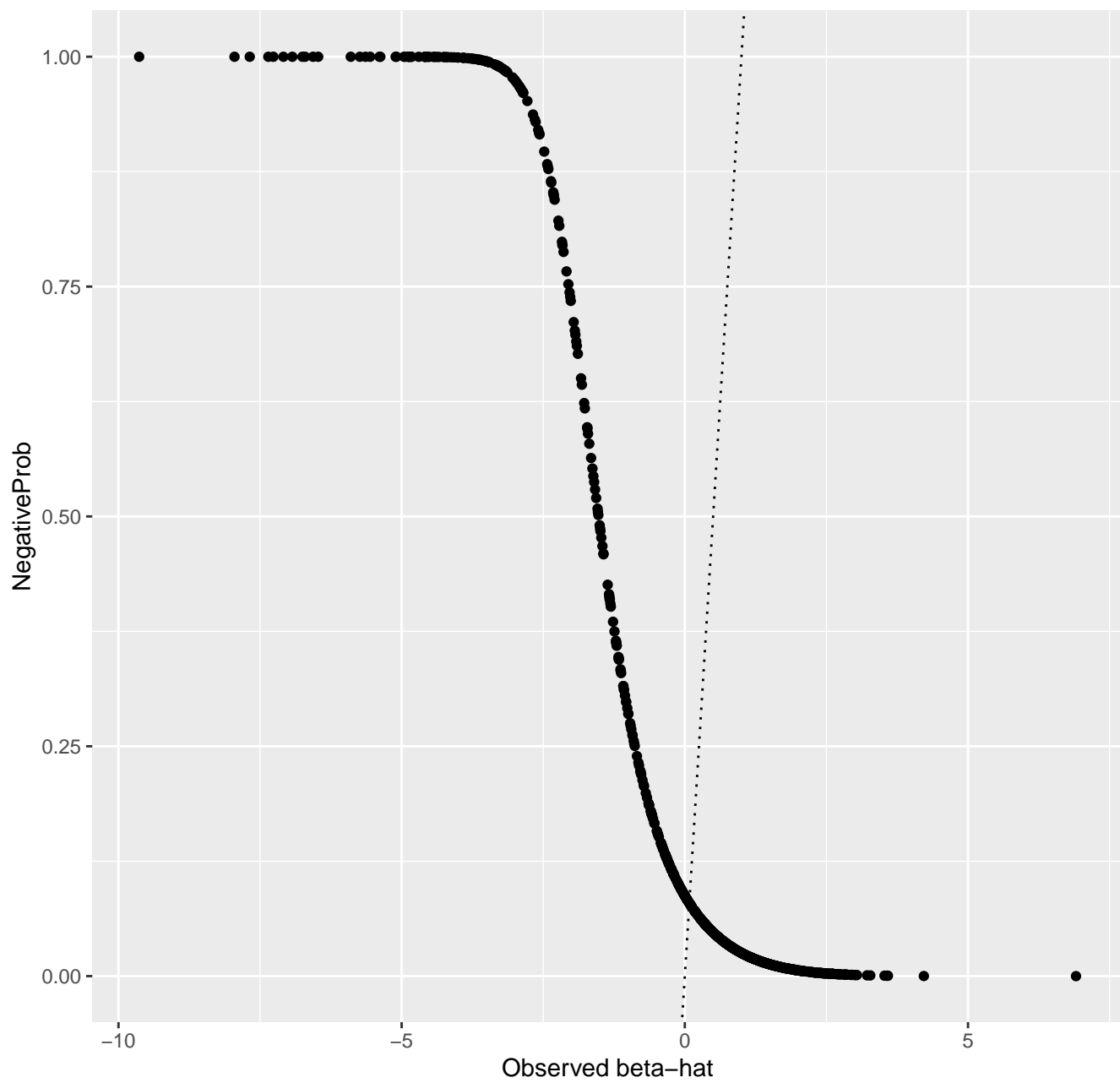
NRAS_(4893) negative



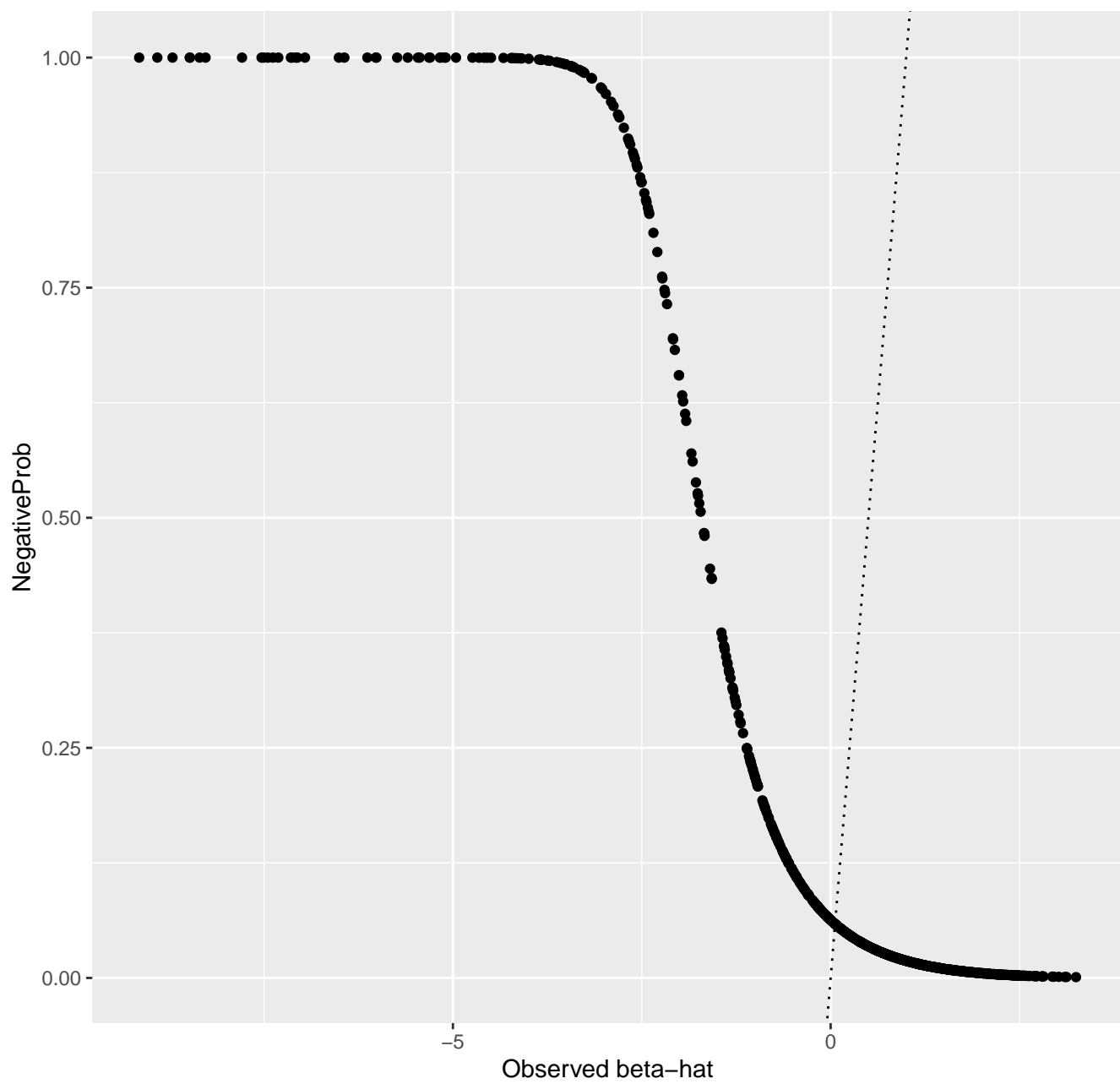
IER3IP1_(51124) negative



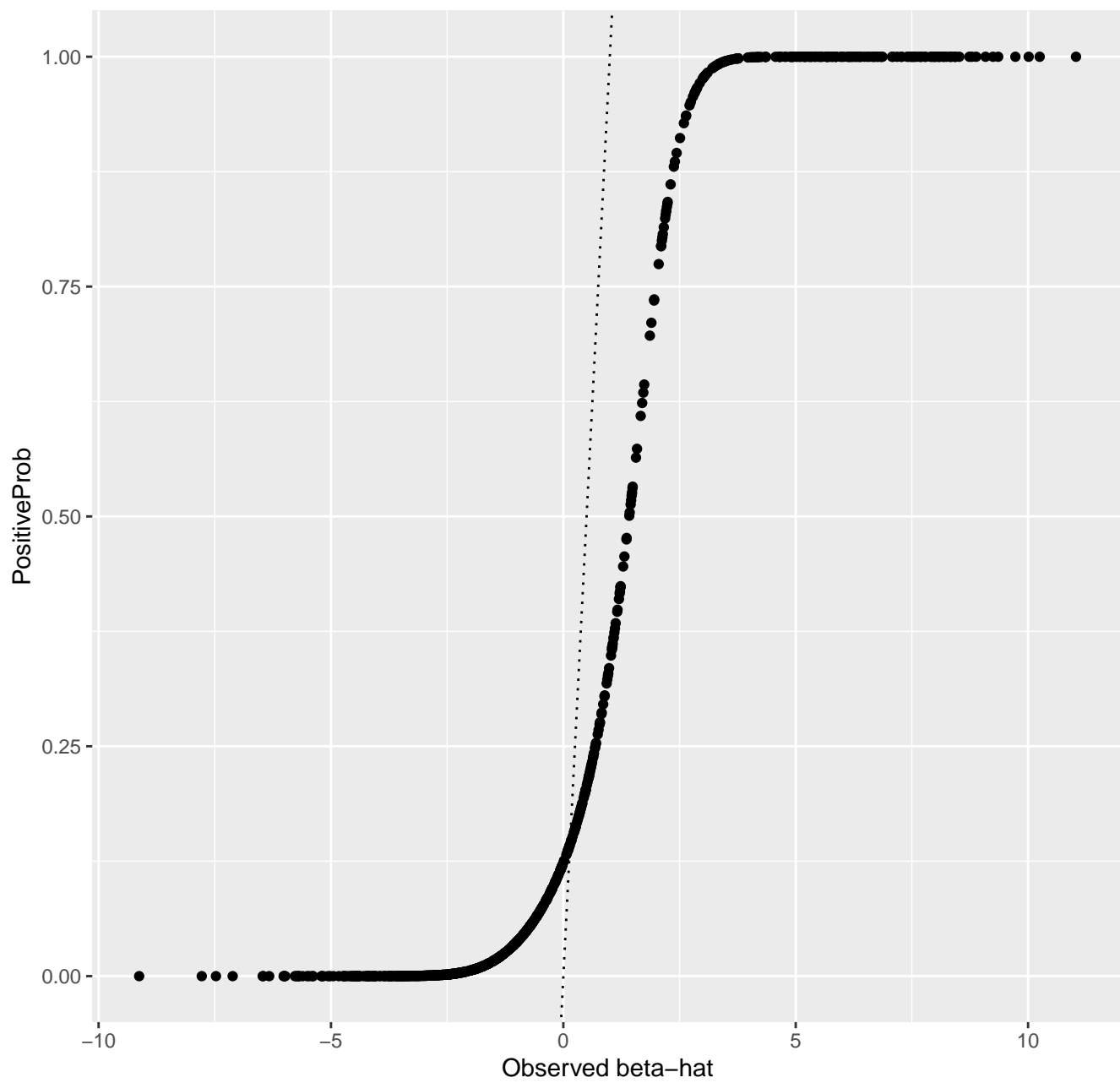
SOX9_(6662) negative



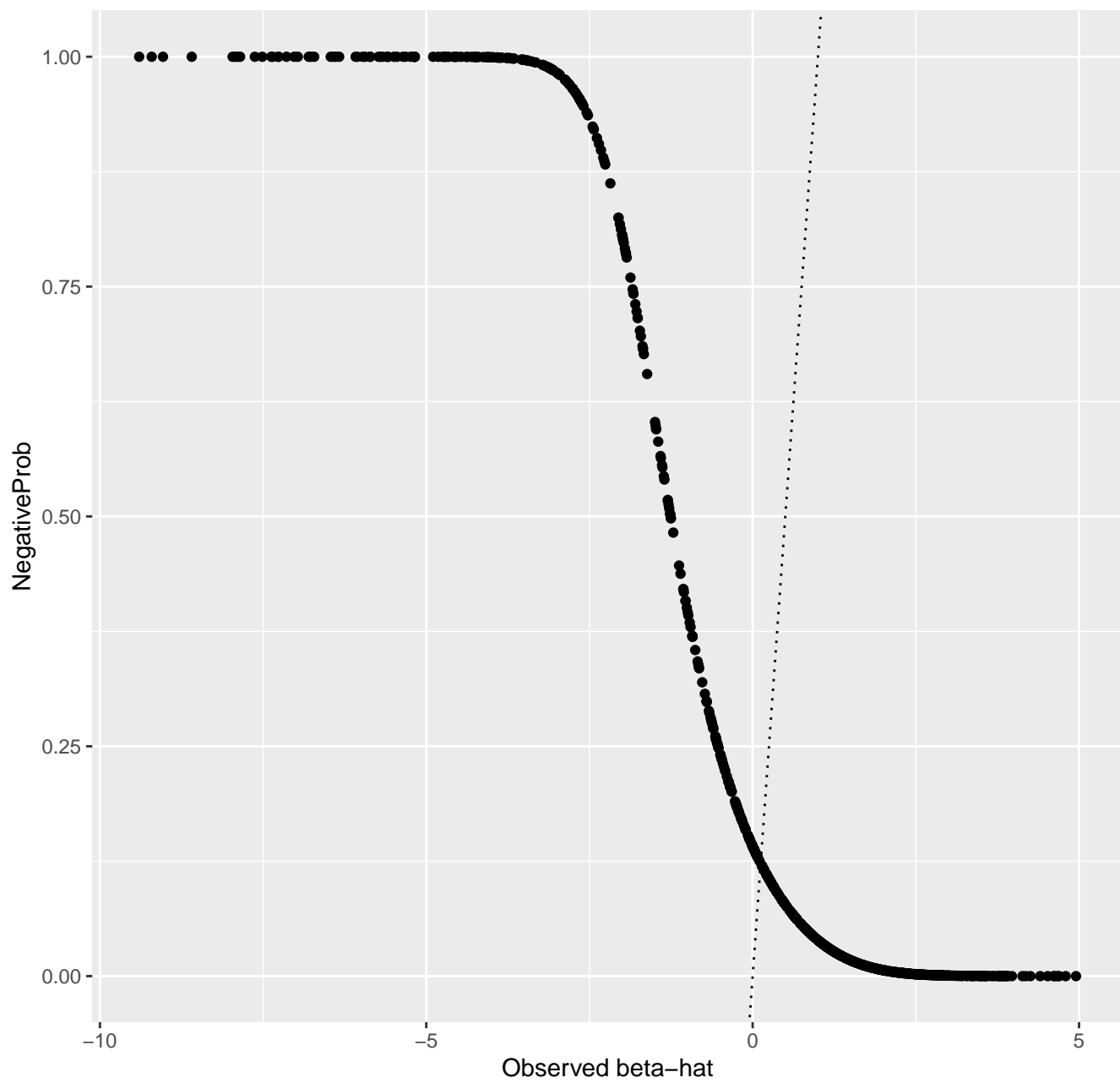
CCDC51_(79714) negative



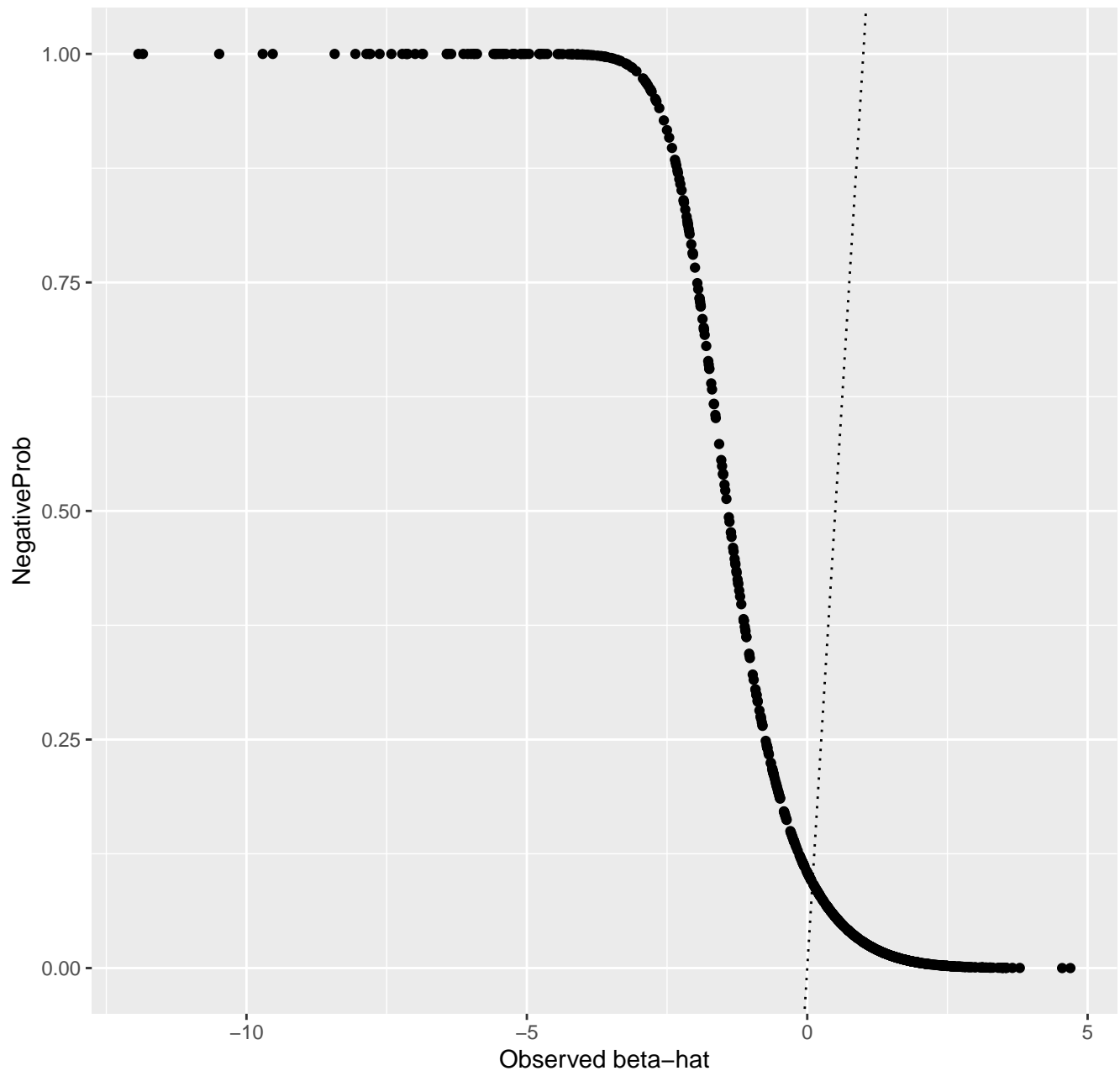
YRDC_(79693) positive



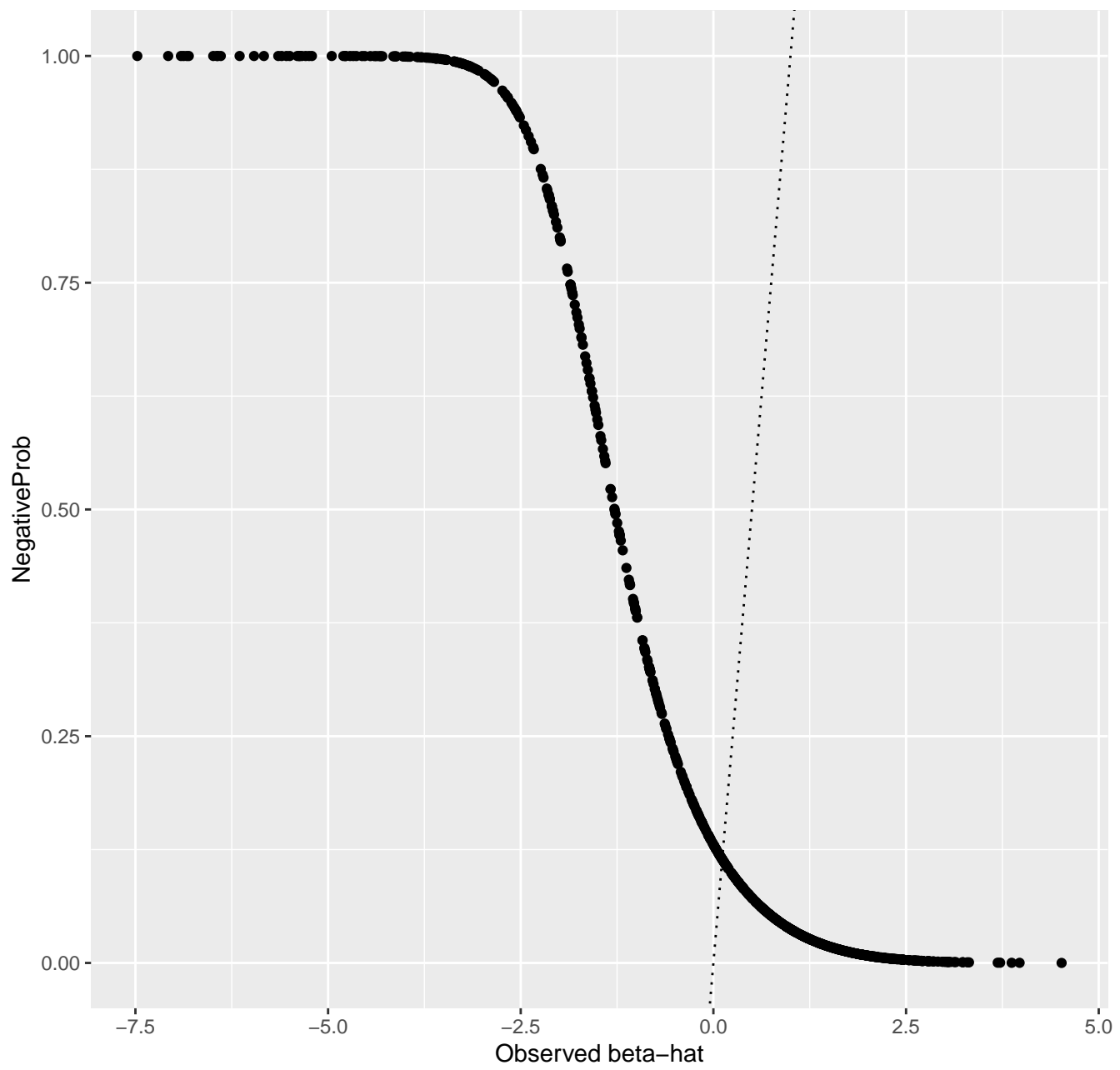
NAMPT_(10135) negative



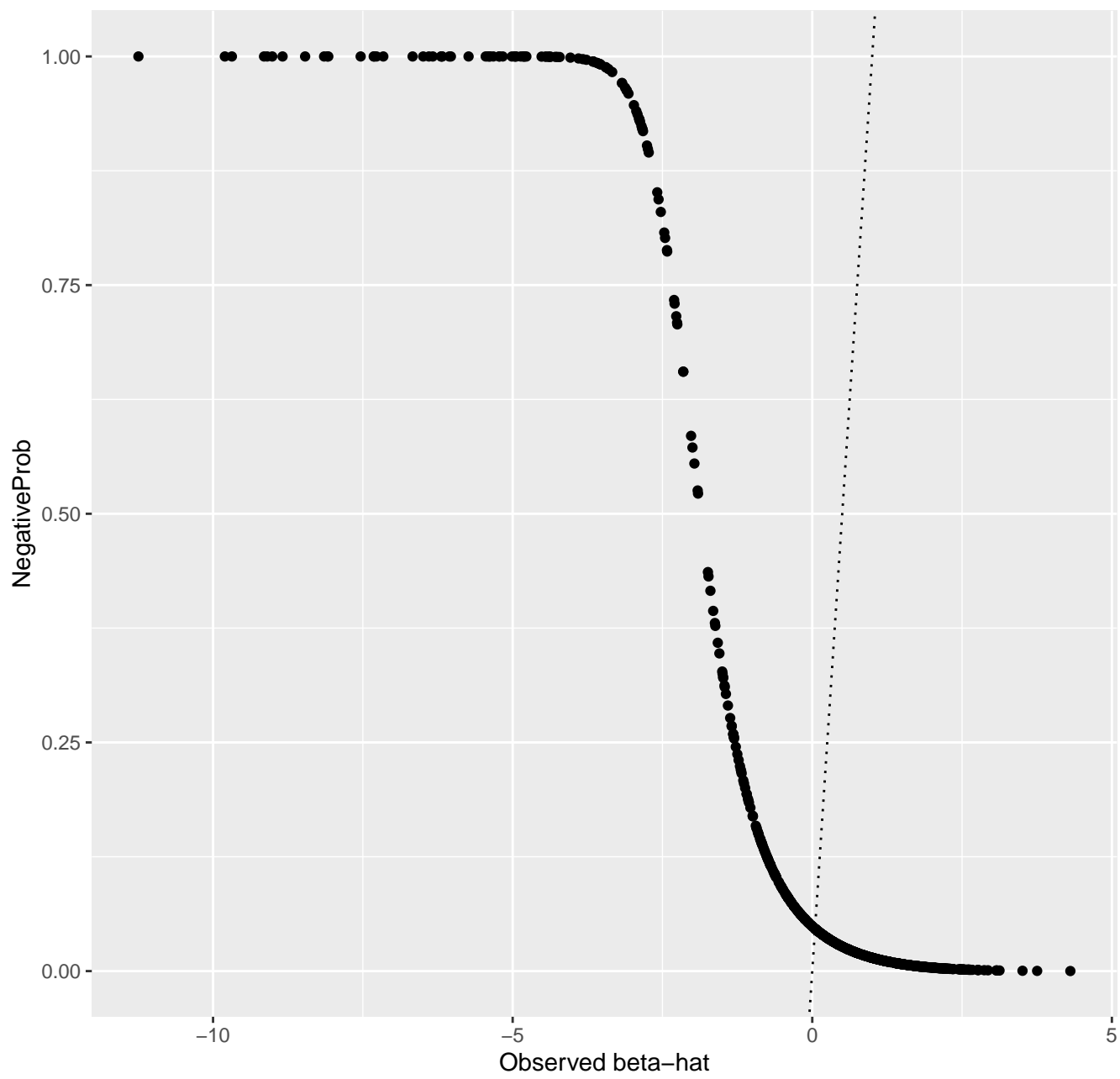
SREBF1_(6720) negative



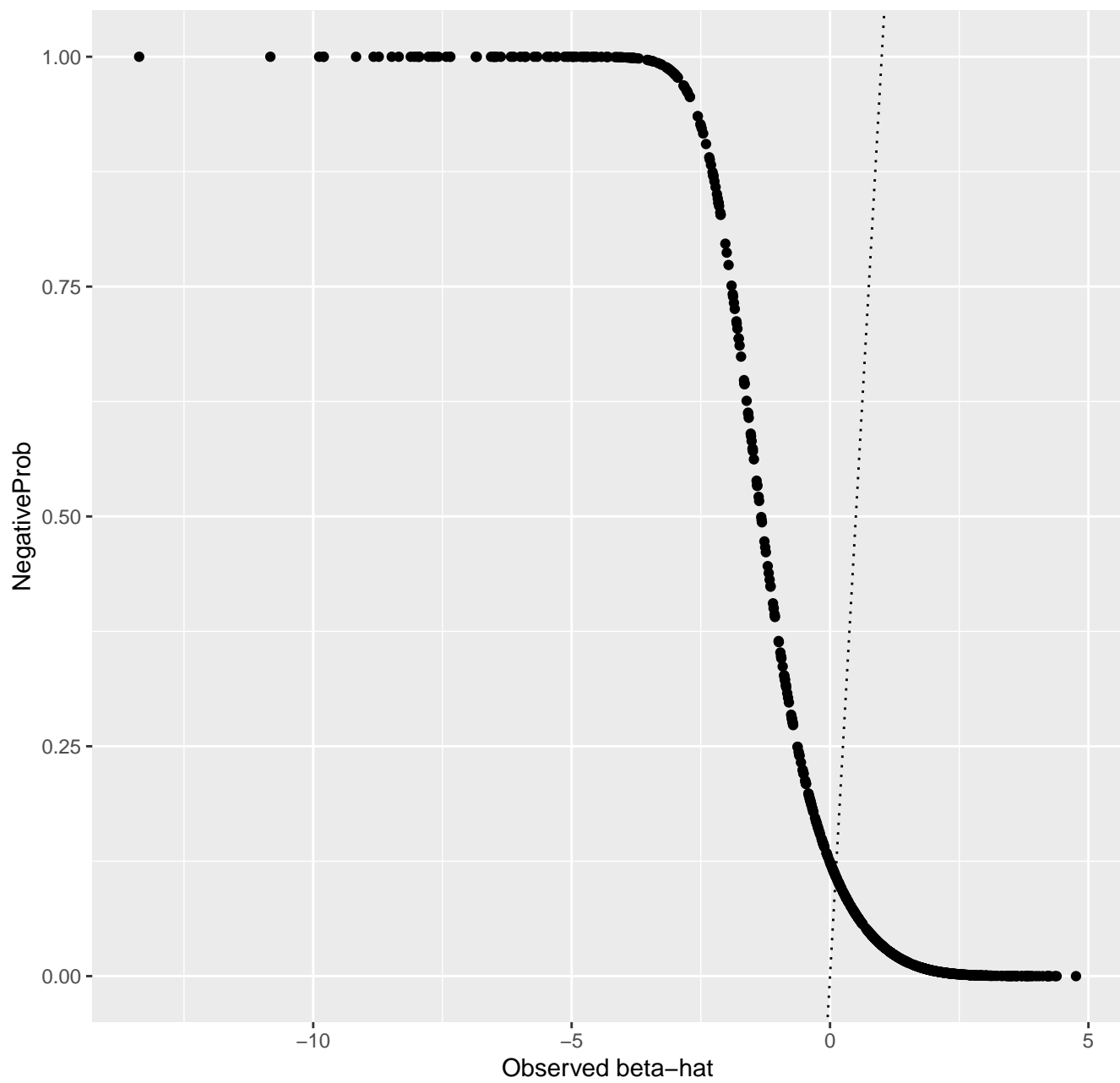
ELMO2_(63916) negative



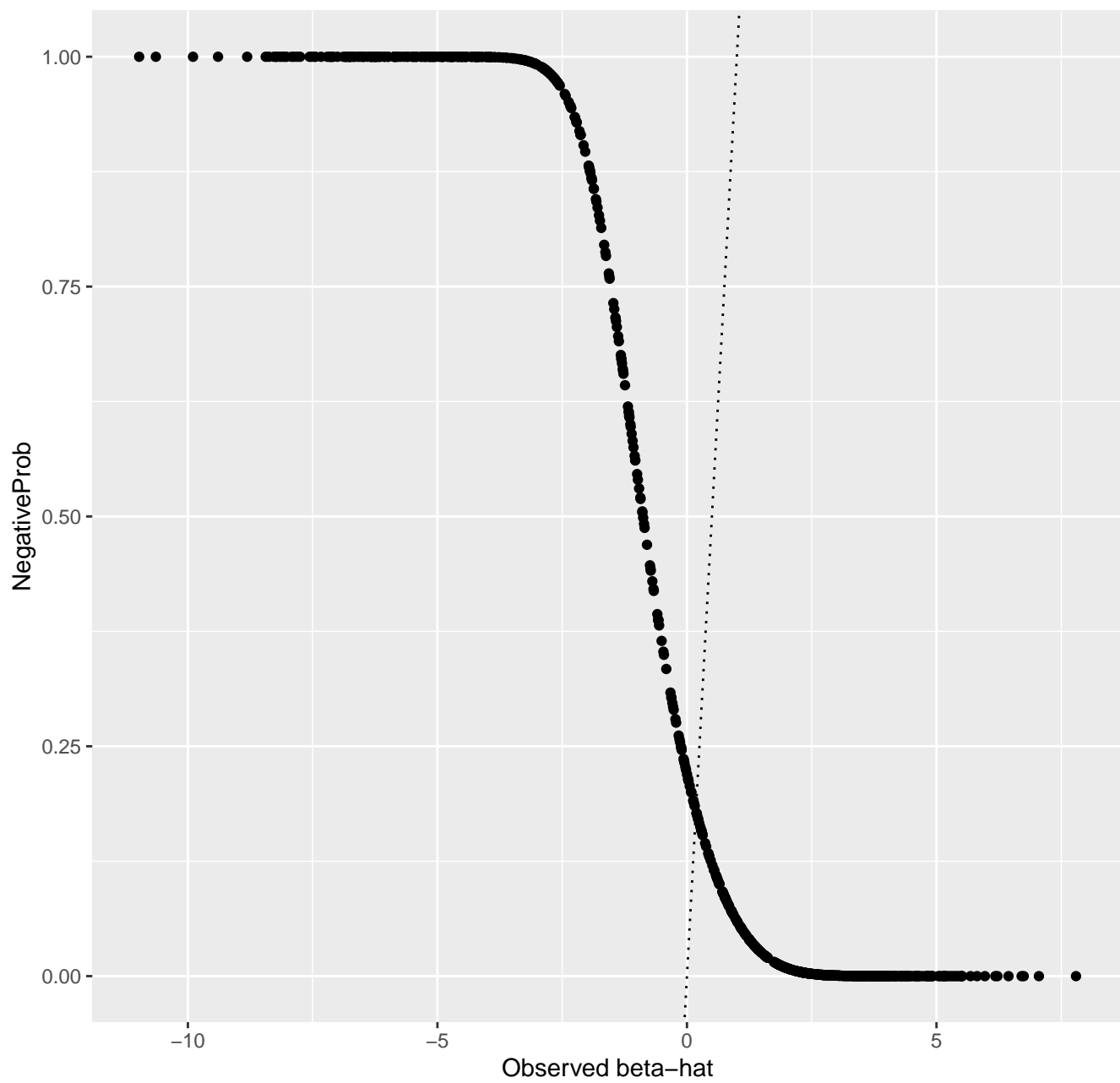
VPS4A_(27183) negative



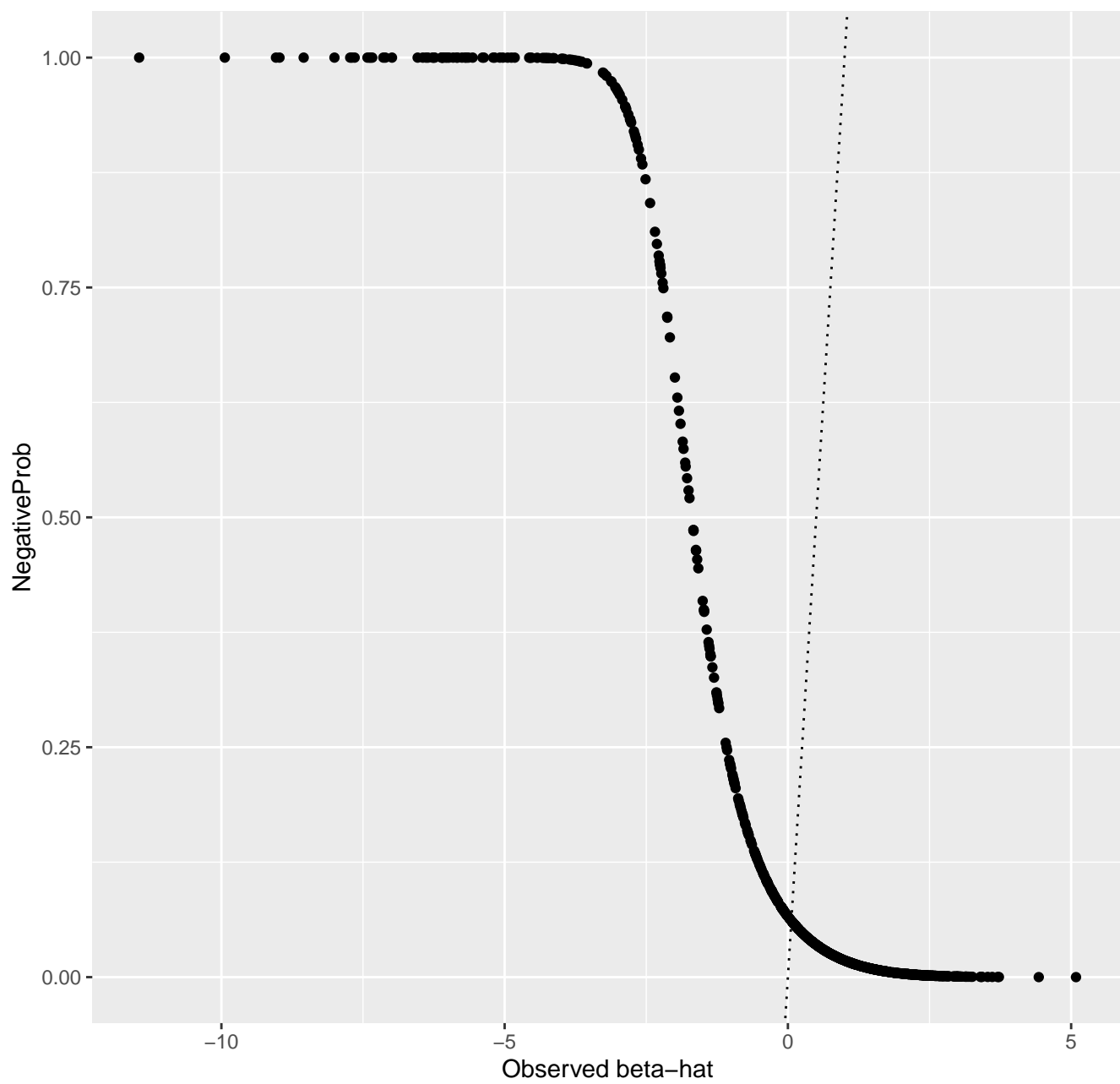
CDK4_(1019) negative



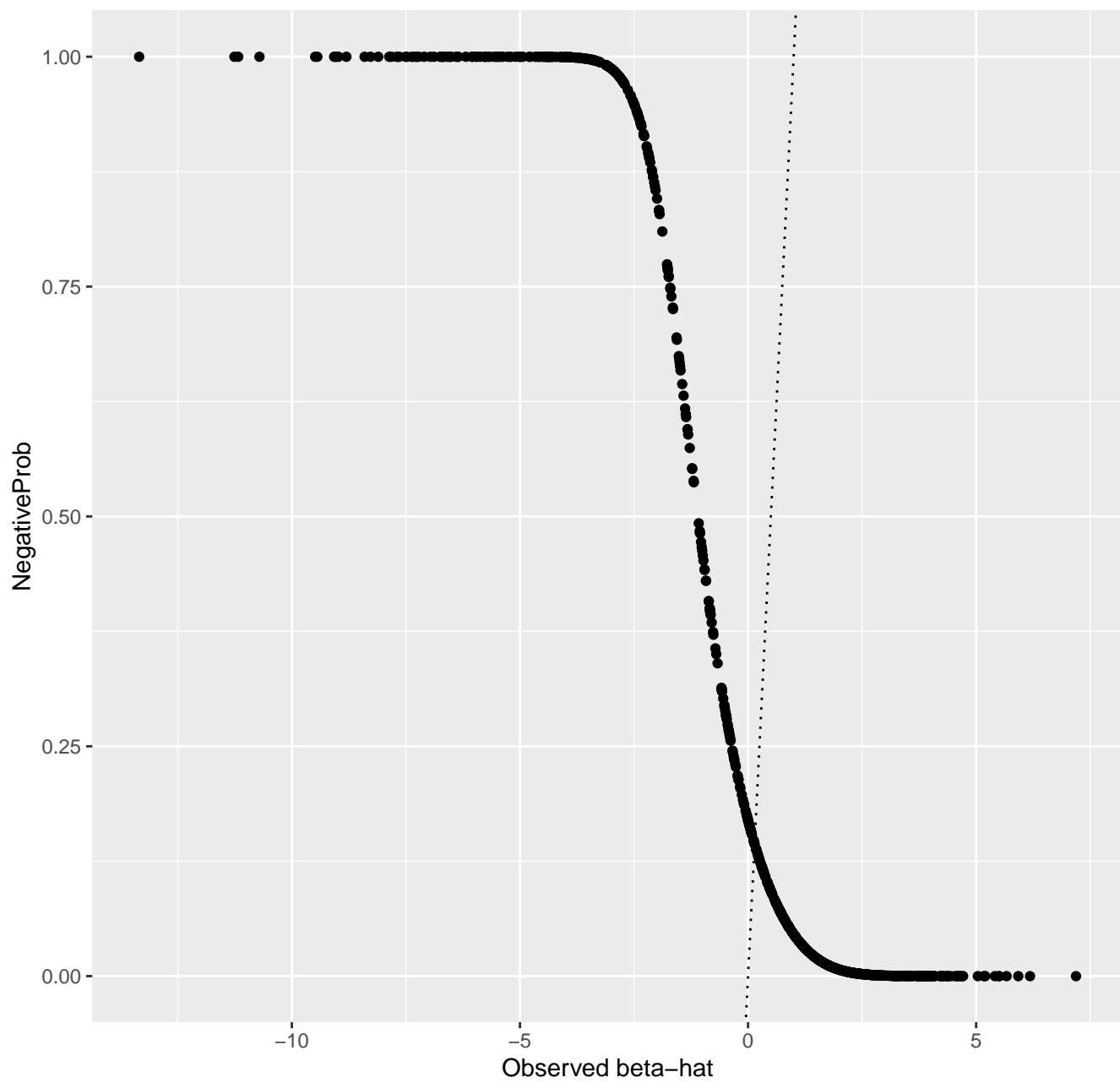
TYMS_(7298) negative



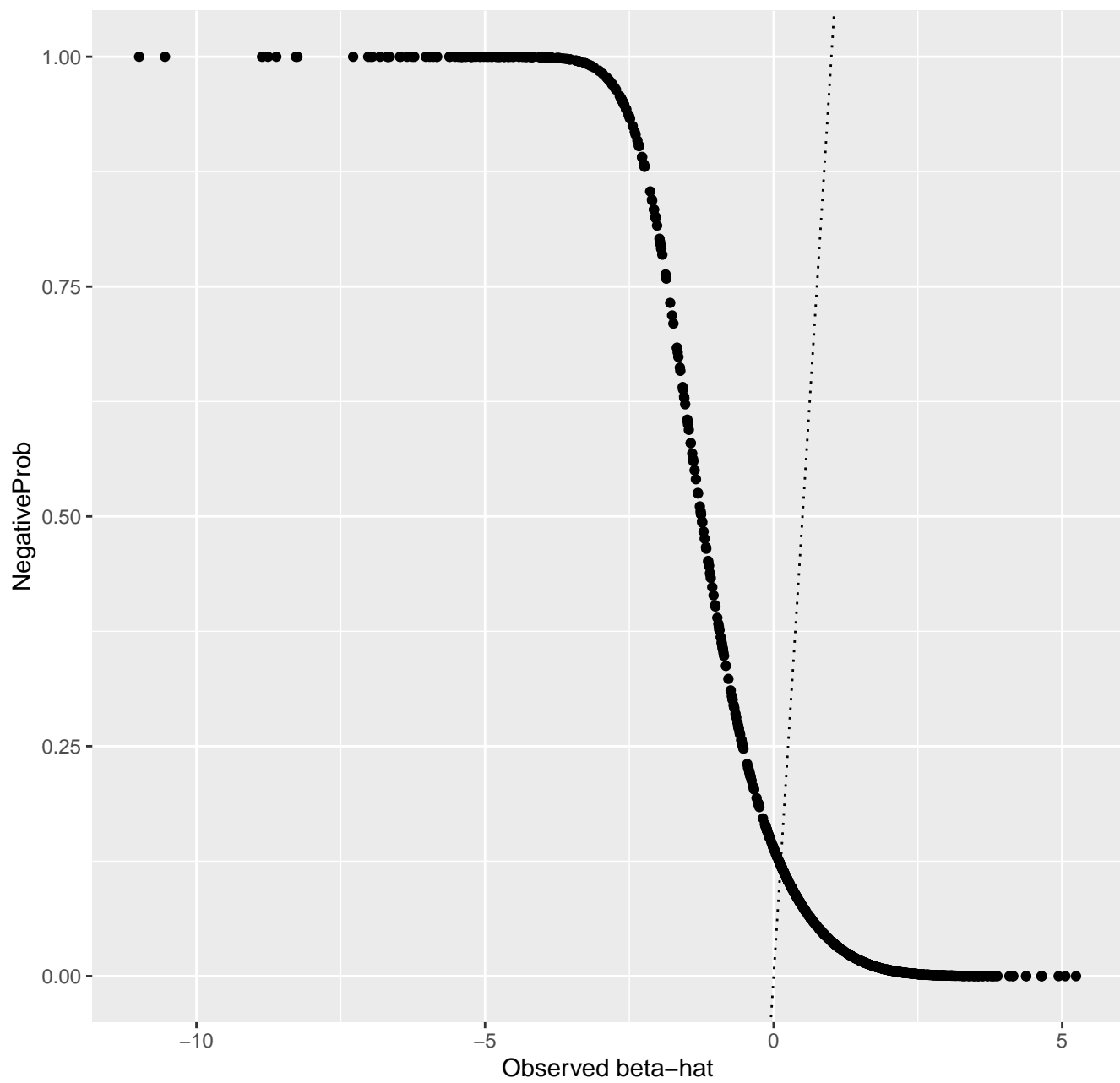
NMNAT1_(64802) negative



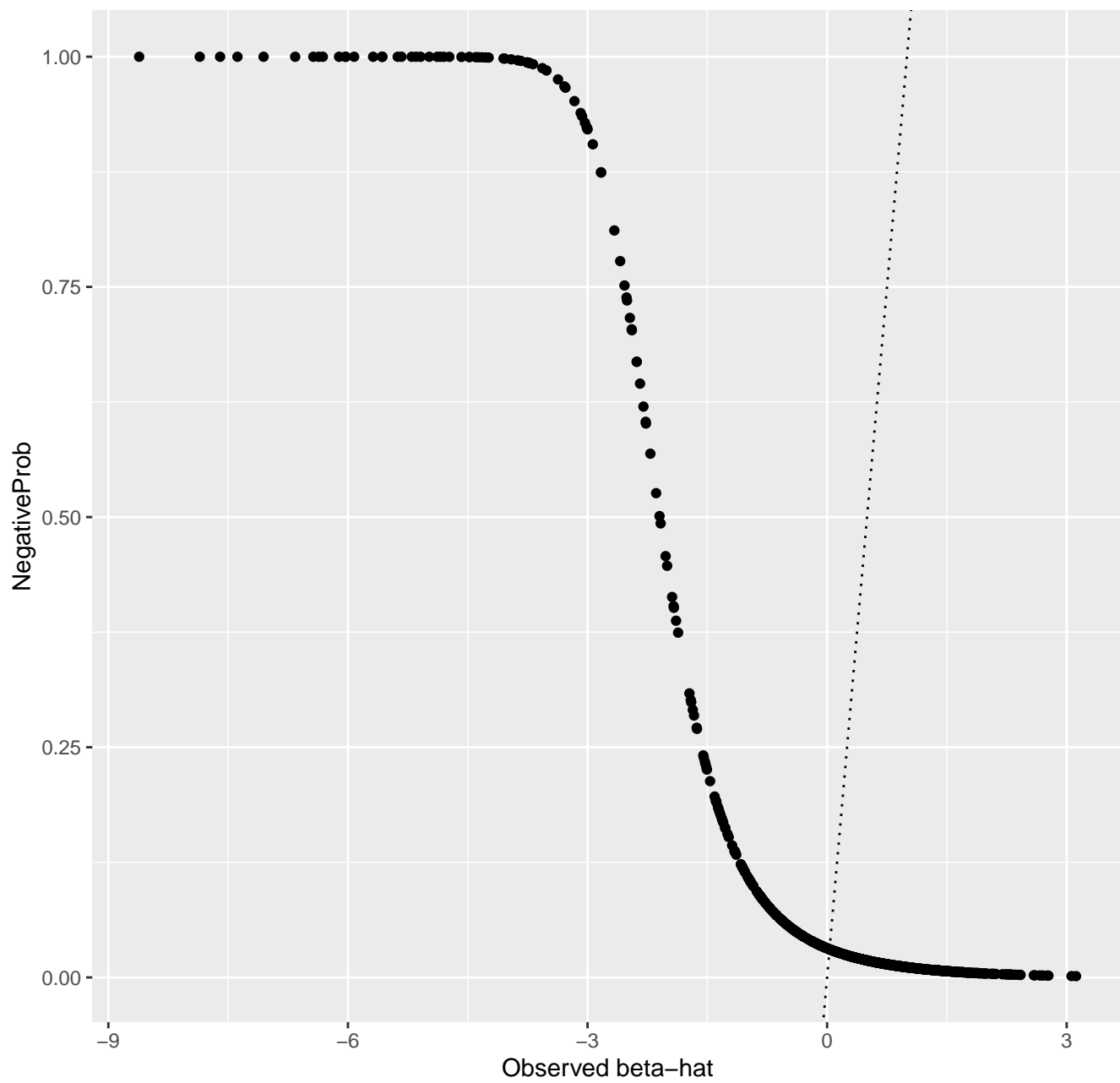
NXT1_(29107) negative



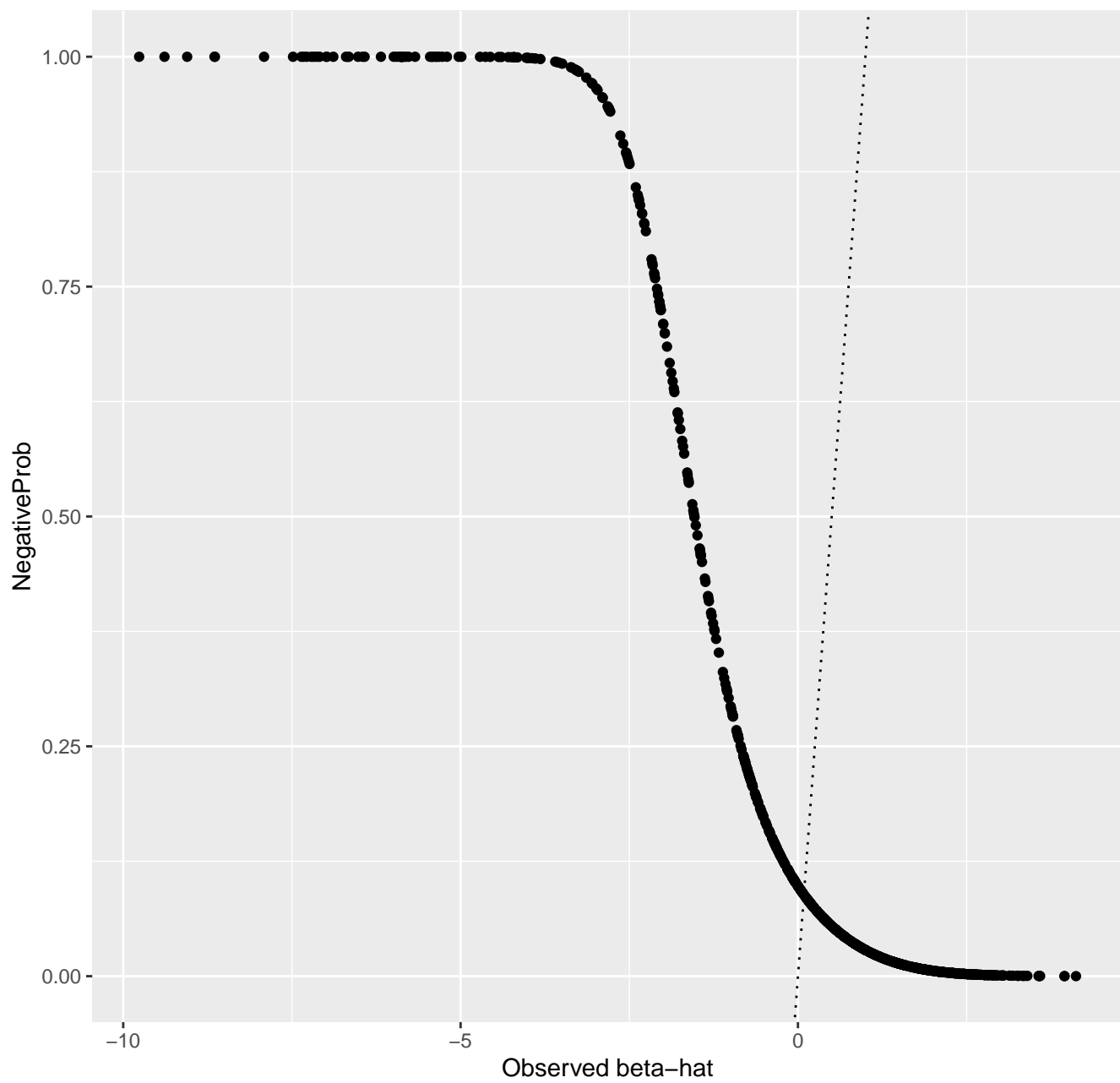
MARCH5_(54708) negative



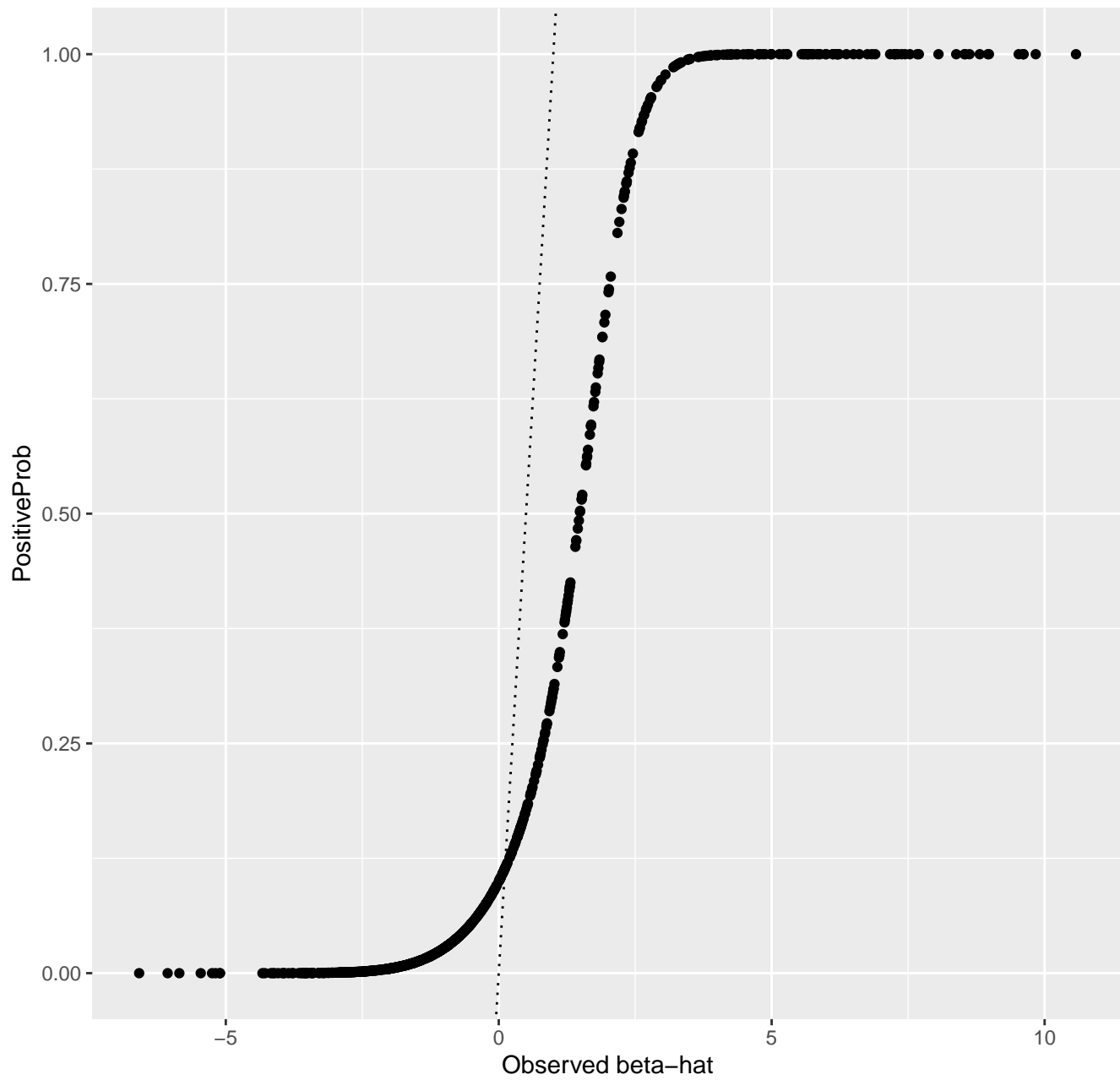
PRDX1_(5052) negative



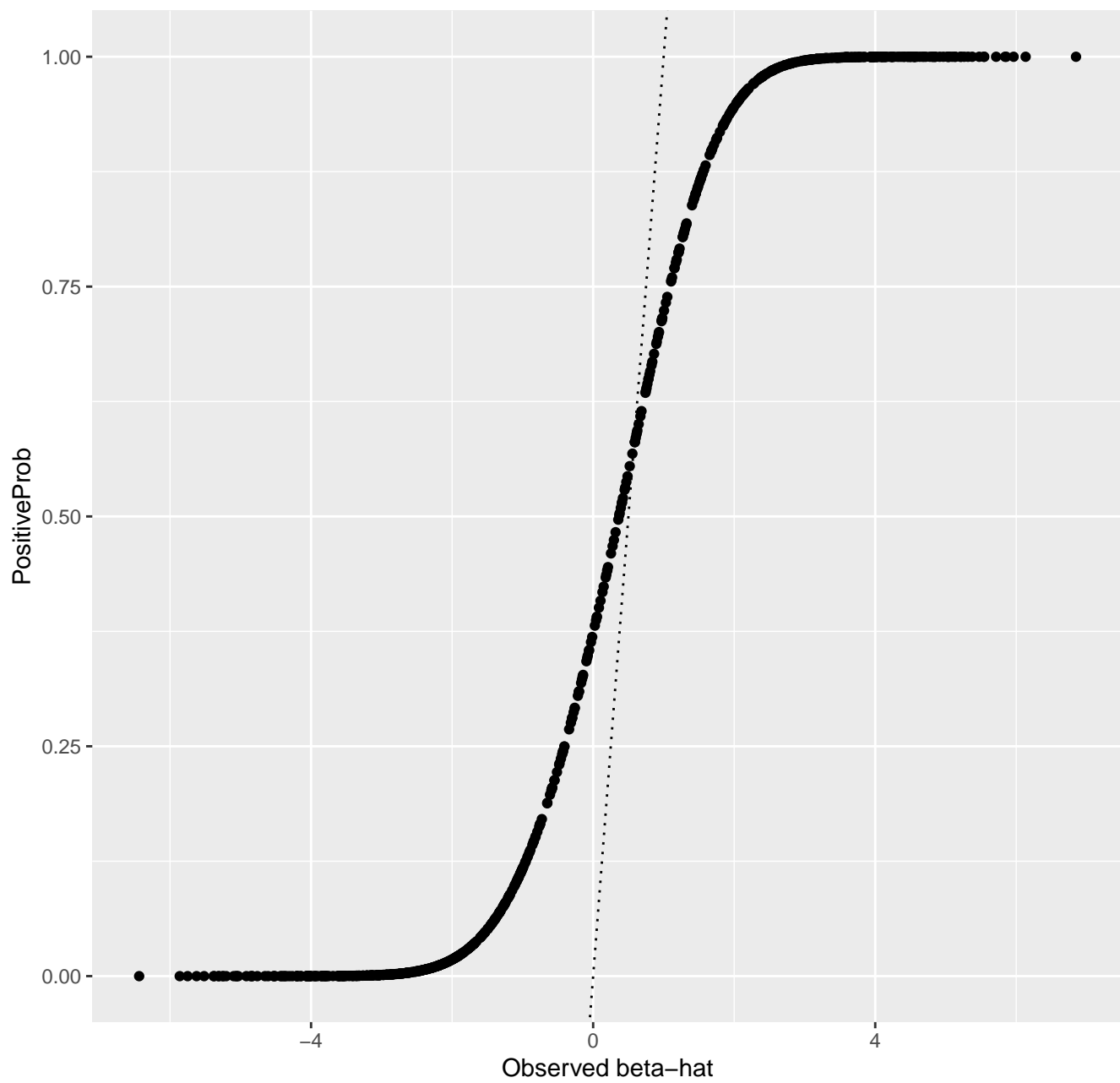
TUBB4B_(10383) negative



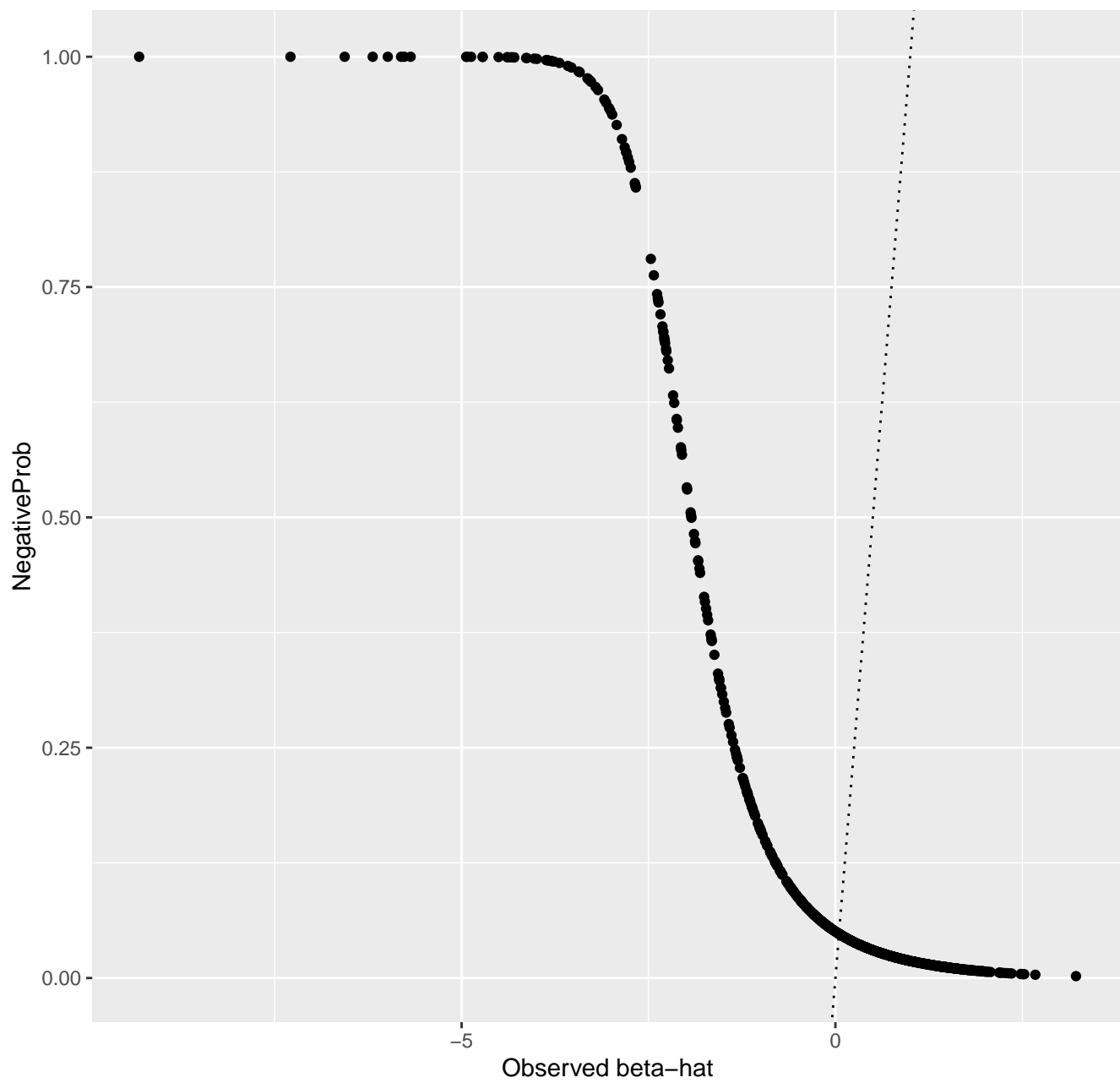
EIF1AX_(1964) positive



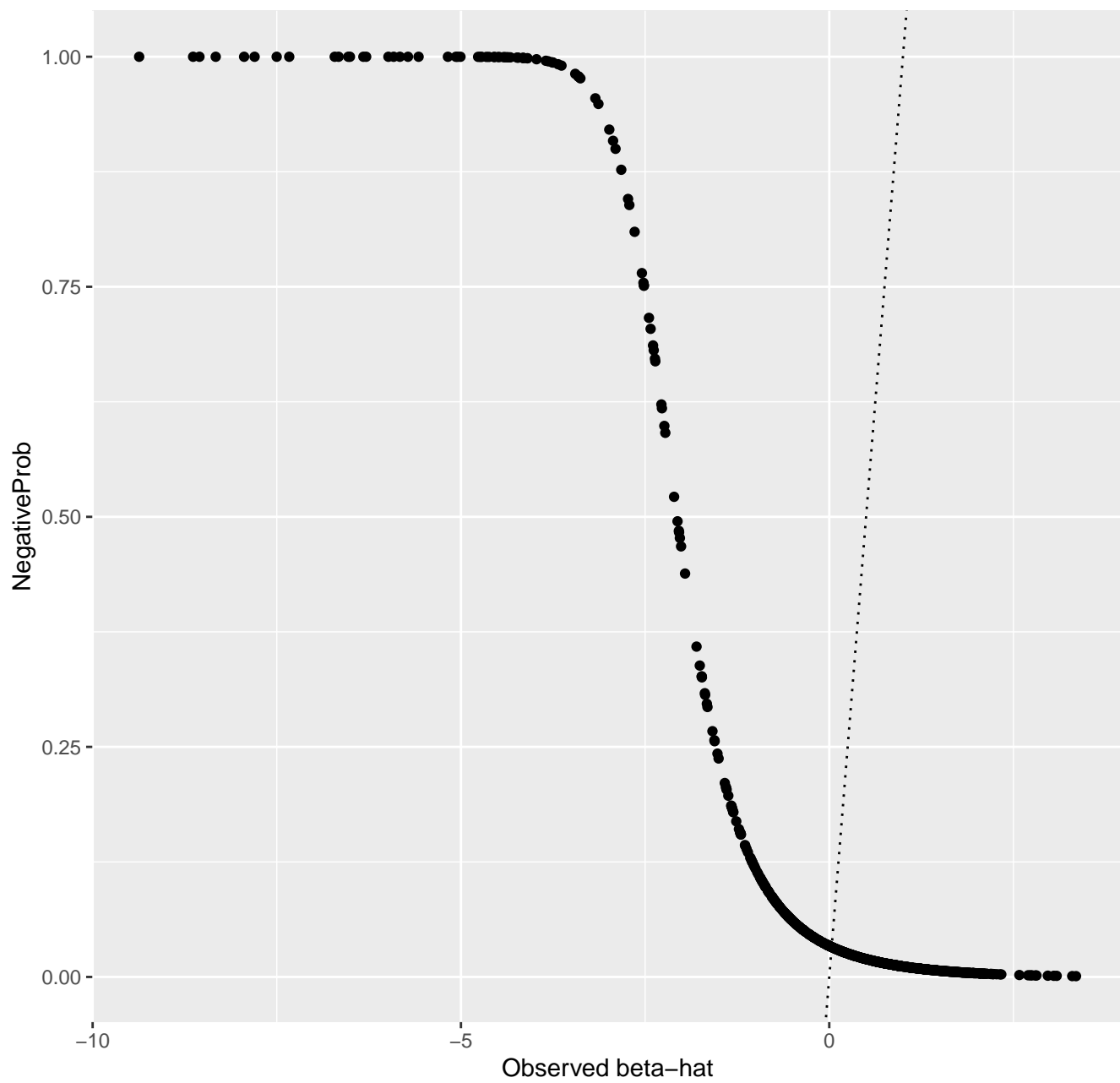
GPX4_(2879) positive



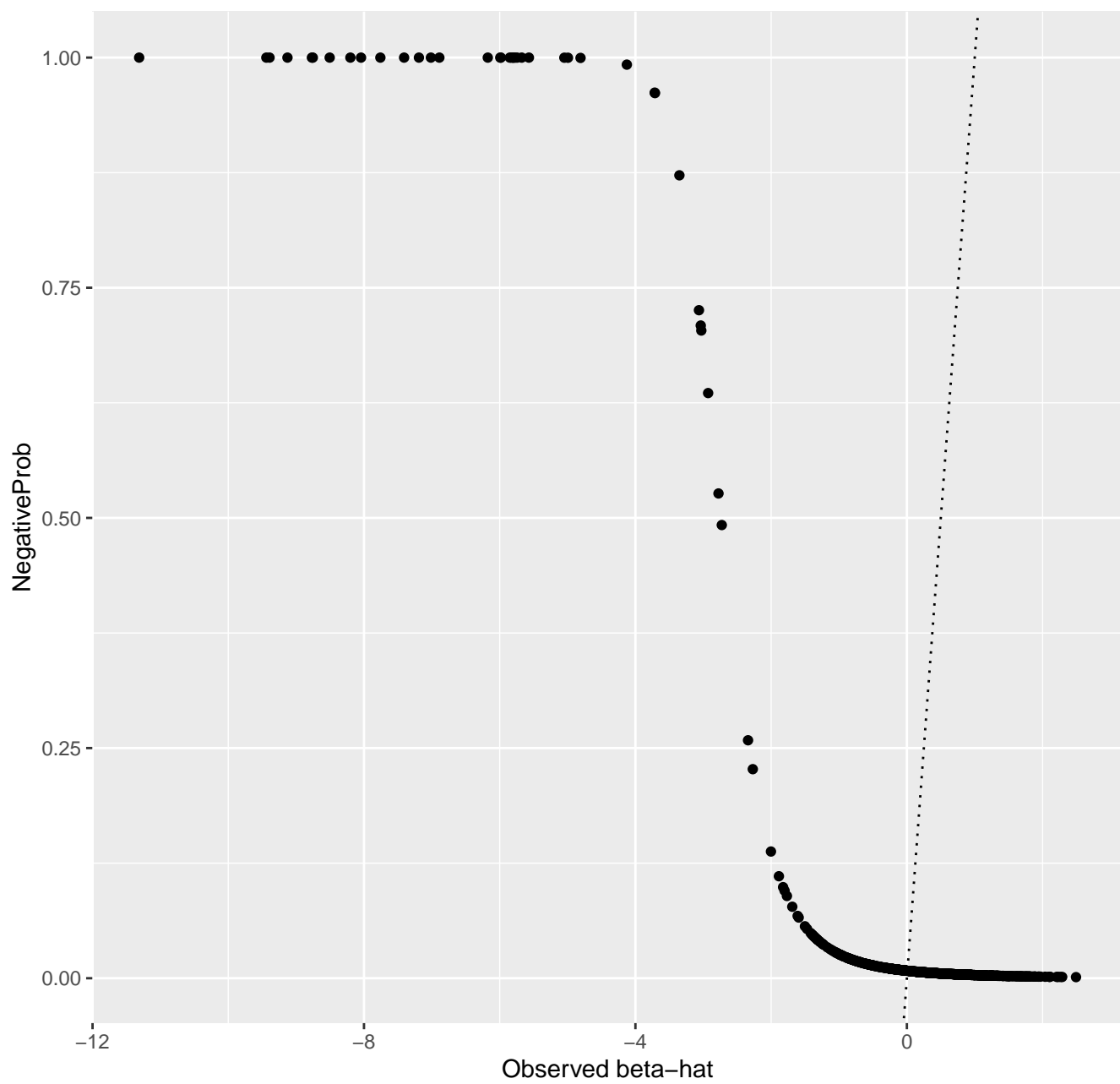
GAB1_(2549) negative



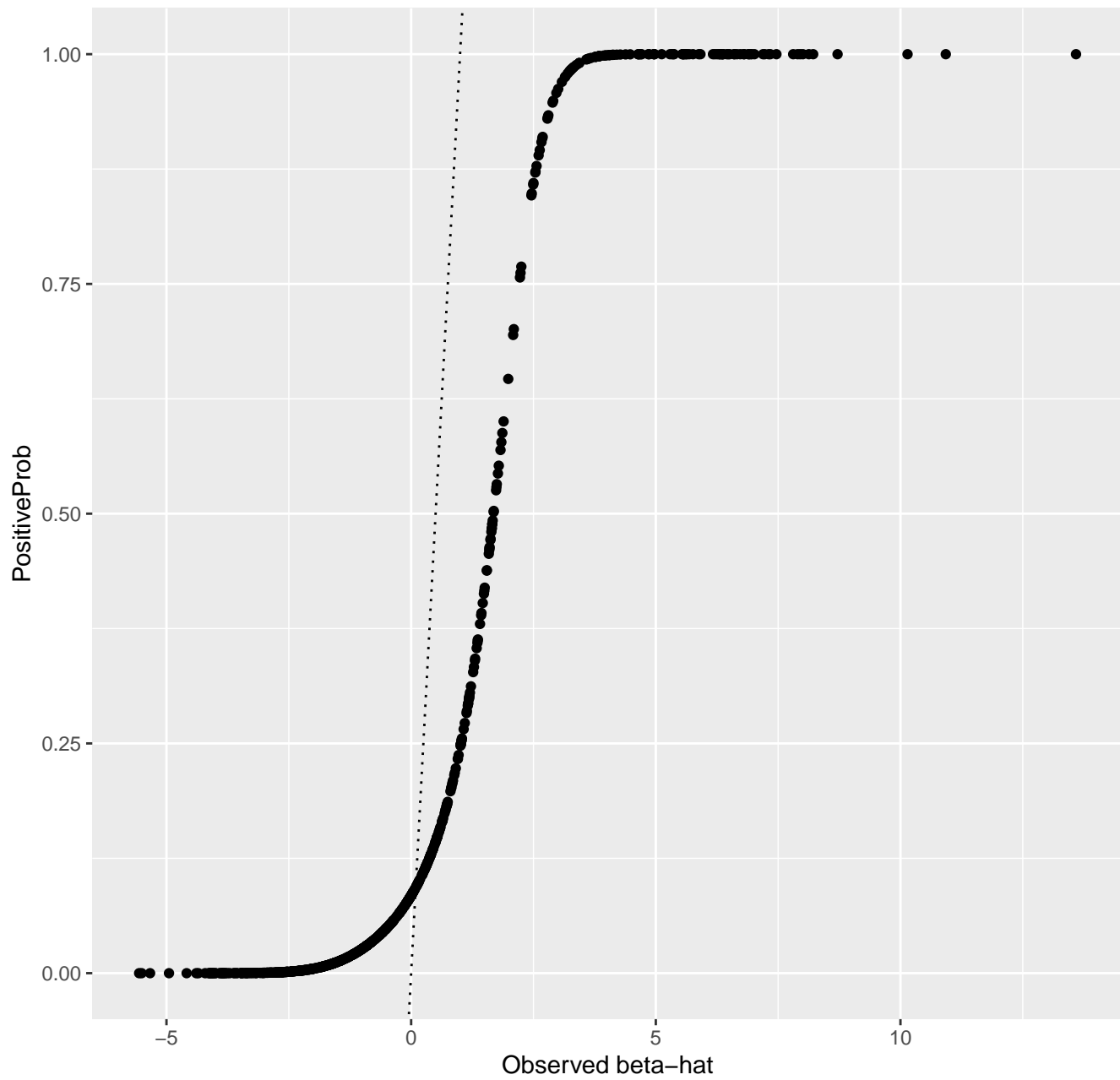
ATP6V0E1_(8992) negative



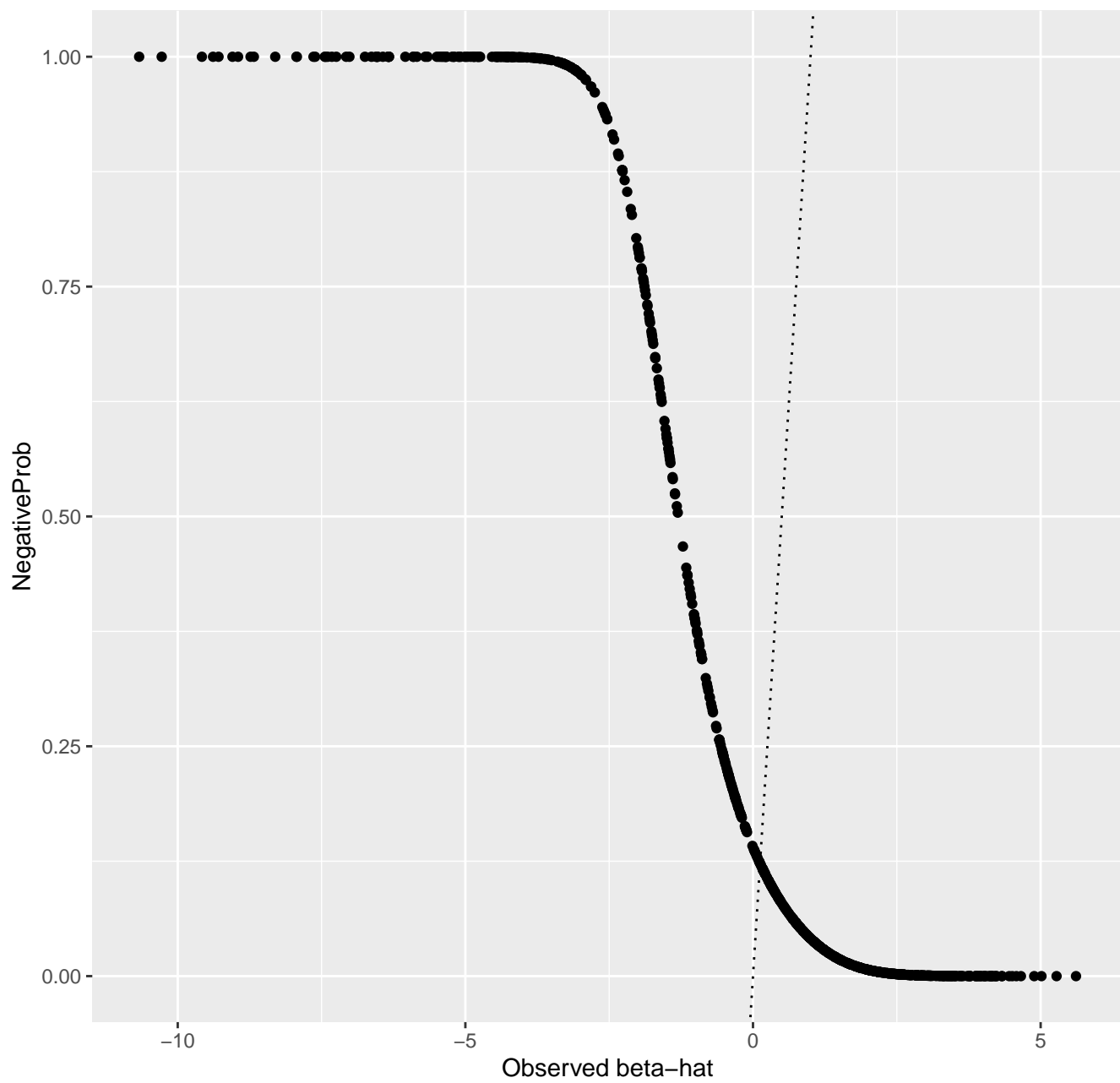
SMARCA2_(6595) negative



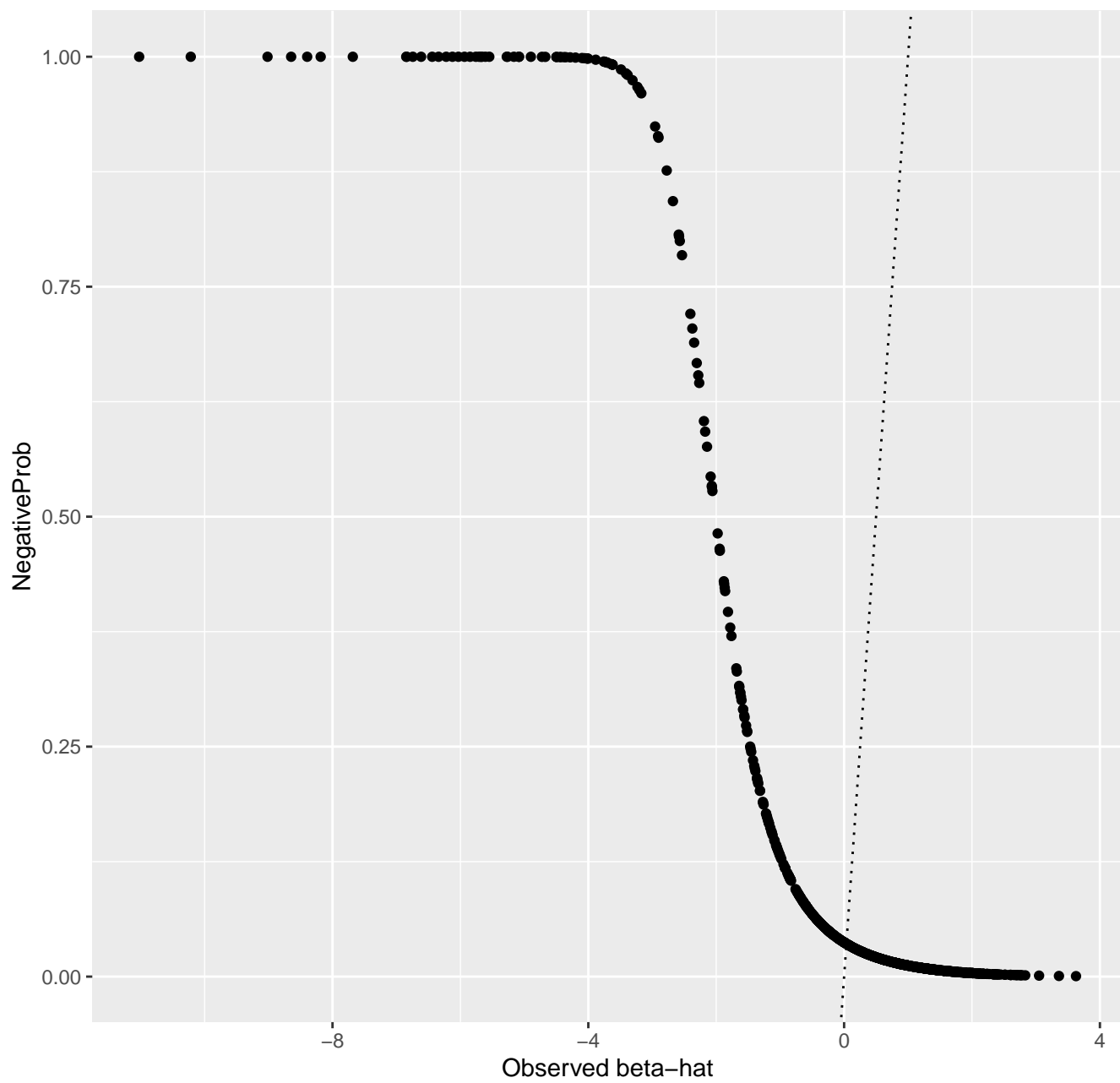
DDX3X_(1654) positive



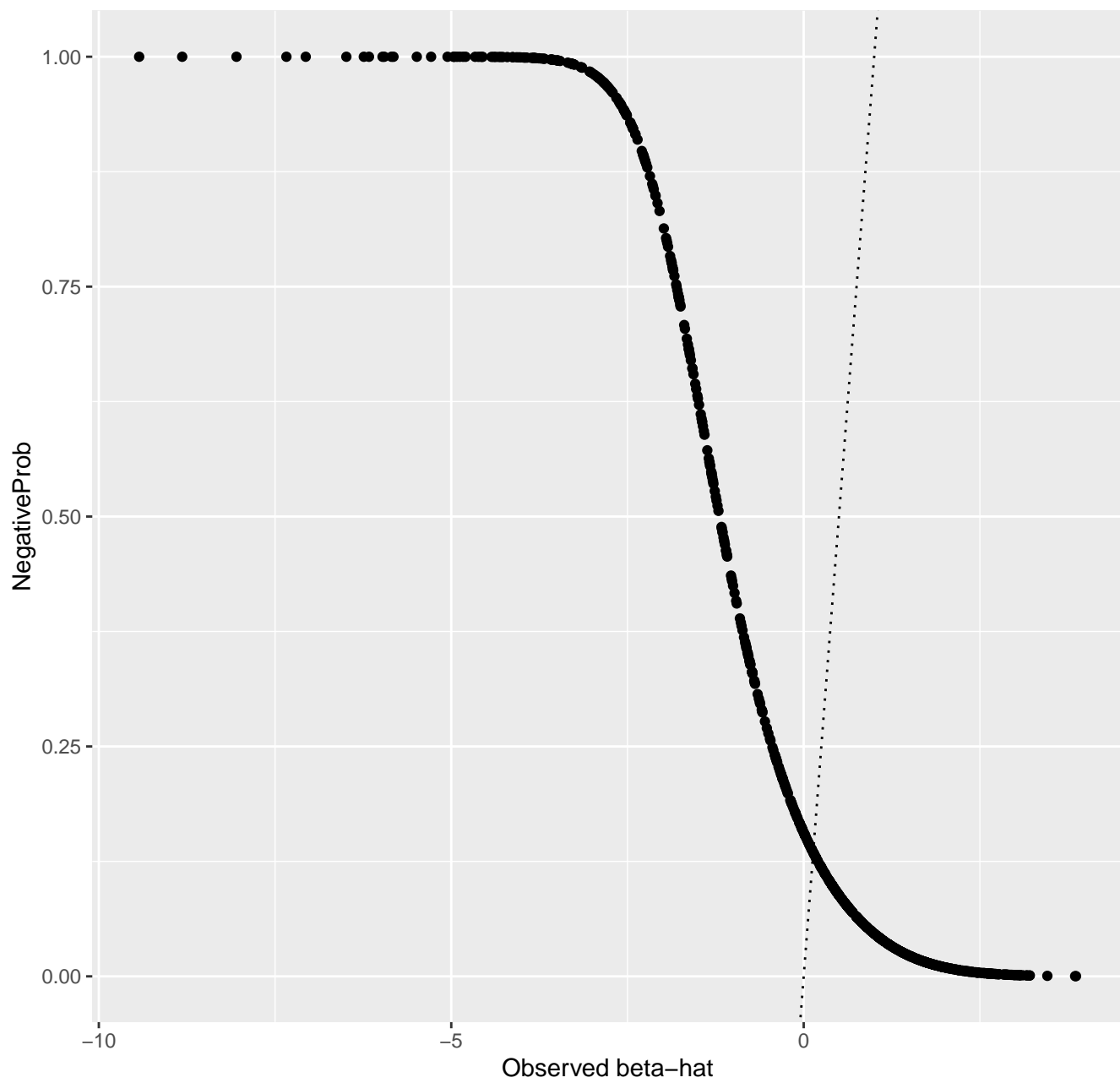
PRKRA_(8575) negative



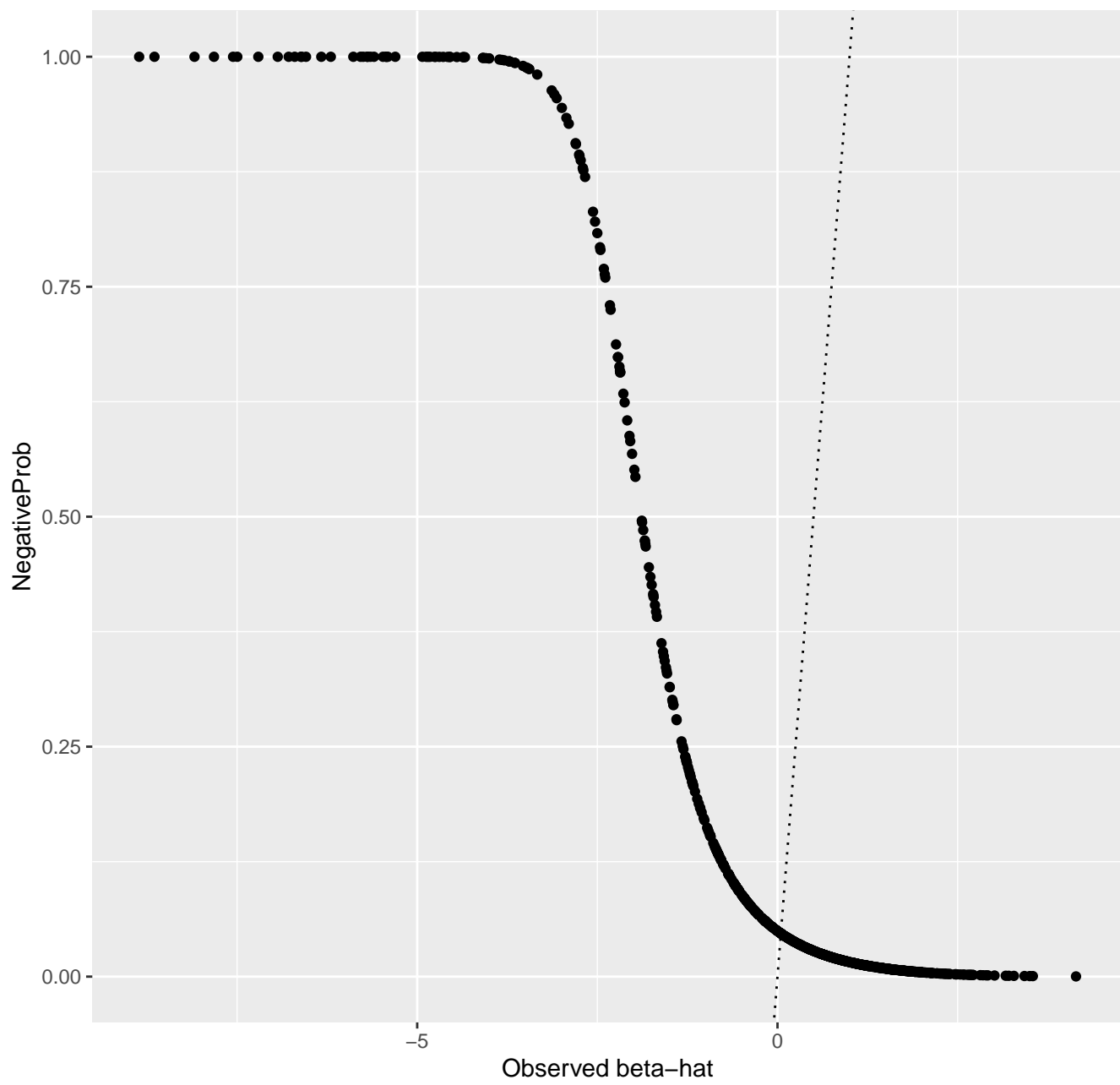
CCND3_(896) negative



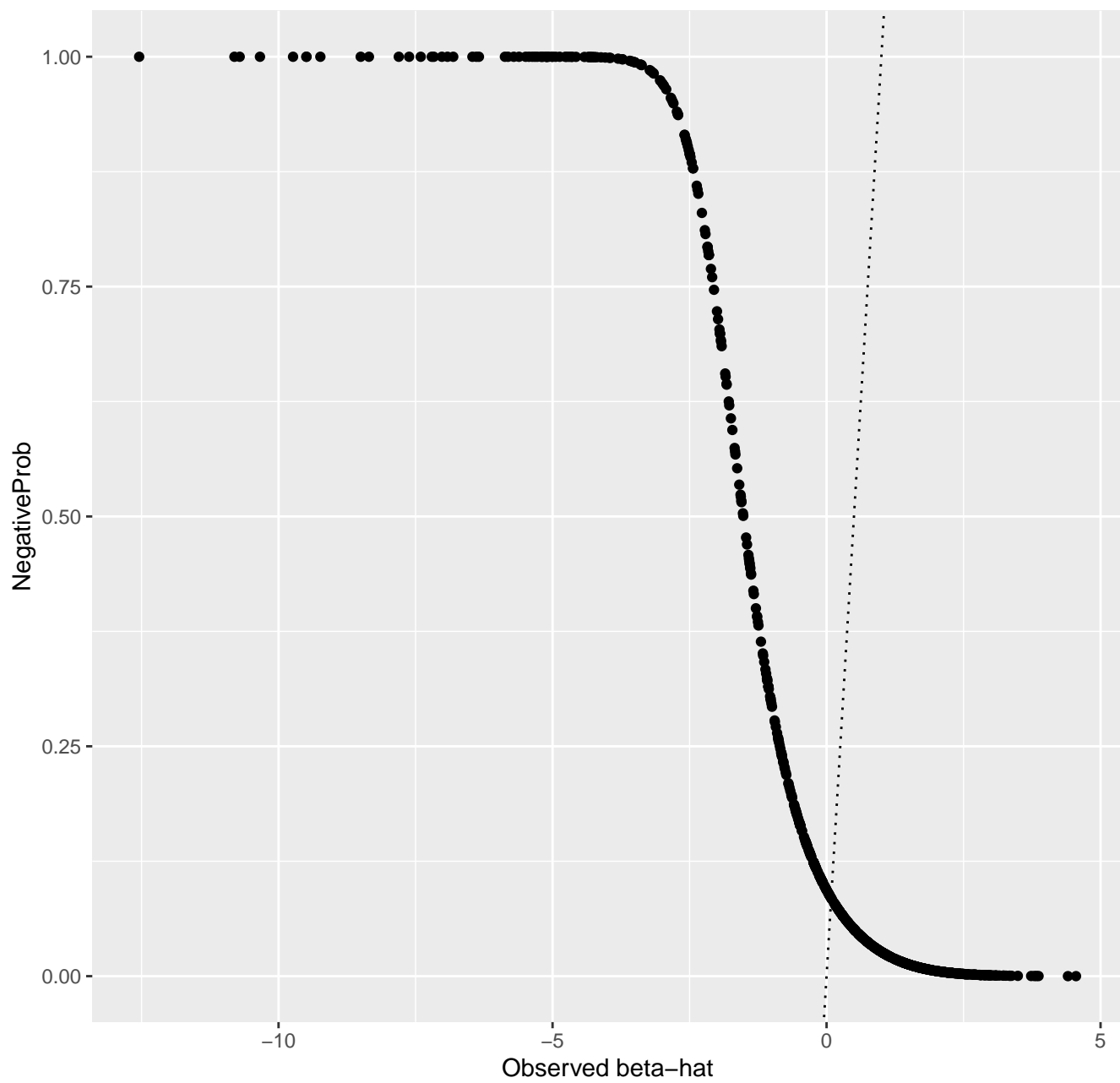
PIK3CA_(5290) negative



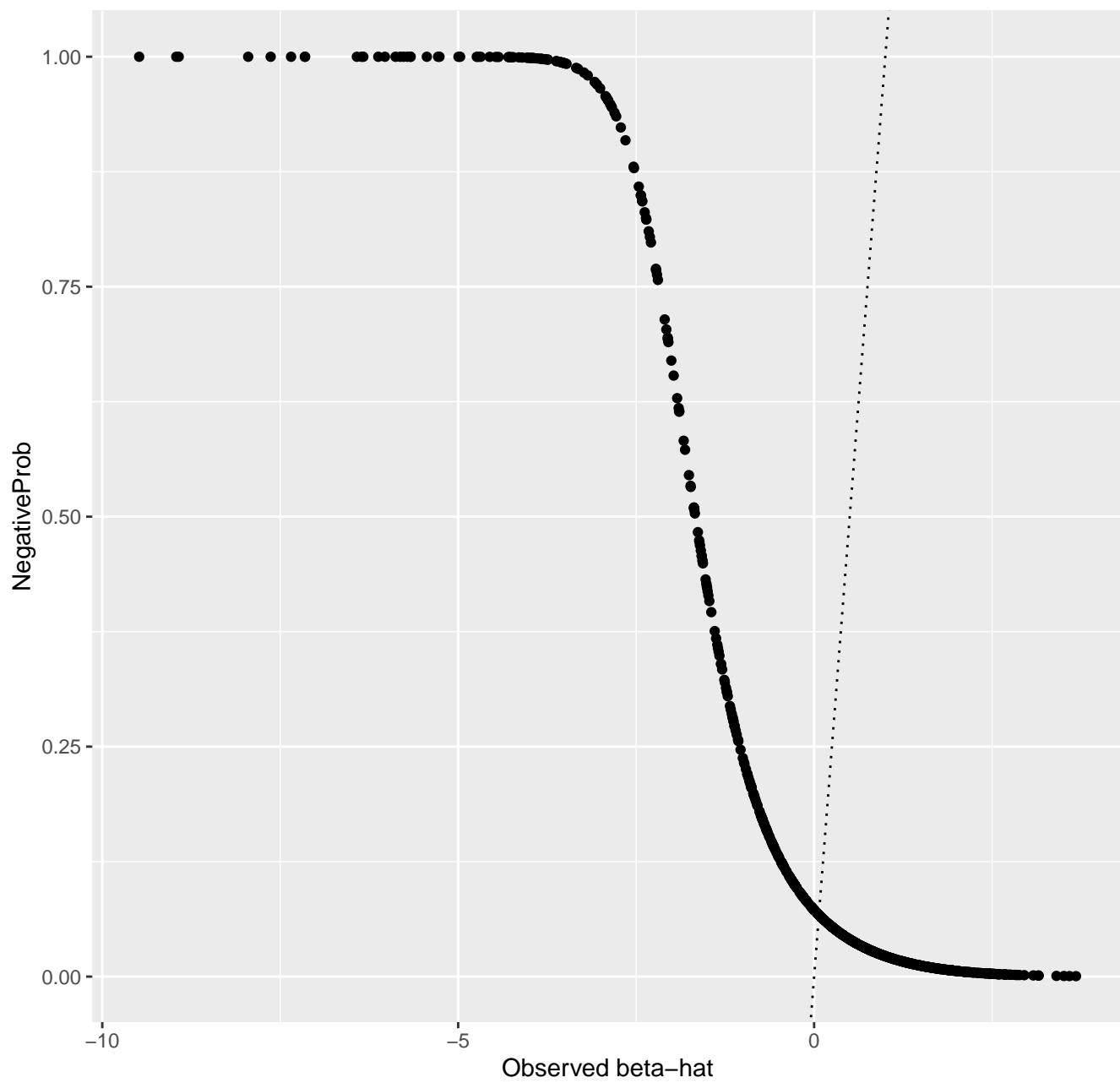
TIMM17A_(10440) negative



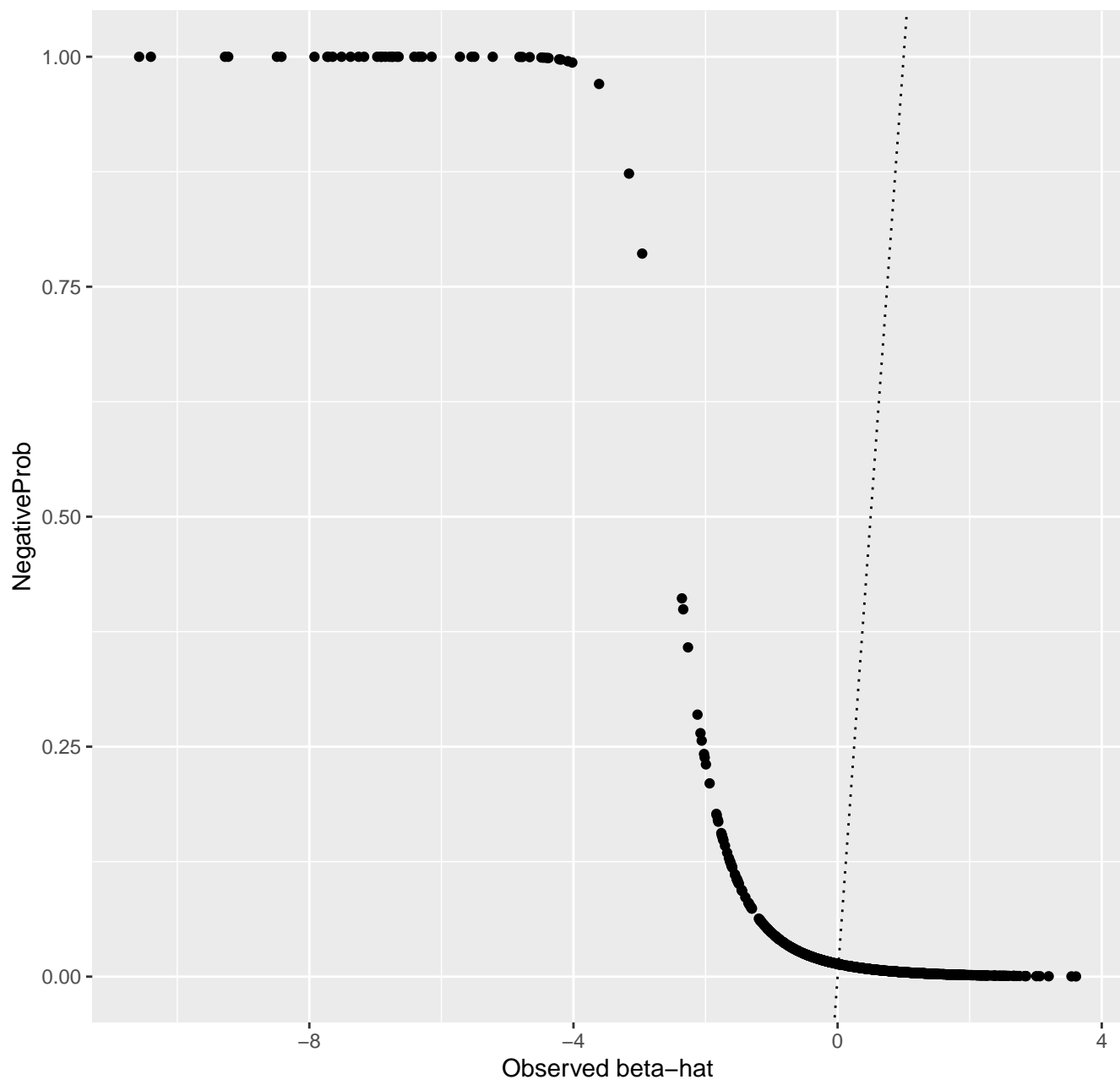
UXS1_(80146) negative



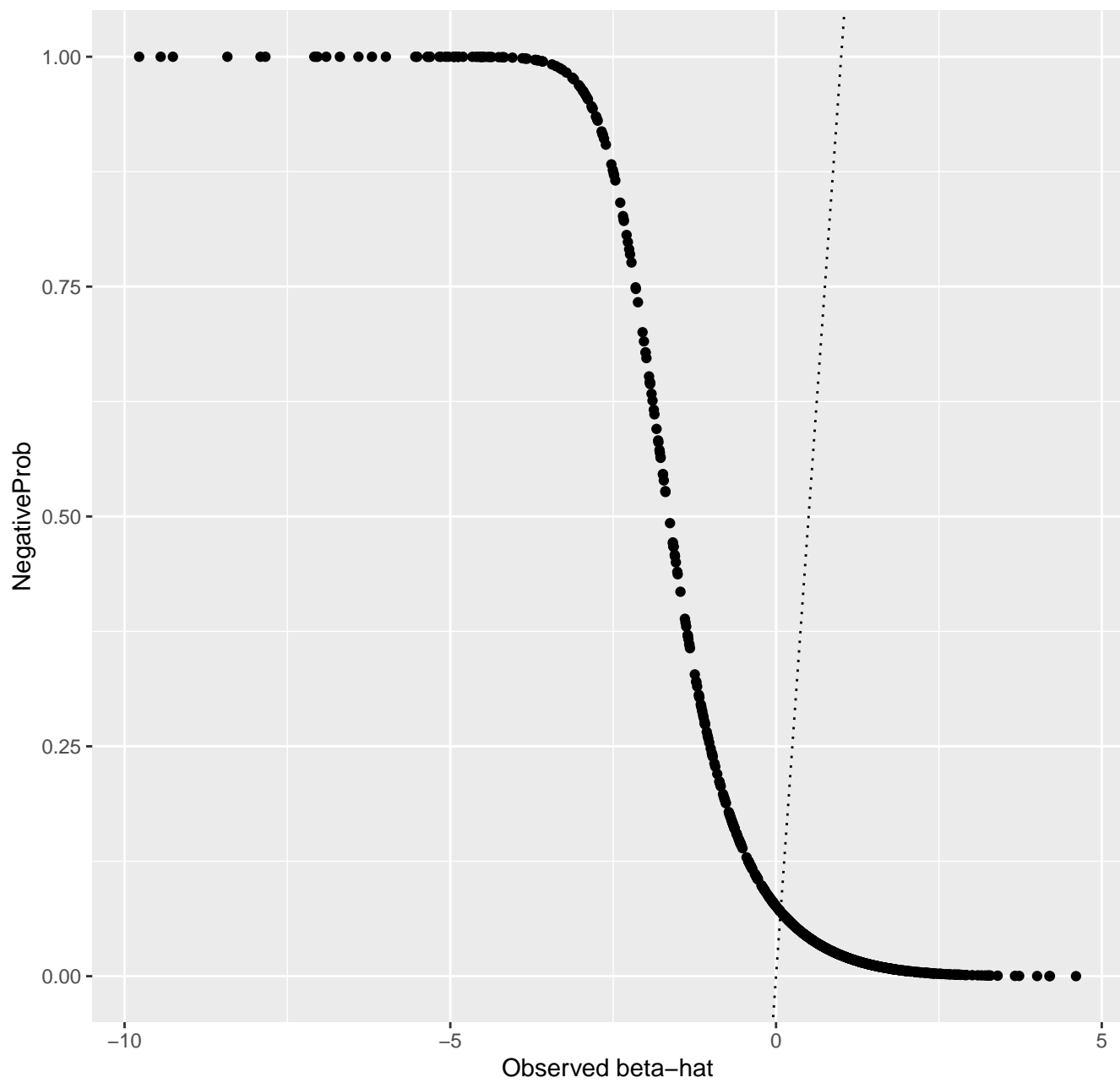
FIS1_(51024) negative



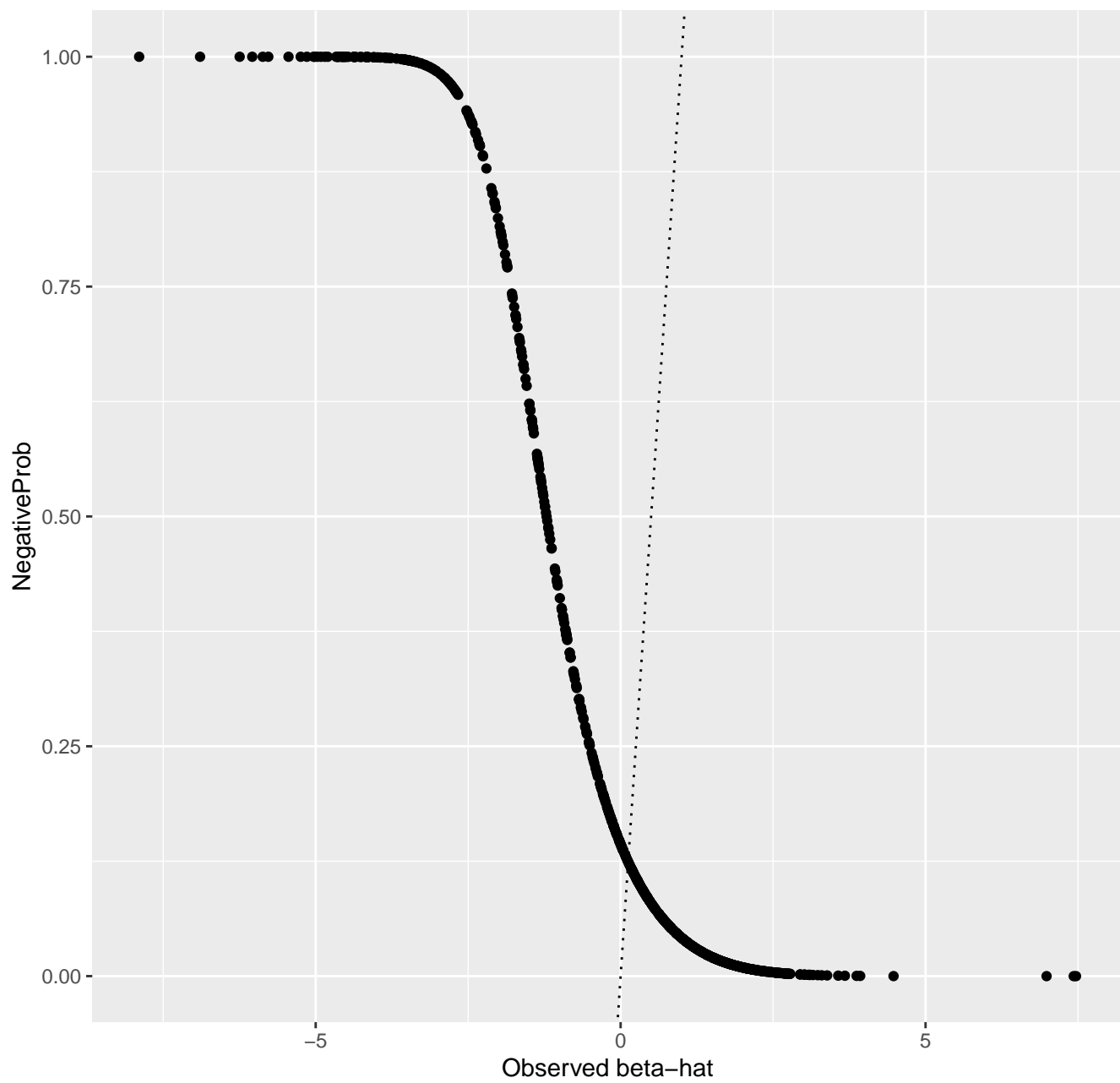
SOX10_(6663) negative



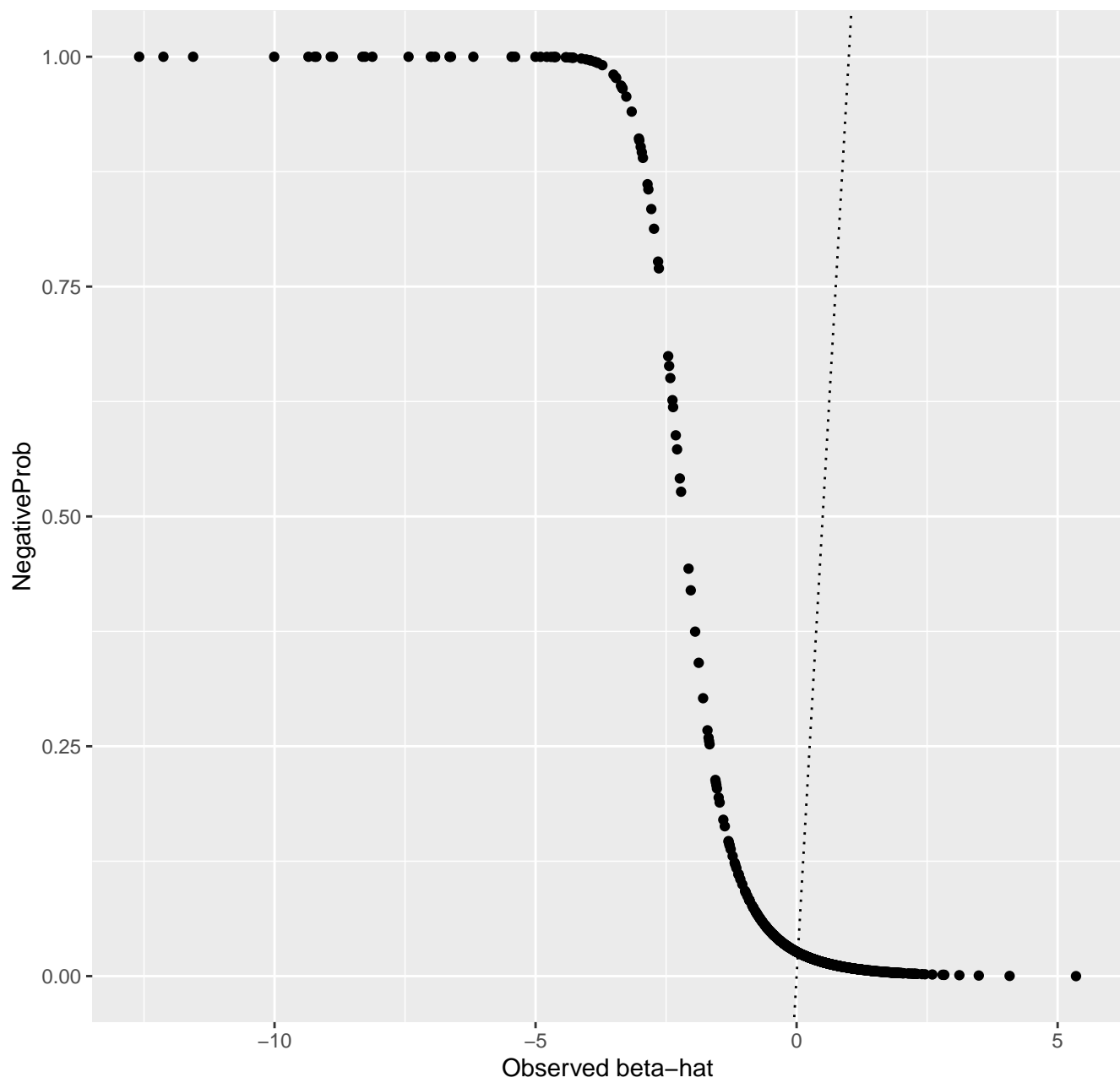
RAF1_(5894) negative



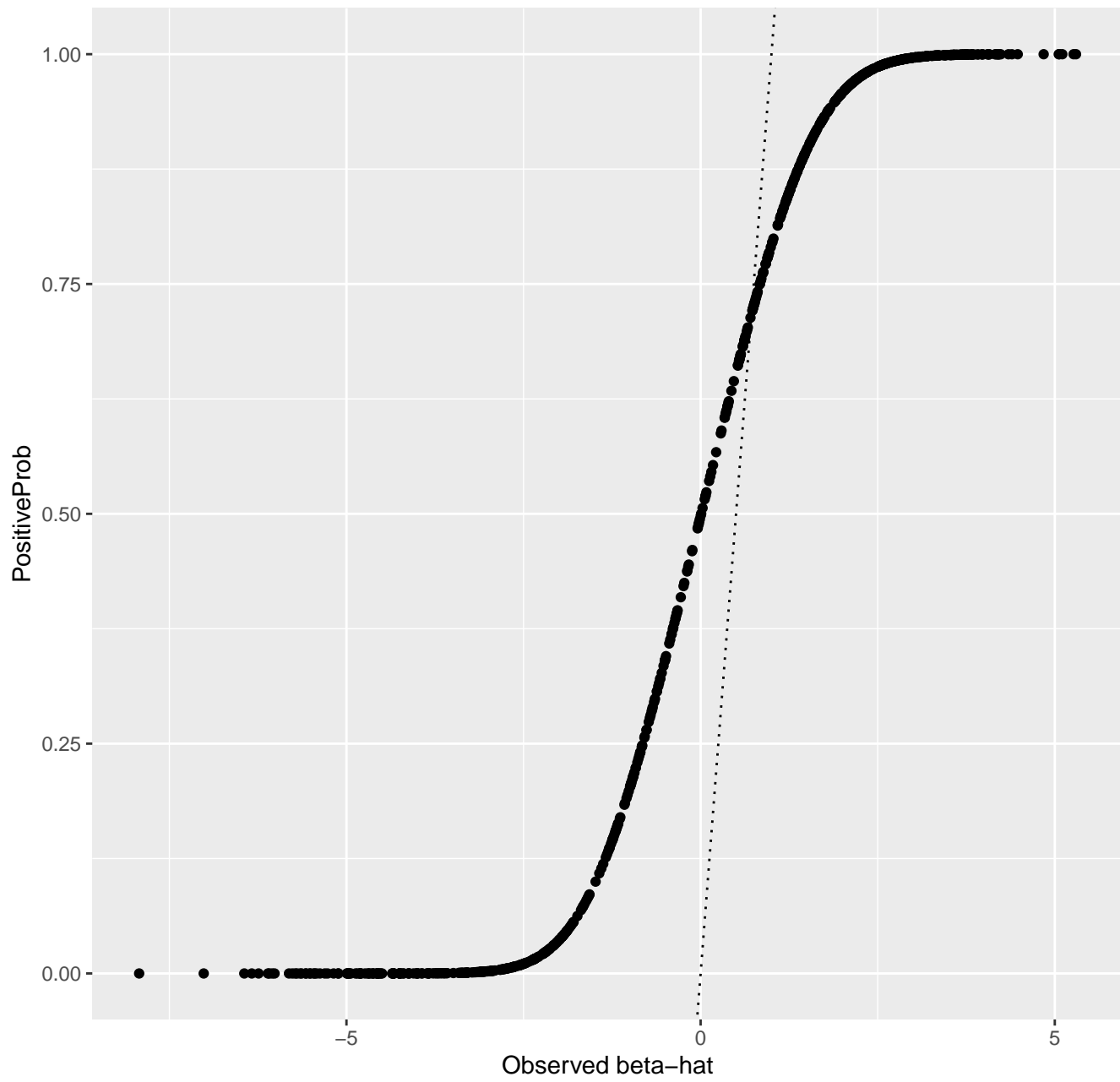
PRKAR1A_(5573) negative



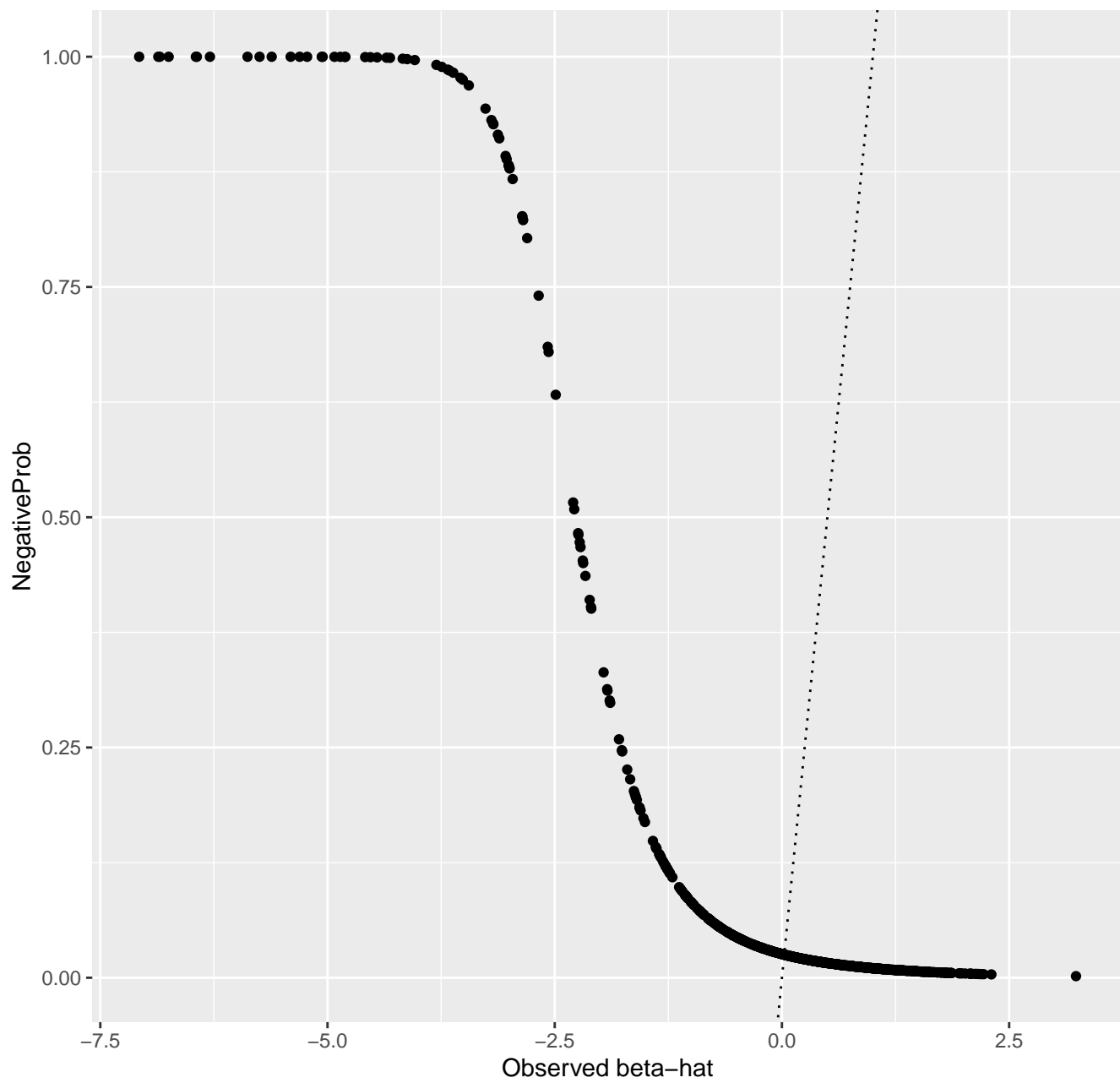
DUSP4_(1846) negative



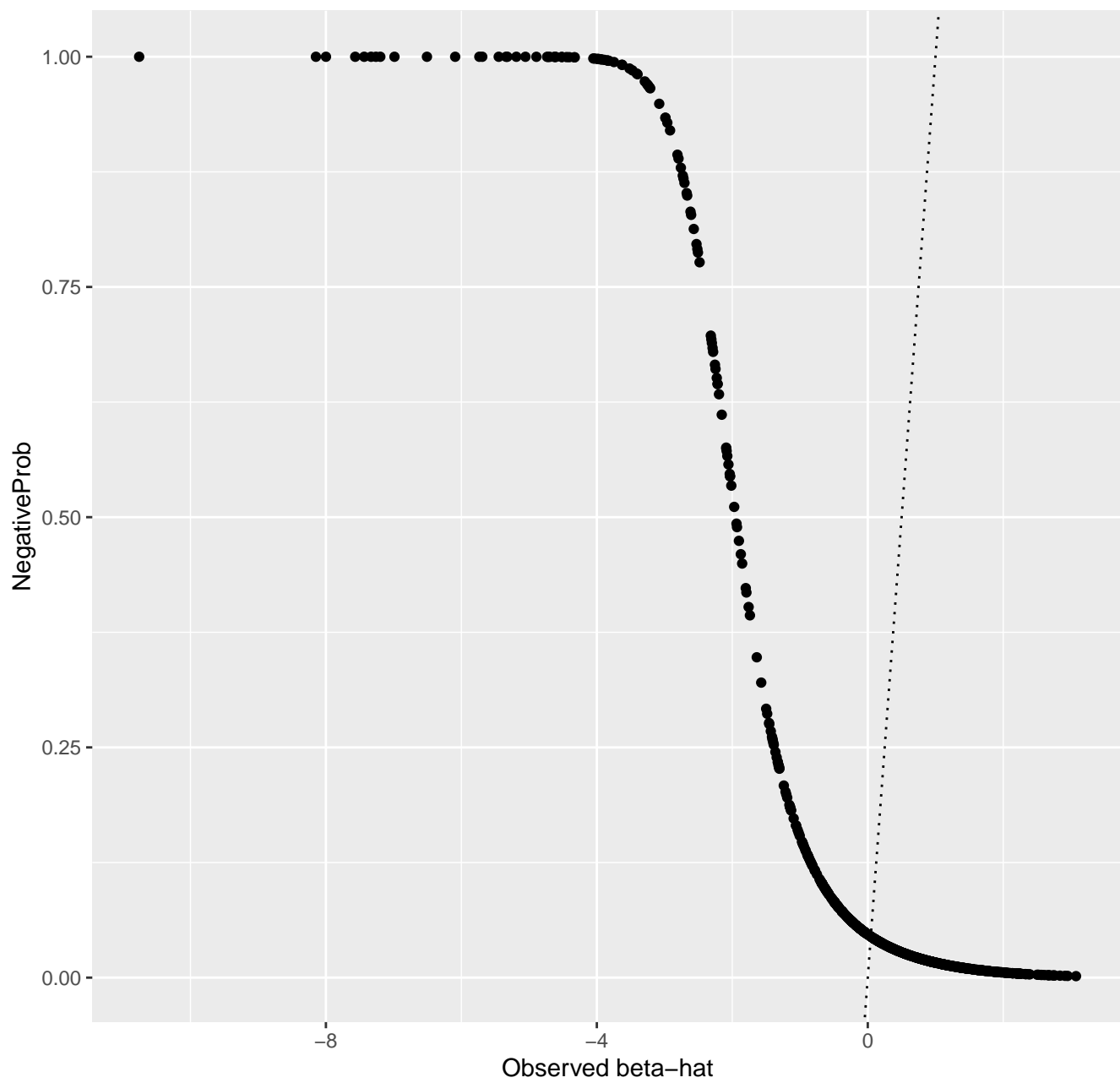
EFR3A_(23167) positive



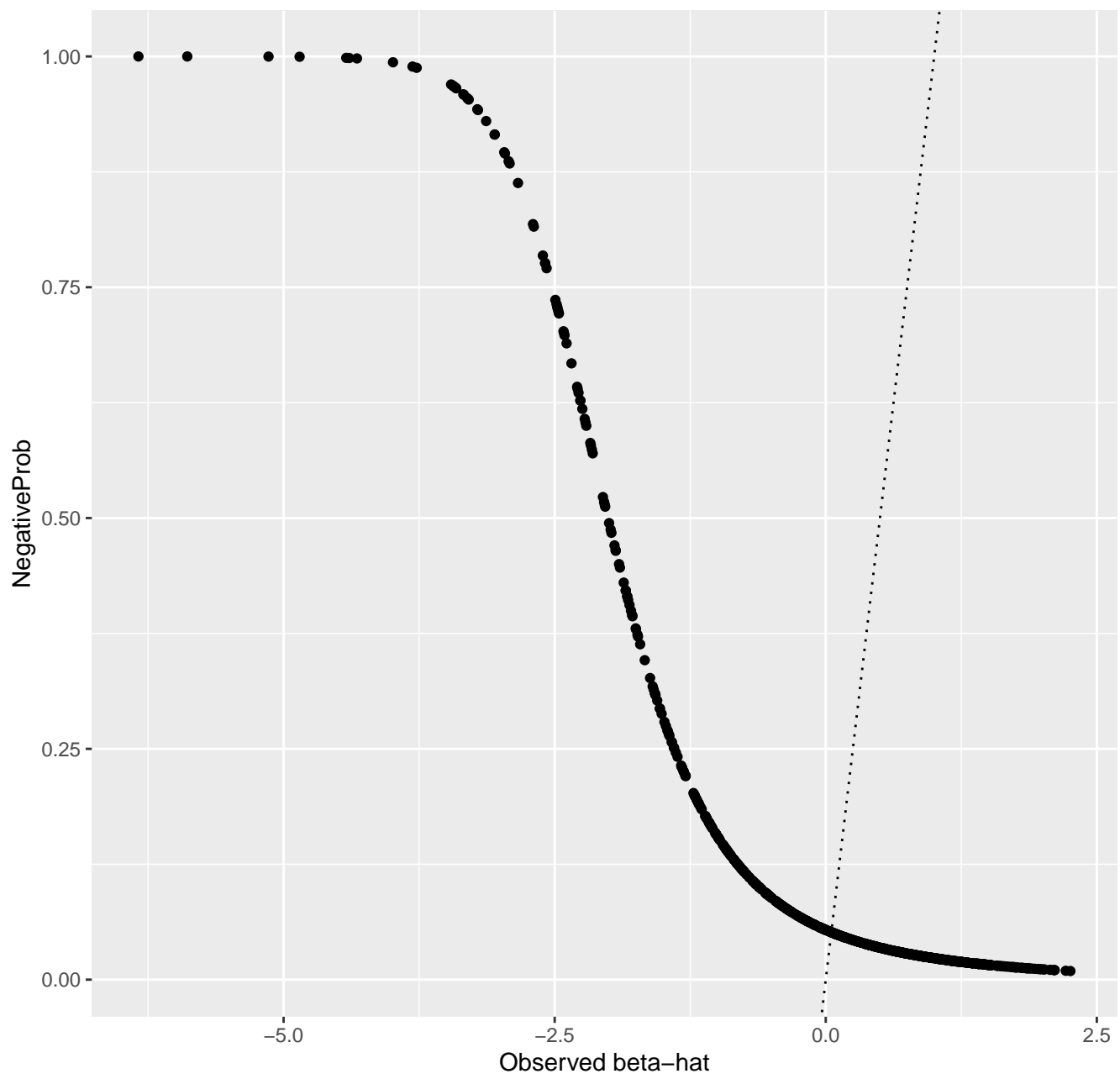
ERBB3_(2065) negative



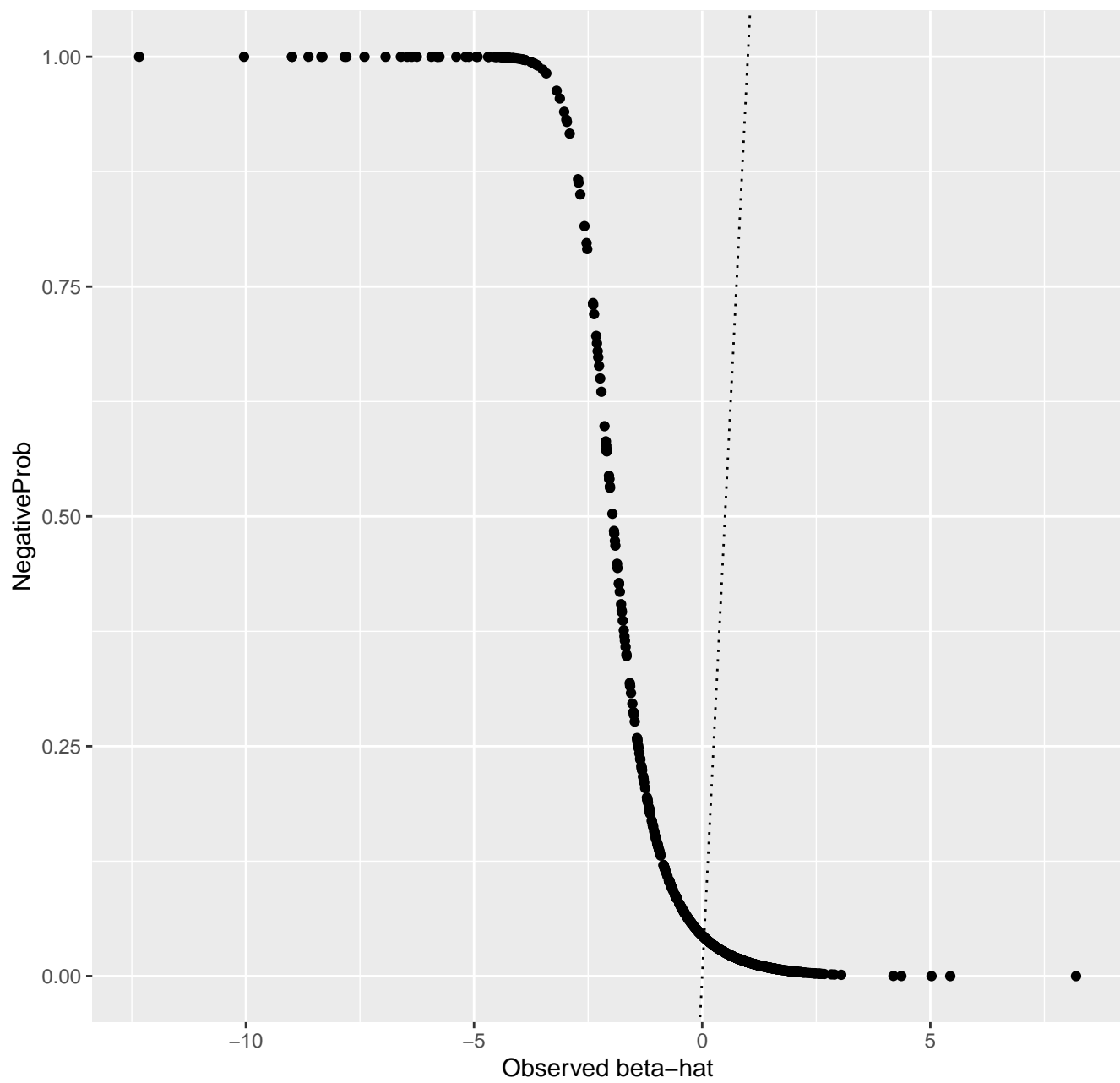
ERBB2_(2064) negative



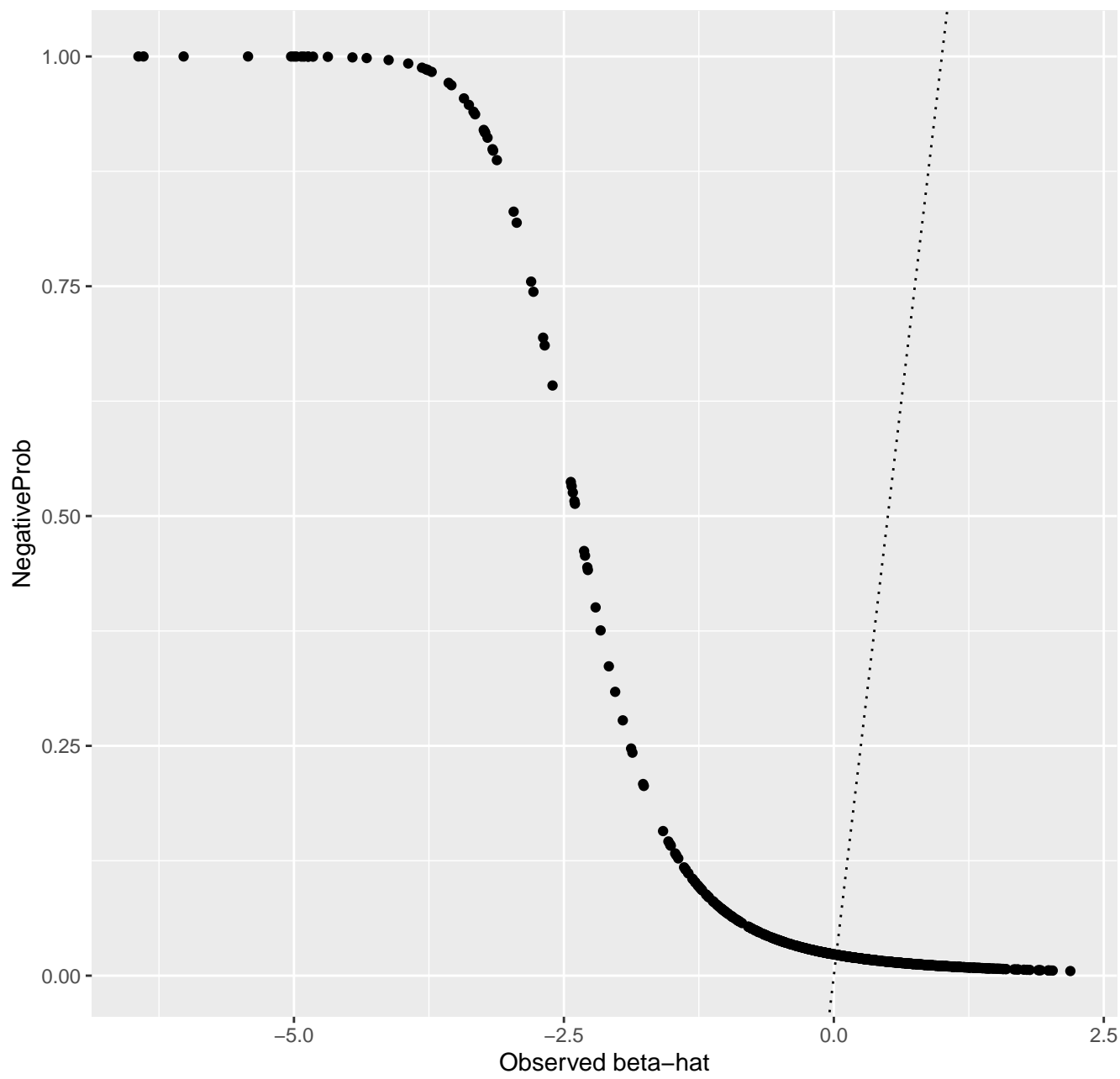
YIPF5_(81555) negative



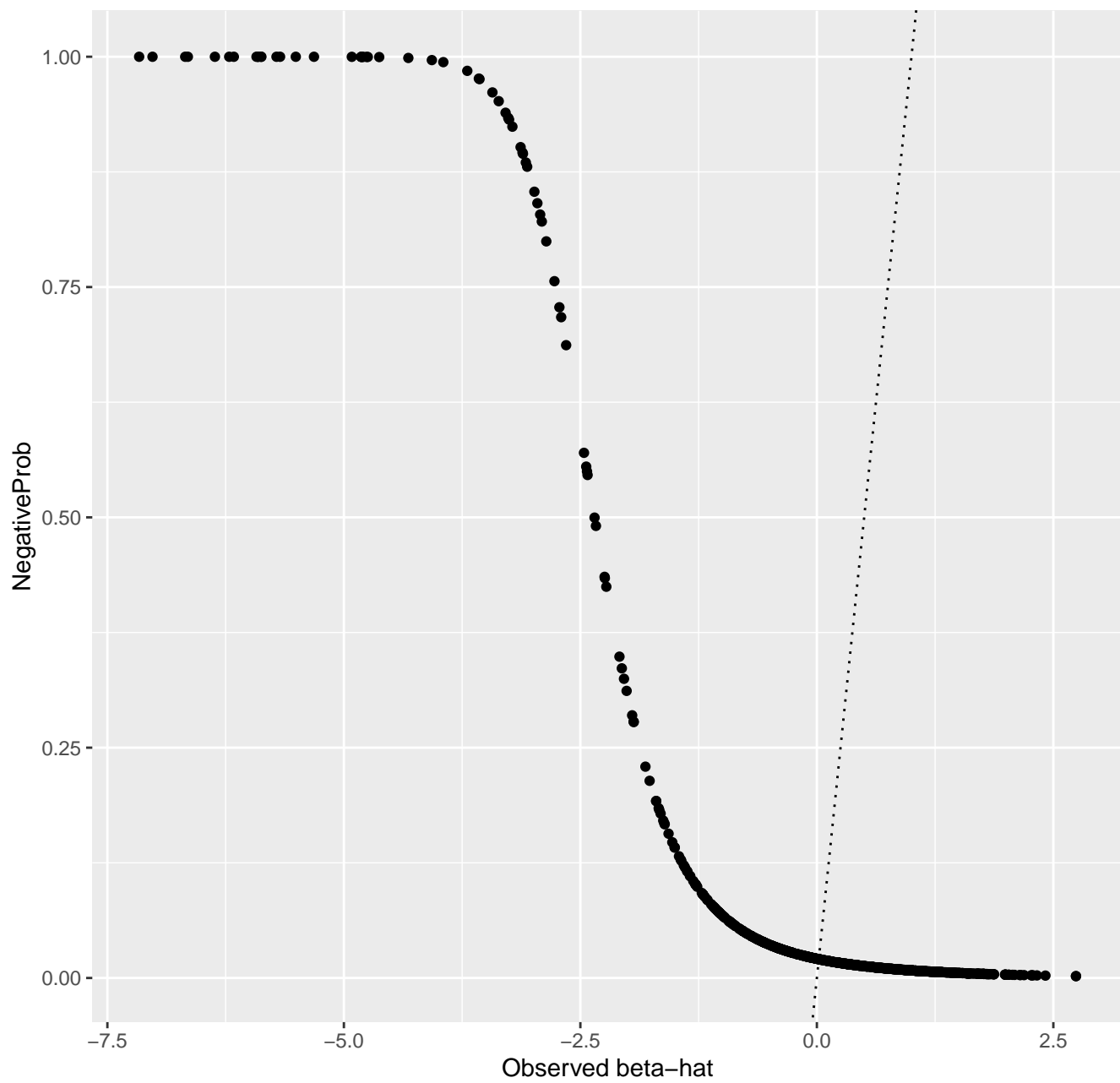
CDH2_(1000) negative



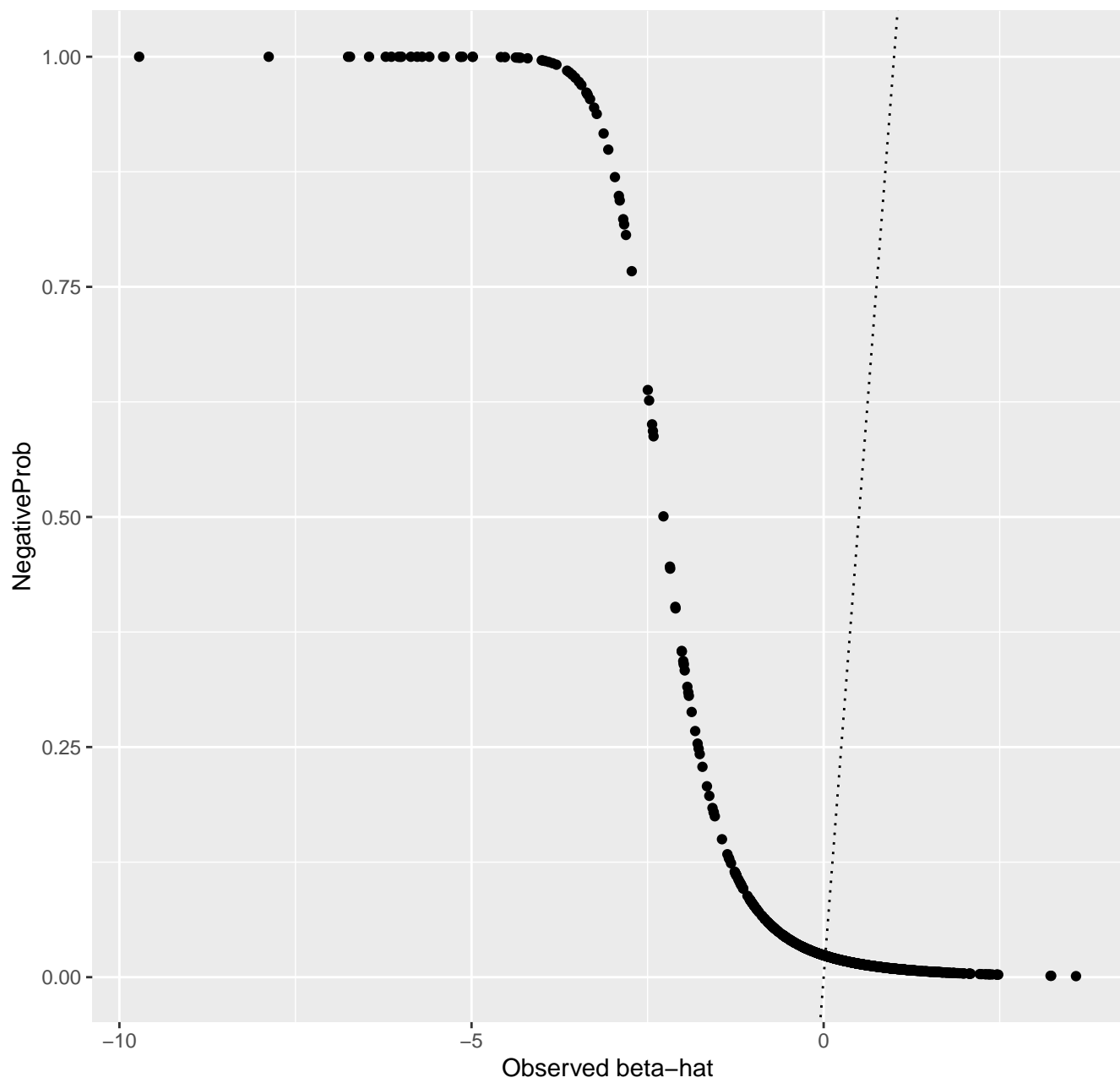
ATP7A_(538) negative



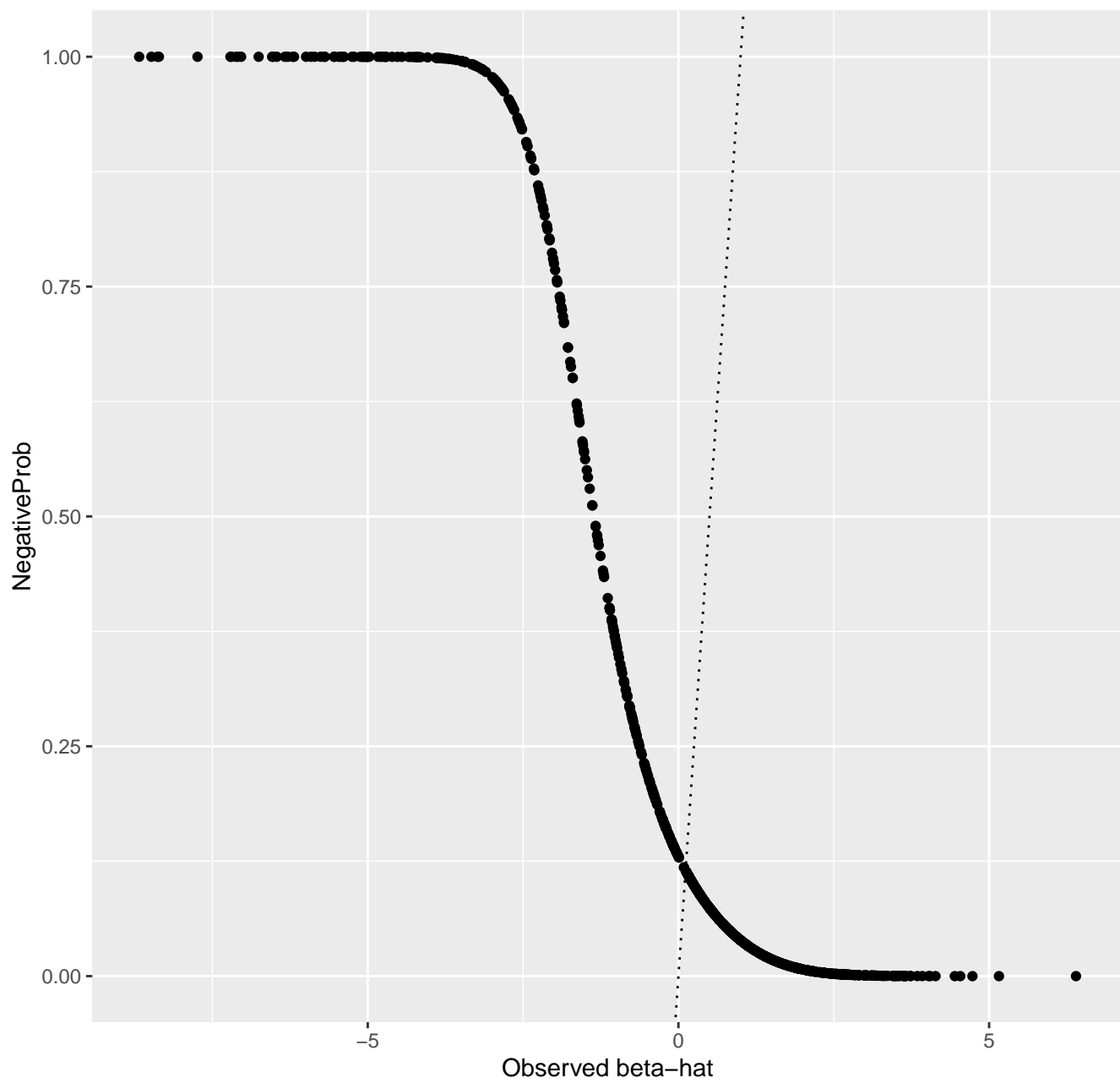
ALDH18A1_(5832) negative



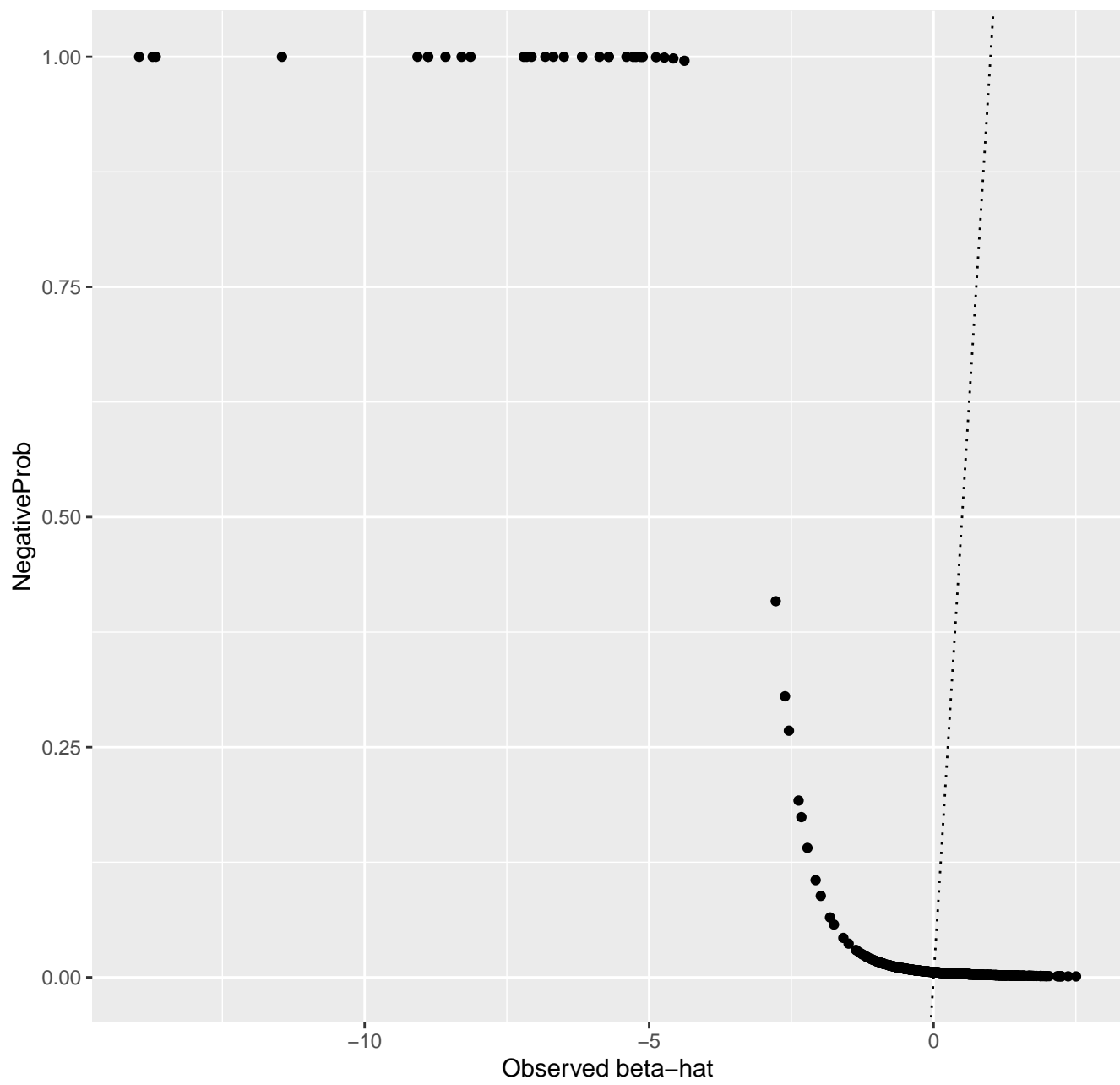
TTC7A_(57217) negative



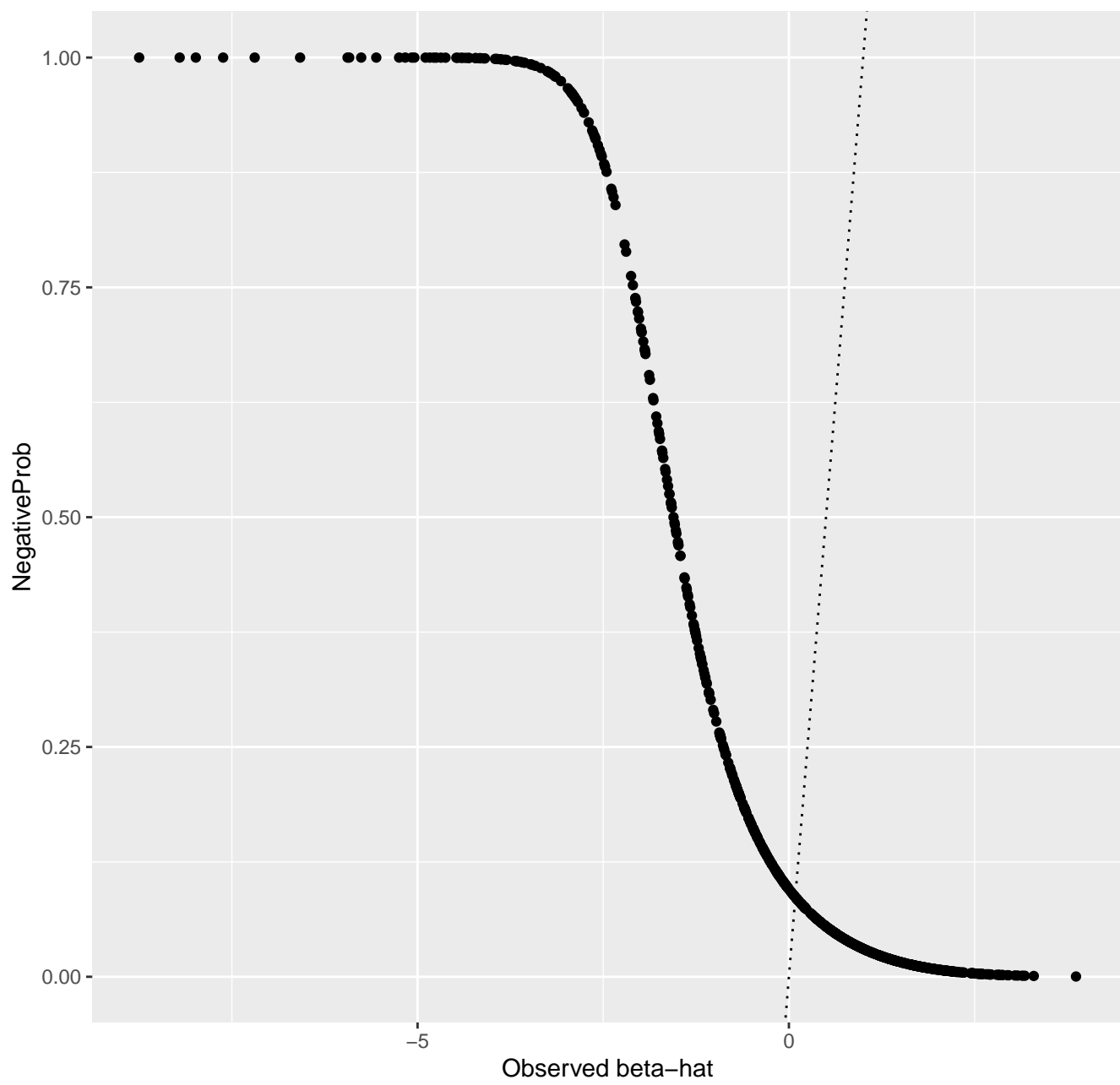
ACSL3_(2181) negative



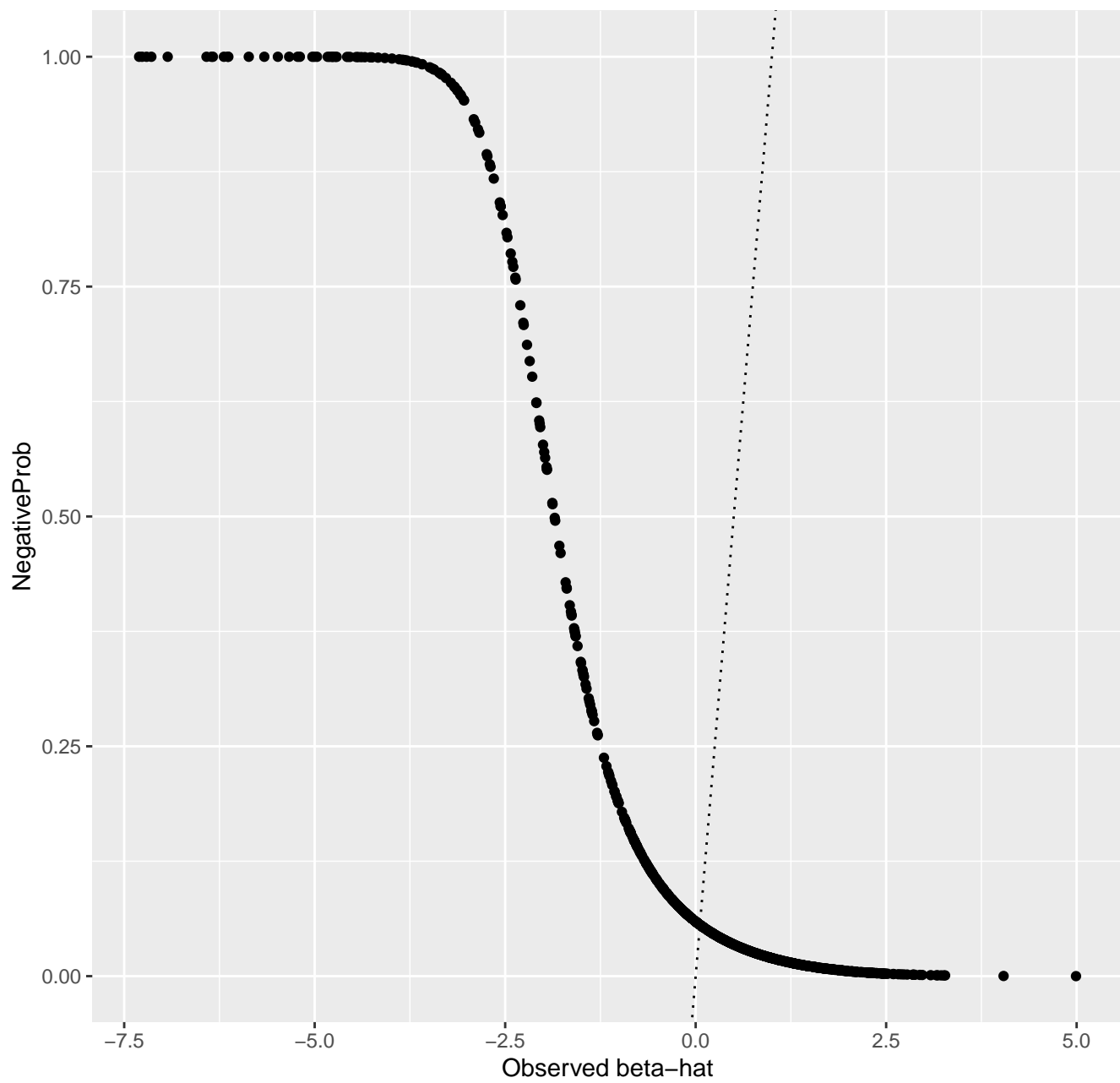
KCNK13_(56659) negative



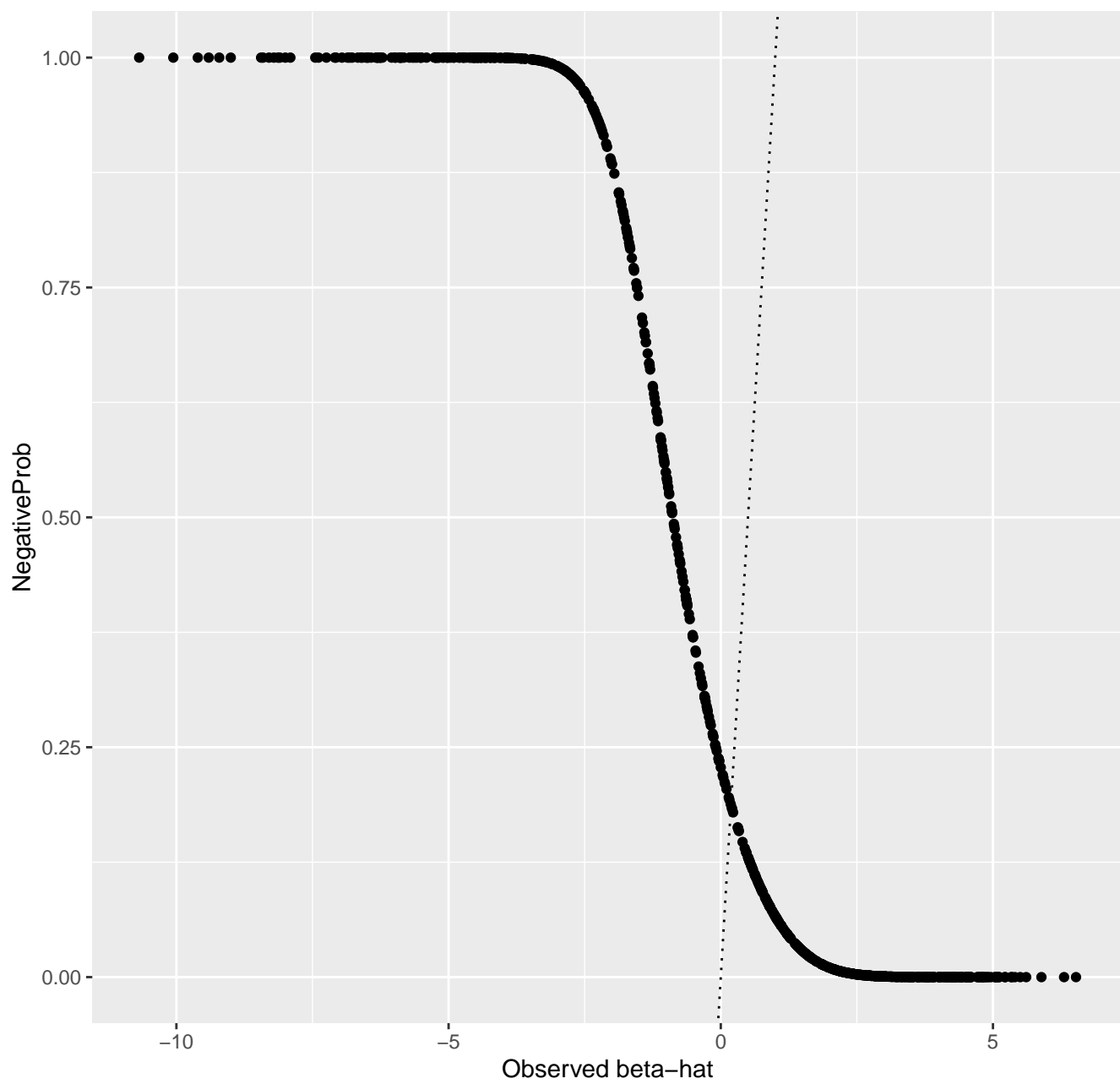
C3orf38_(285237) negative



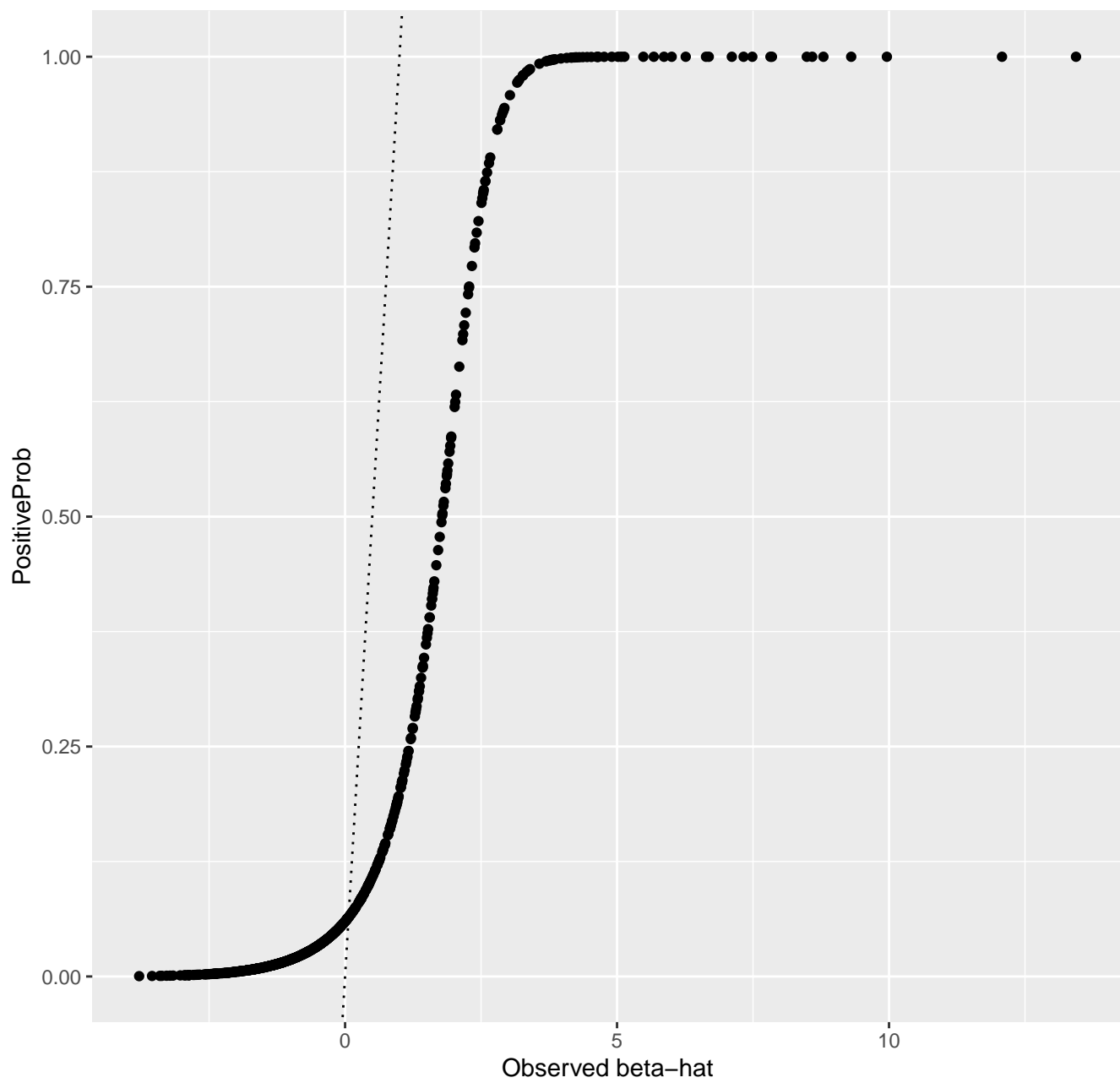
CEPT1_(10390) negative



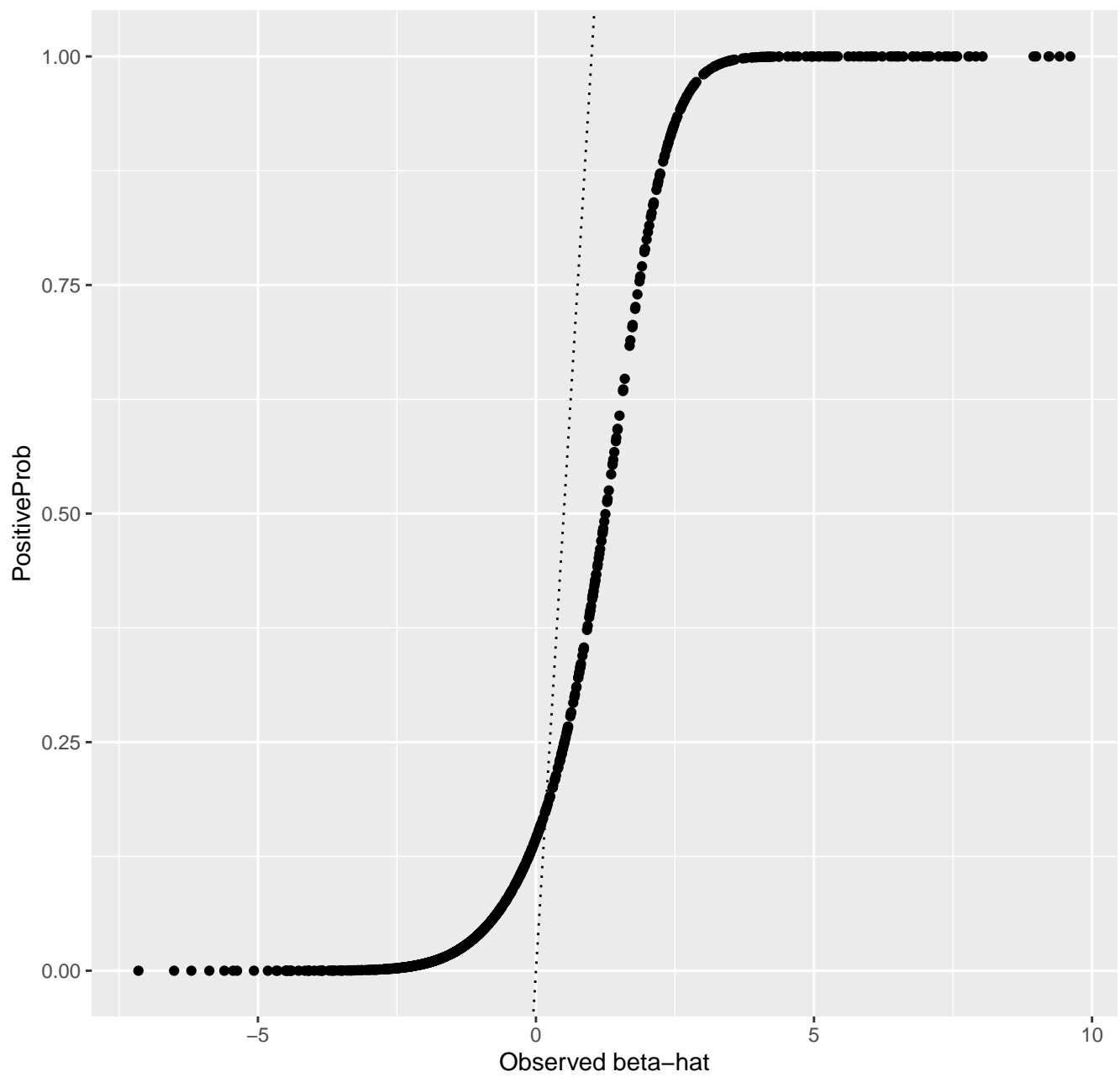
SEPHS2_(22928) negative



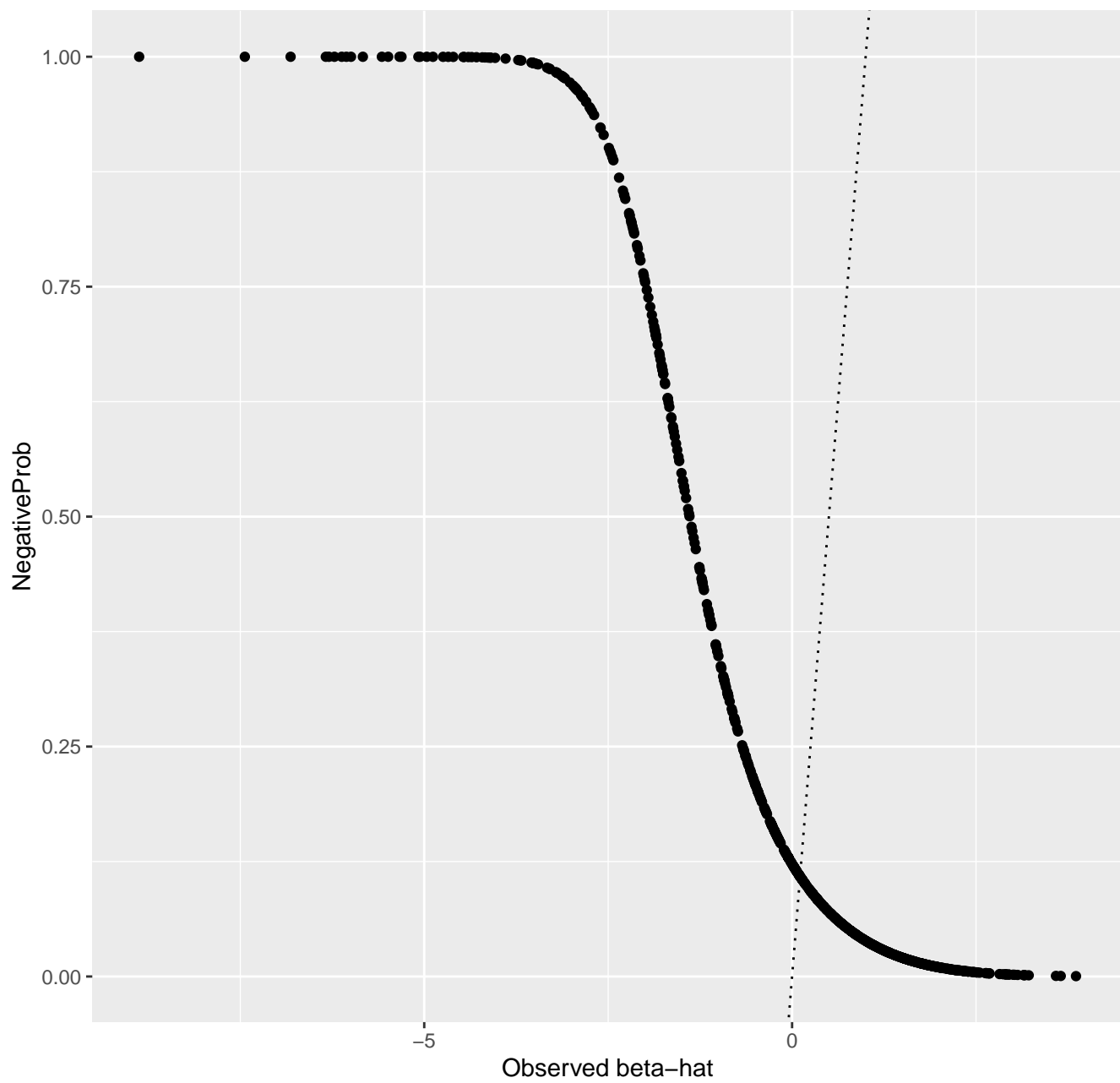
CDKN1A_(1026) positive



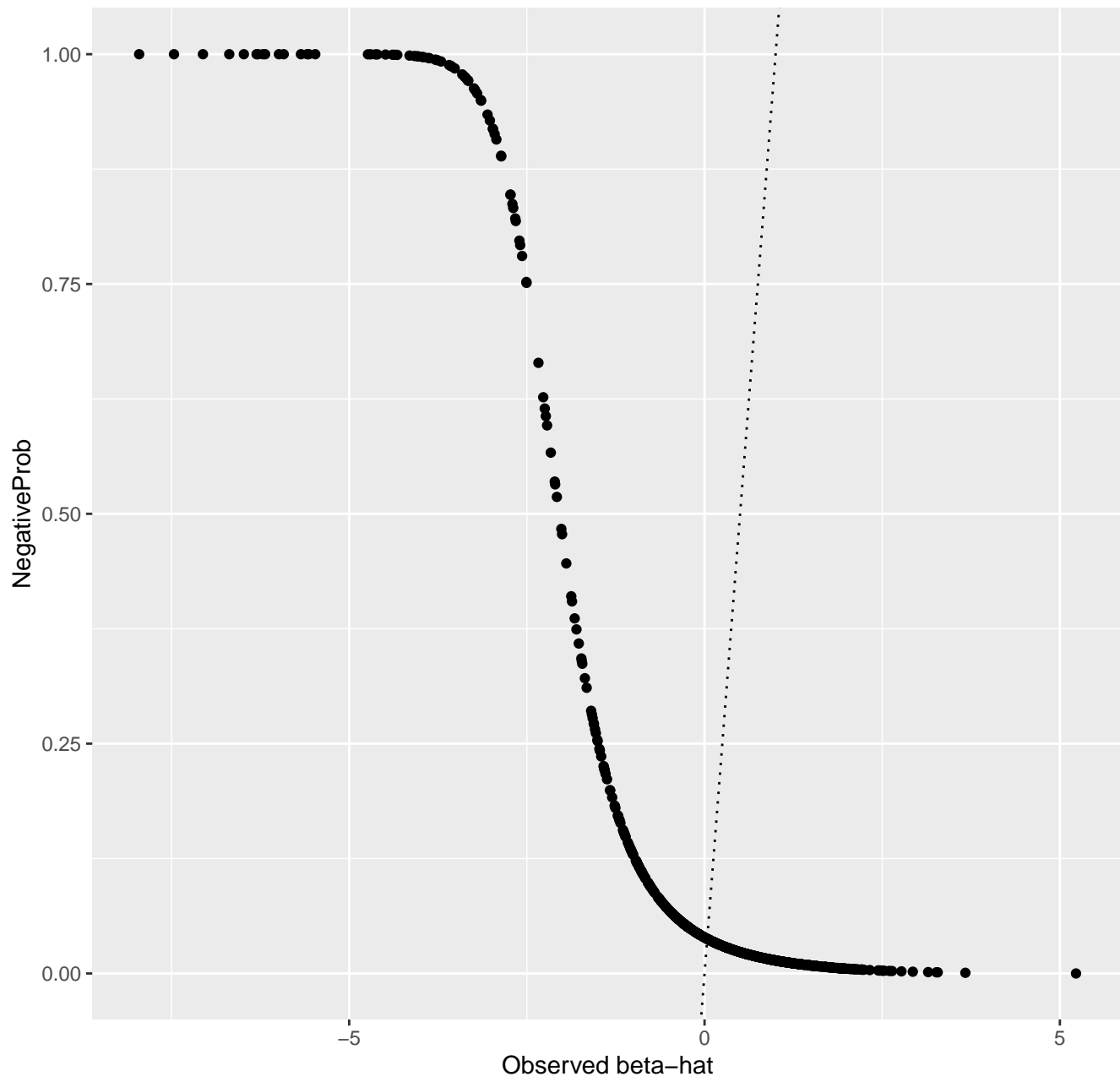
PPP2R1A_(5518) positive



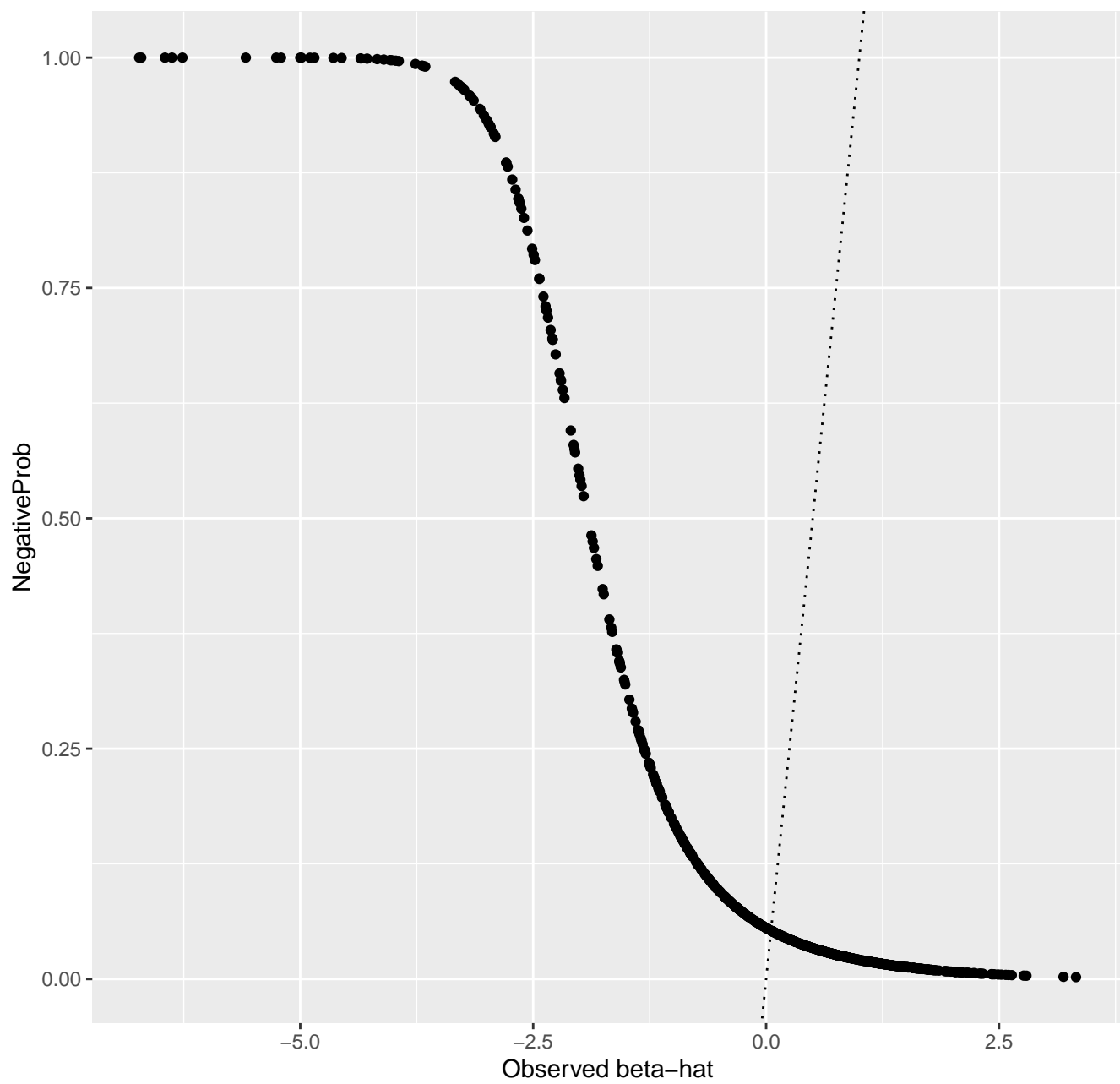
YAP1_(10413) negative



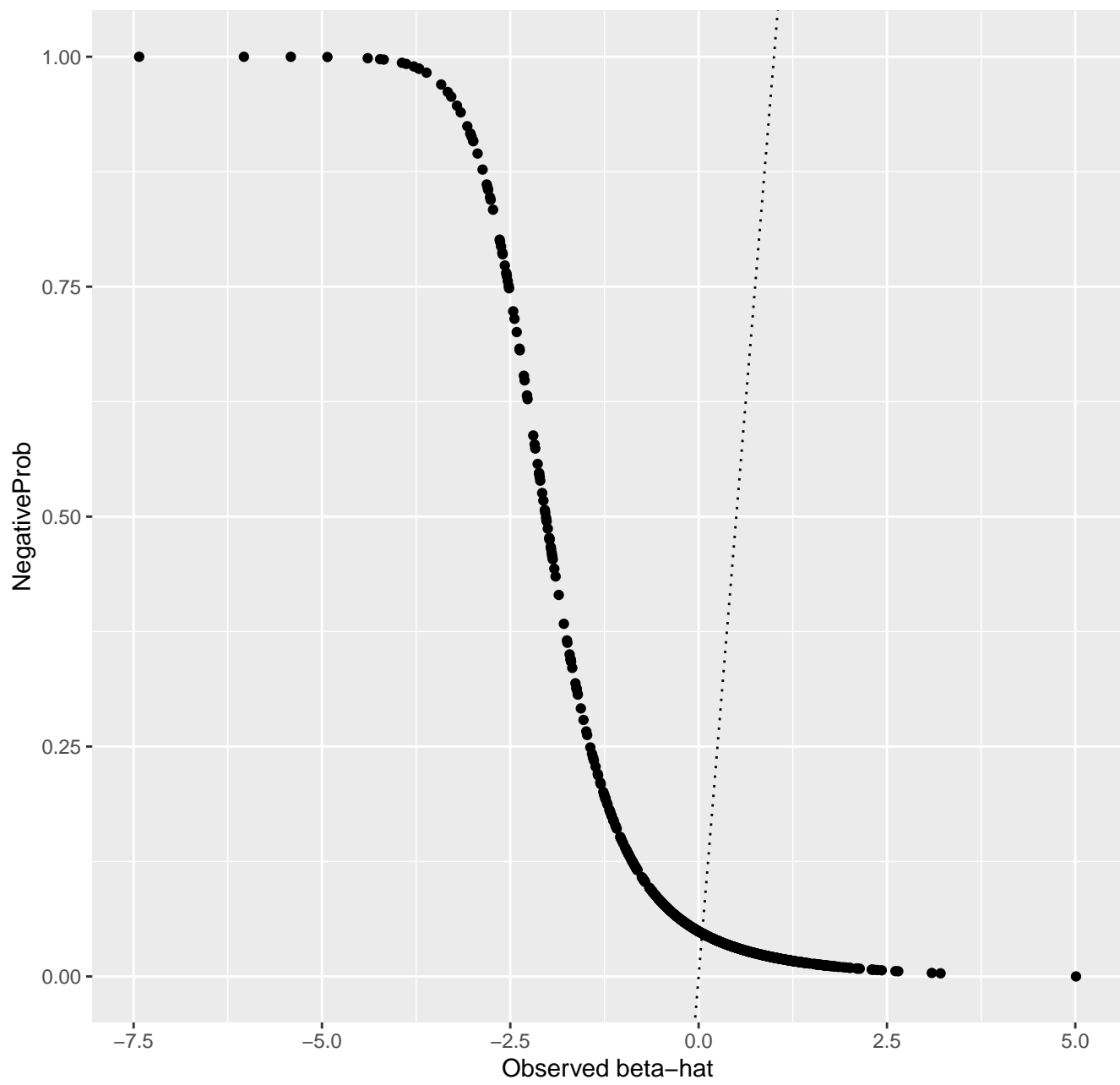
FDFT1_(2222) negative



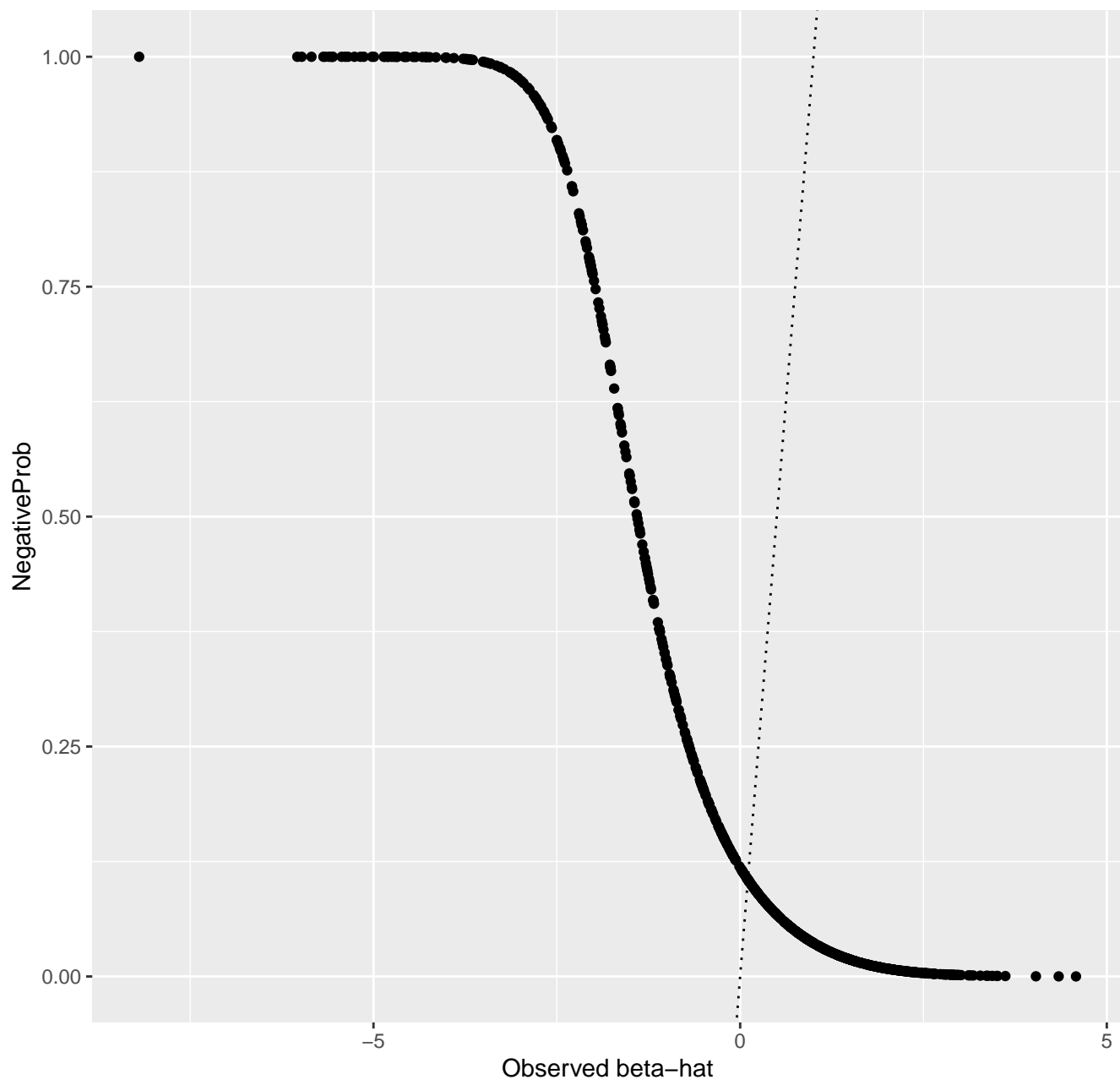
DYNC1LI2_(1783) negative



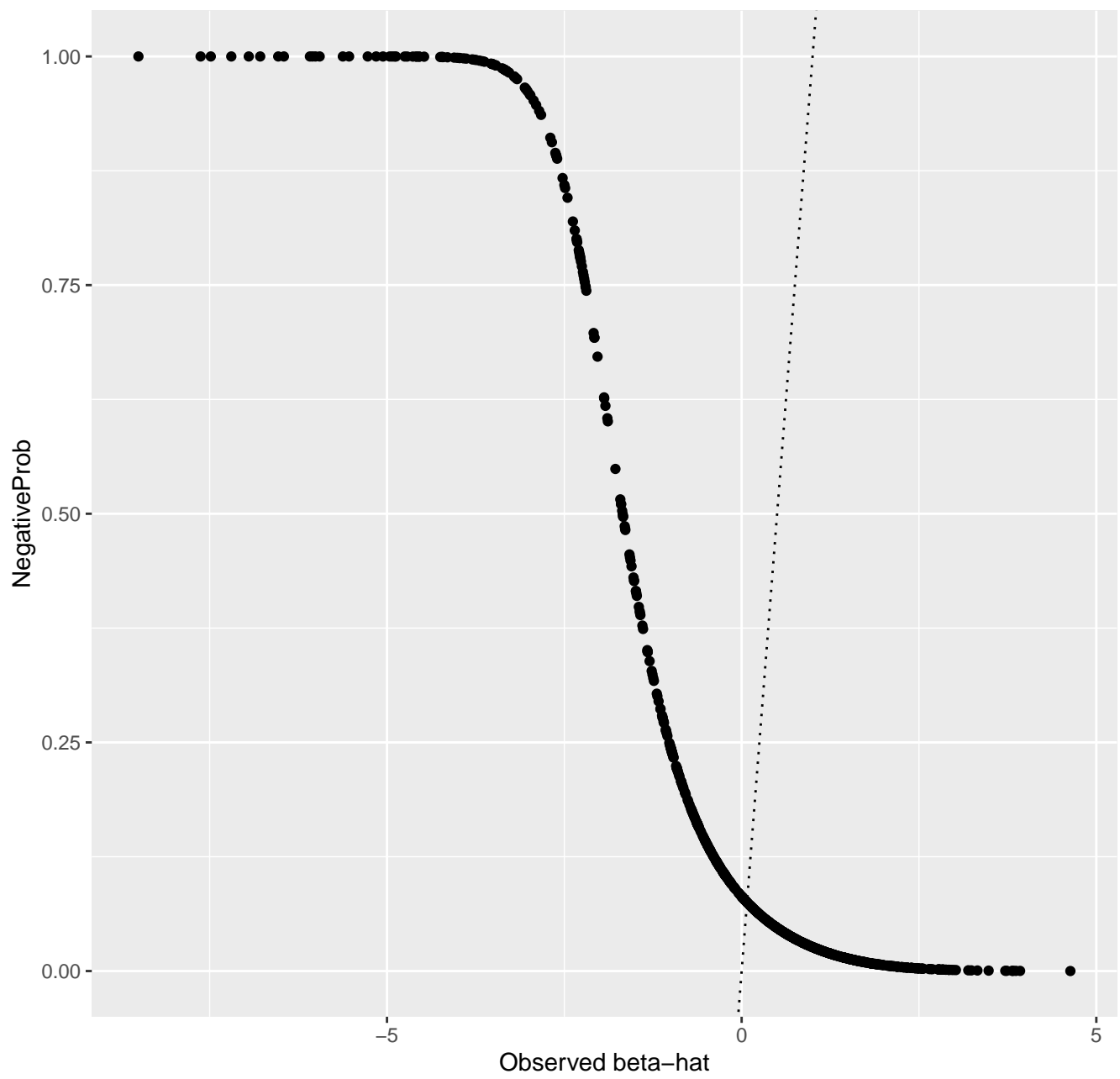
FRS2_(10818) negative



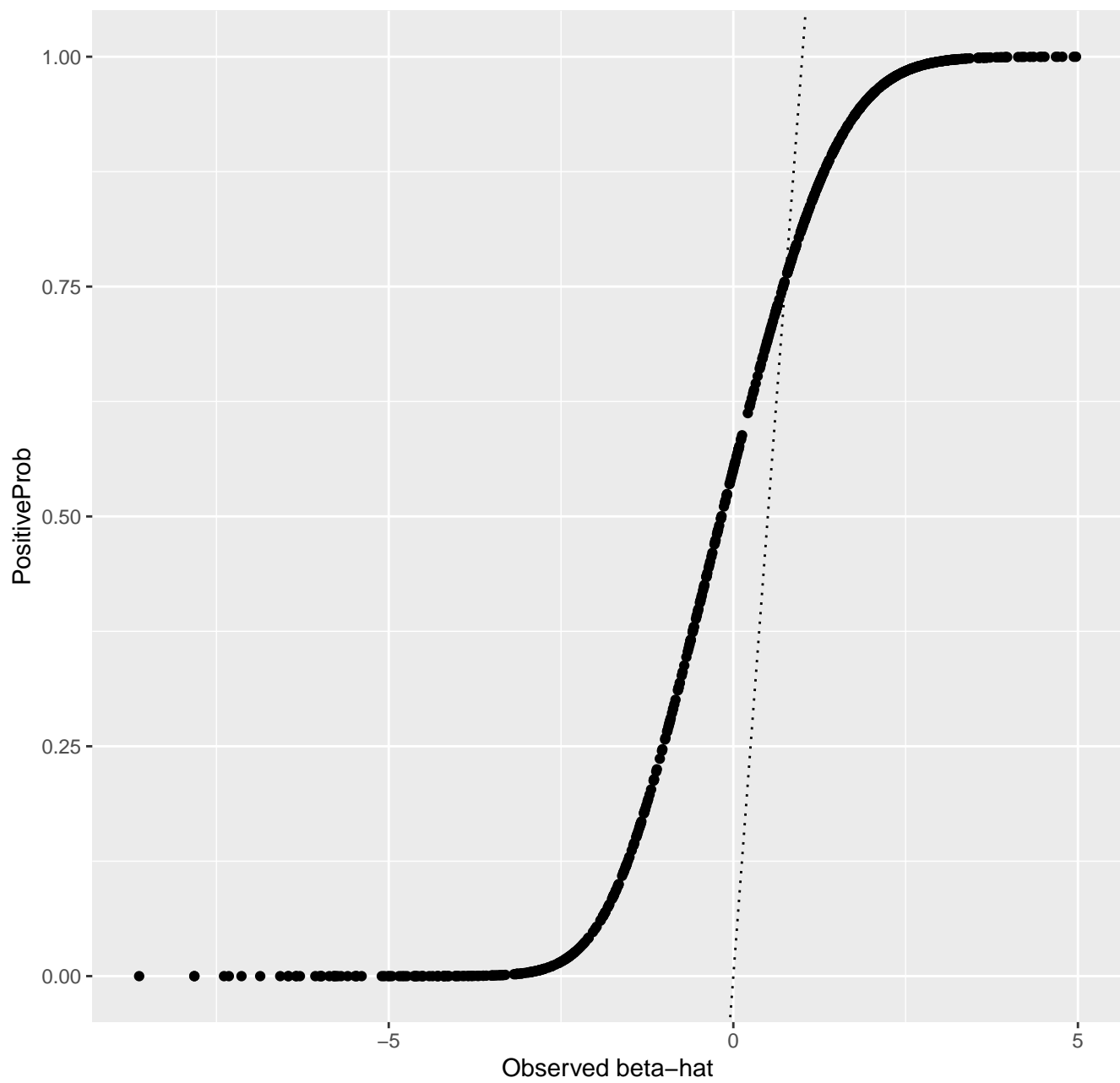
FCHO2_(115548) negative



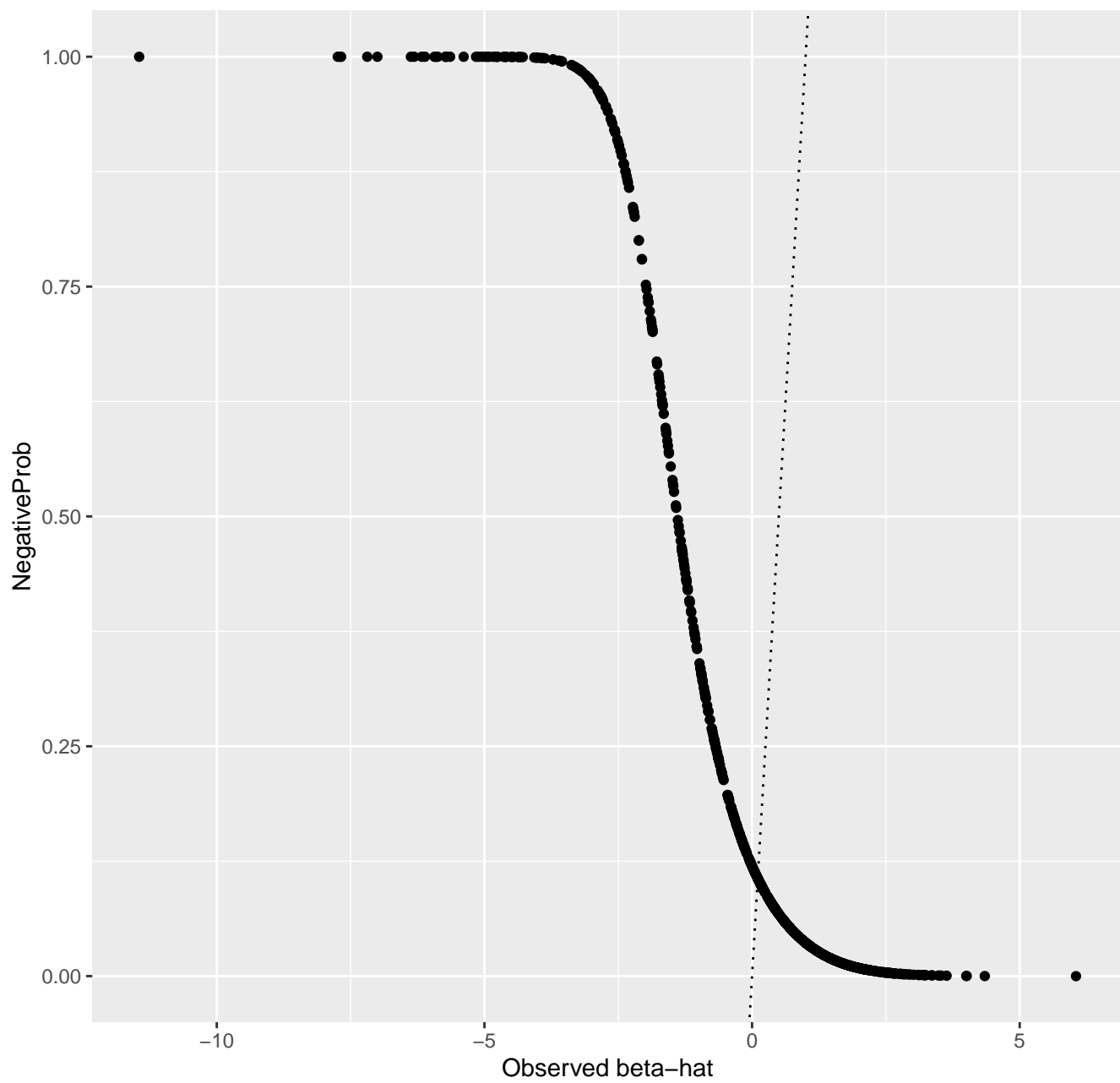
IRS2_(8660) negative



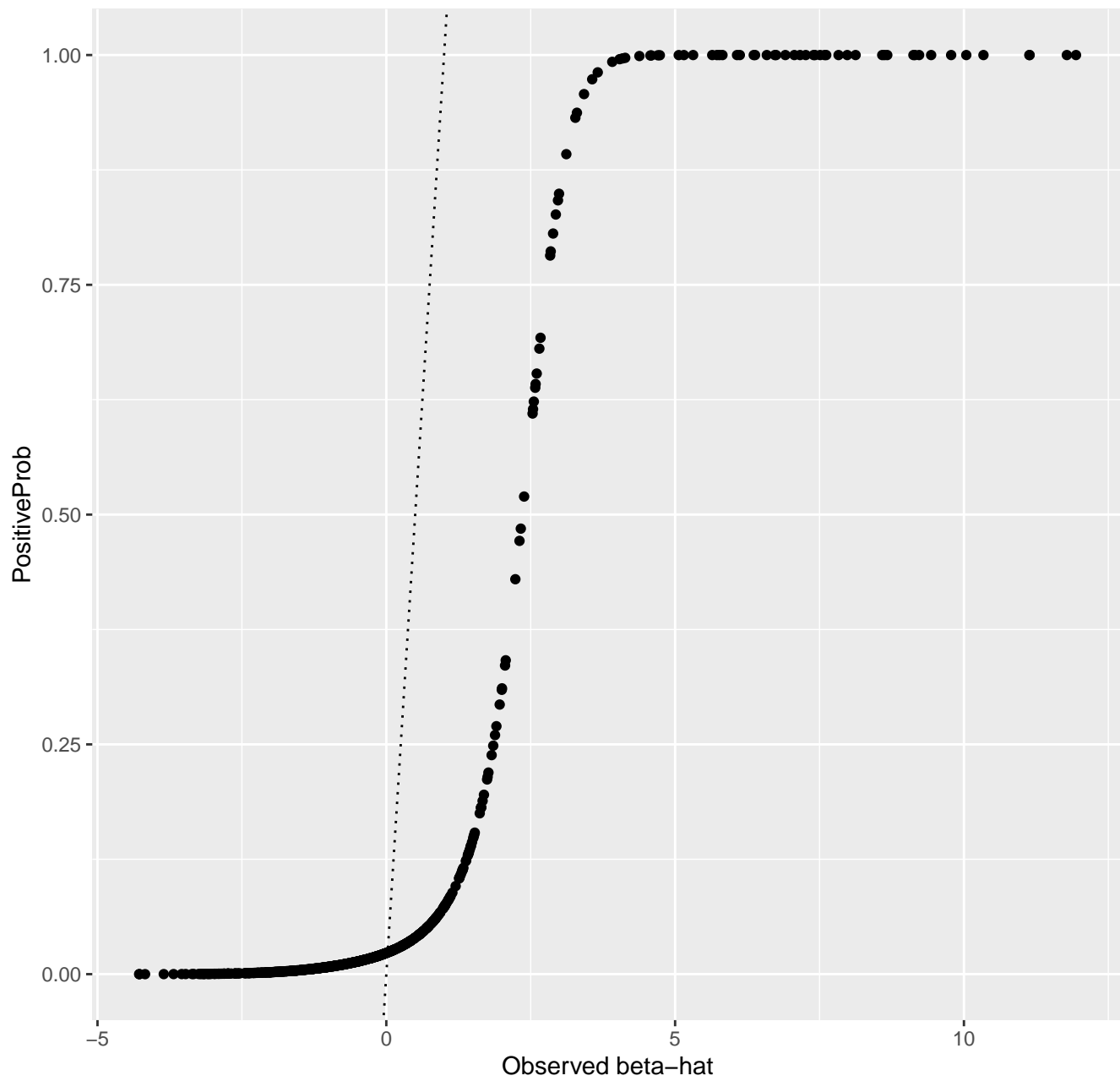
MCL1_(4170) positive



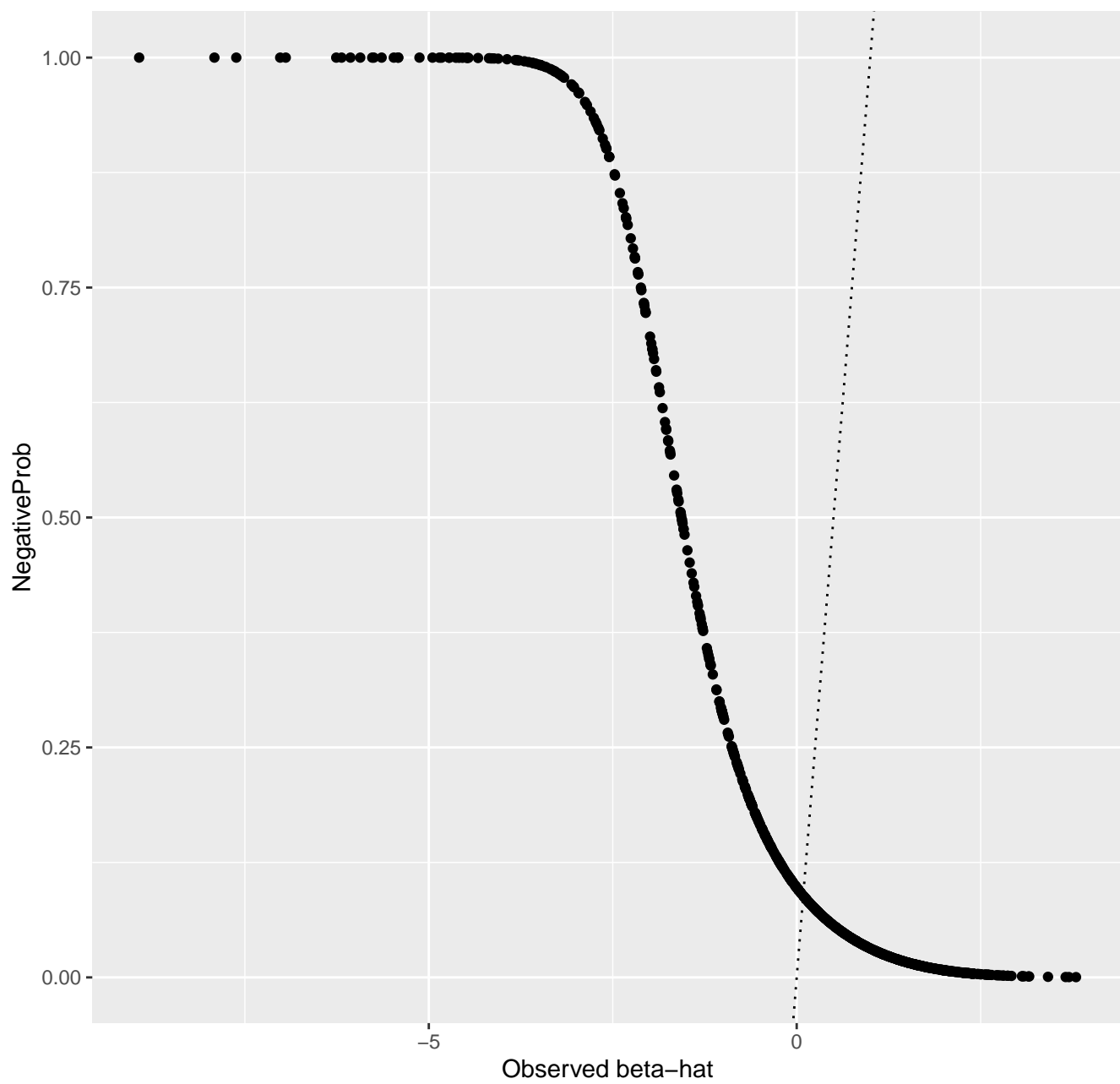
MAP3K7_(6885) negative



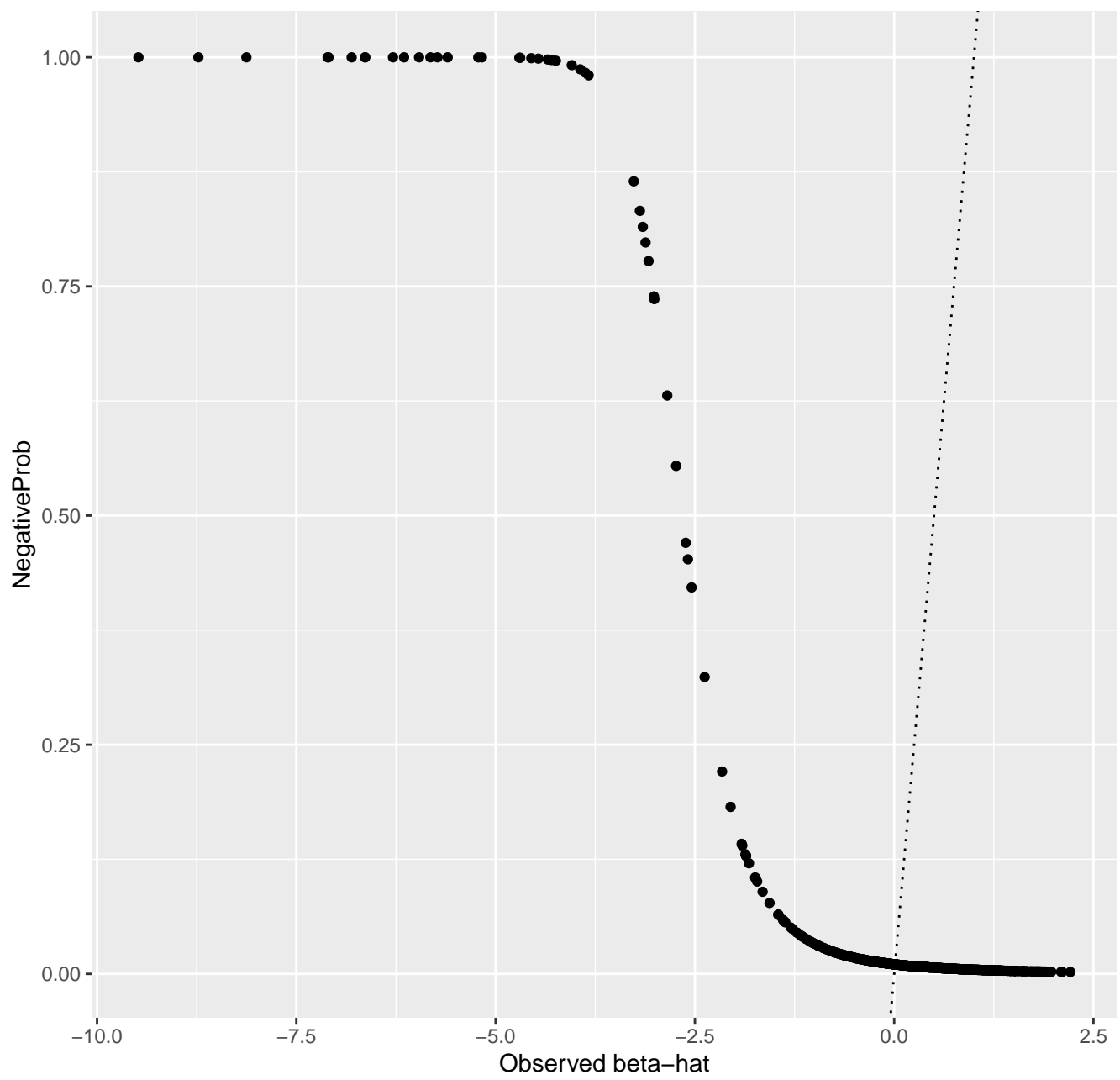
NAA10_(8260) positive



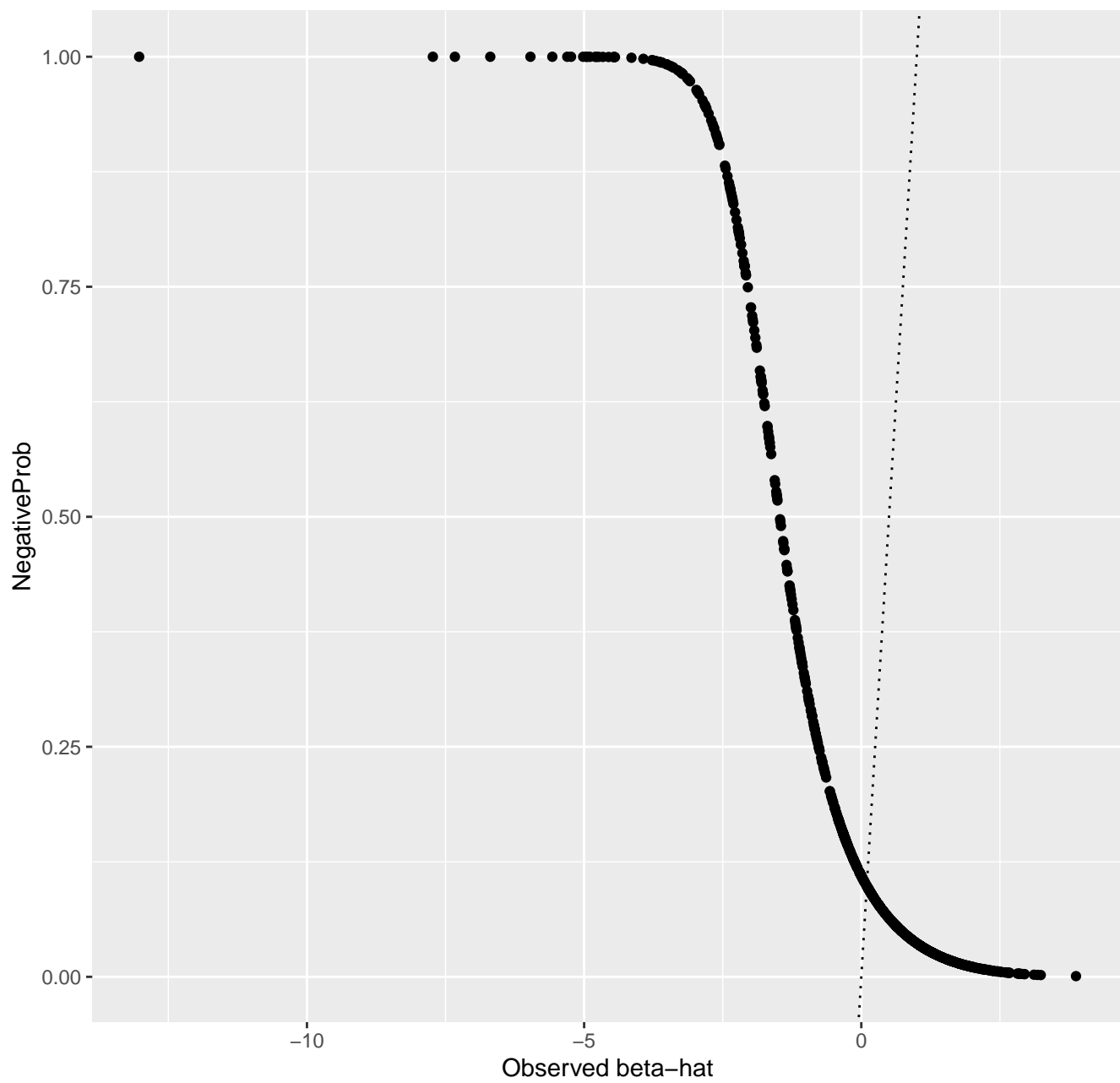
EGLN1_(54583) negative



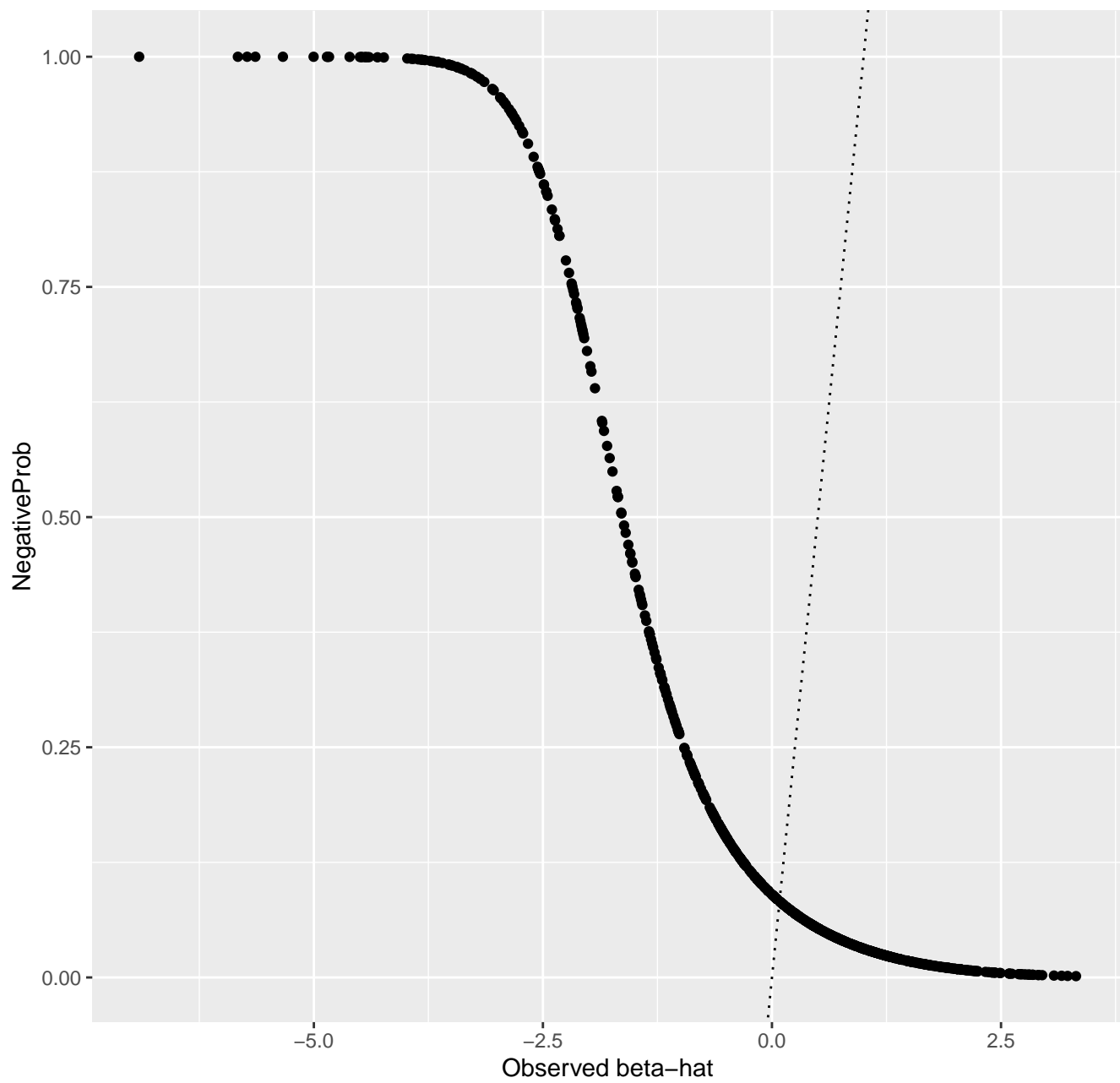
MITF_(4286) negative



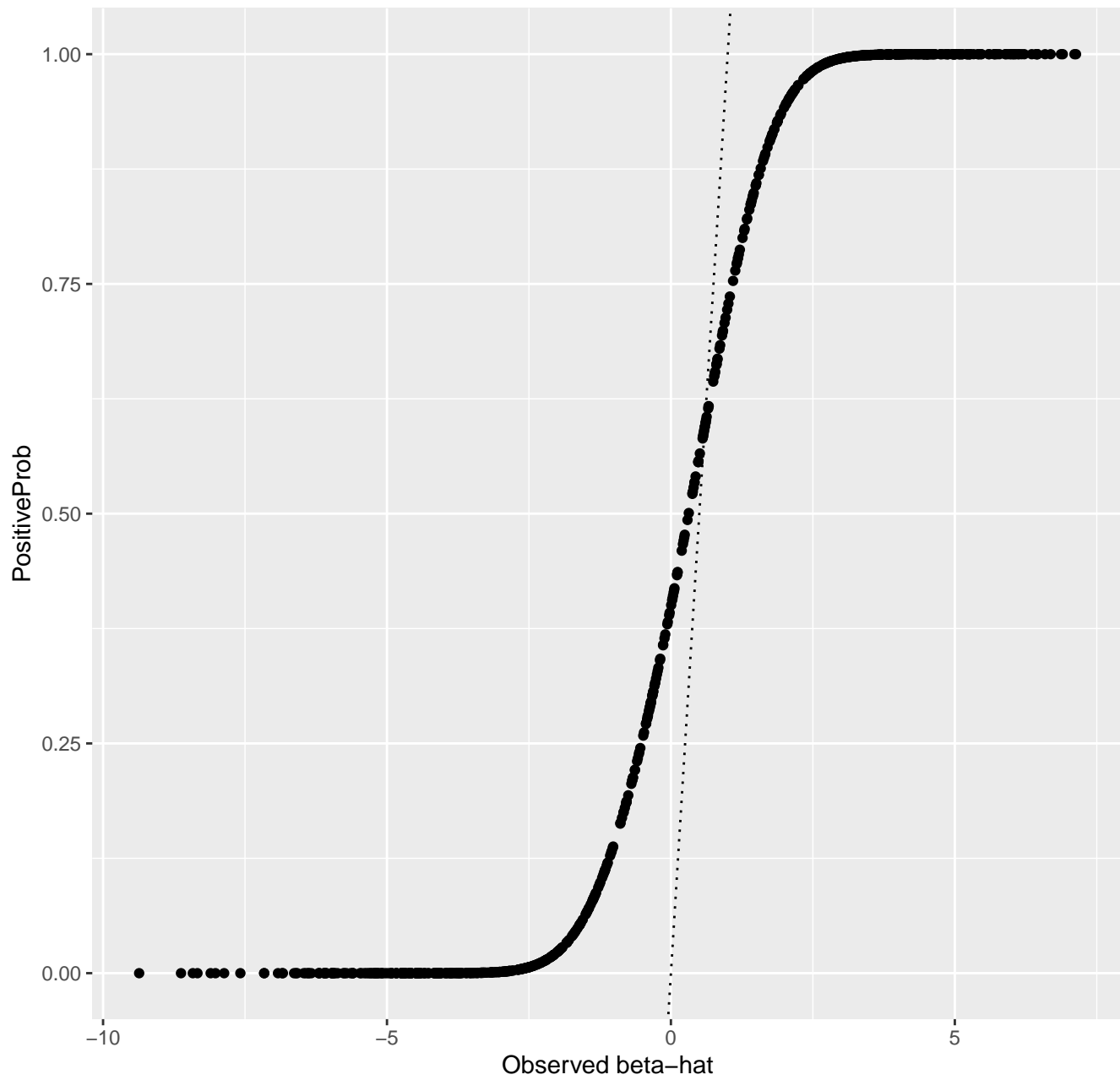
VPS36_(51028) negative



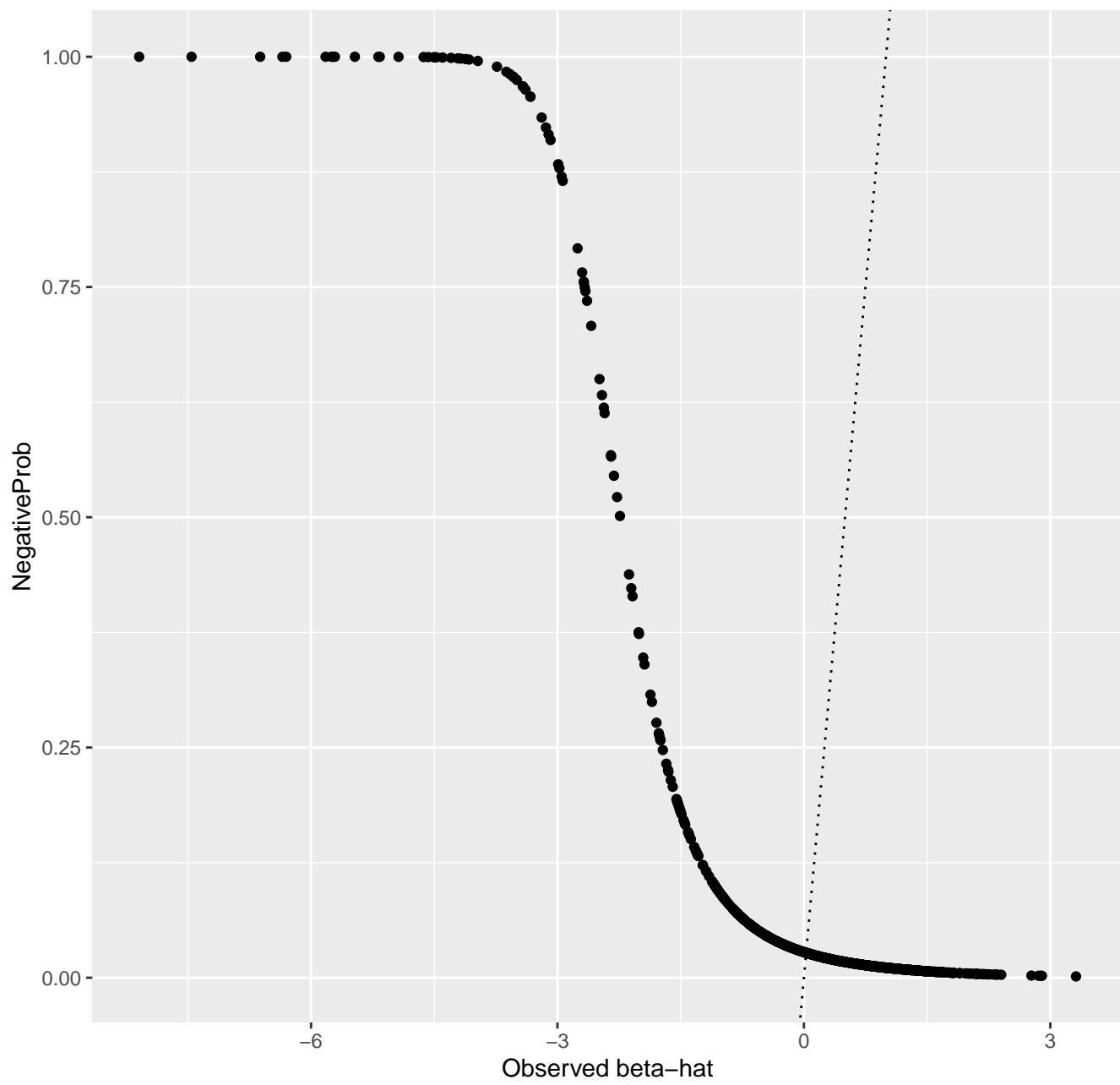
HSPA13_(6782) negative



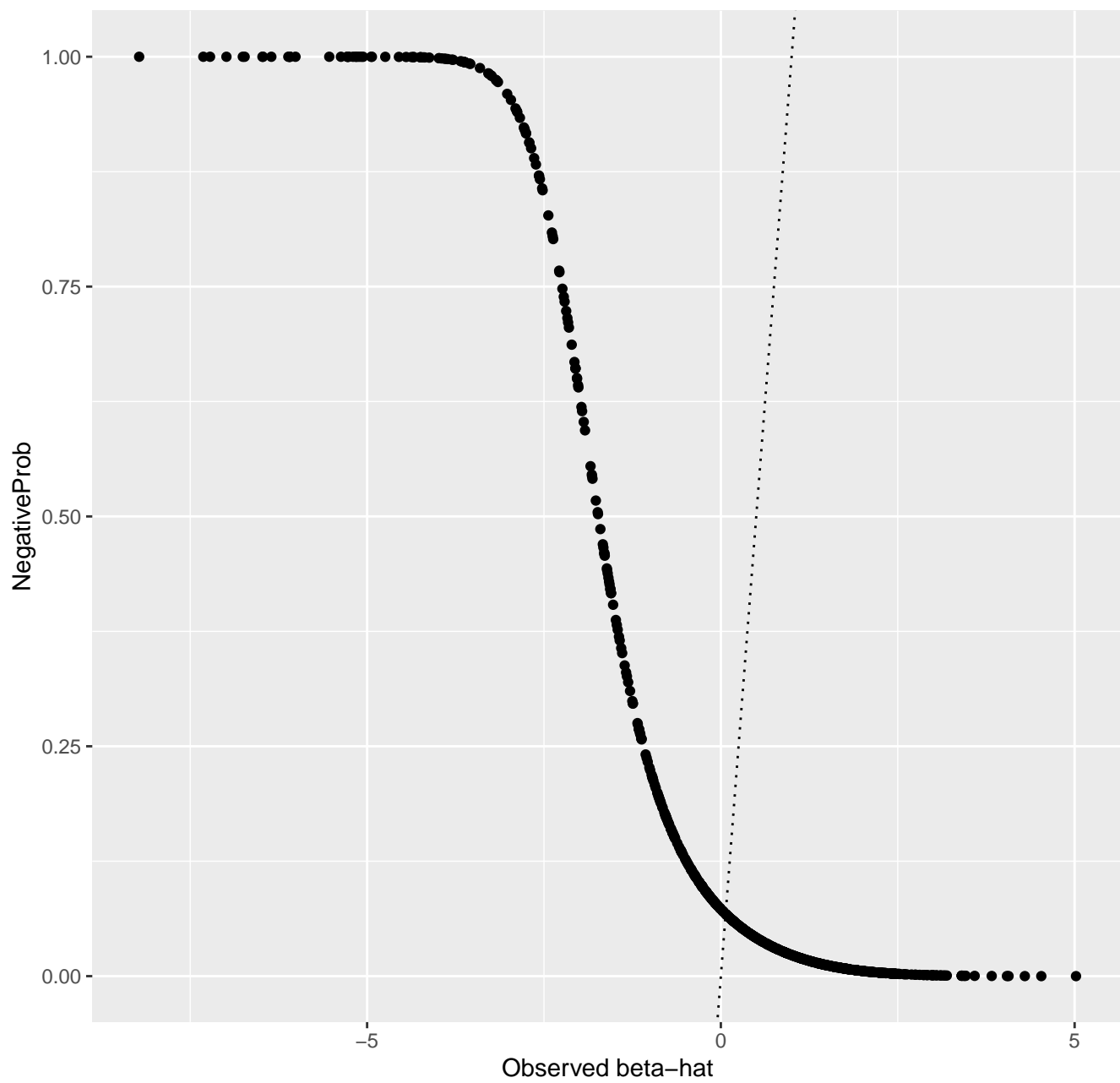
CCND1_(595) positive



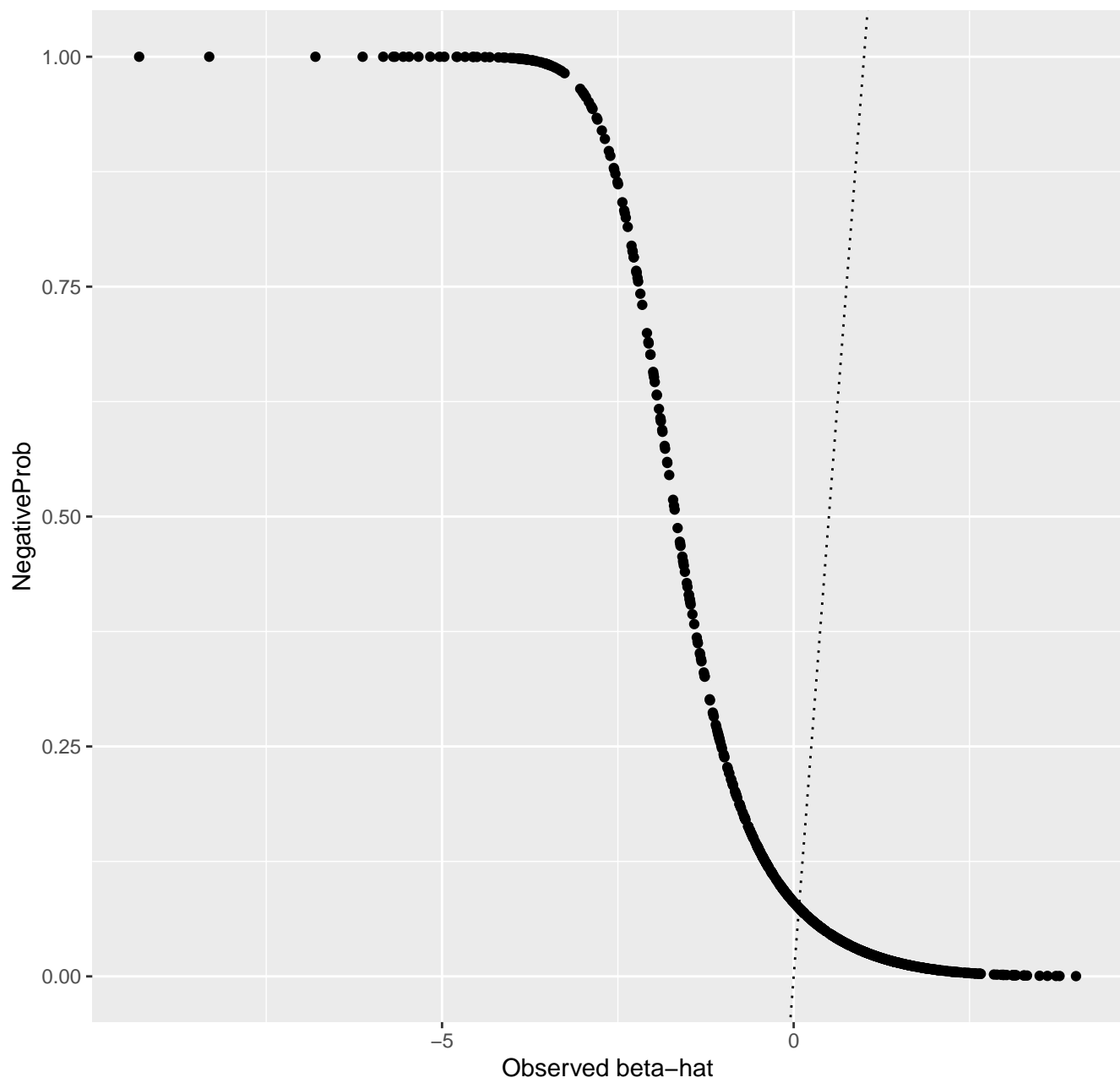
TPK1_(27010) negative



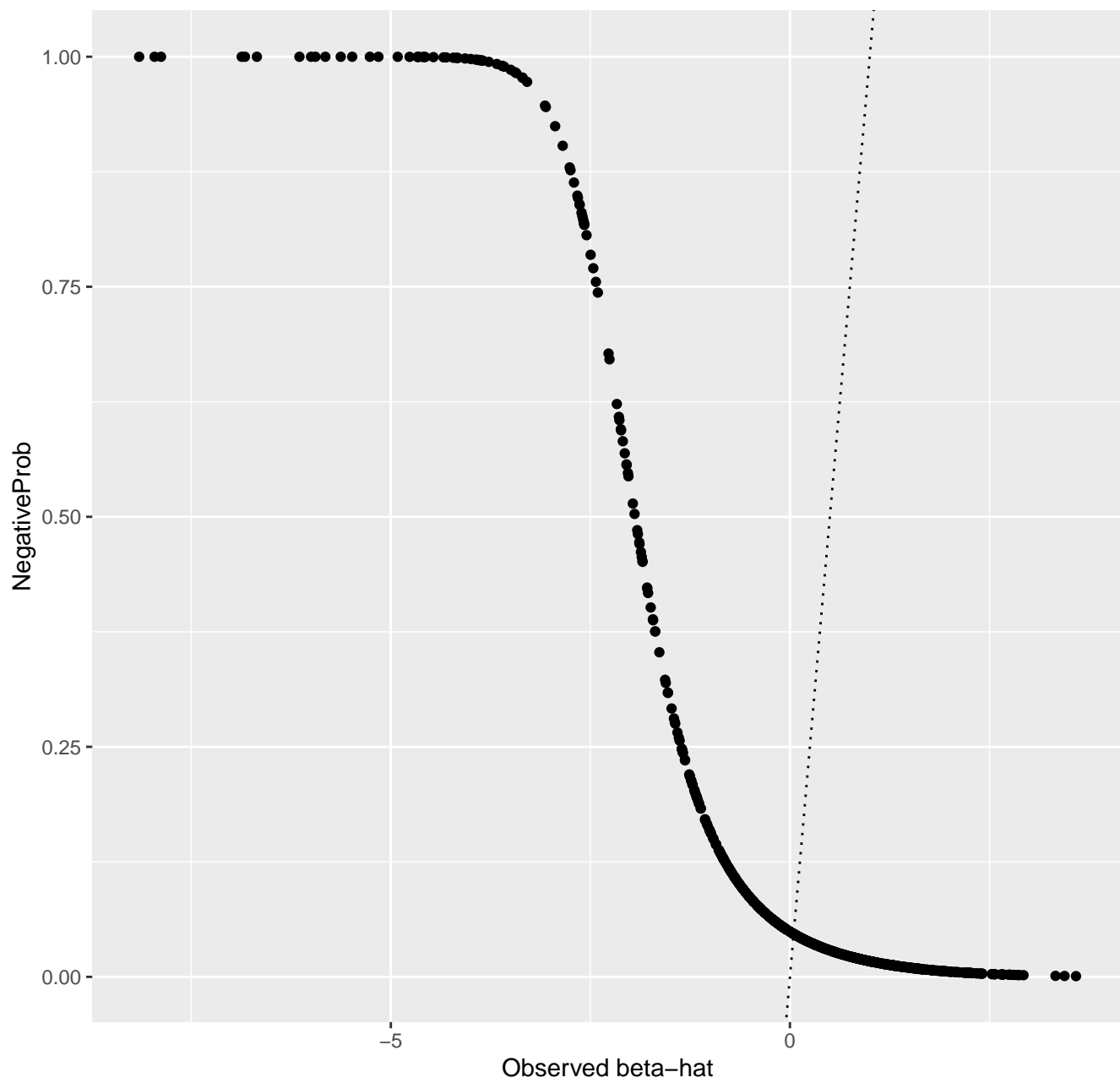
PPCDC_(60490) negative



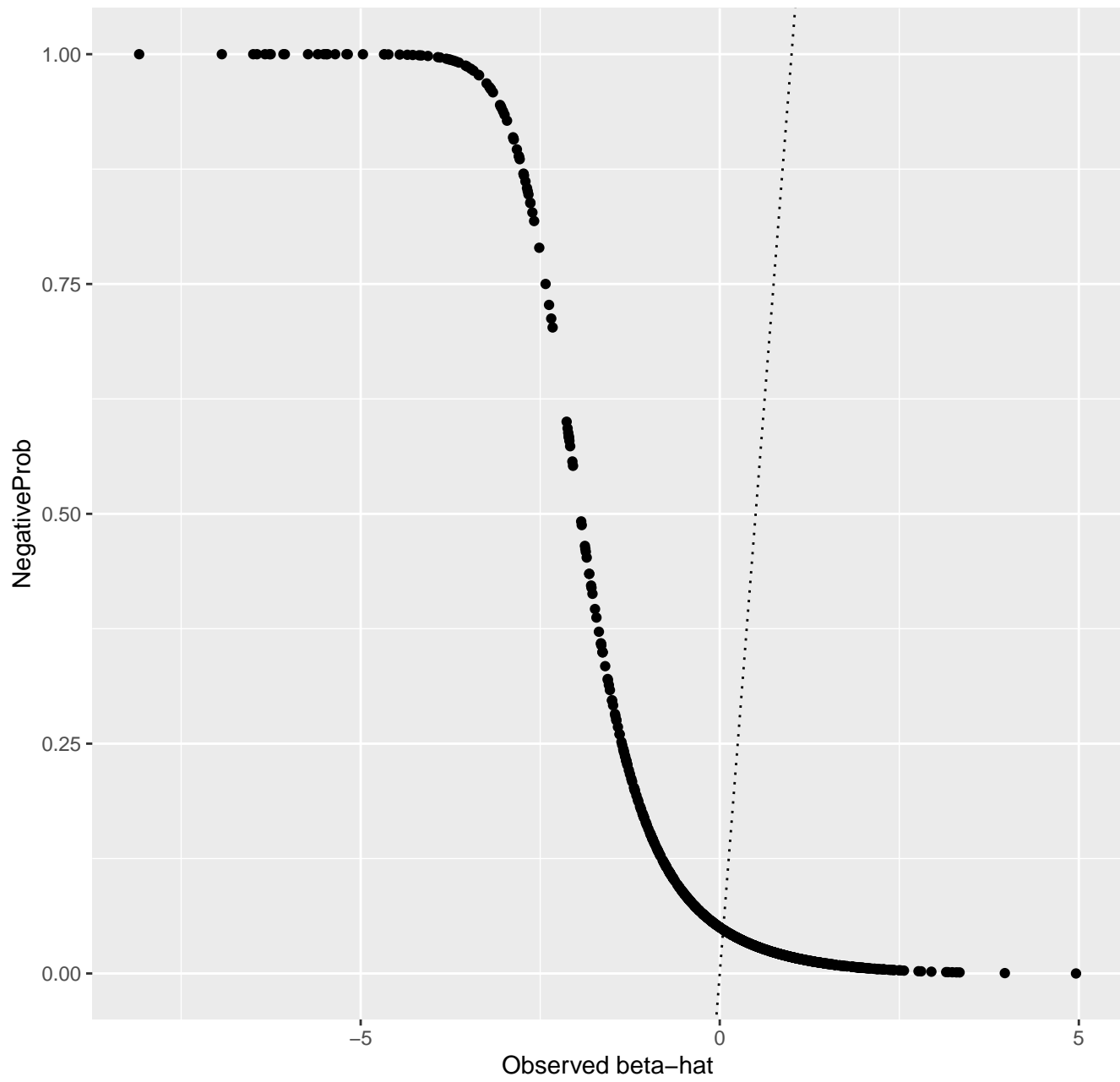
DNAJC19_(131118) negative



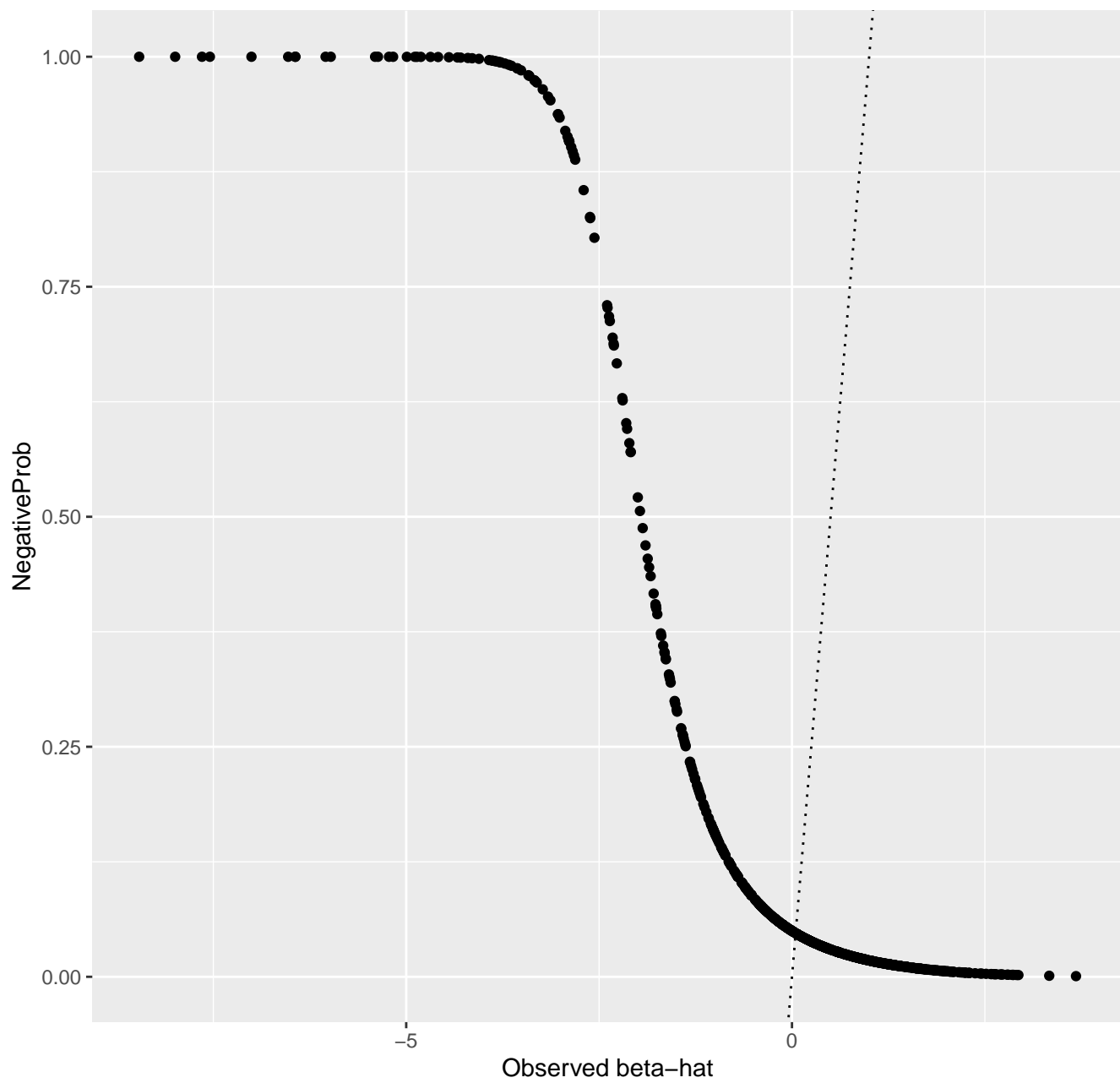
ASNS_(440) negative



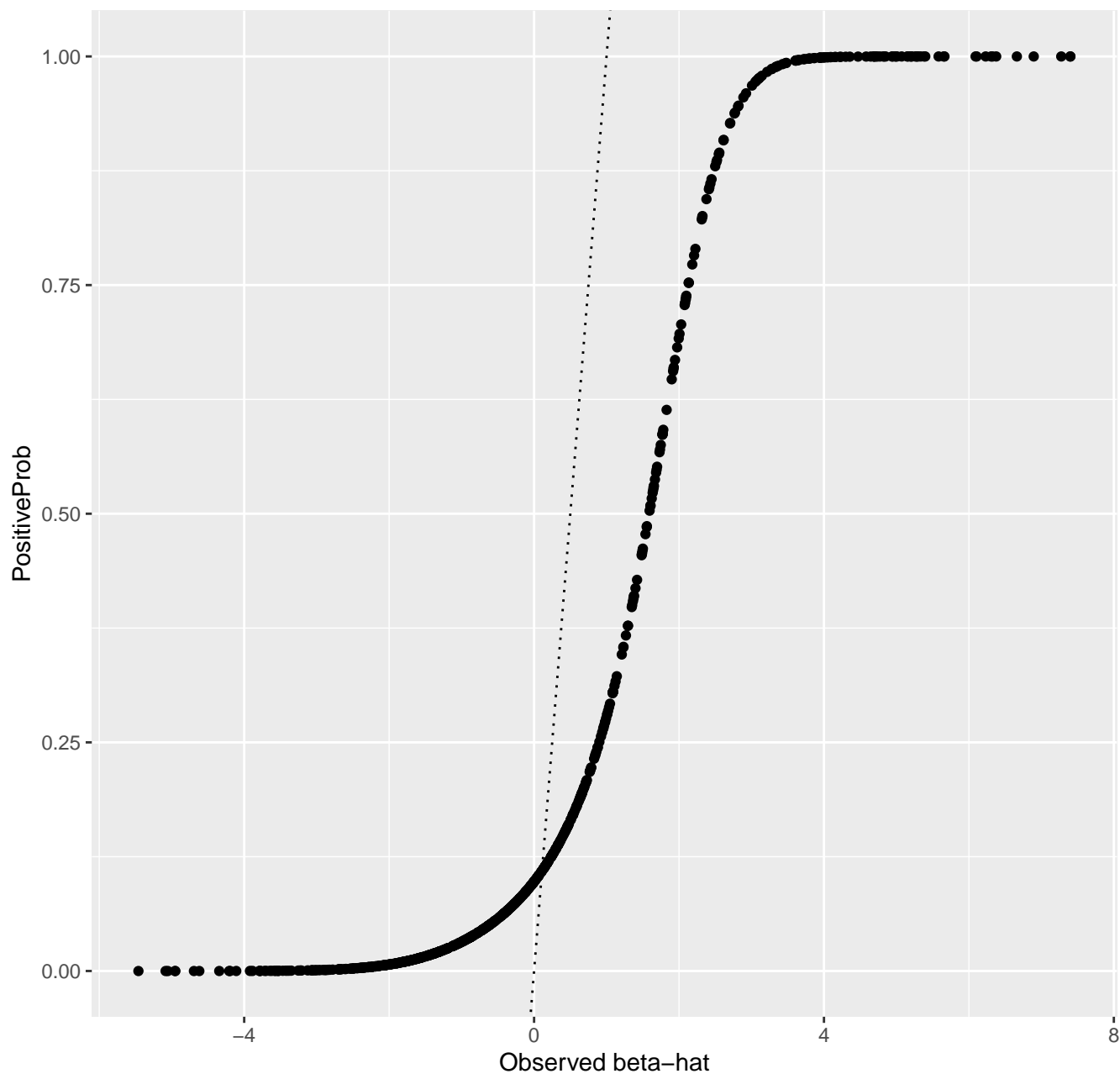
UAP1_(6675) negative



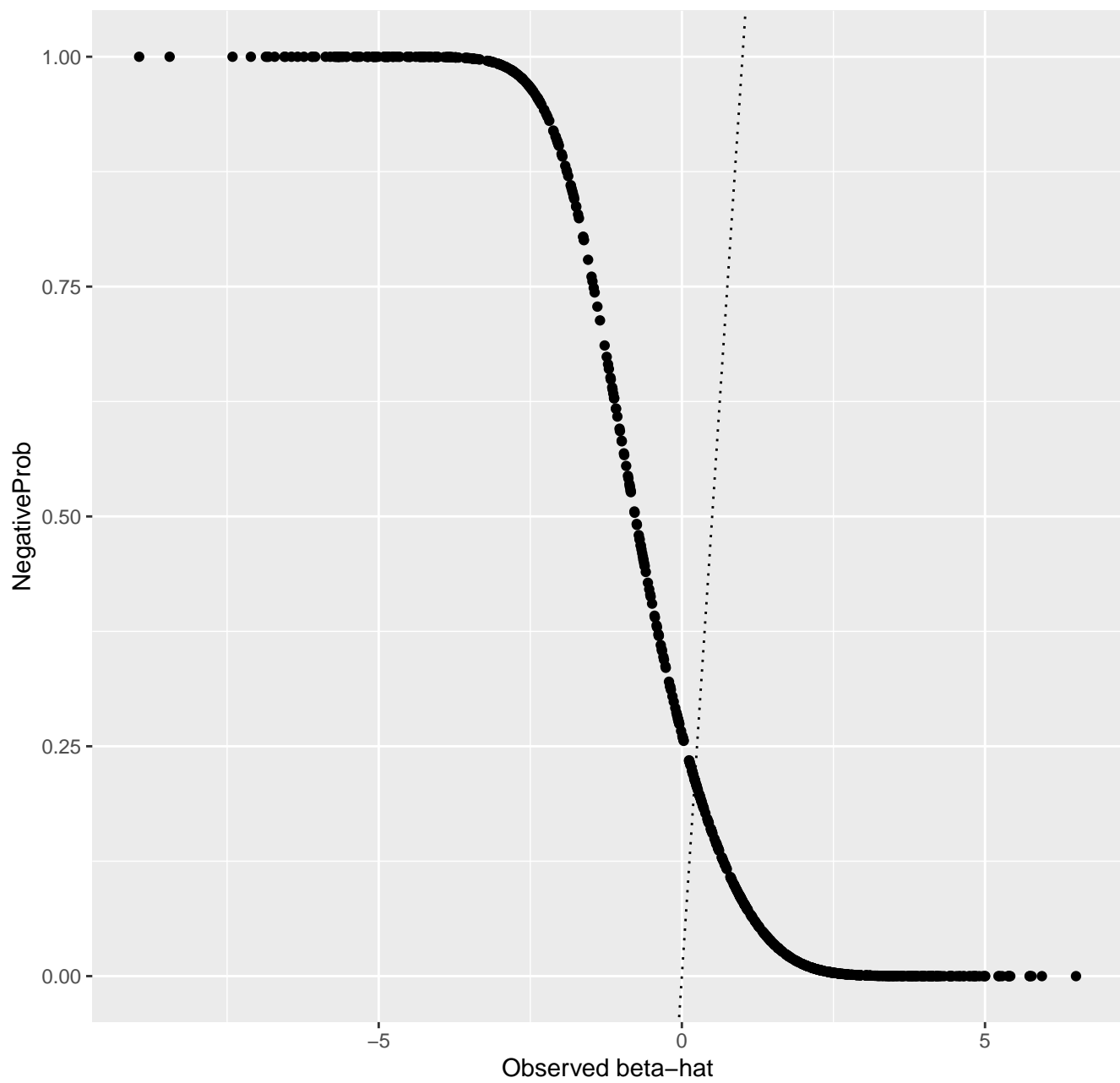
SNAI2_(6591) negative



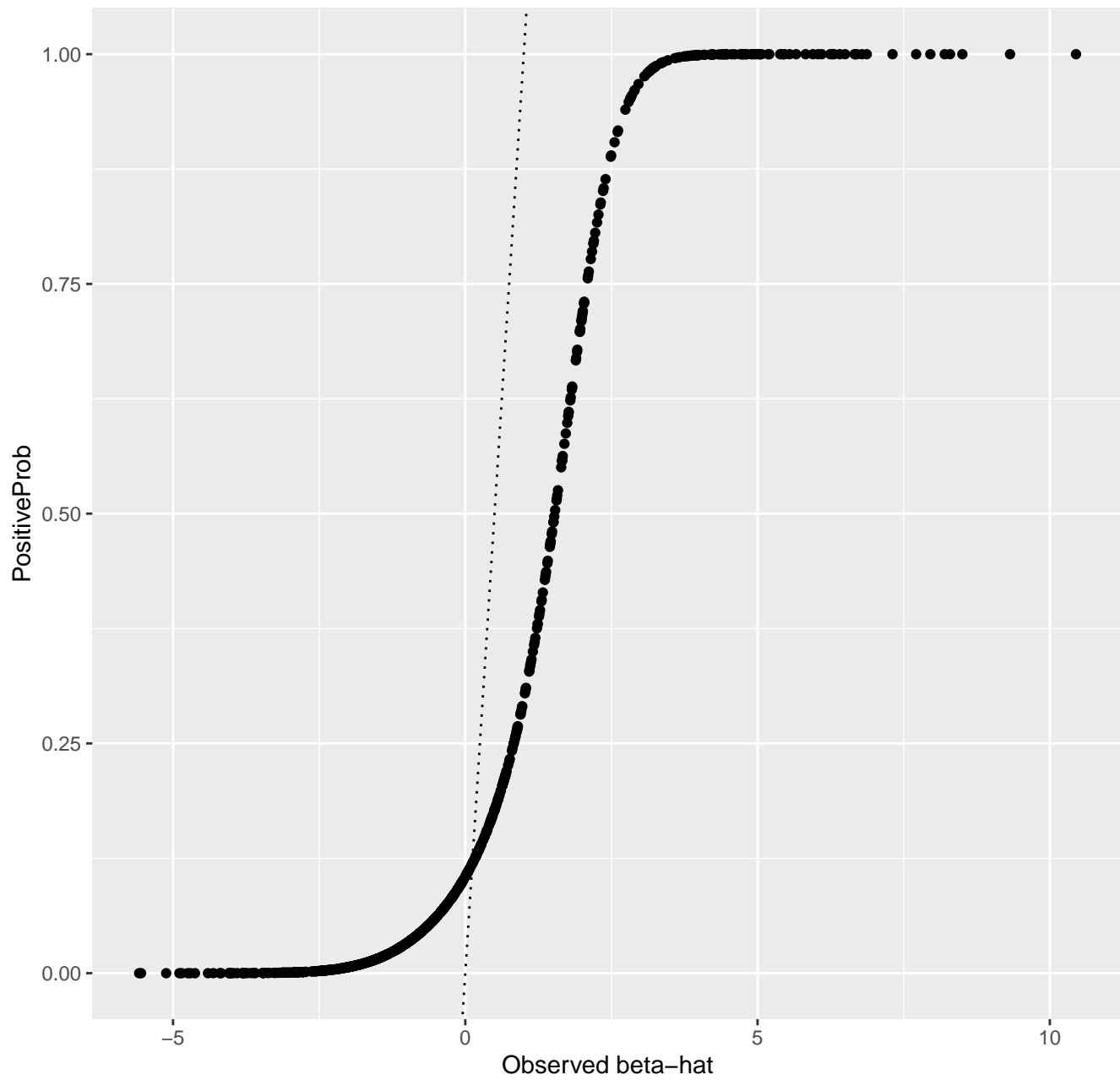
MVK_(4598) positive



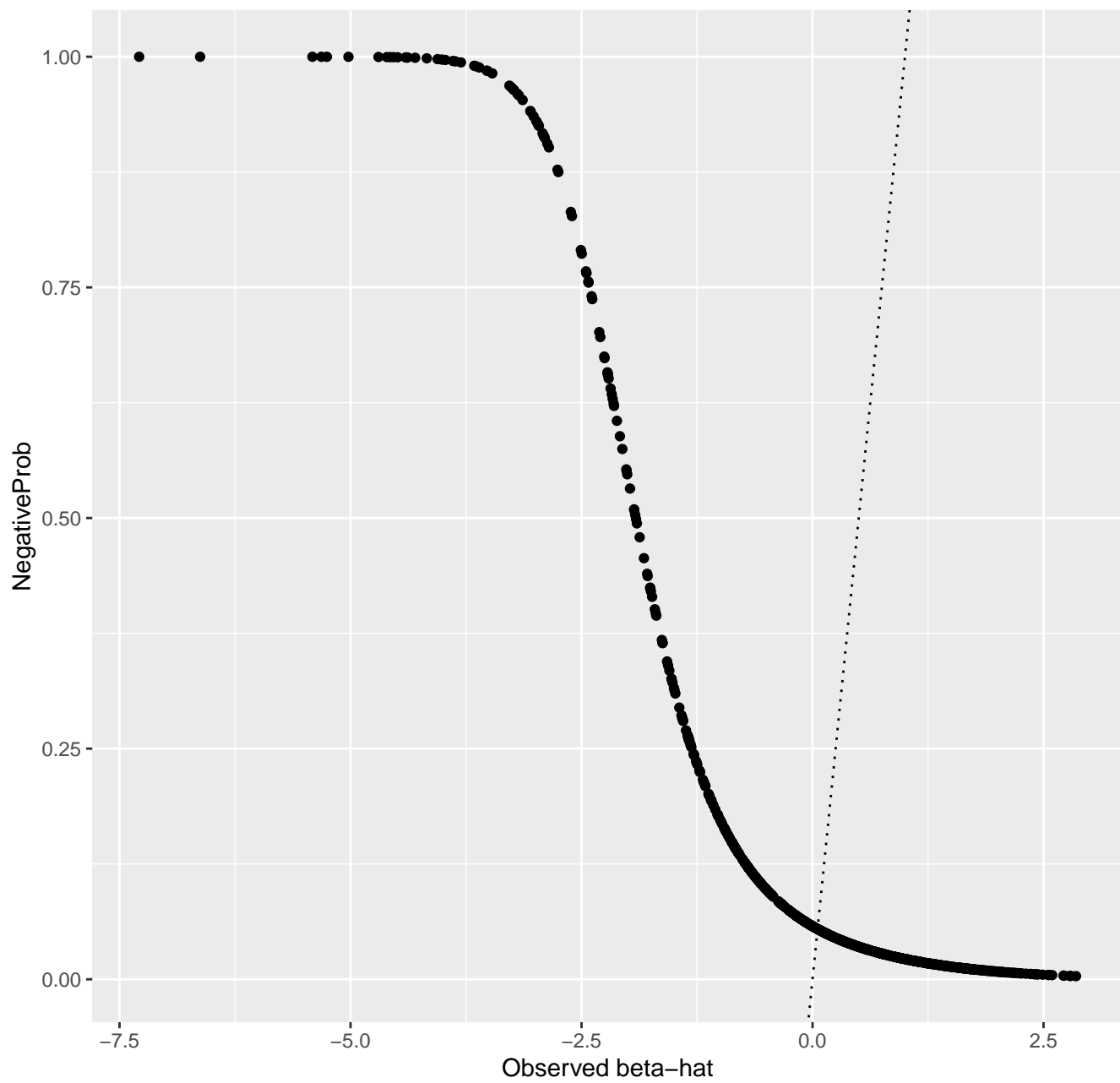
FERMT2_(10979) negative



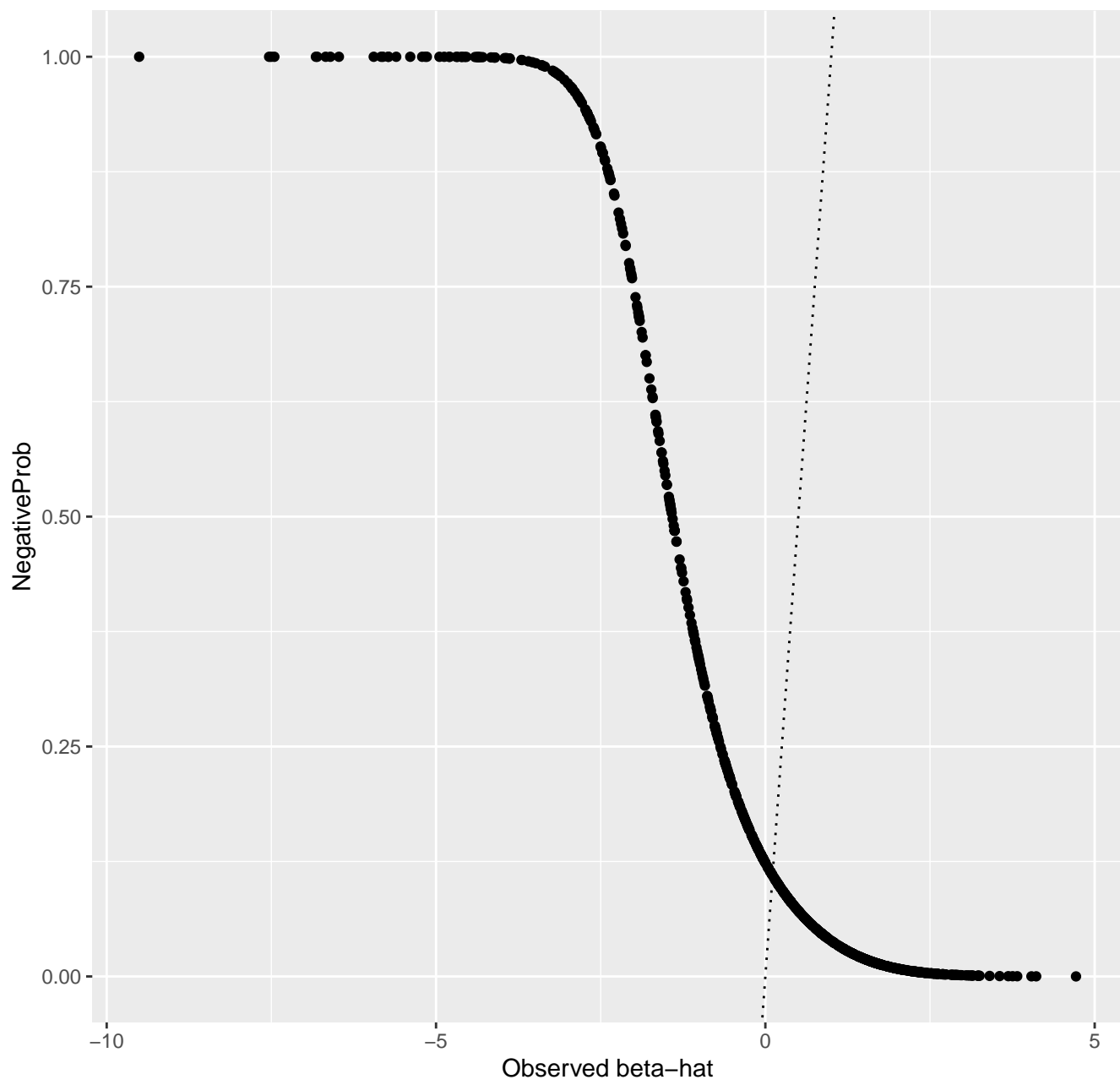
PPP1R12A_(4659) positive



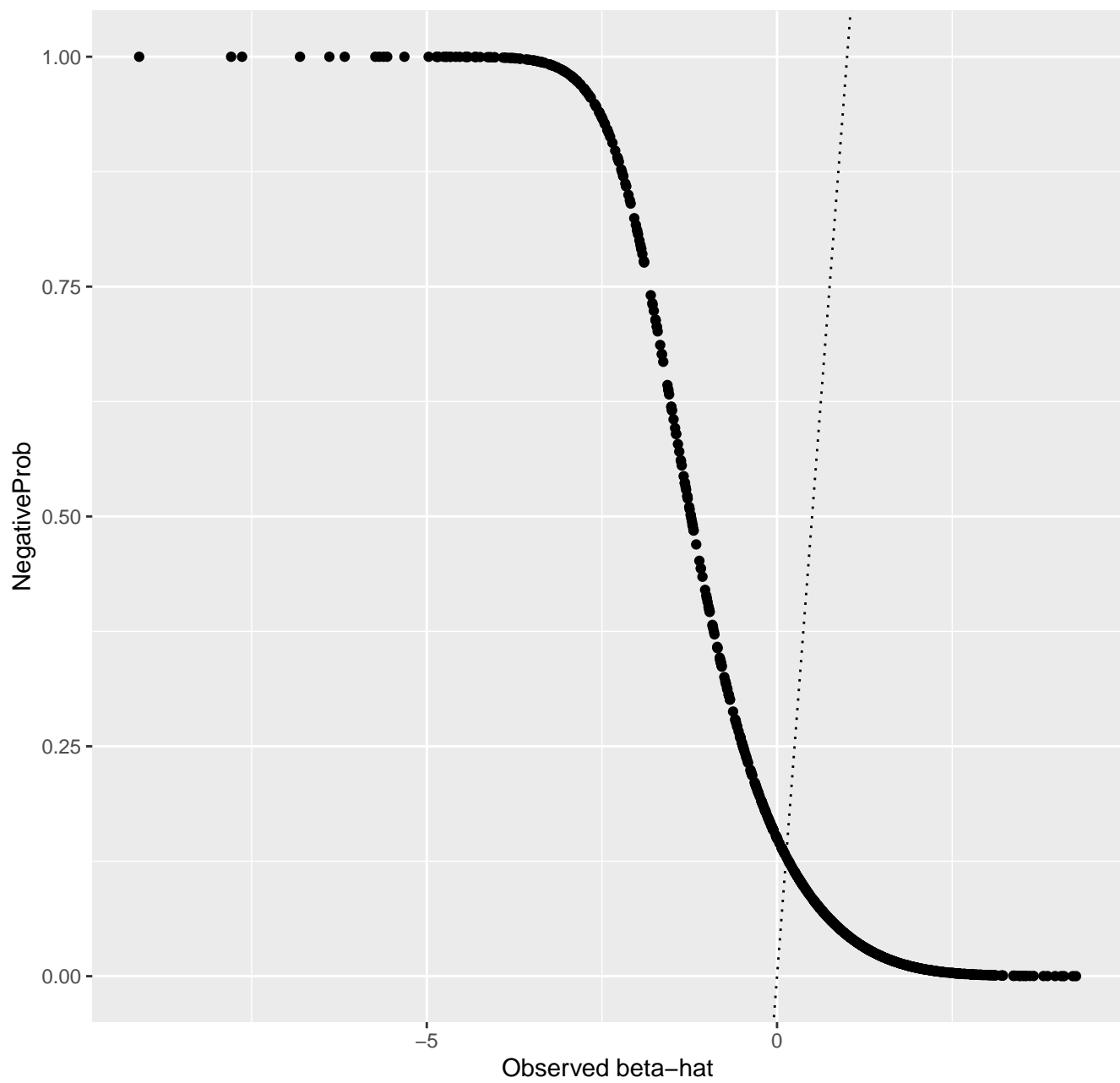
REXO2_(25996) negative



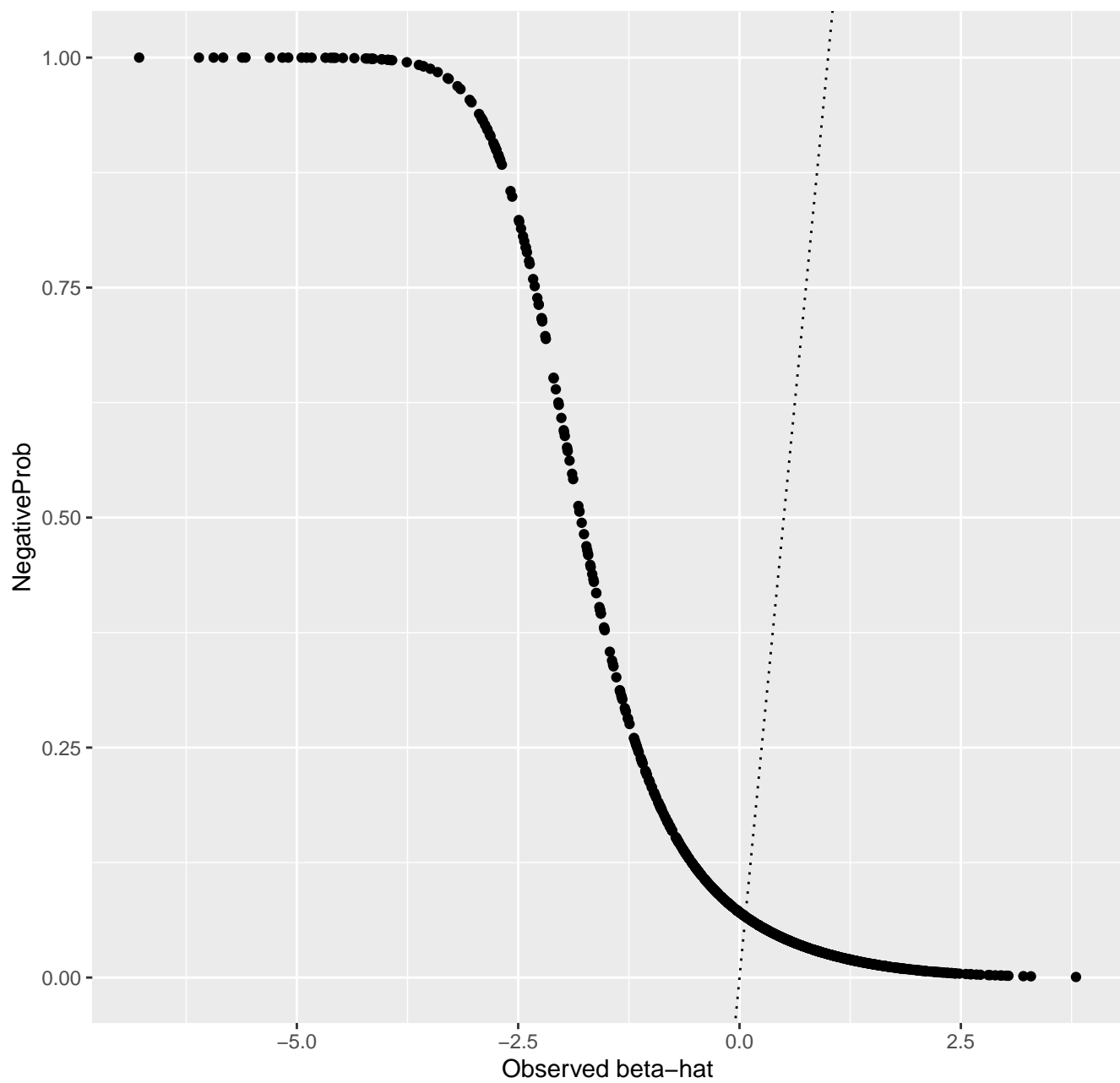
ITGB5_(3693) negative



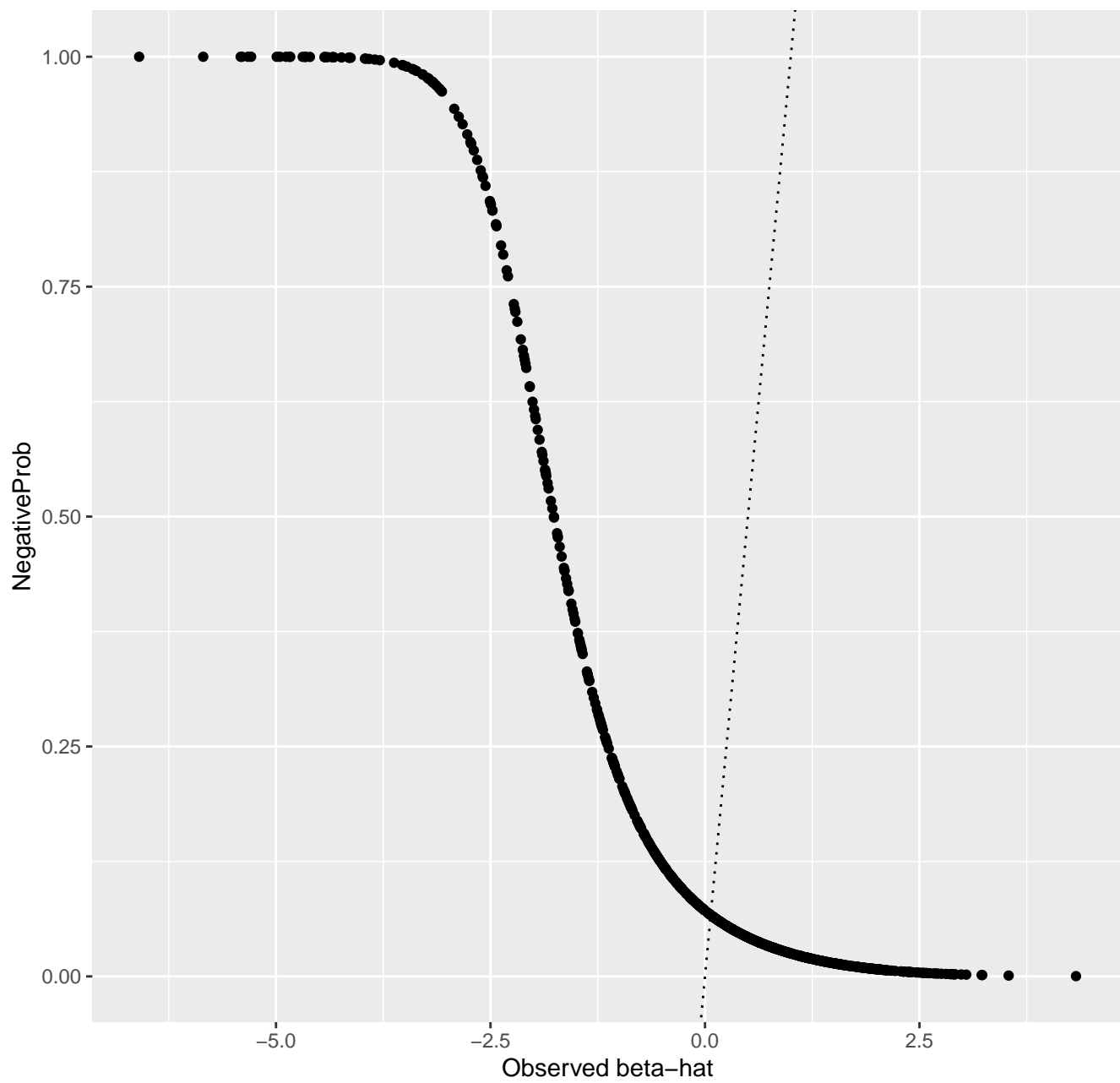
IPPK_(64768) negative



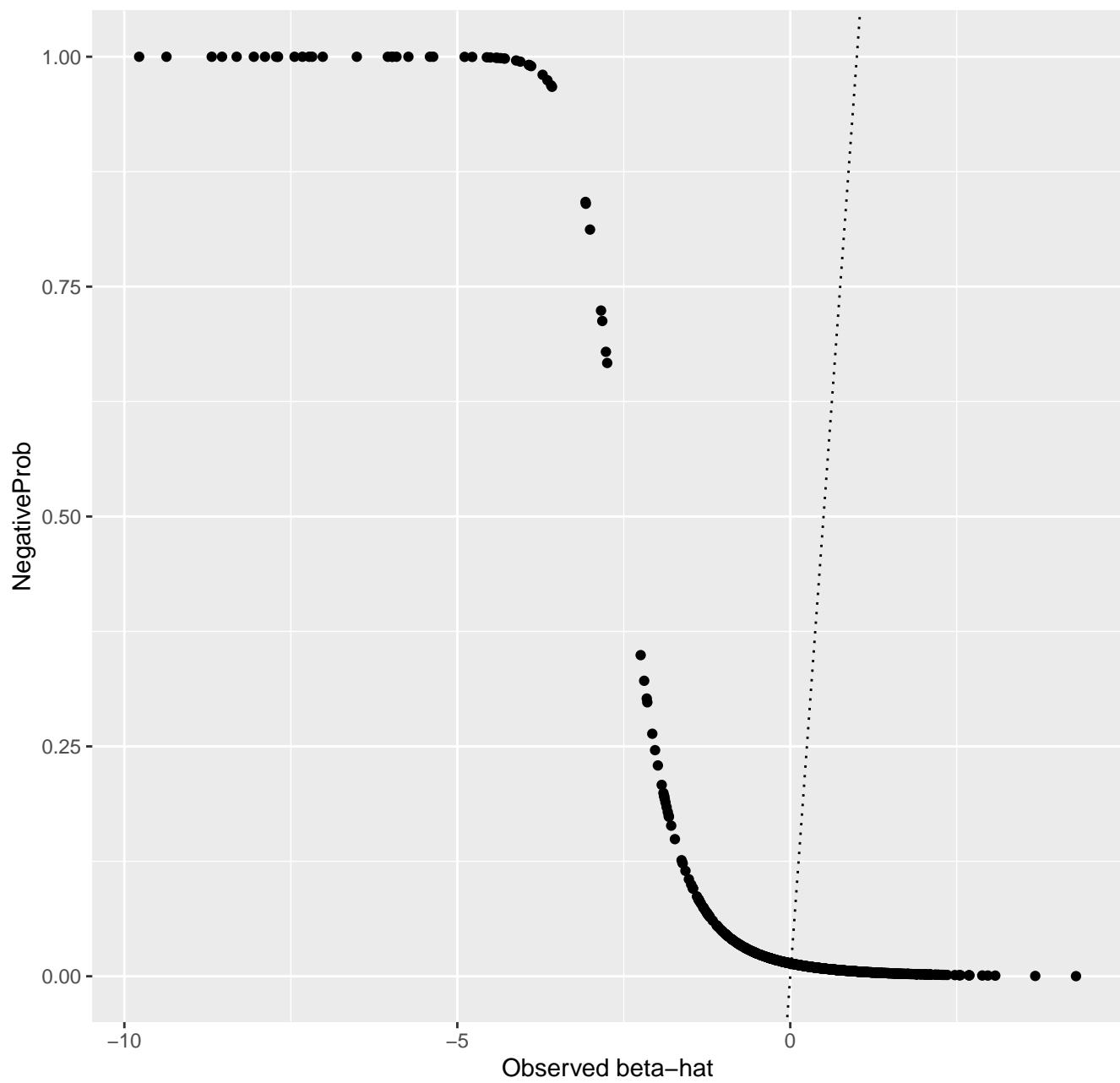
PDE12_(201626) negative



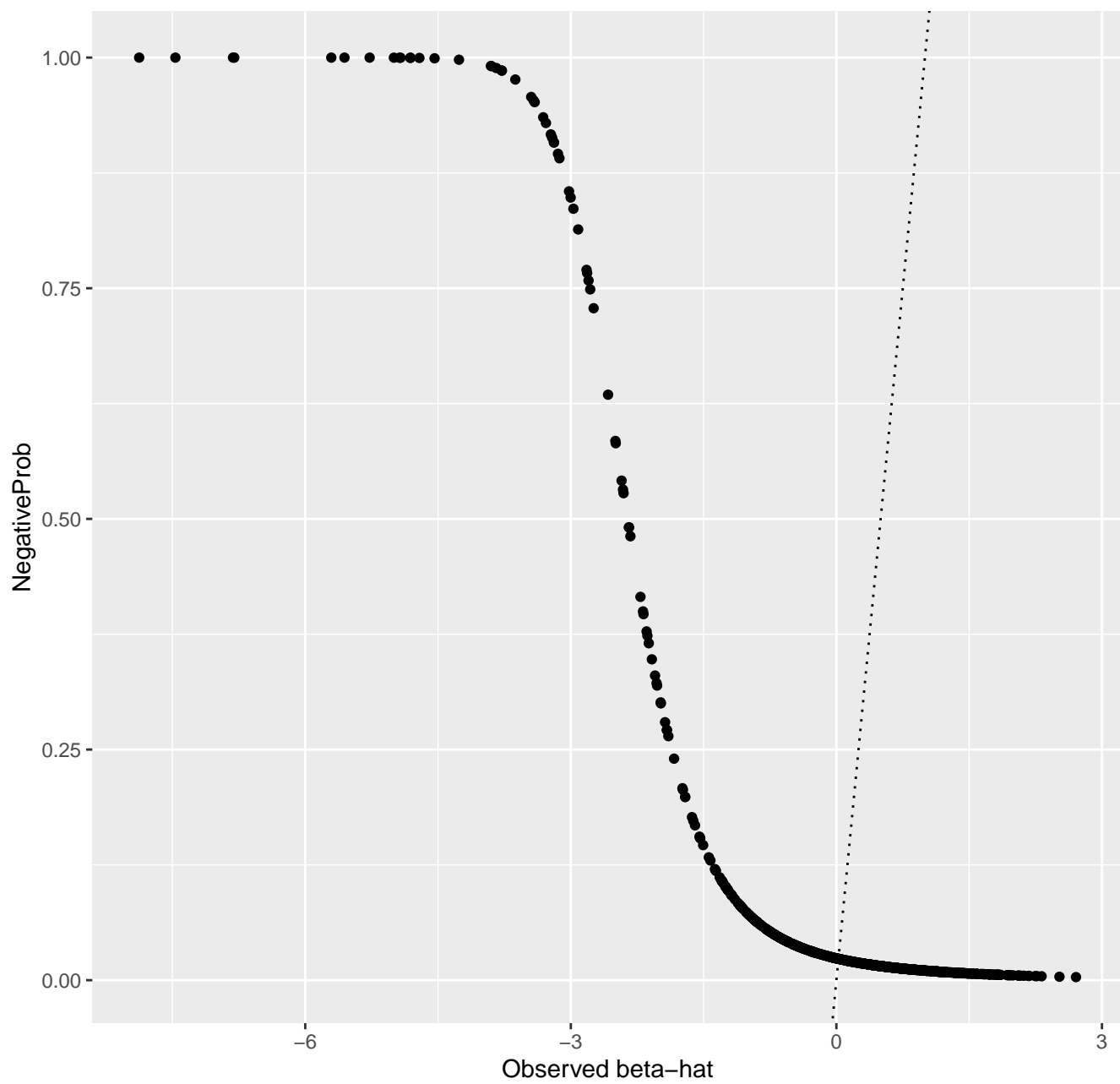
TDP2_(51567) negative



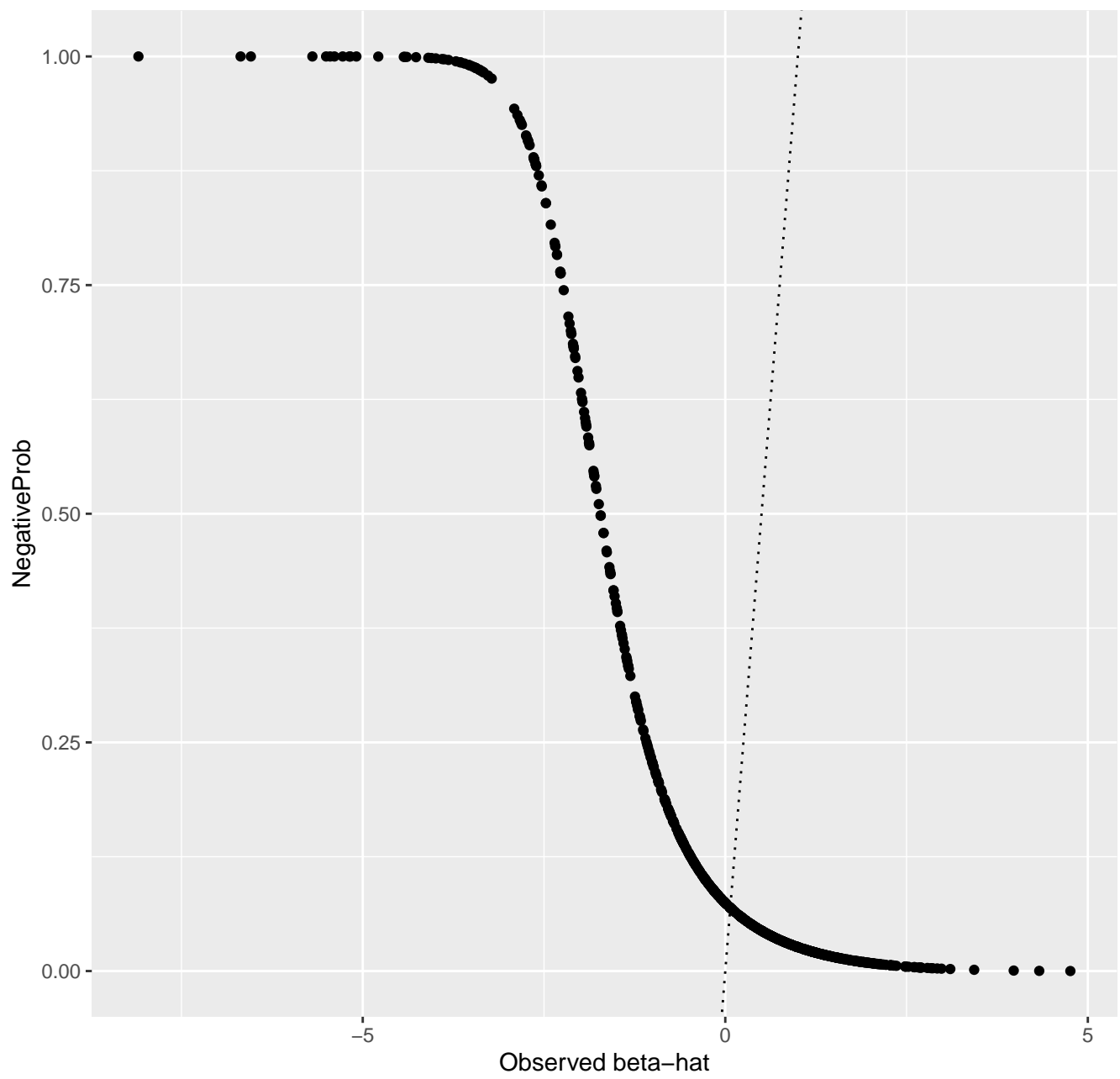
MYB_(4602) negative



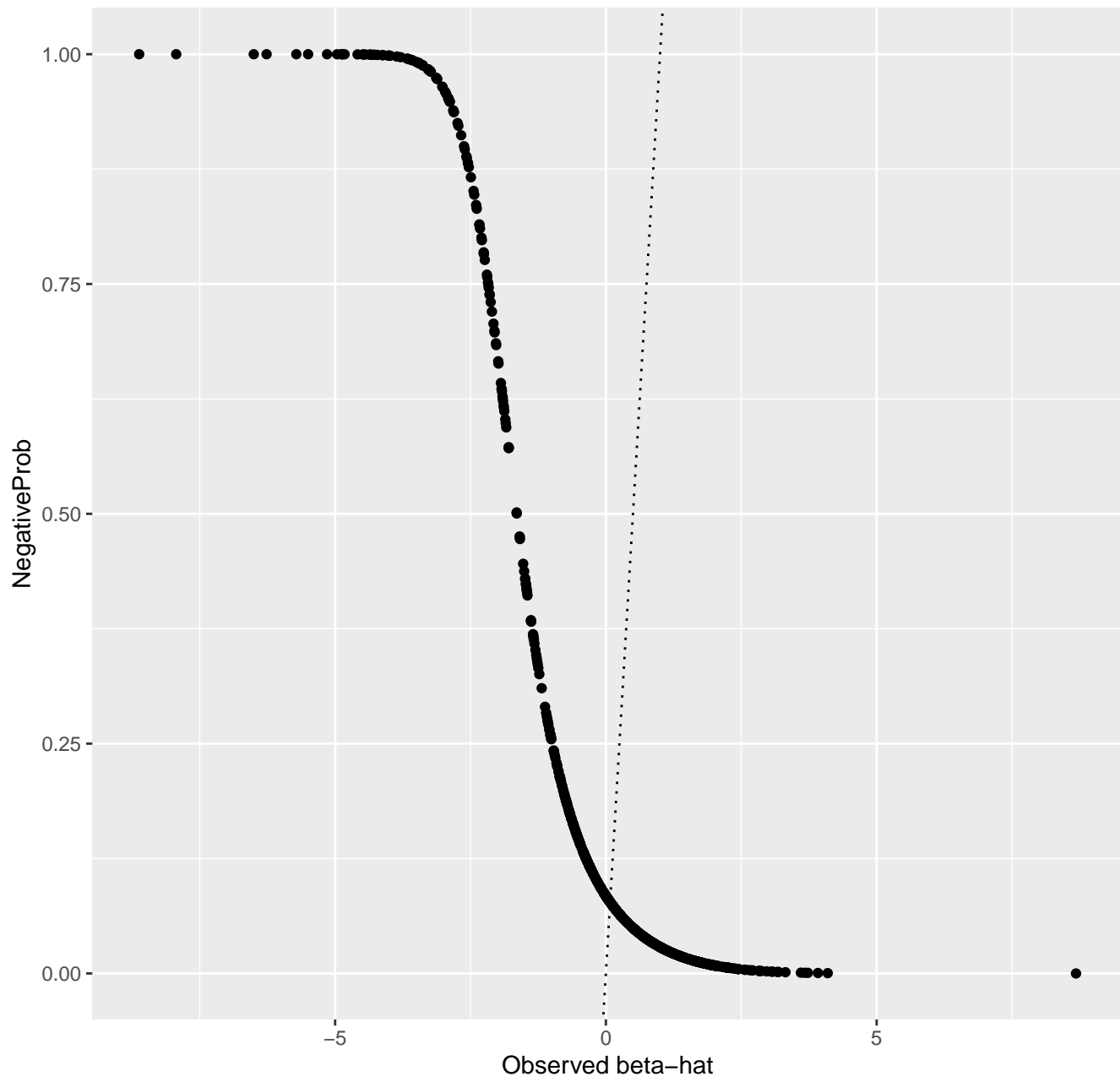
FITM2_(128486) negative



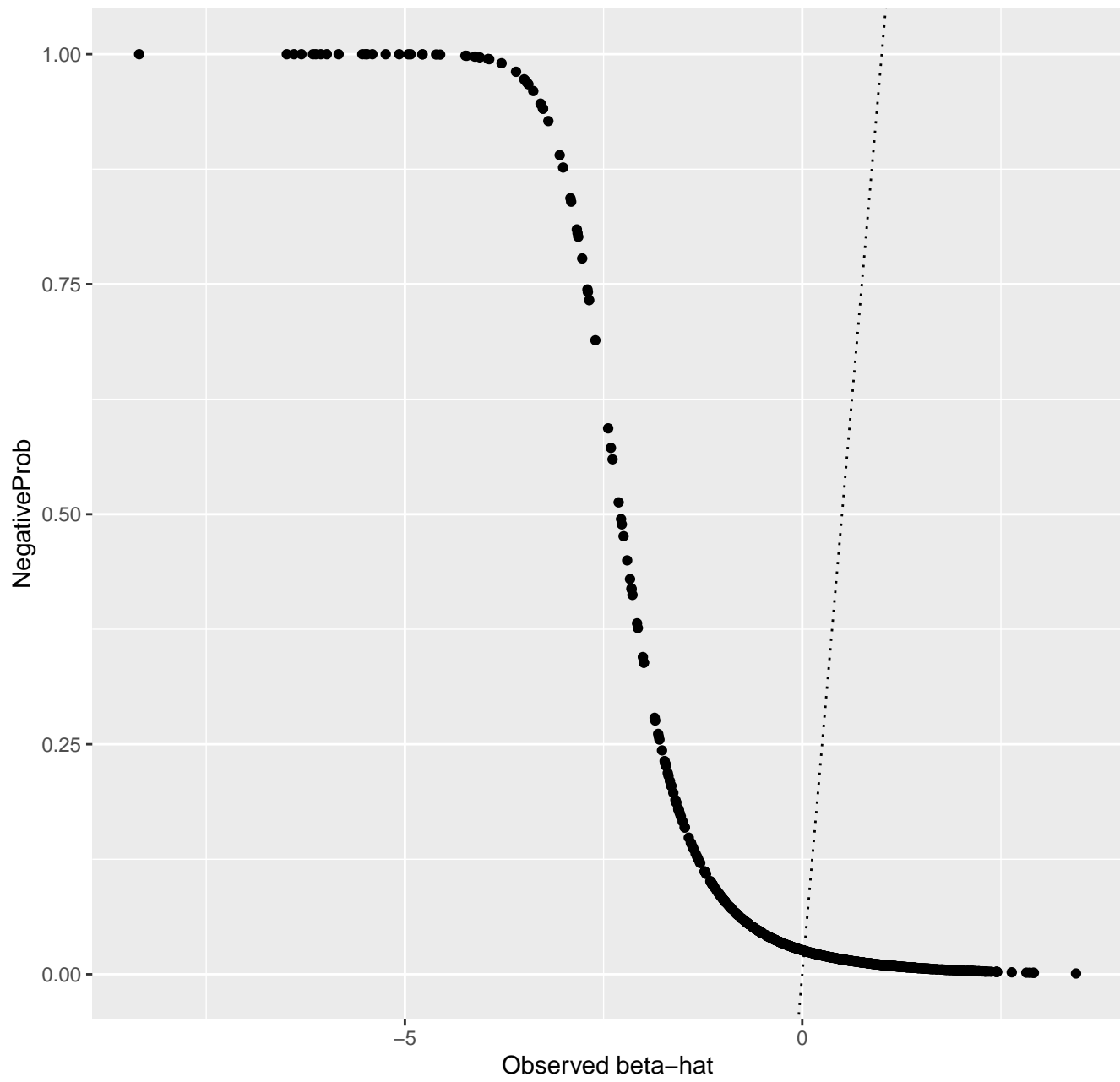
ZEB1_(6935) negative



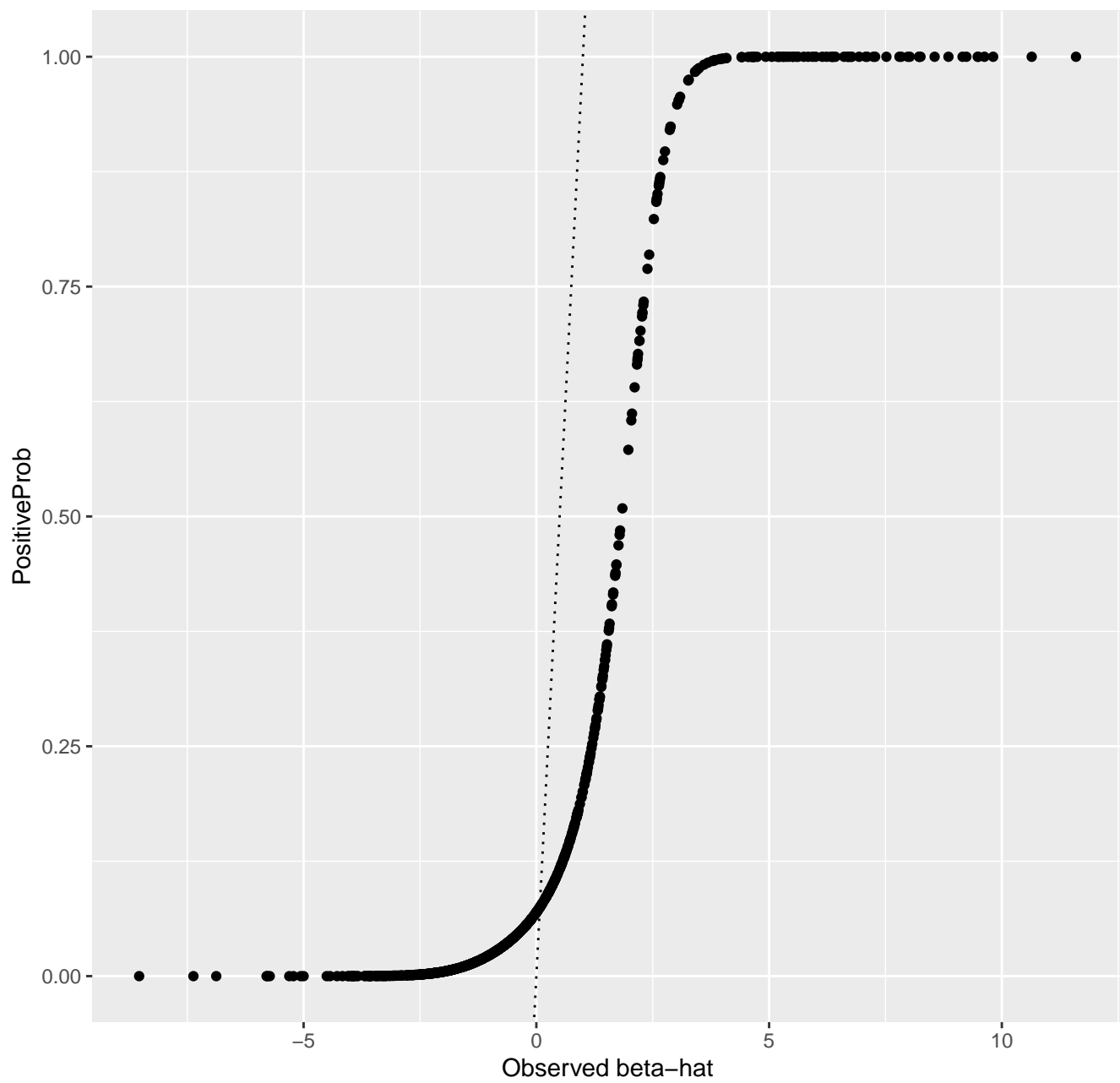
RNF31_(55072) negative



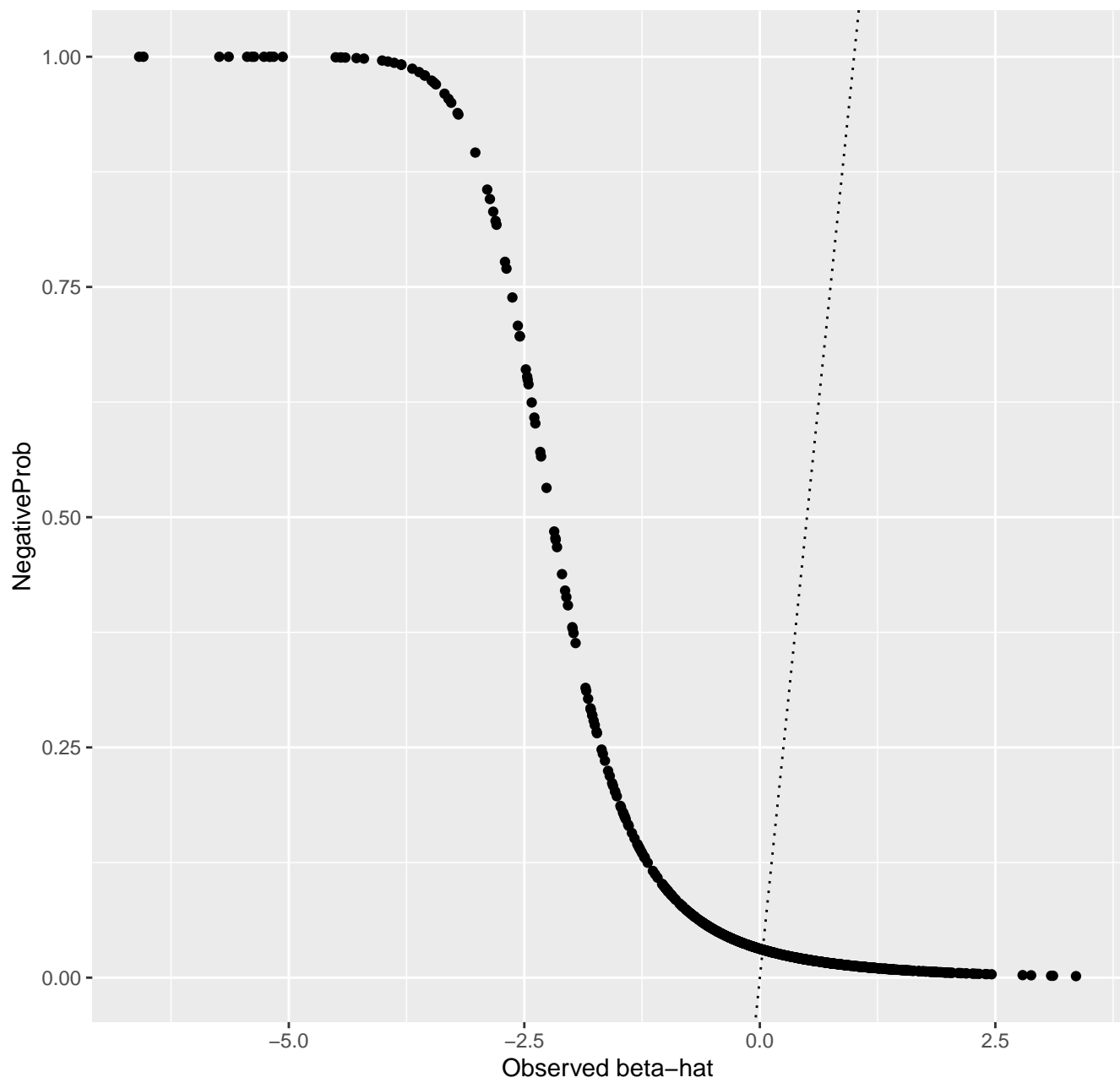
GOT2_(2806) negative



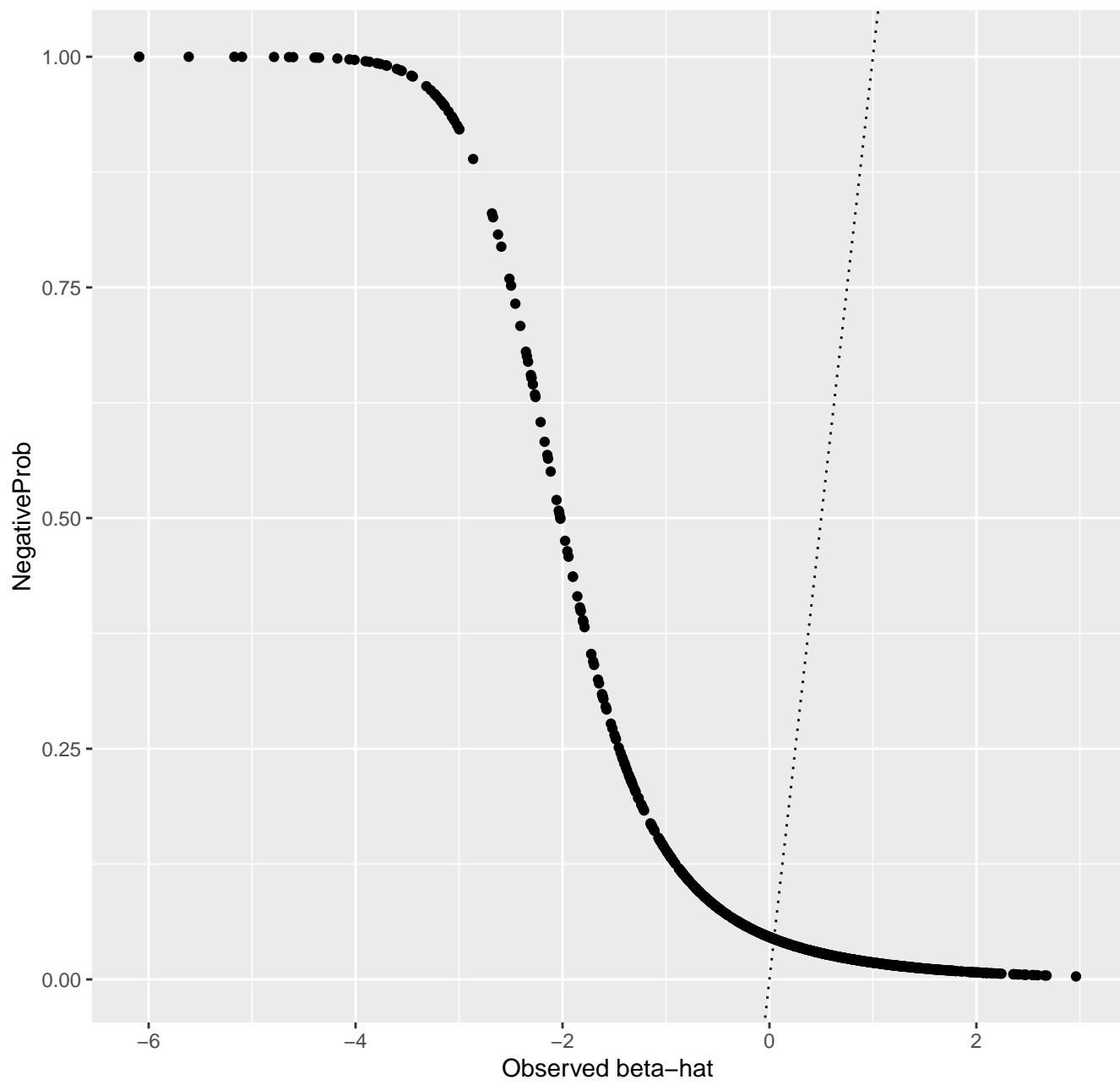
PSMB5_(5693) positive



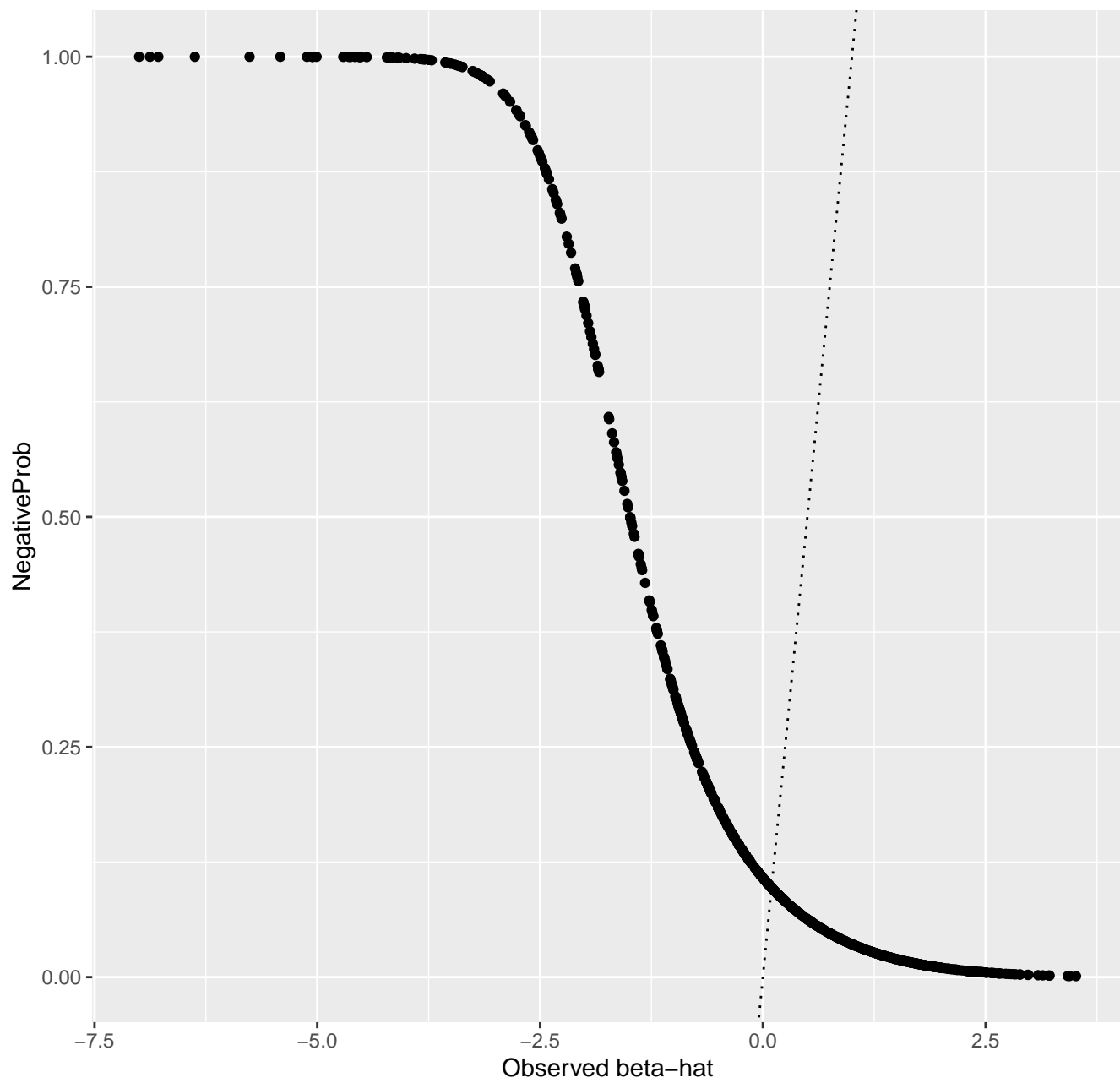
PRKCI_(5584) negative



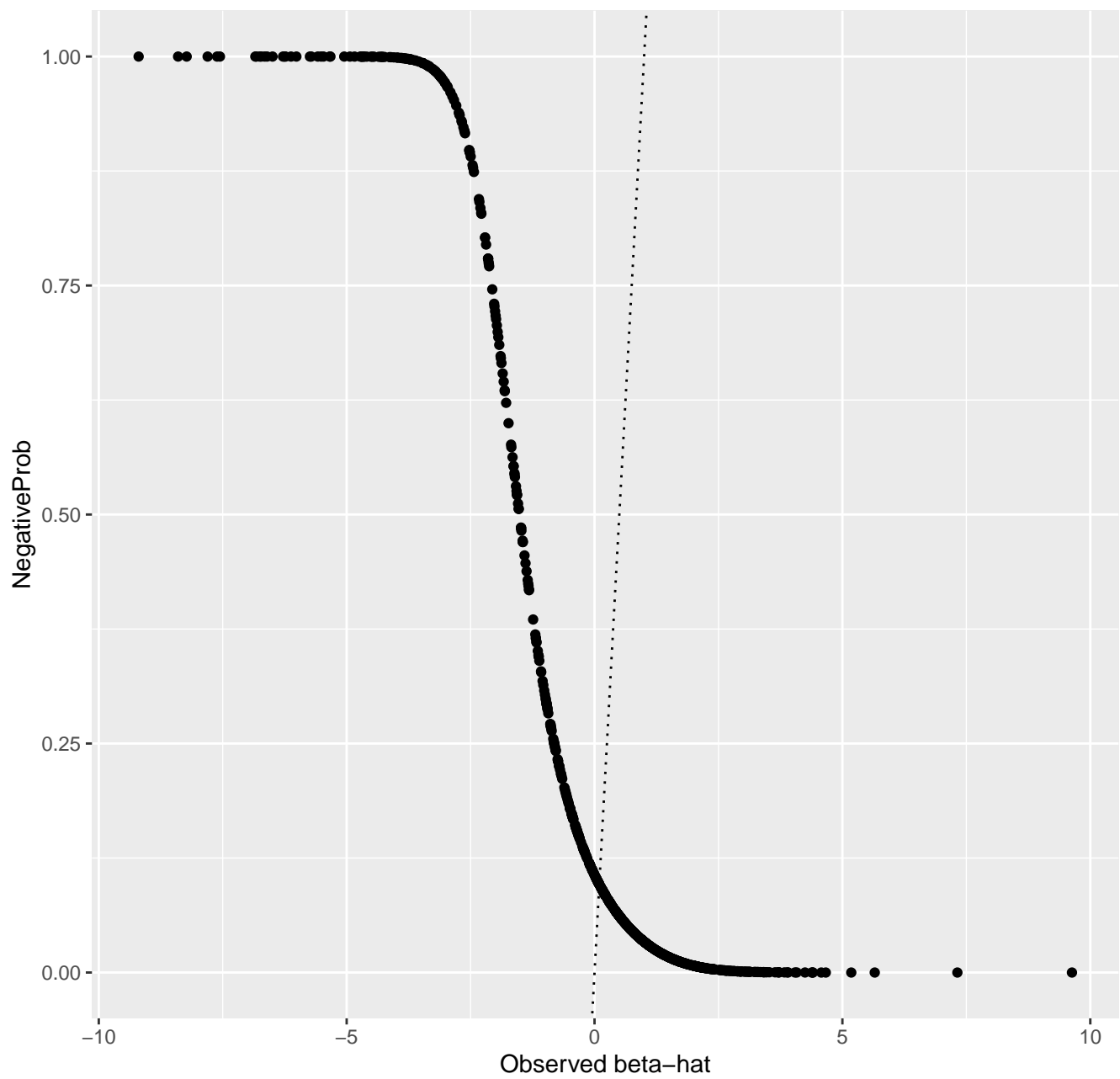
RNF25_(64320) negative



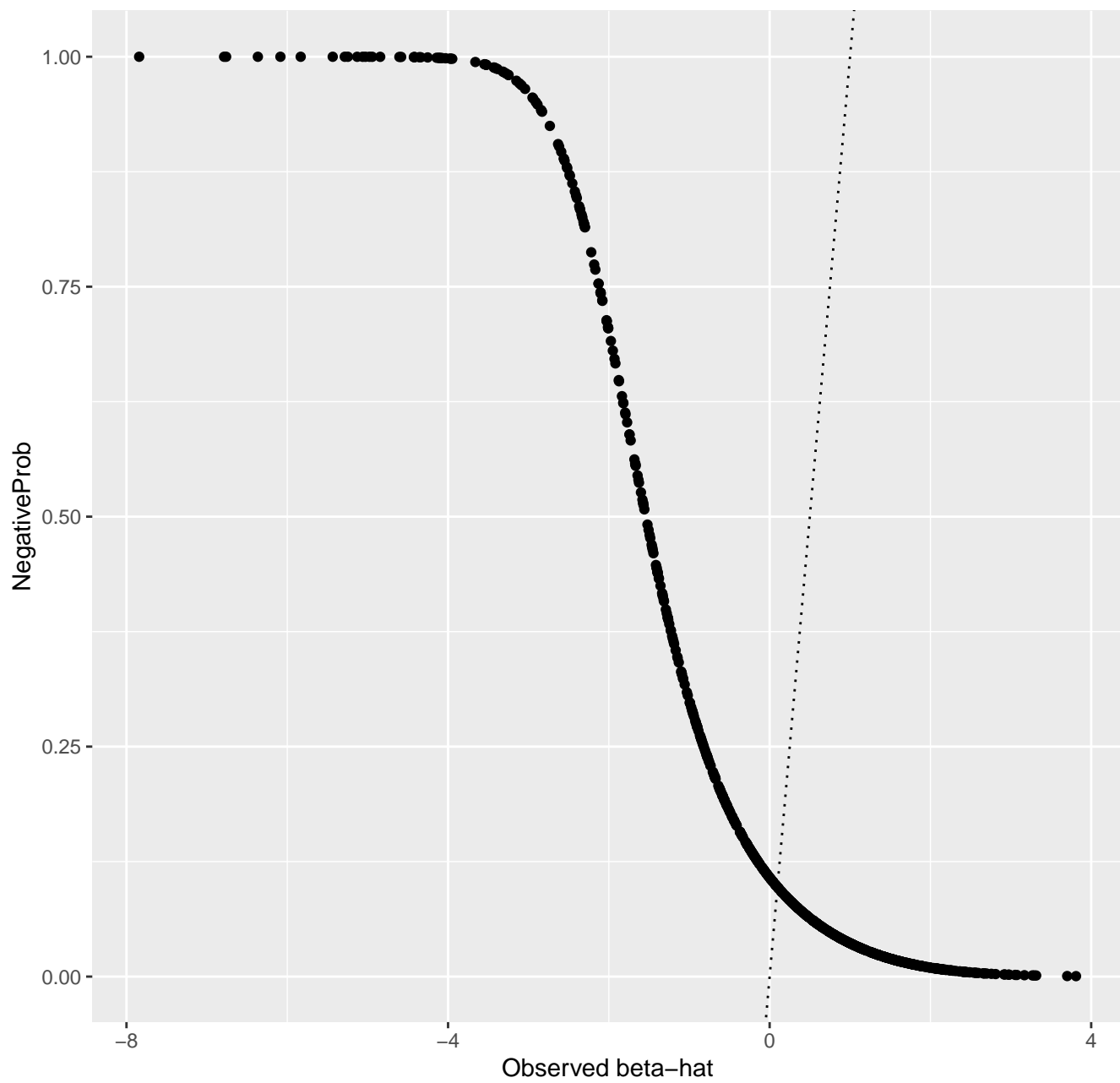
SLC30A9_(10463) negative



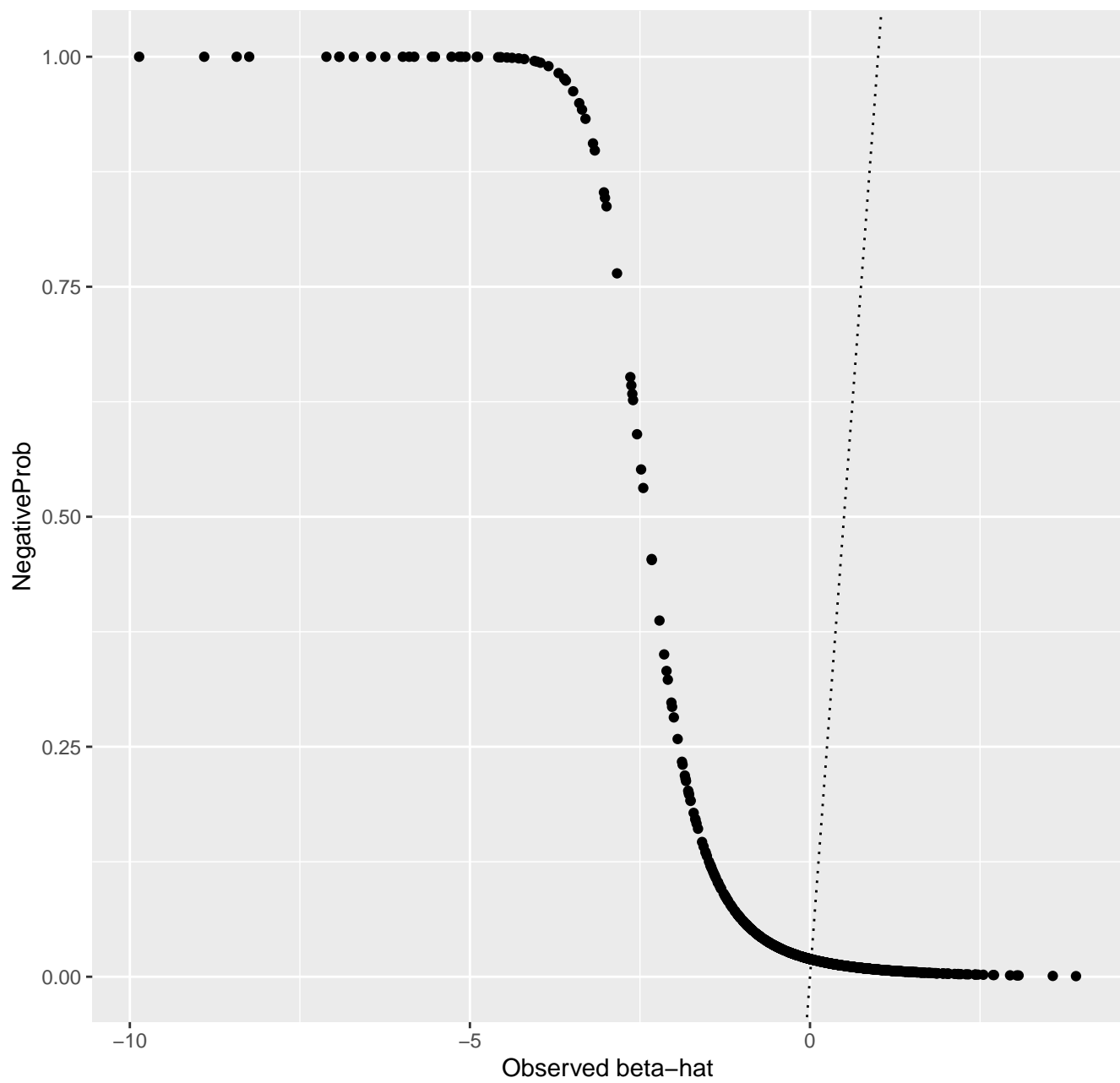
SOCS3_(9021) negative



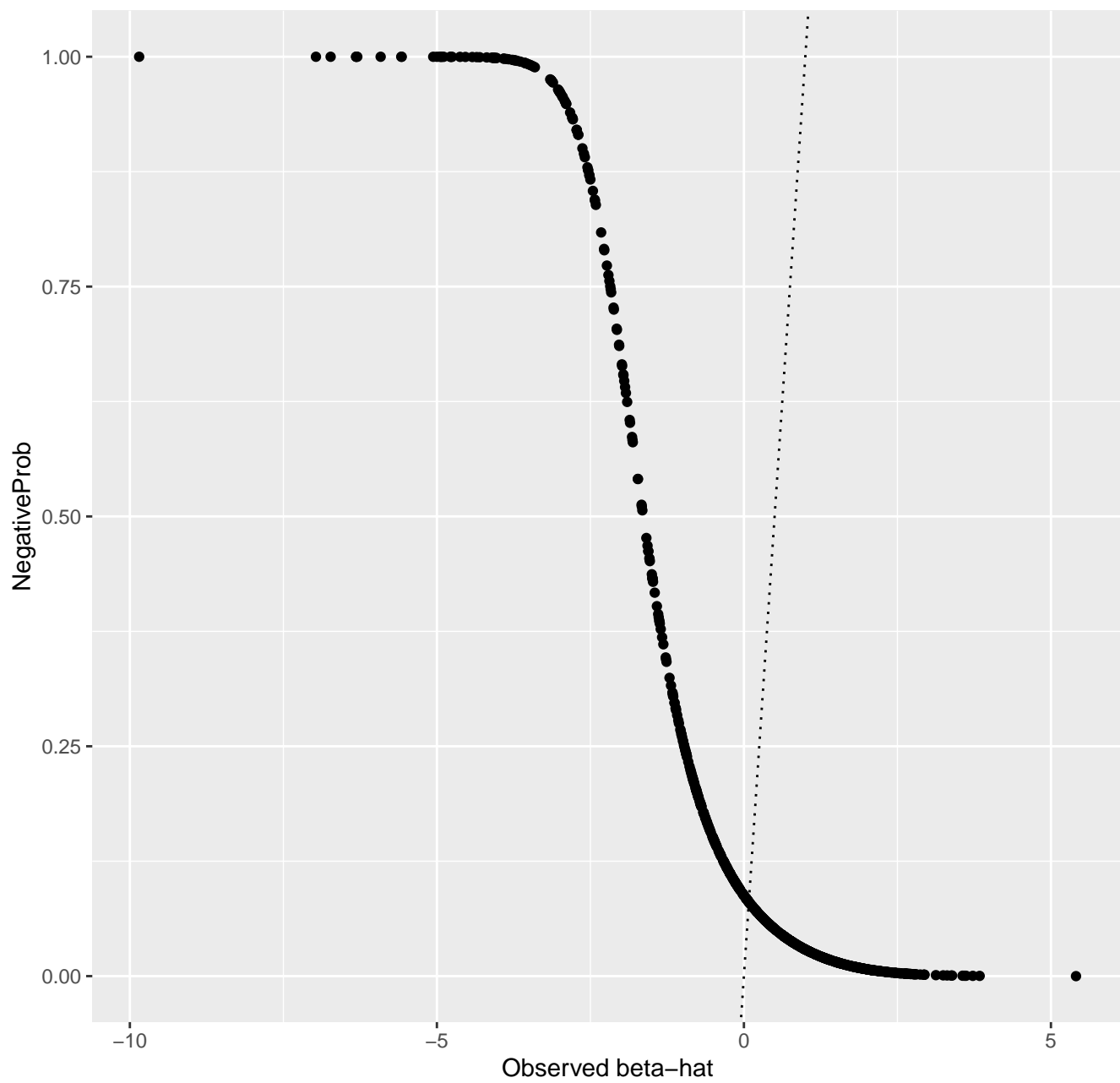
SLC33A1_(9197) negative



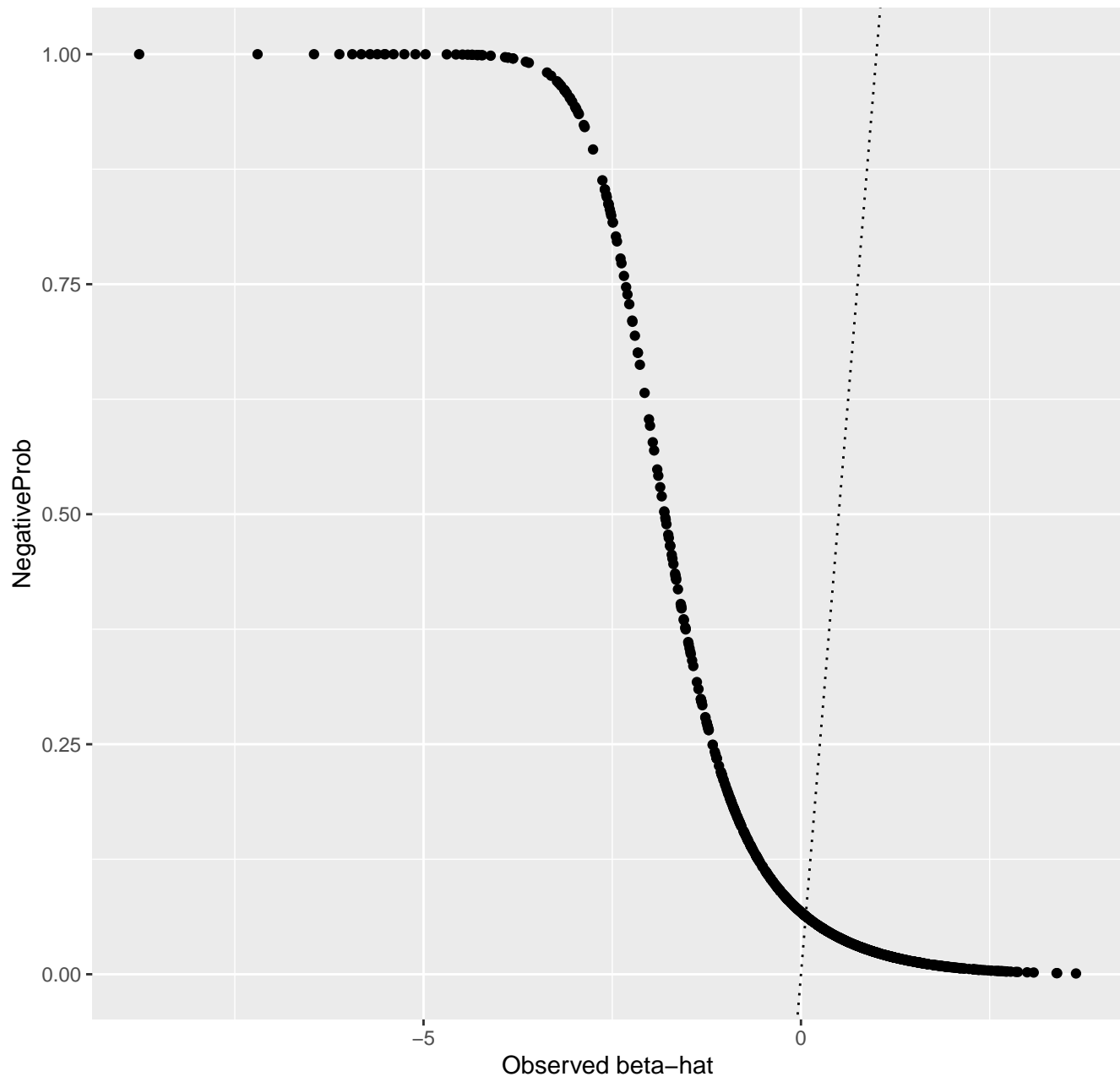
NFE2L2_(4780) negative



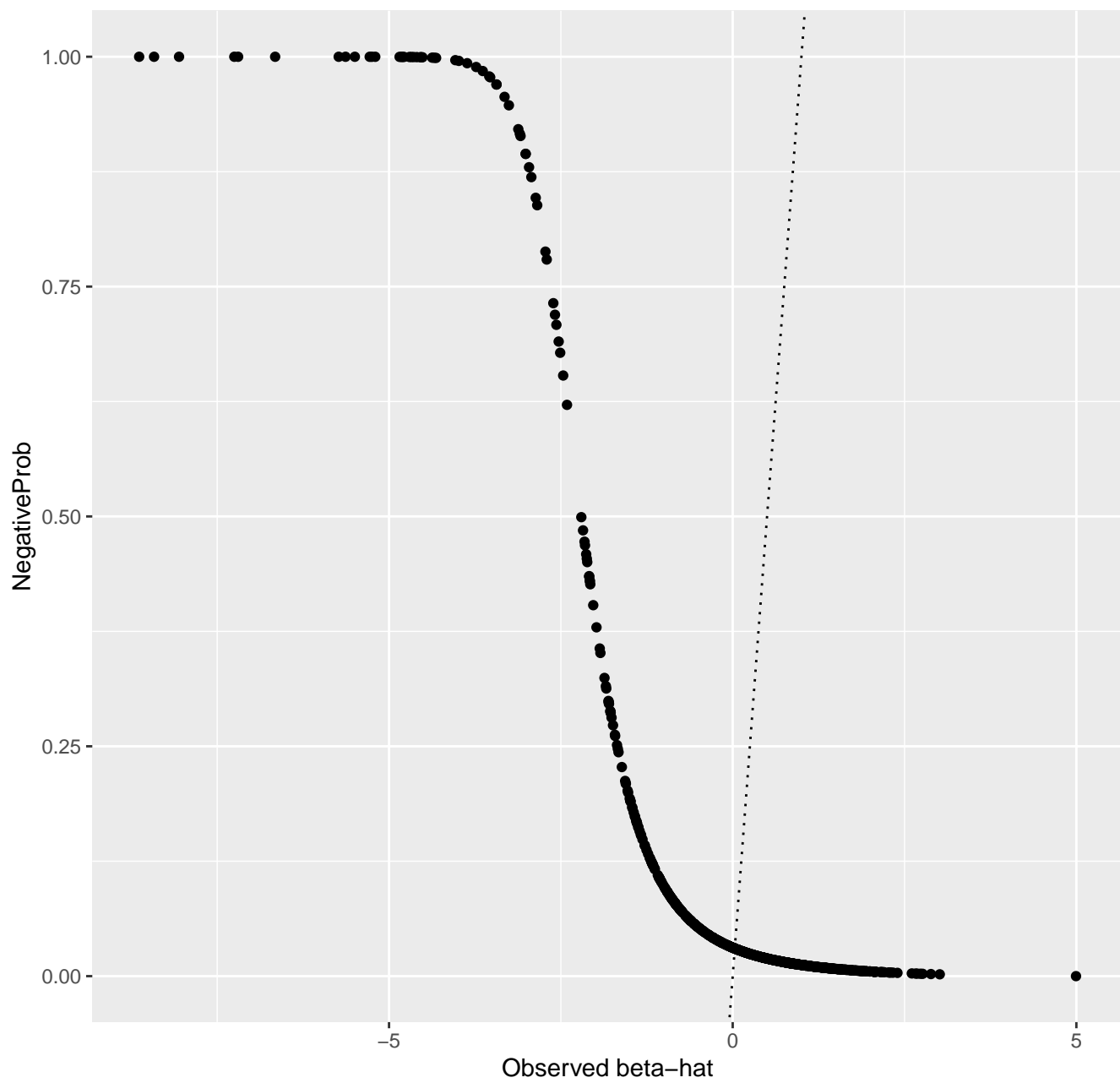
DERL2_(51009) negative



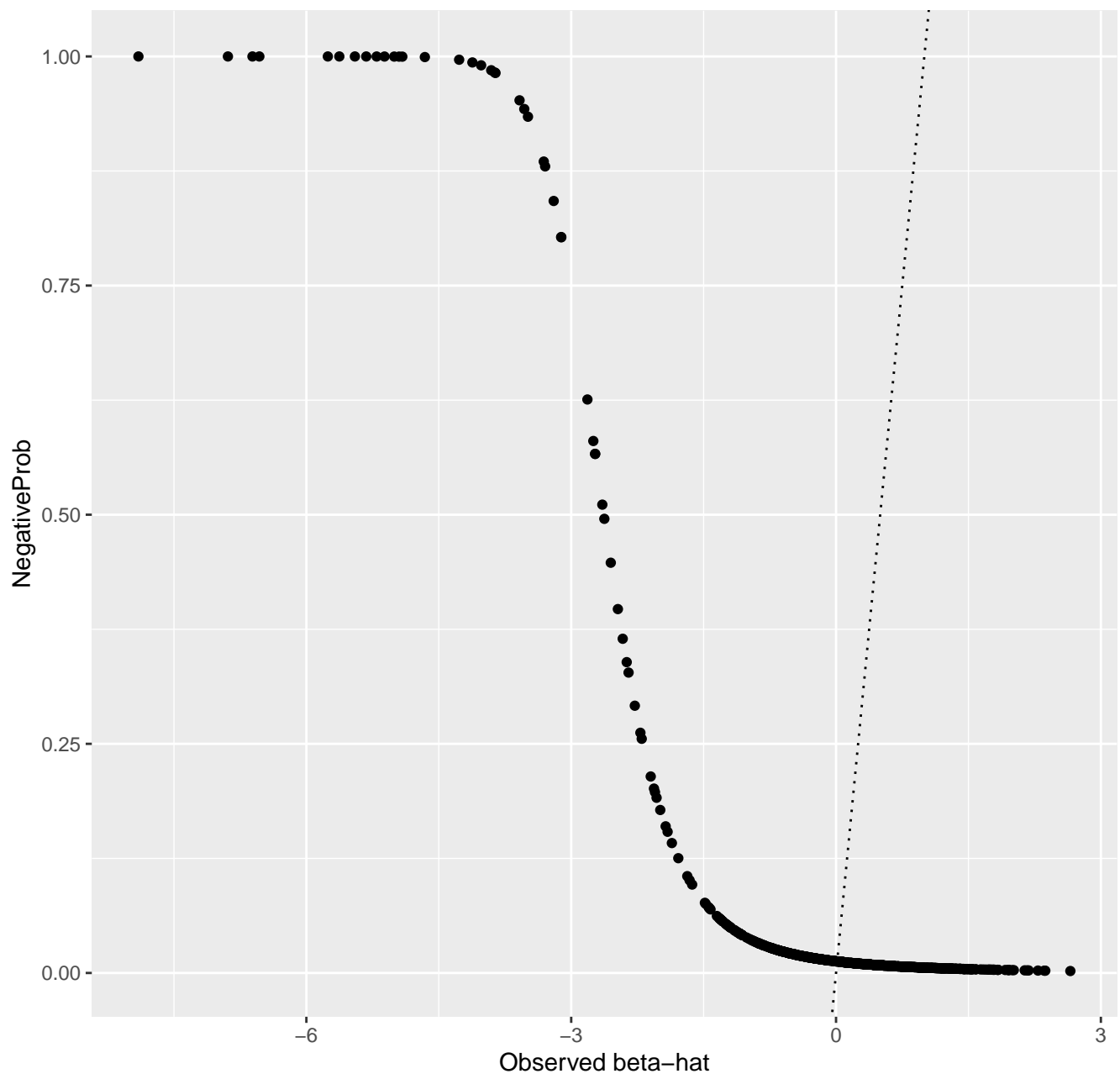
RAB18_(22931) negative



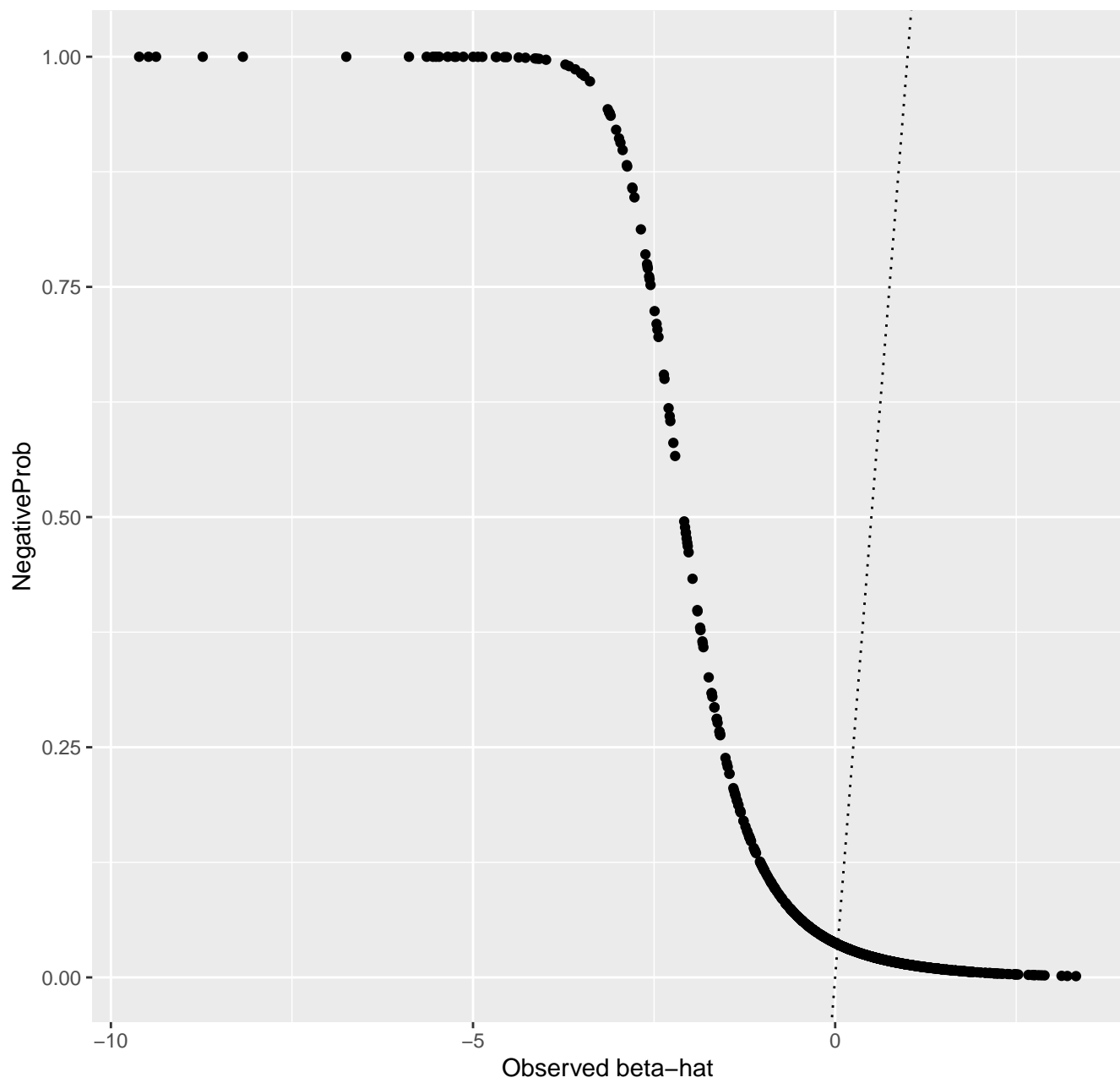
MTX2_(10651) negative



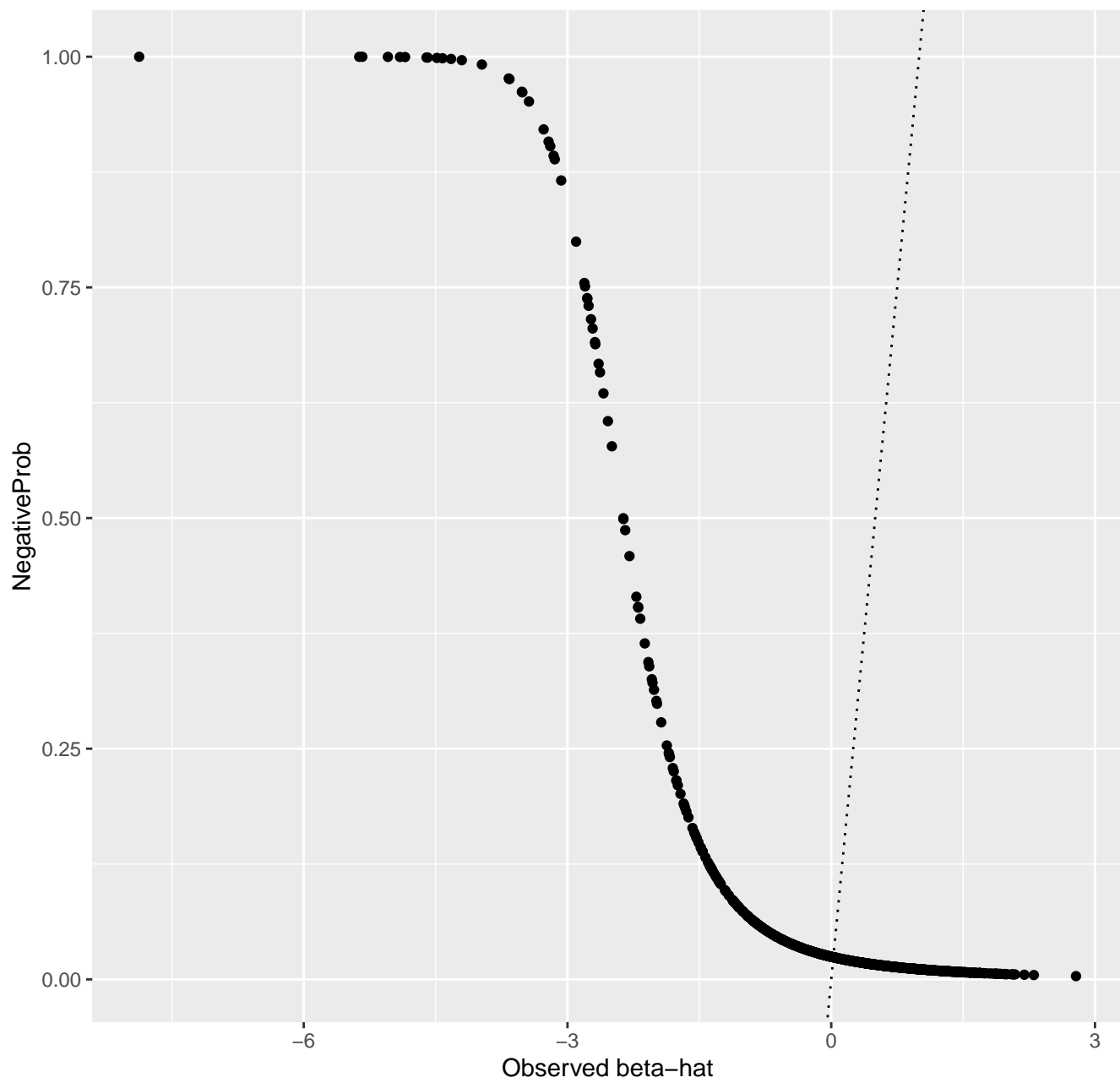
BCL2_(596) negative



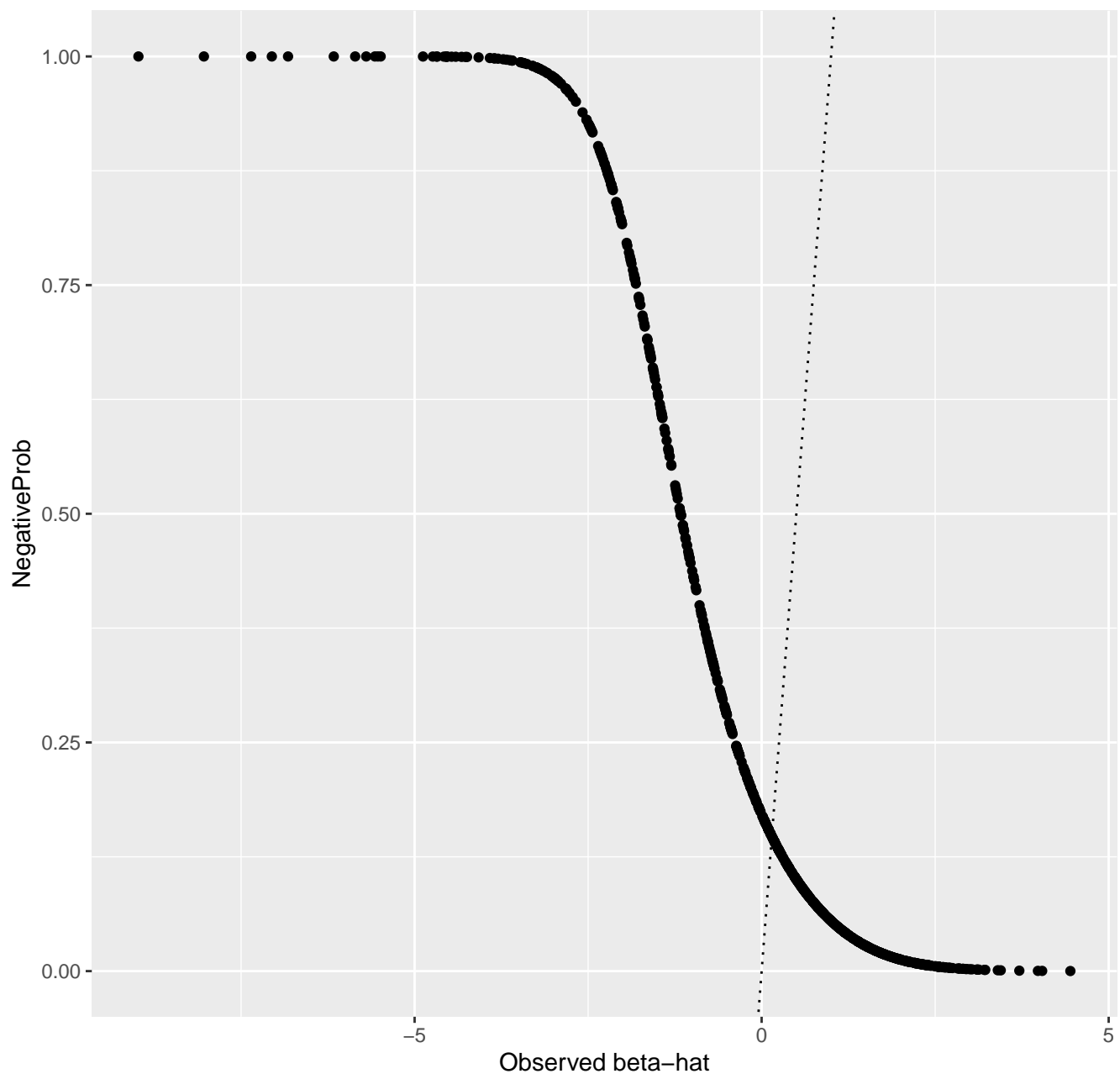
DOCK5_(80005) negative



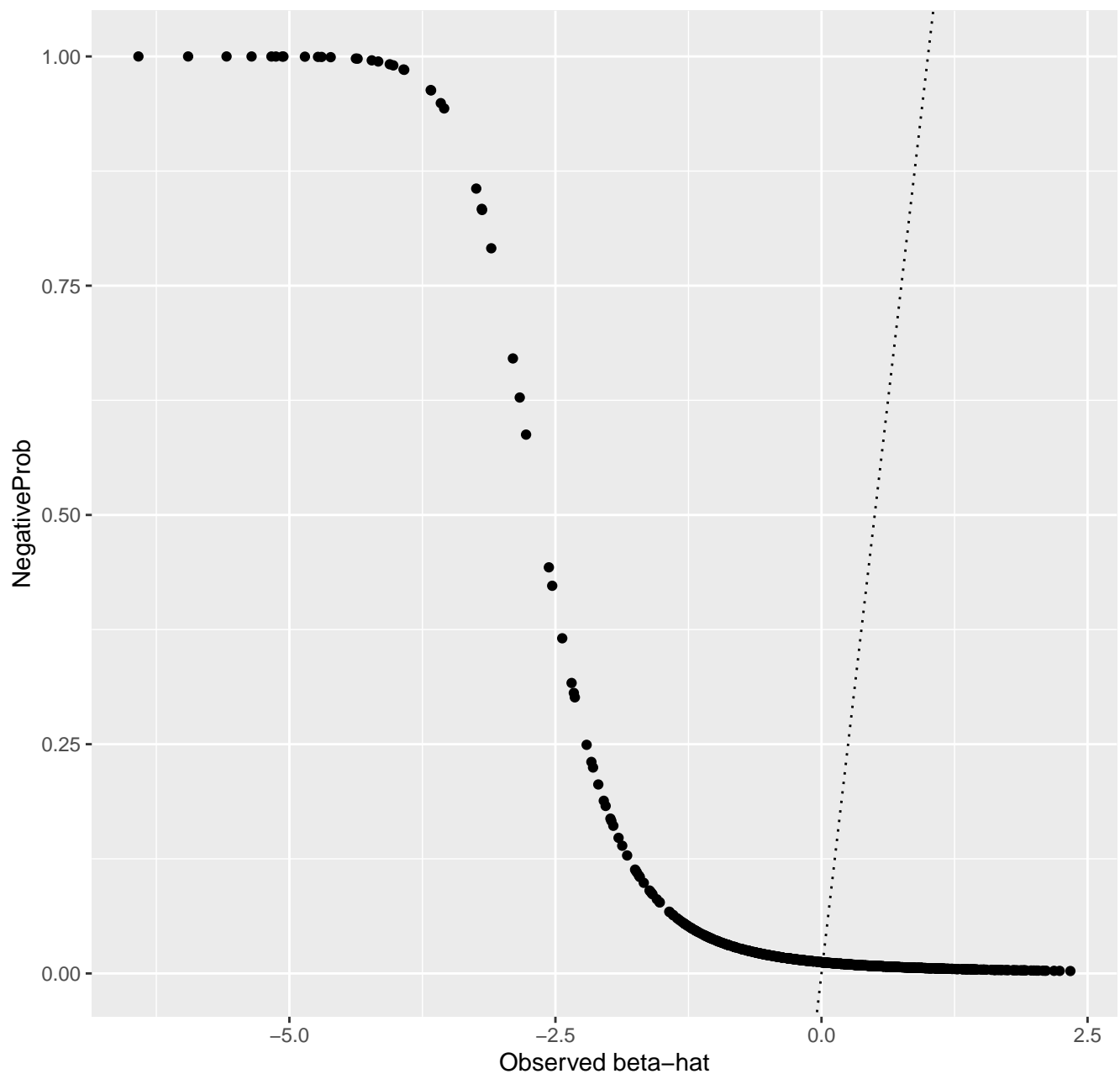
STEAP3_(55240) negative



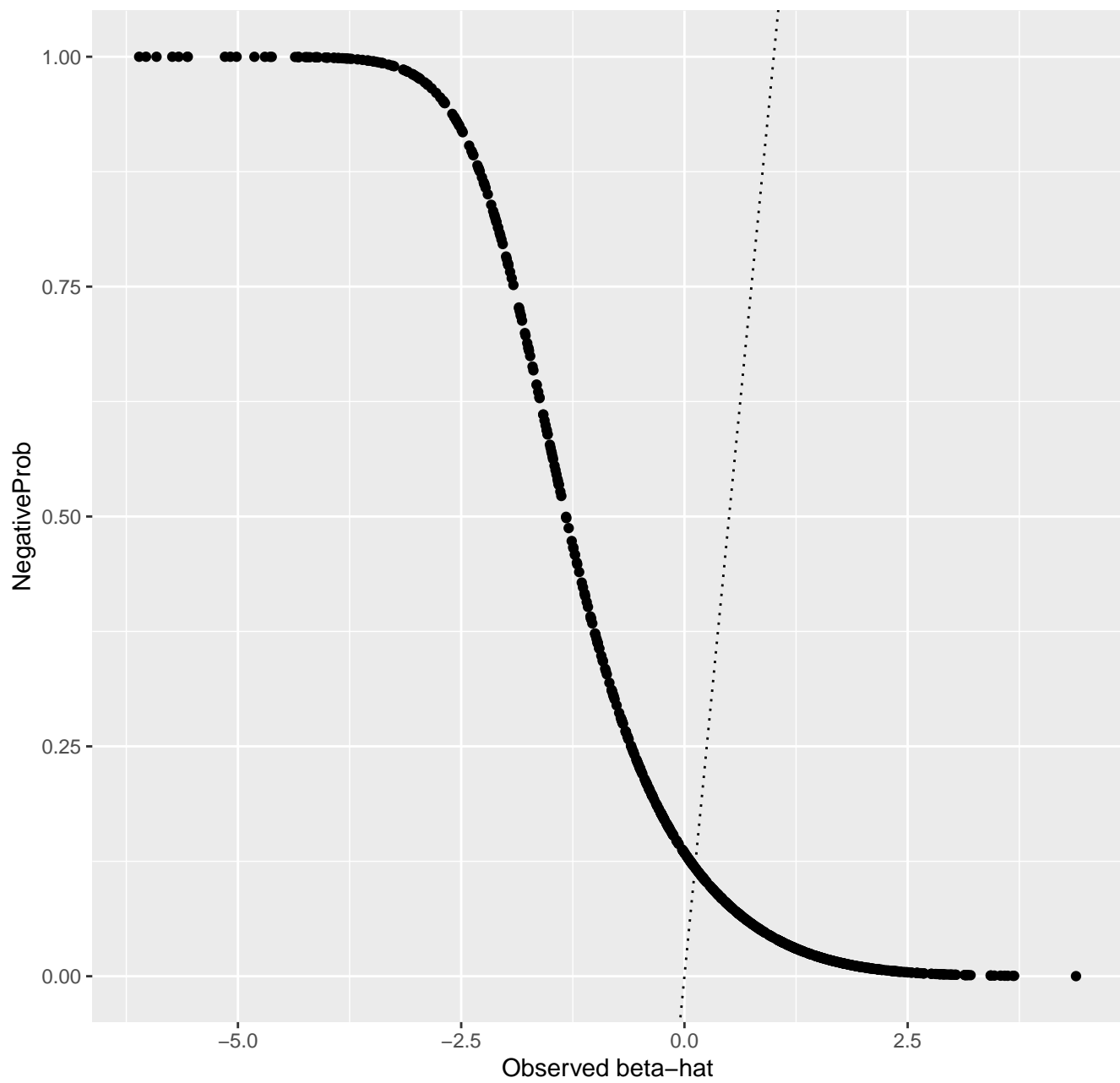
SLC35B2_(347734) negative



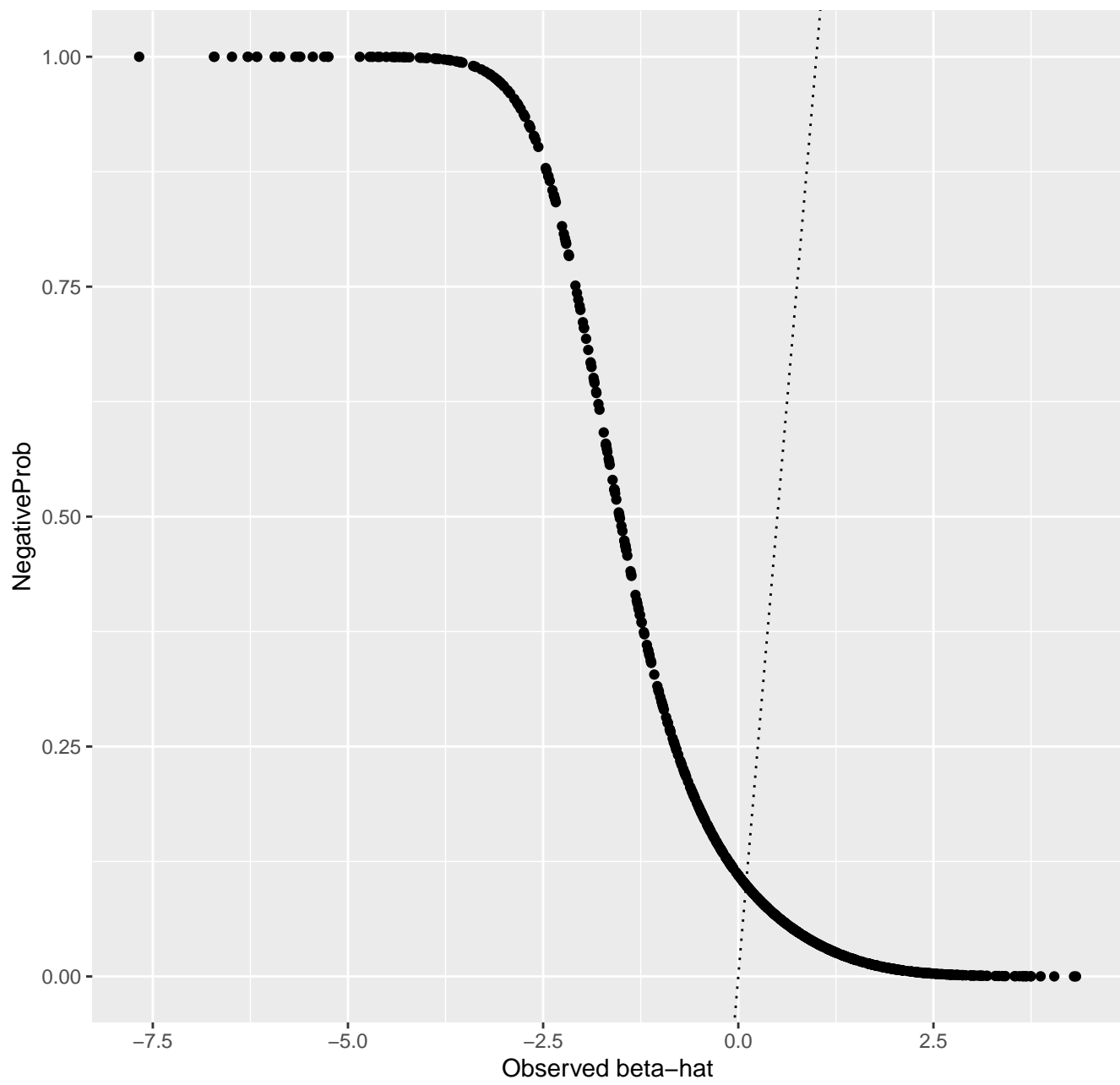
PPARG_(5468) negative



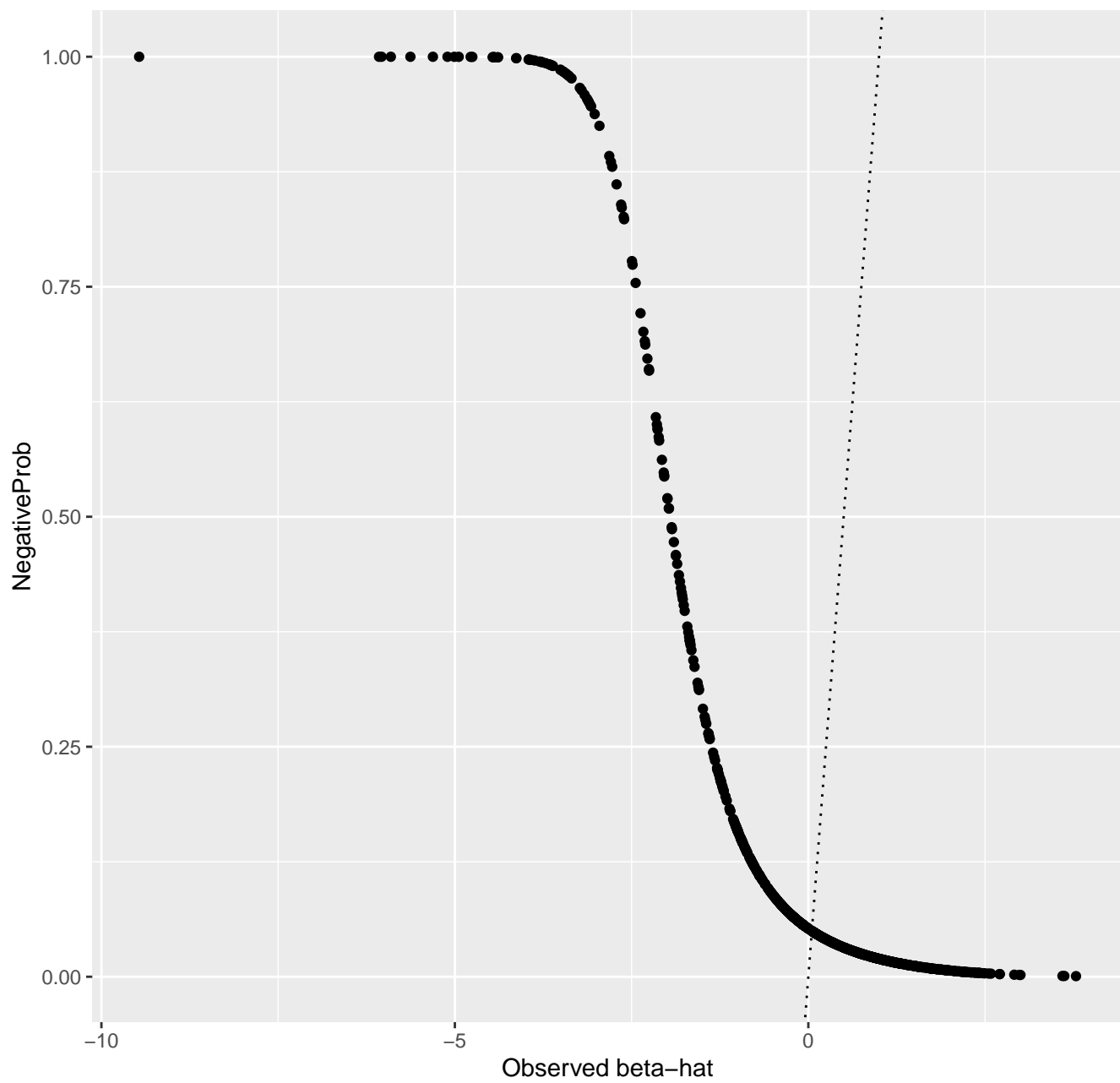
SLC7A5_(8140) negative



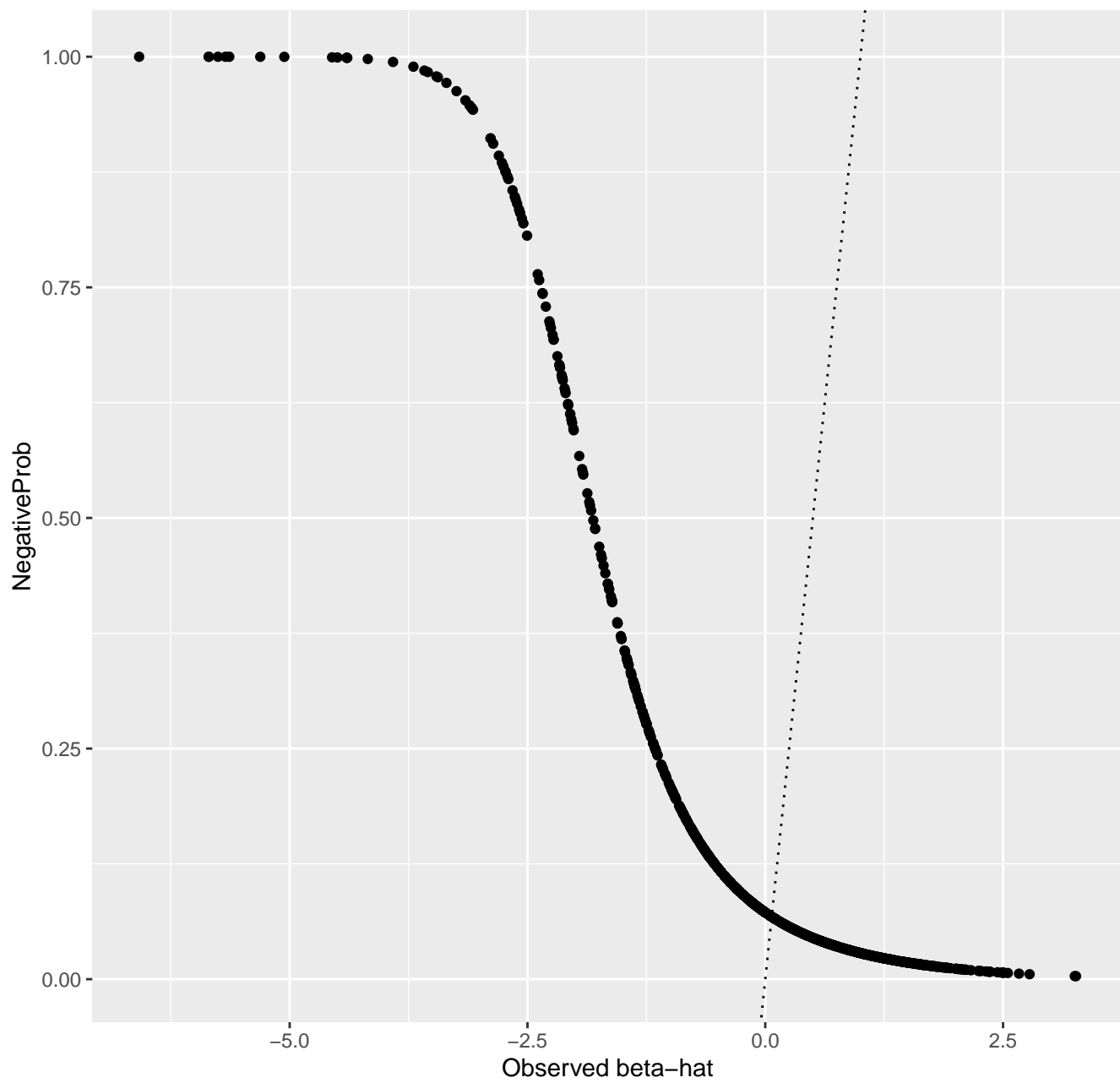
FDPS_(2224) negative



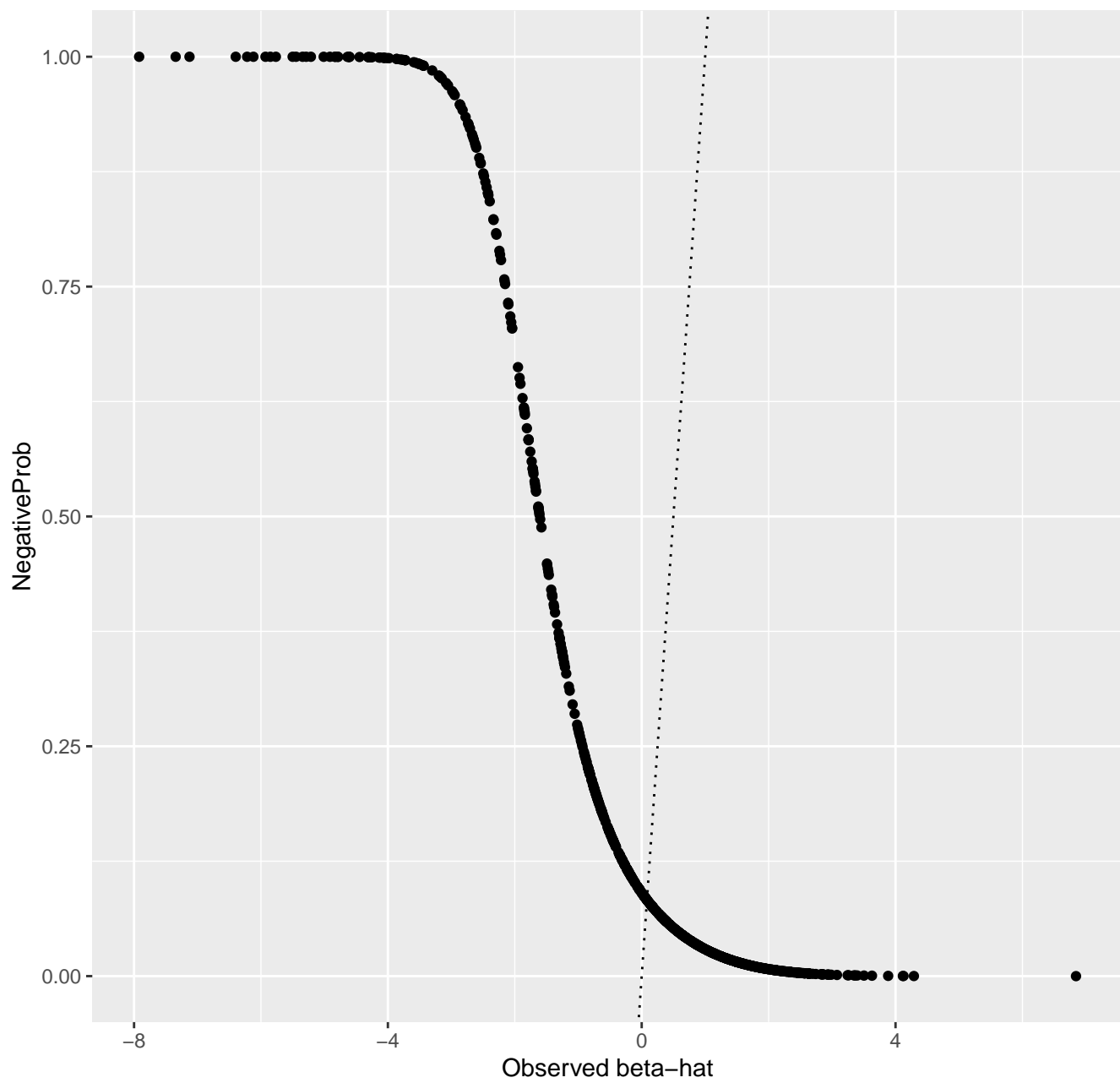
TAPT1_(202018) negative



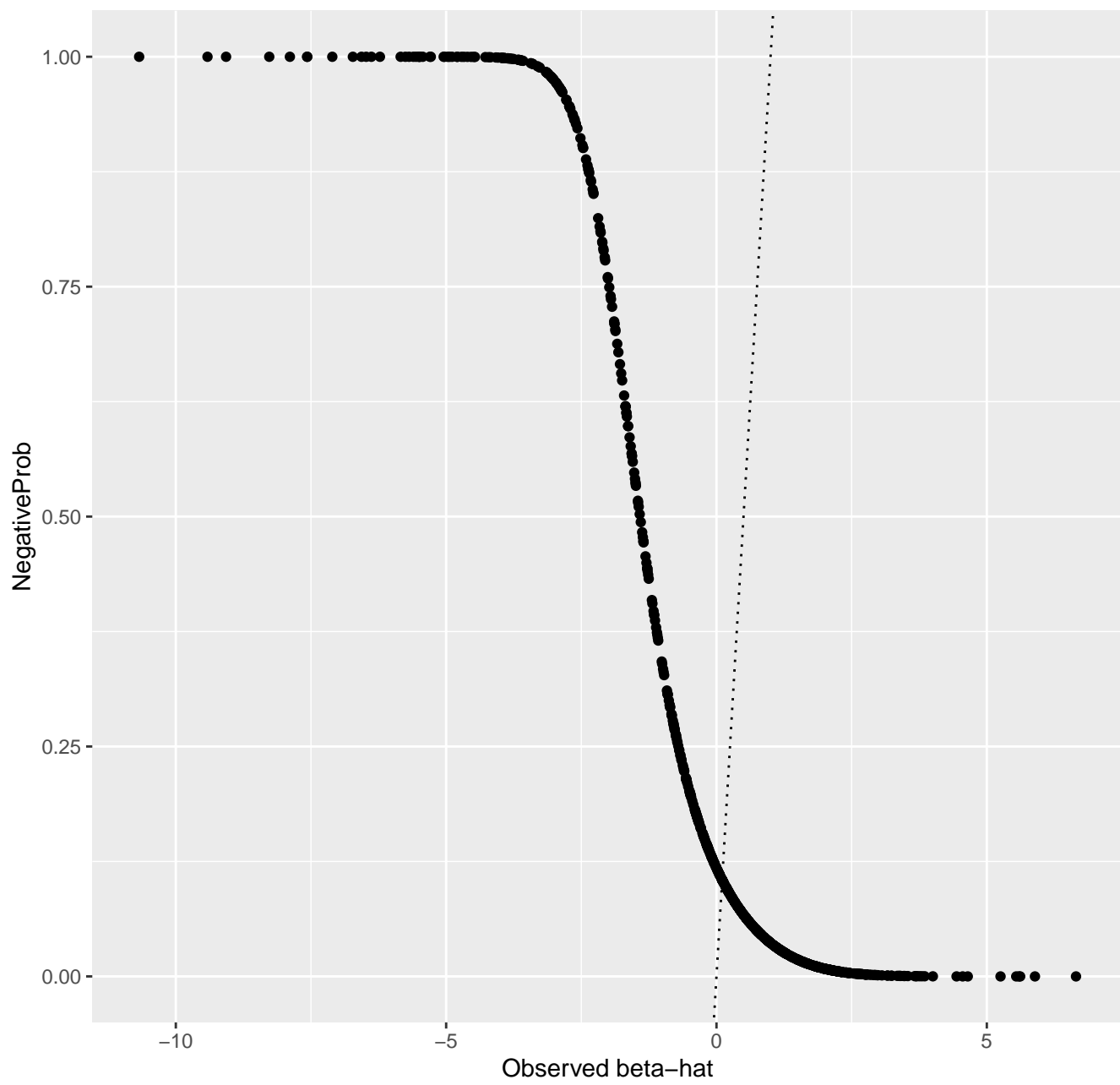
GTSE1_(51512) negative



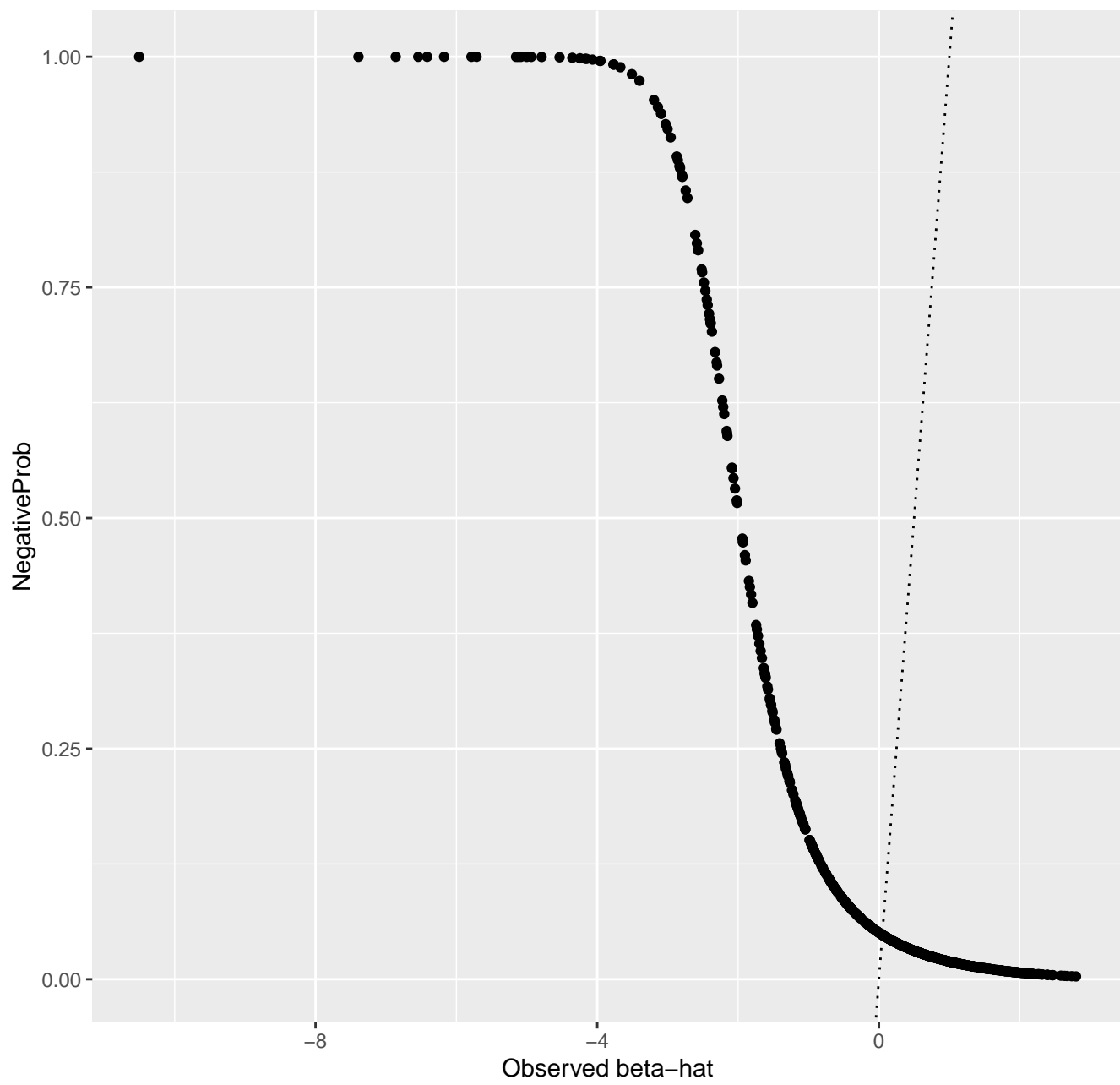
SLC25A28_(81894) negative



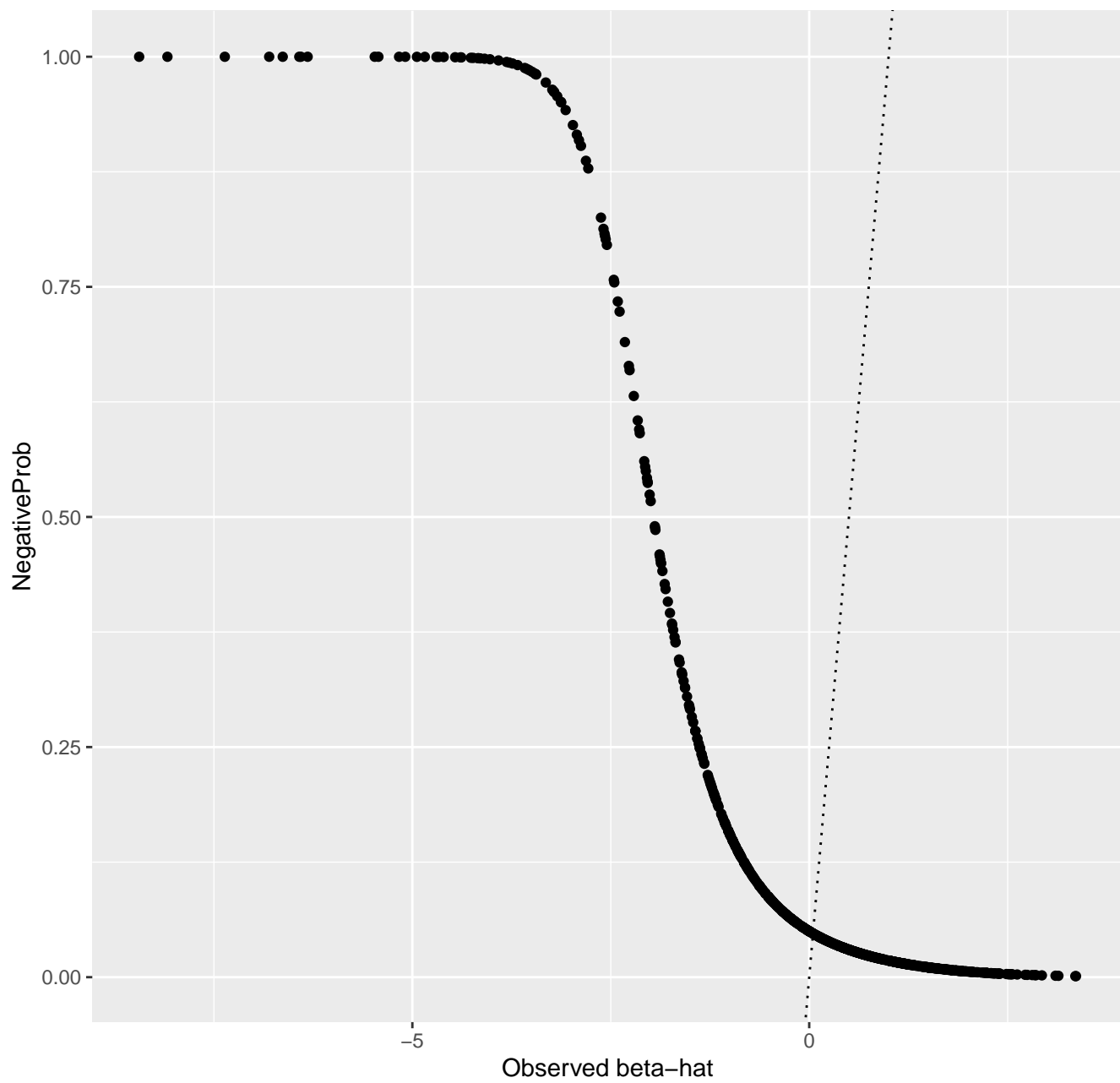
MIB1_(57534) negative



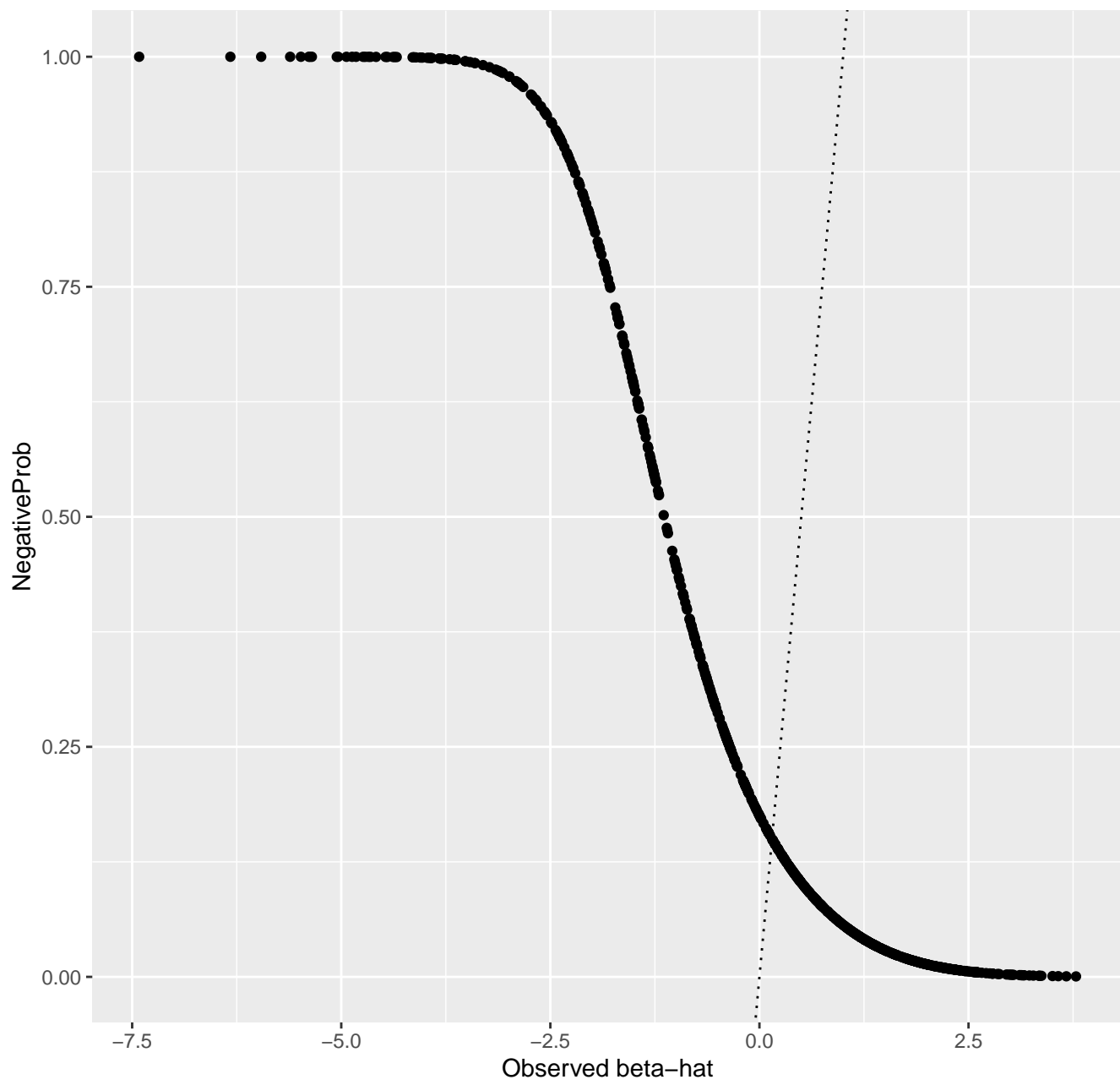
SCYL1_(57410) negative



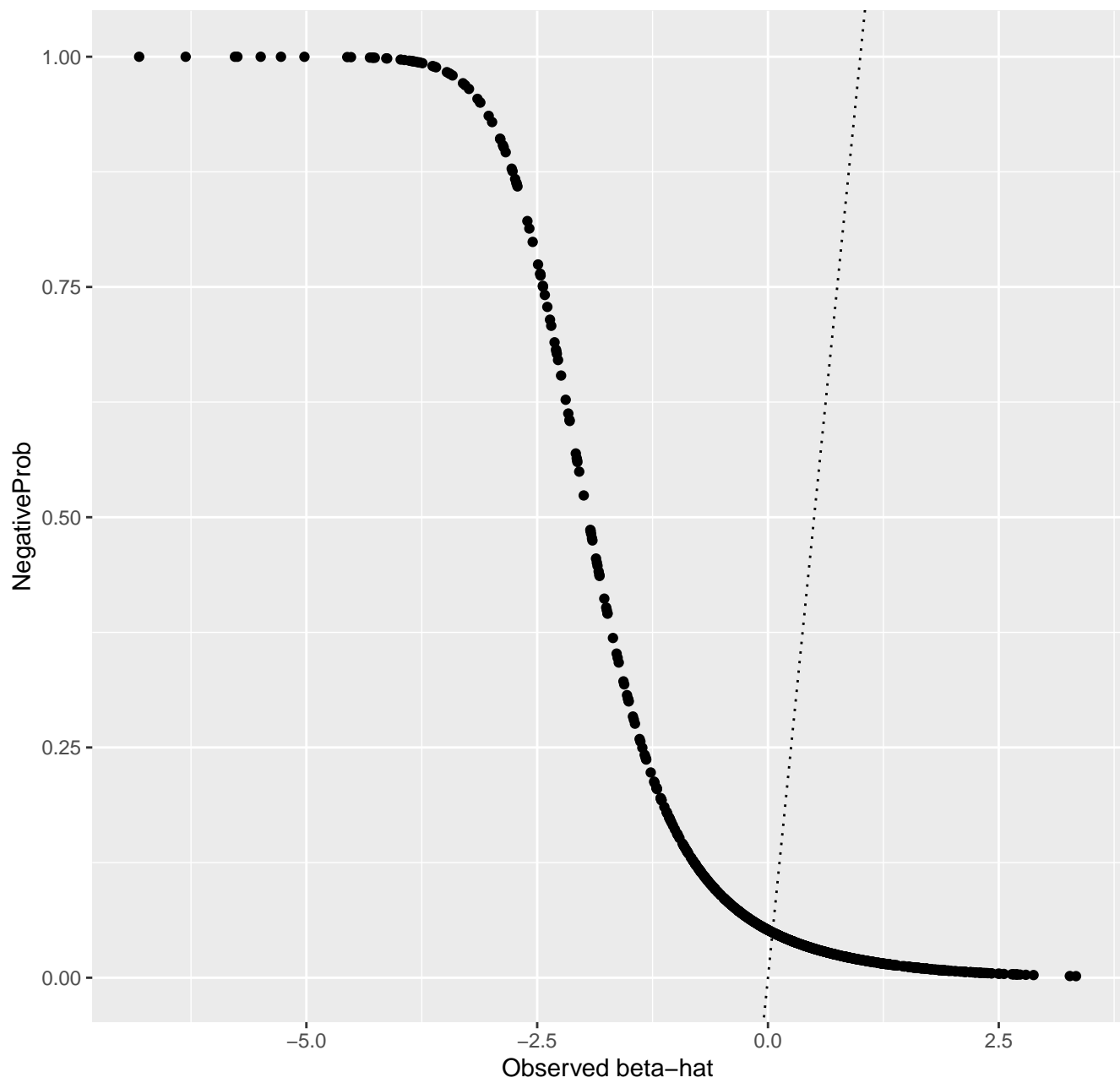
SLC20A1_(6574) negative



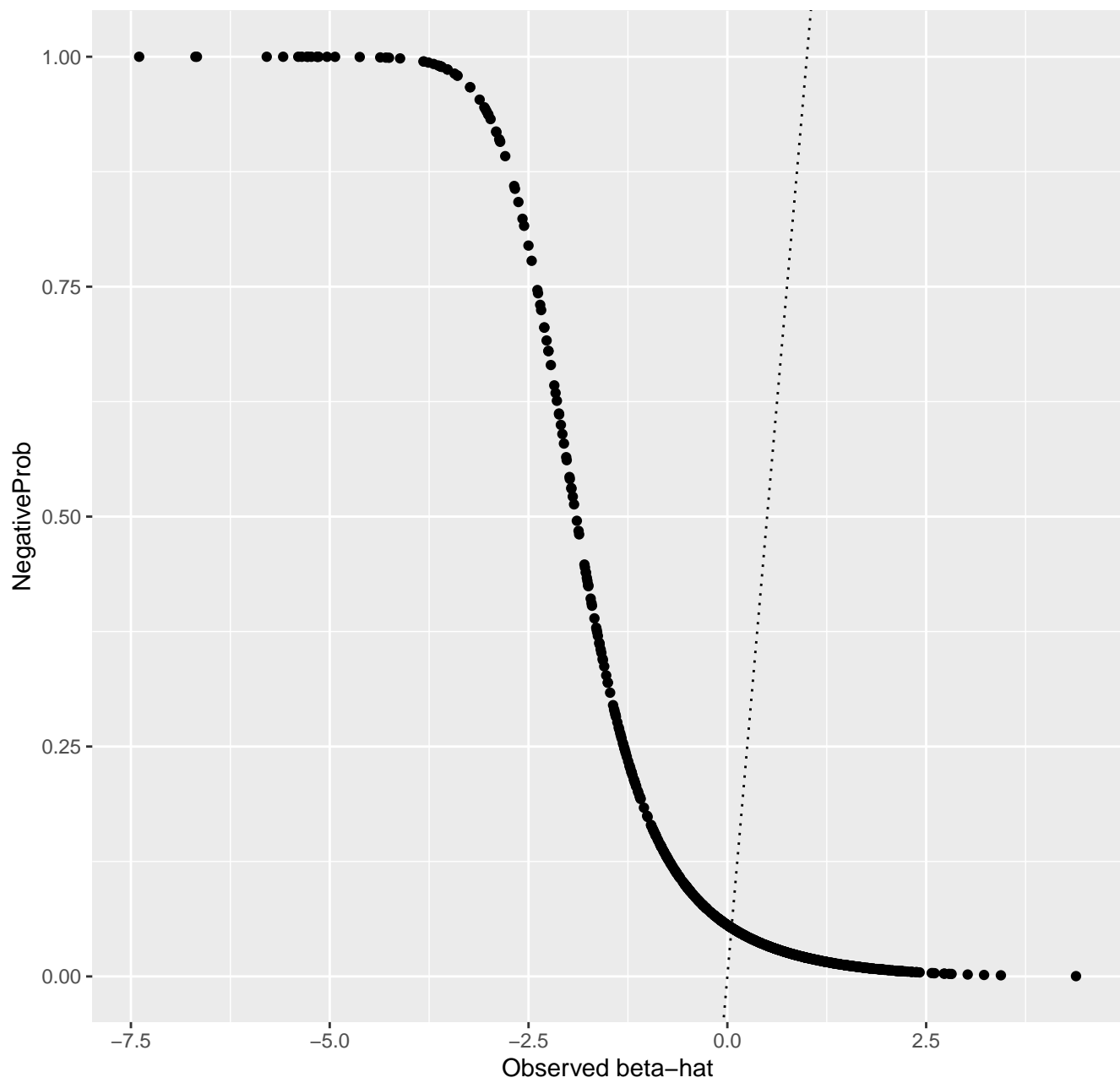
TRIM37_(4591) negative



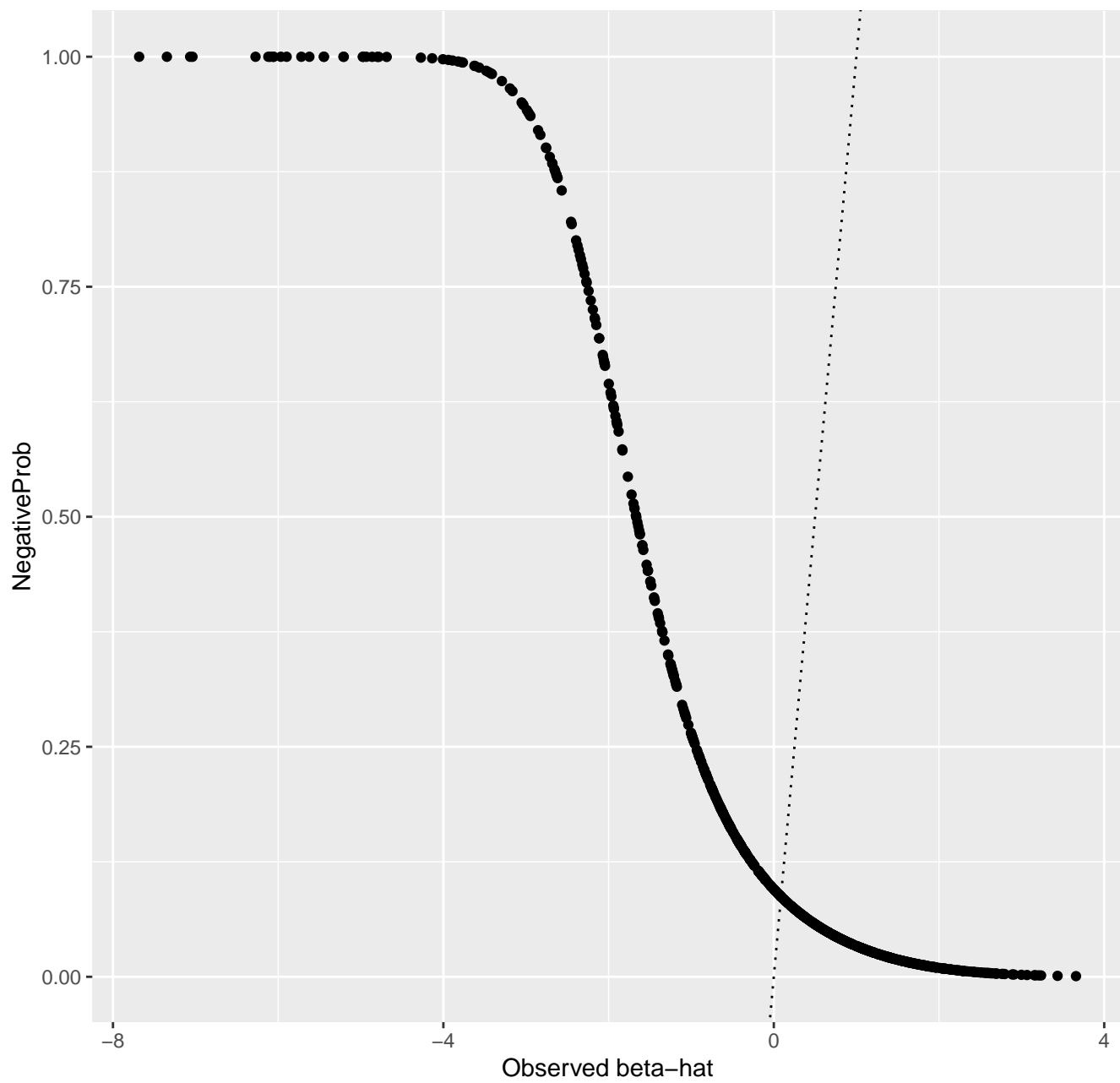
NGDN_(25983) negative



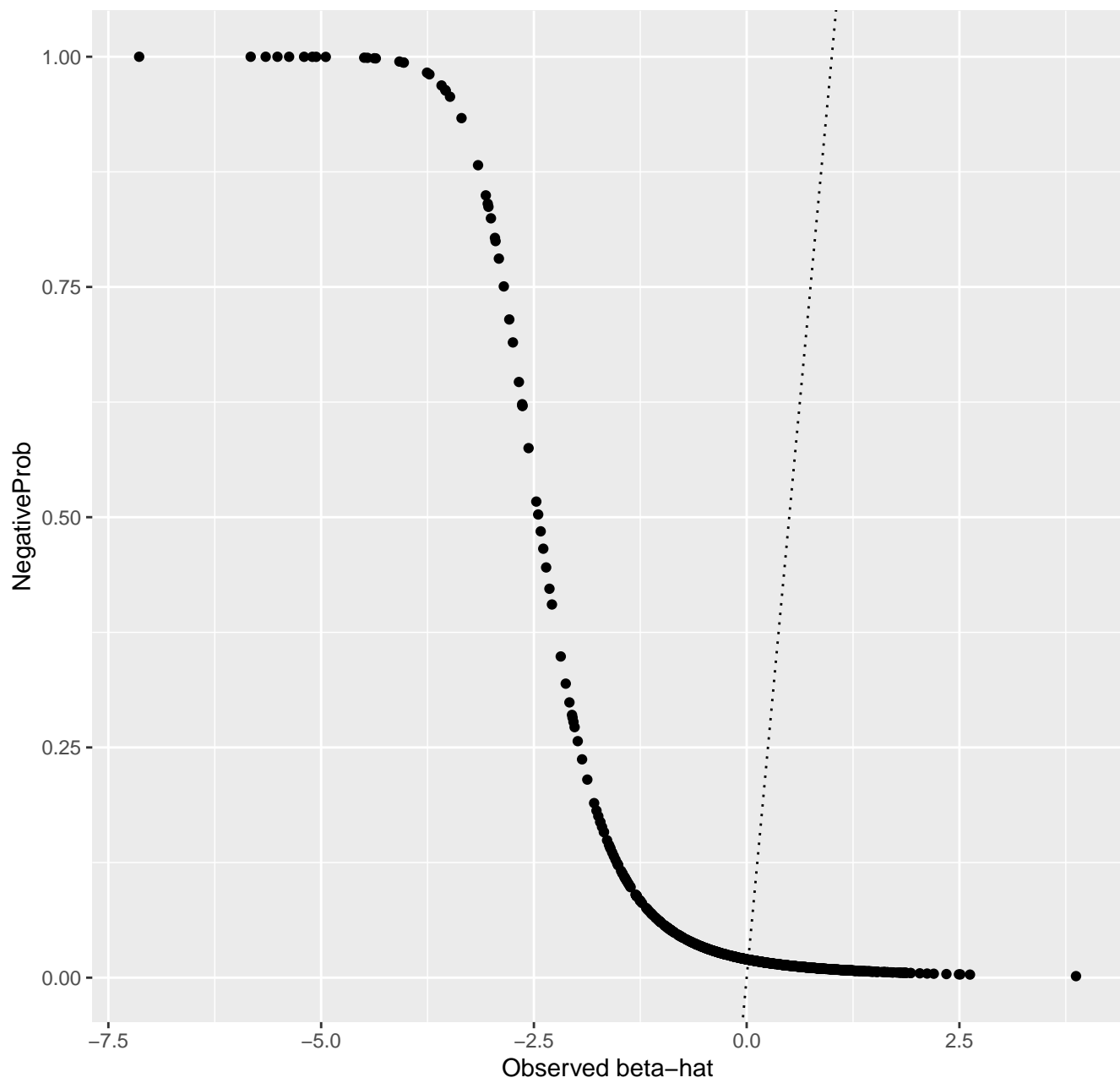
ITPK1_(3705) negative



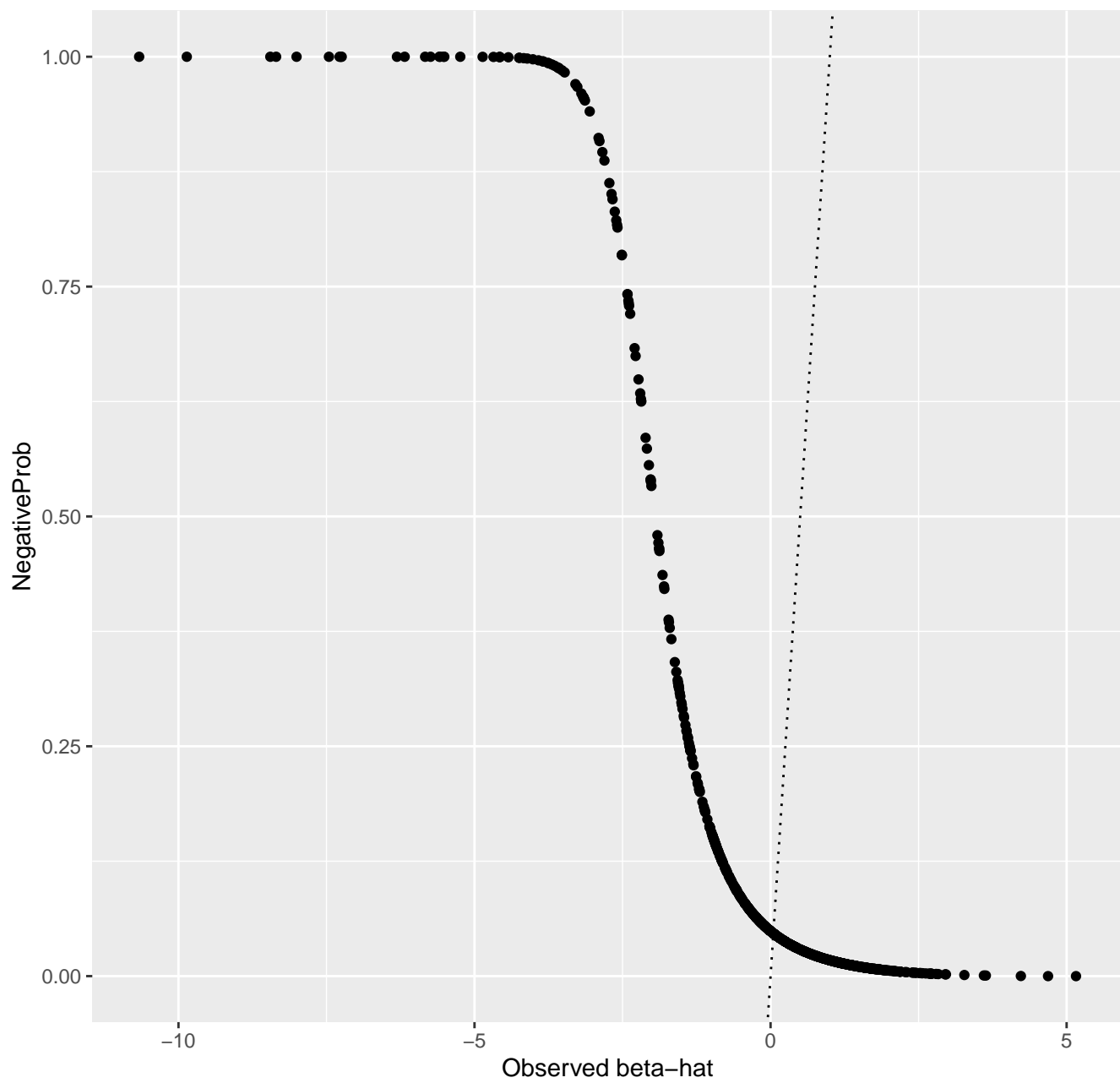
USP14_(9097) negative



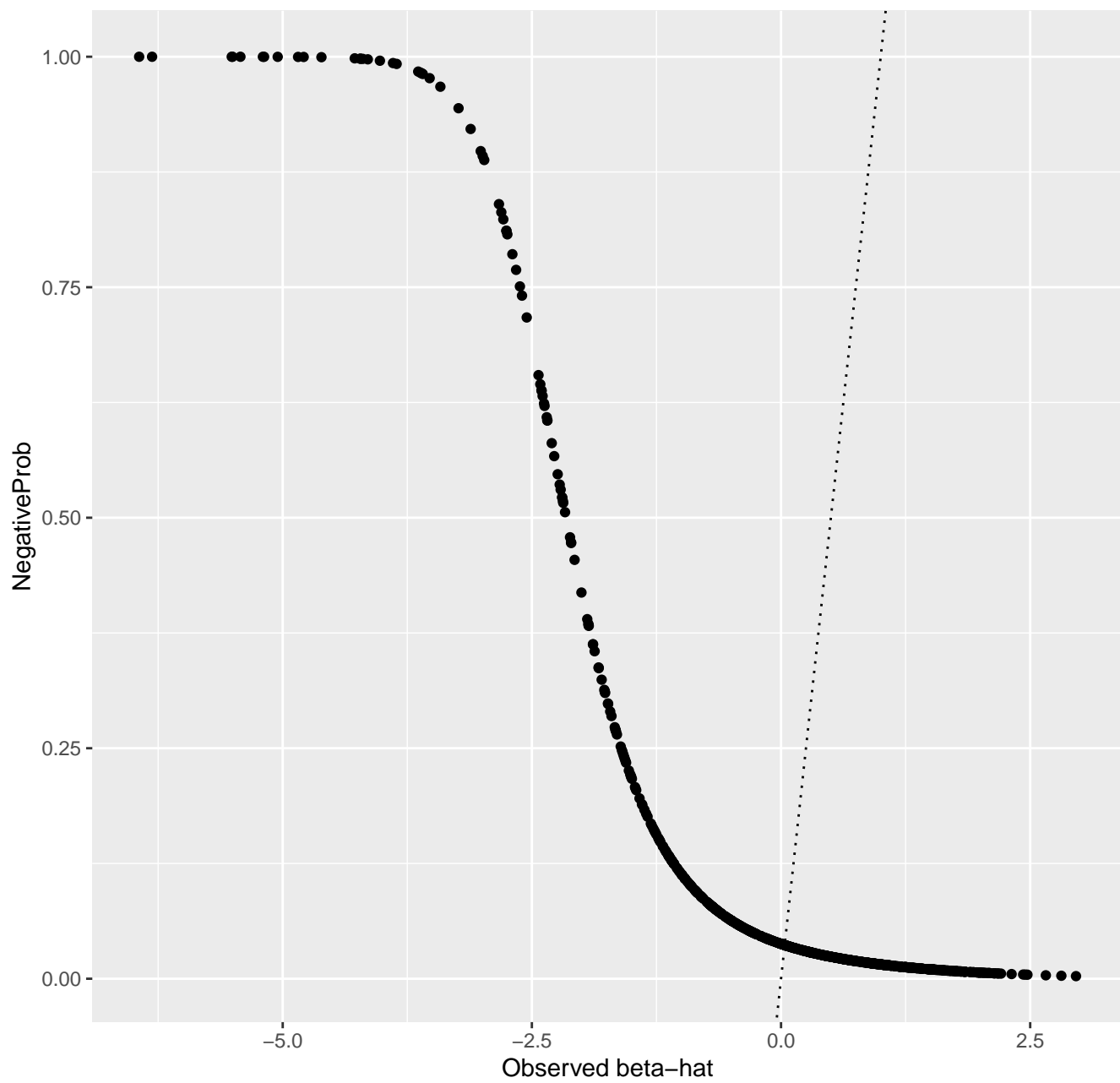
SLF1_(84250) negative



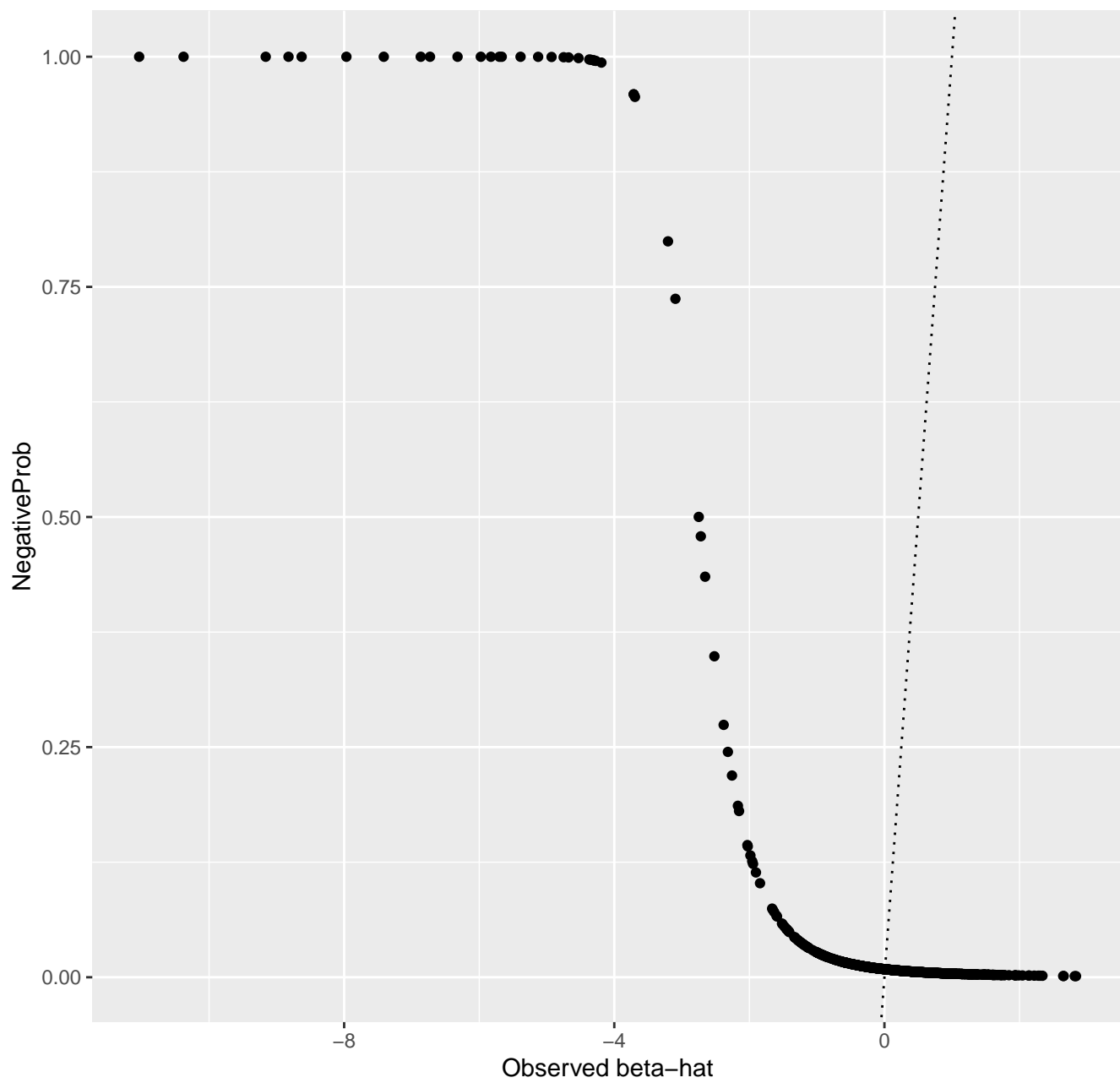
XPR1_(9213) negative



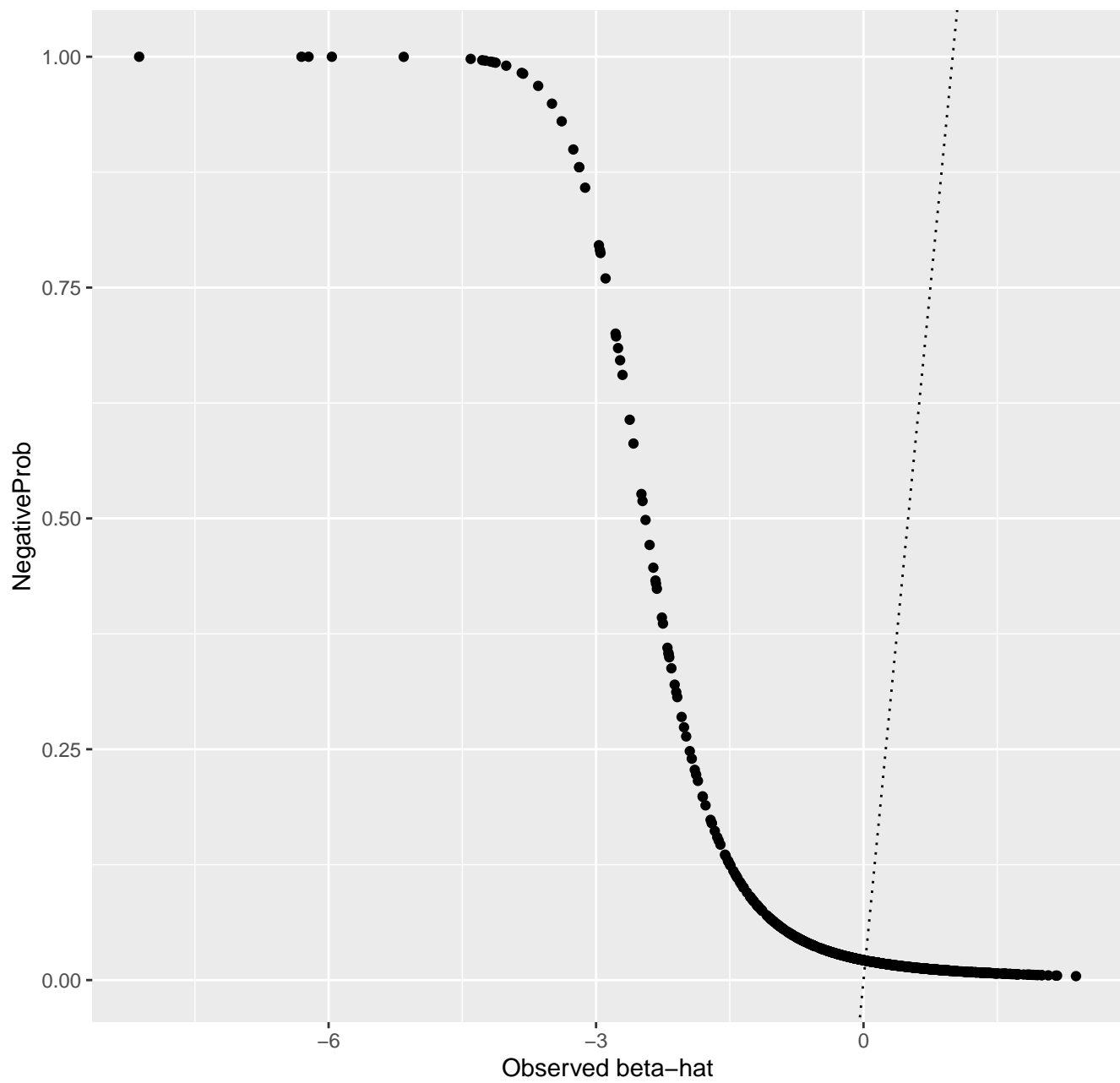
C12orf49_(79794) negative



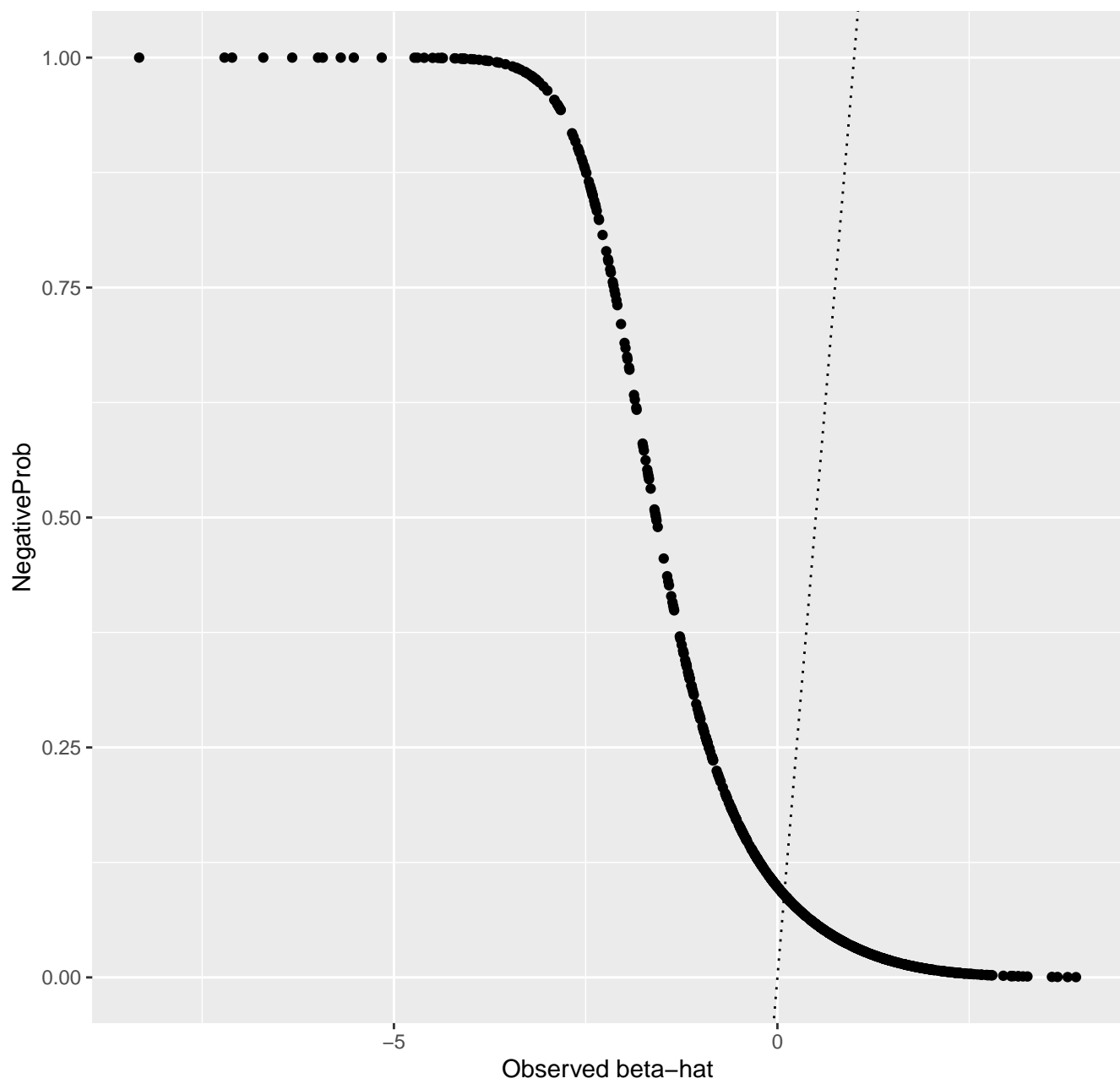
SOX2_(6657) negative



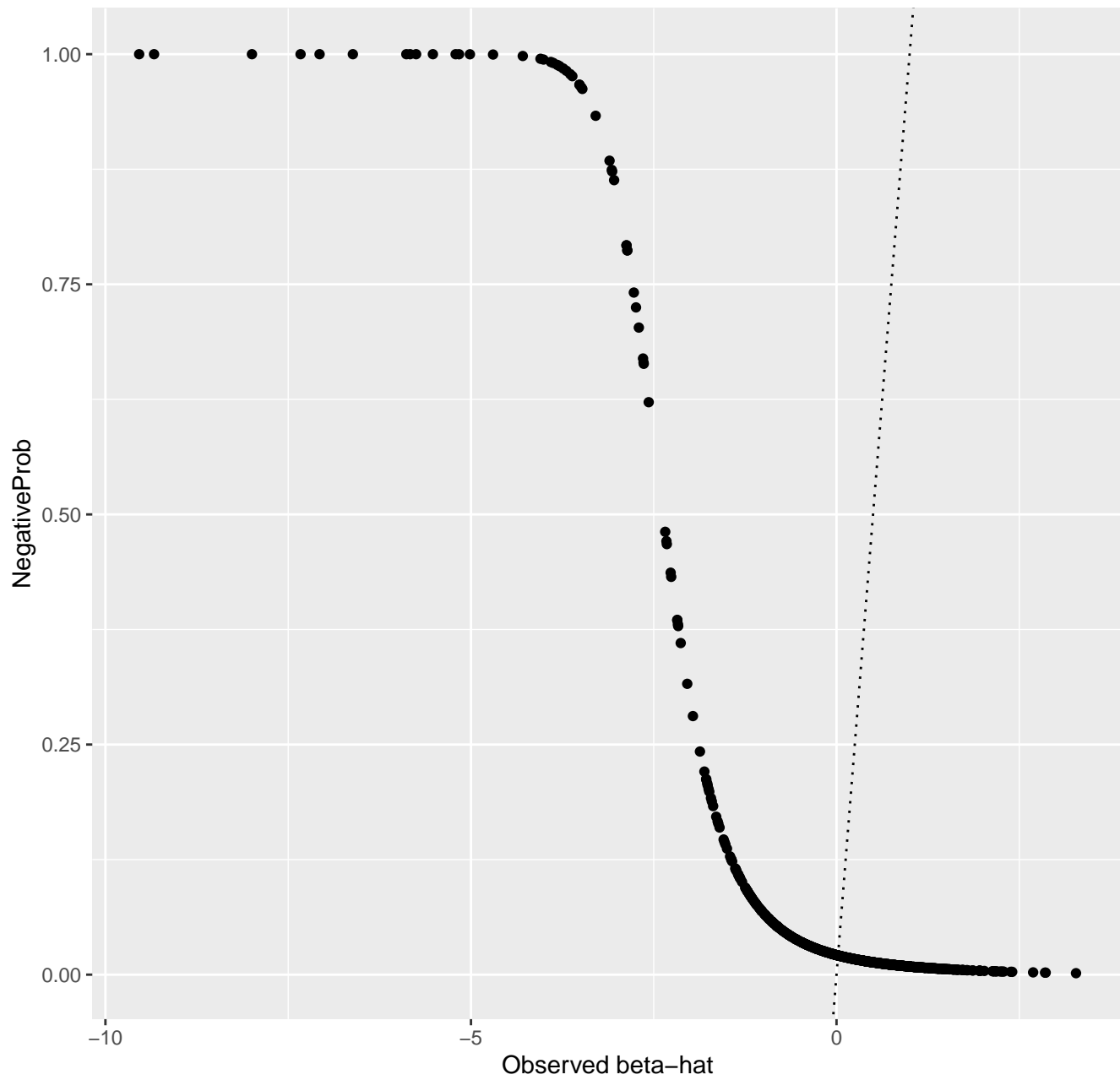
TMEM41A_(90407) negative



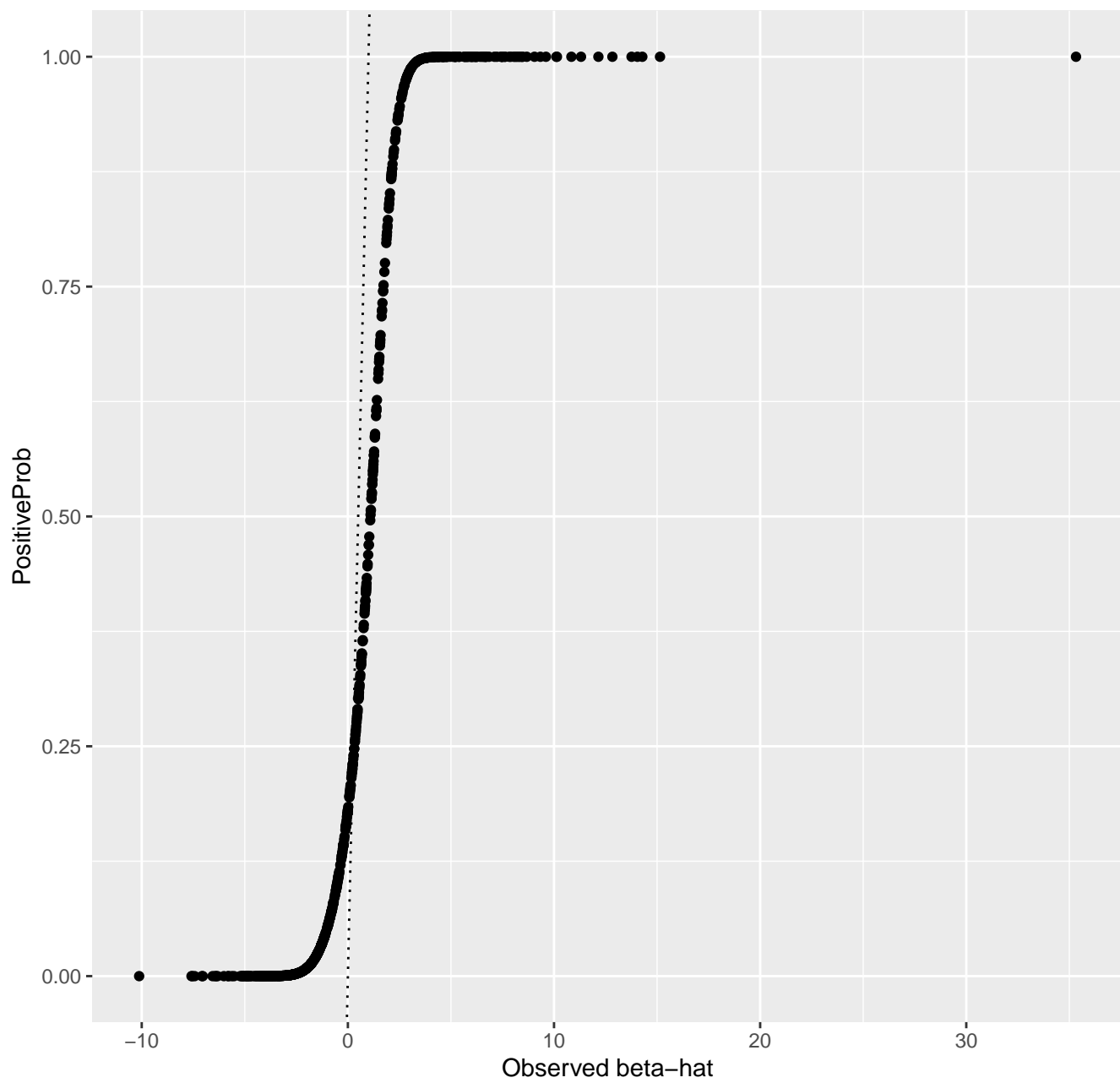
JTB_(10899) negative



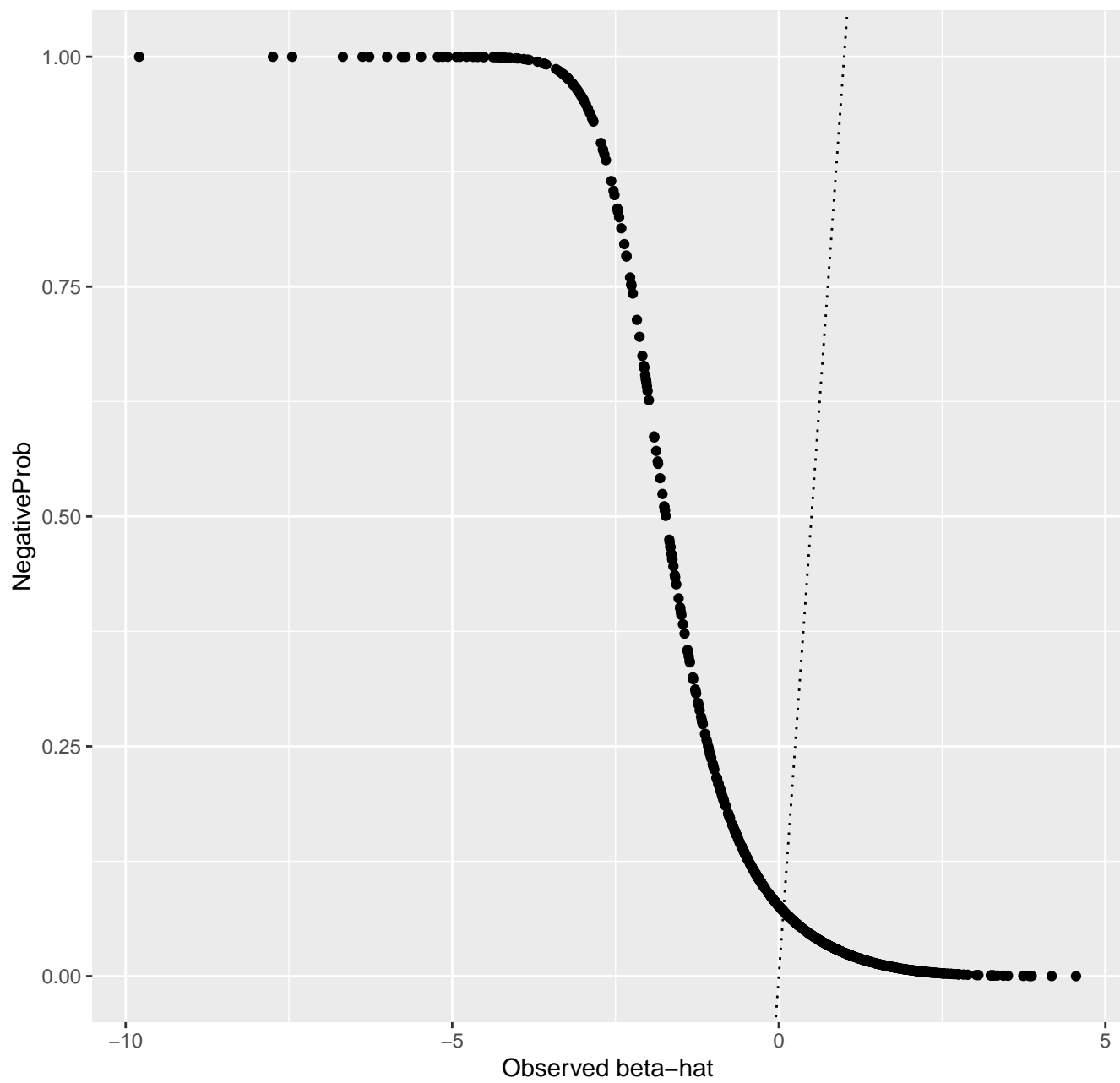
IRS1_(3667) negative



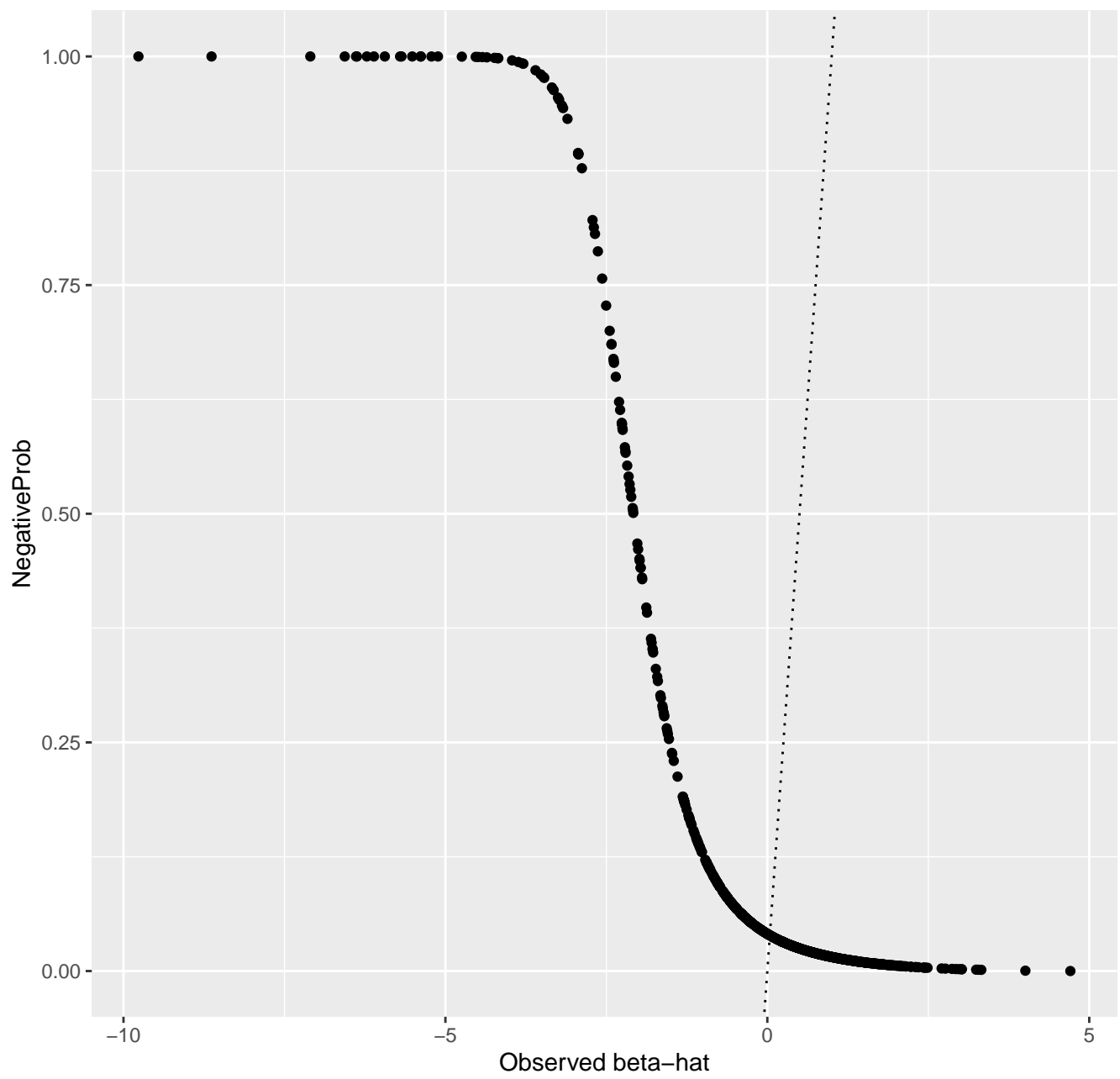
PTEN_(5728) positive



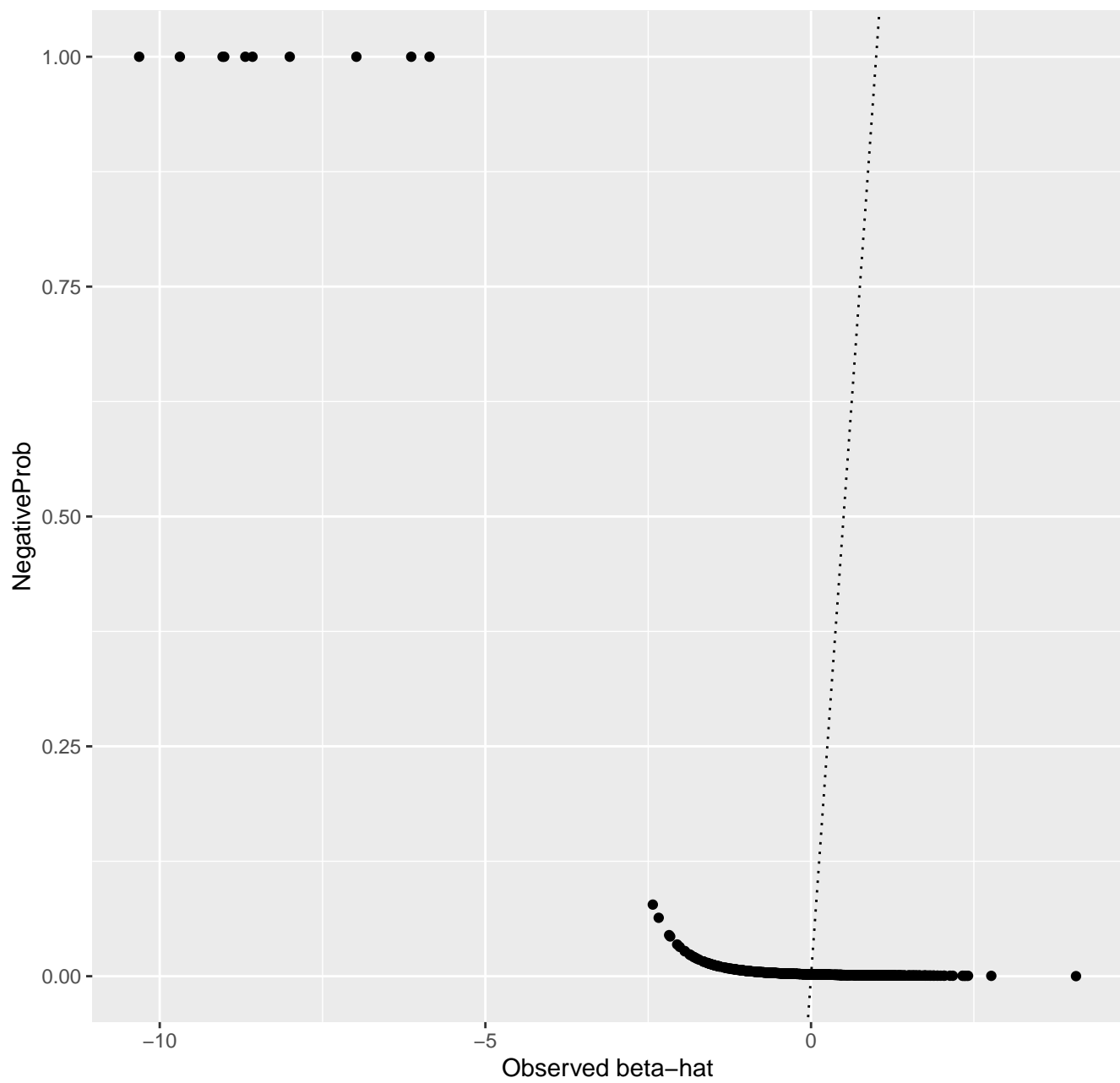
ATP1B3_(483) negative



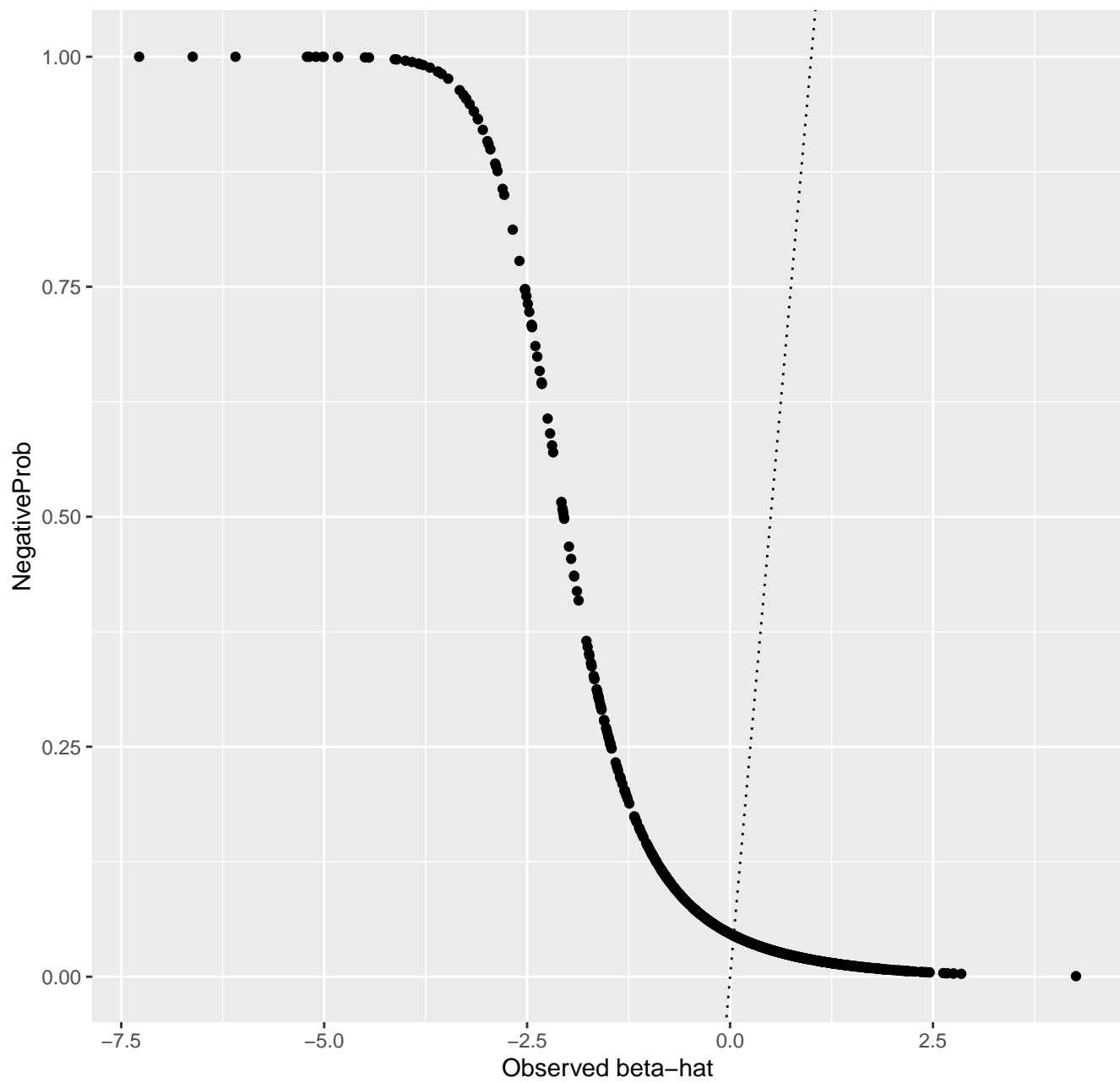
ARHGAP29_(9411) negative



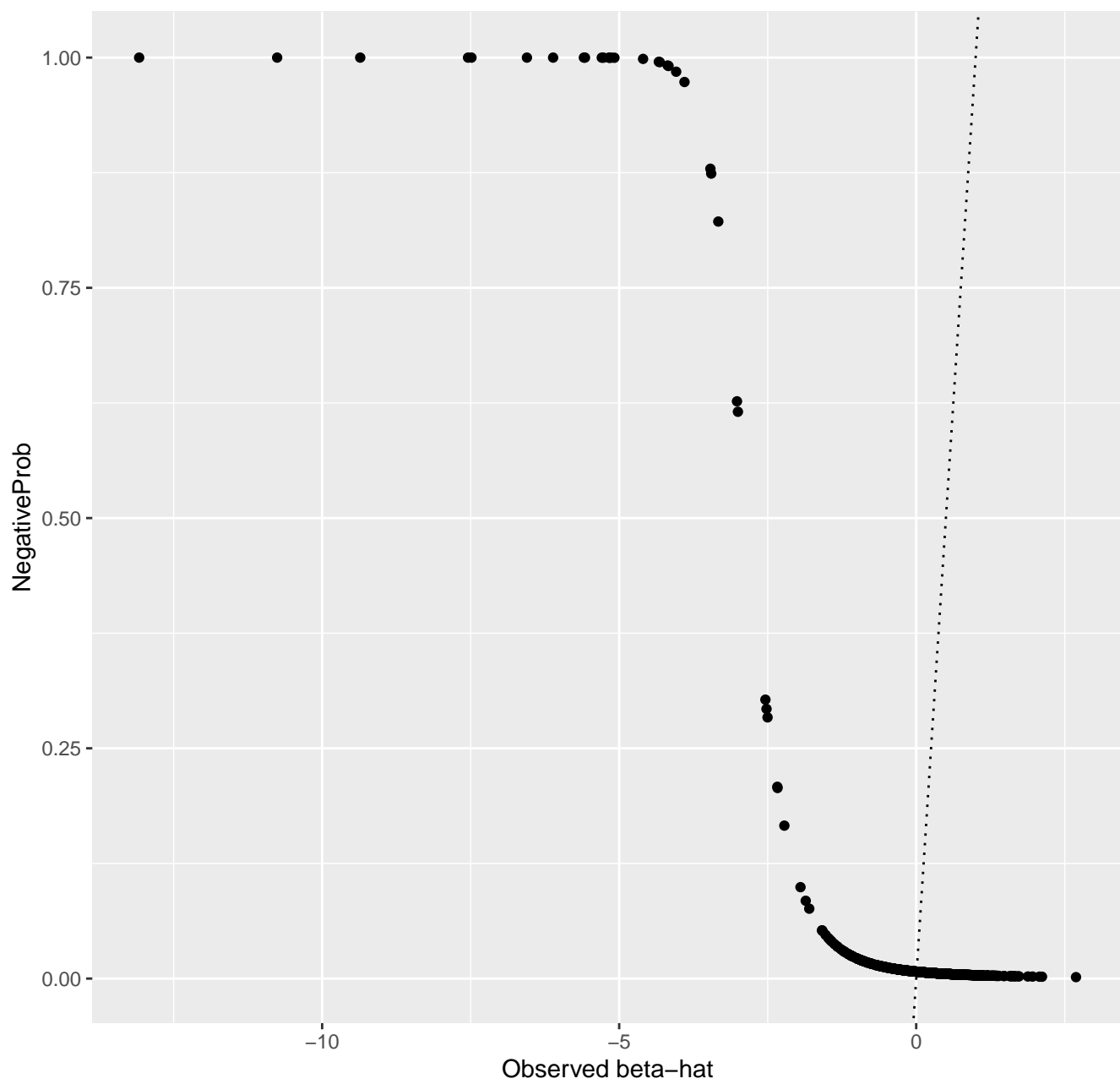
MYOD1_(4654) negative



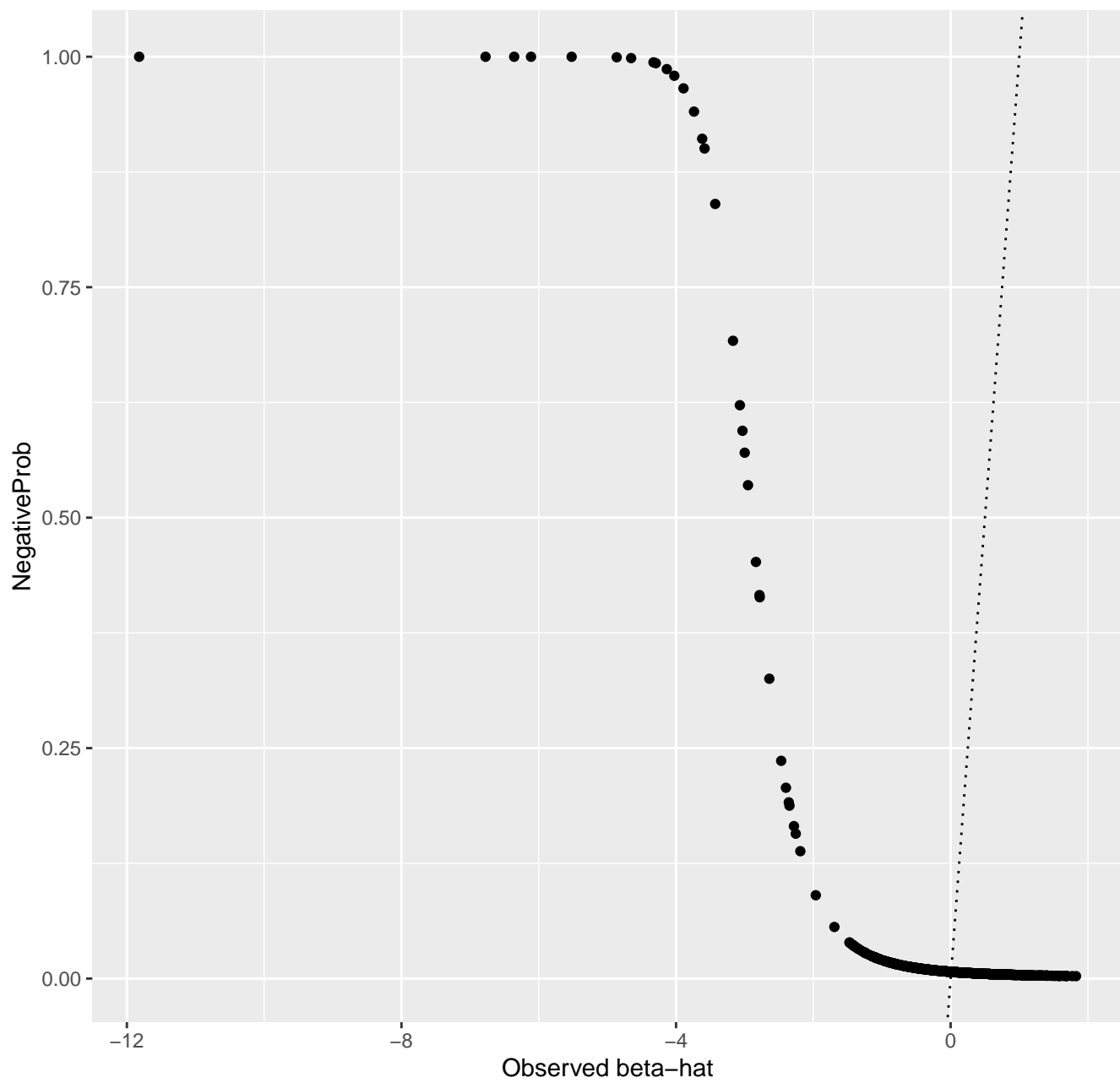
C21orf2_(755) negative



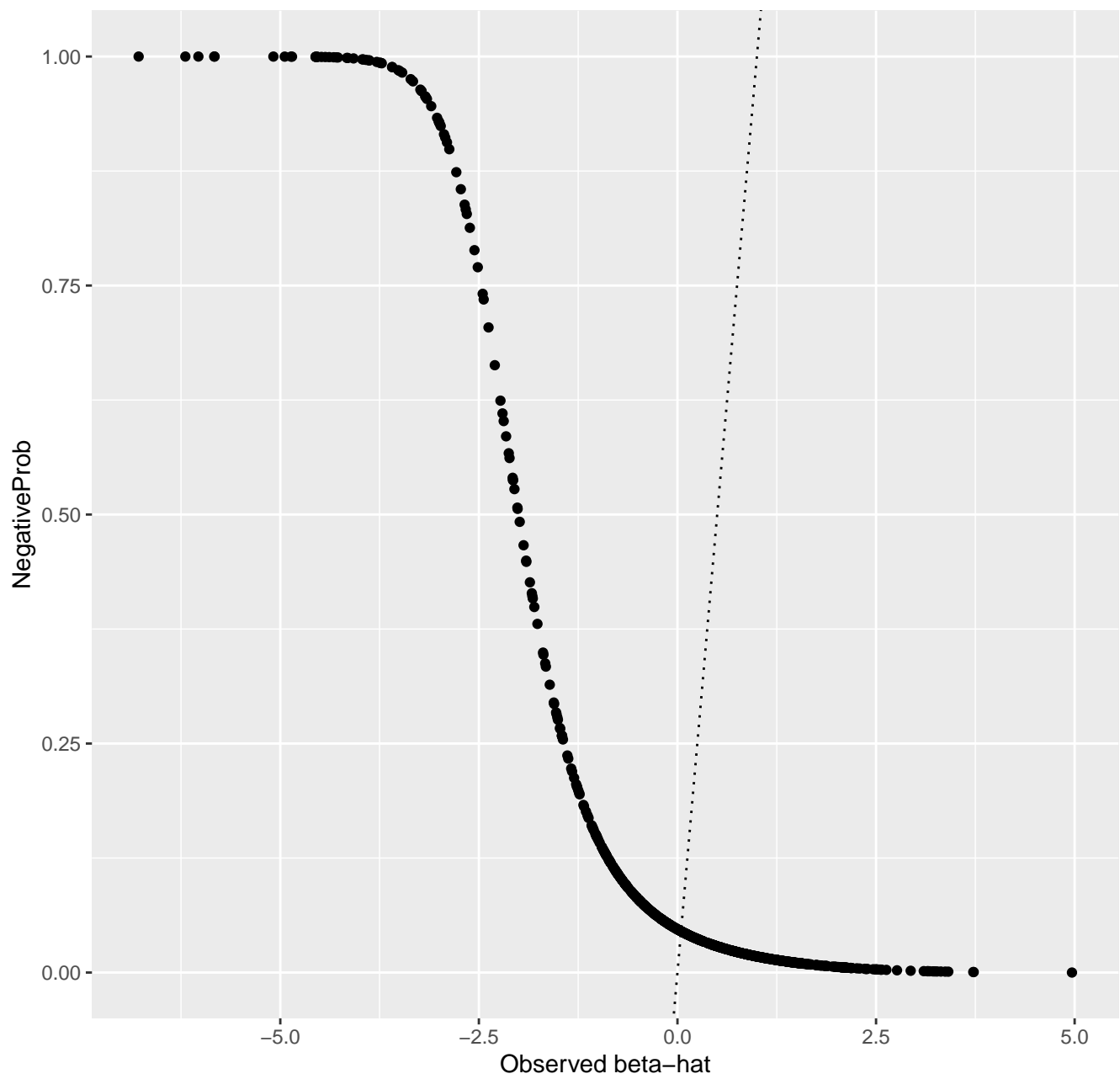
MET_(4233) negative



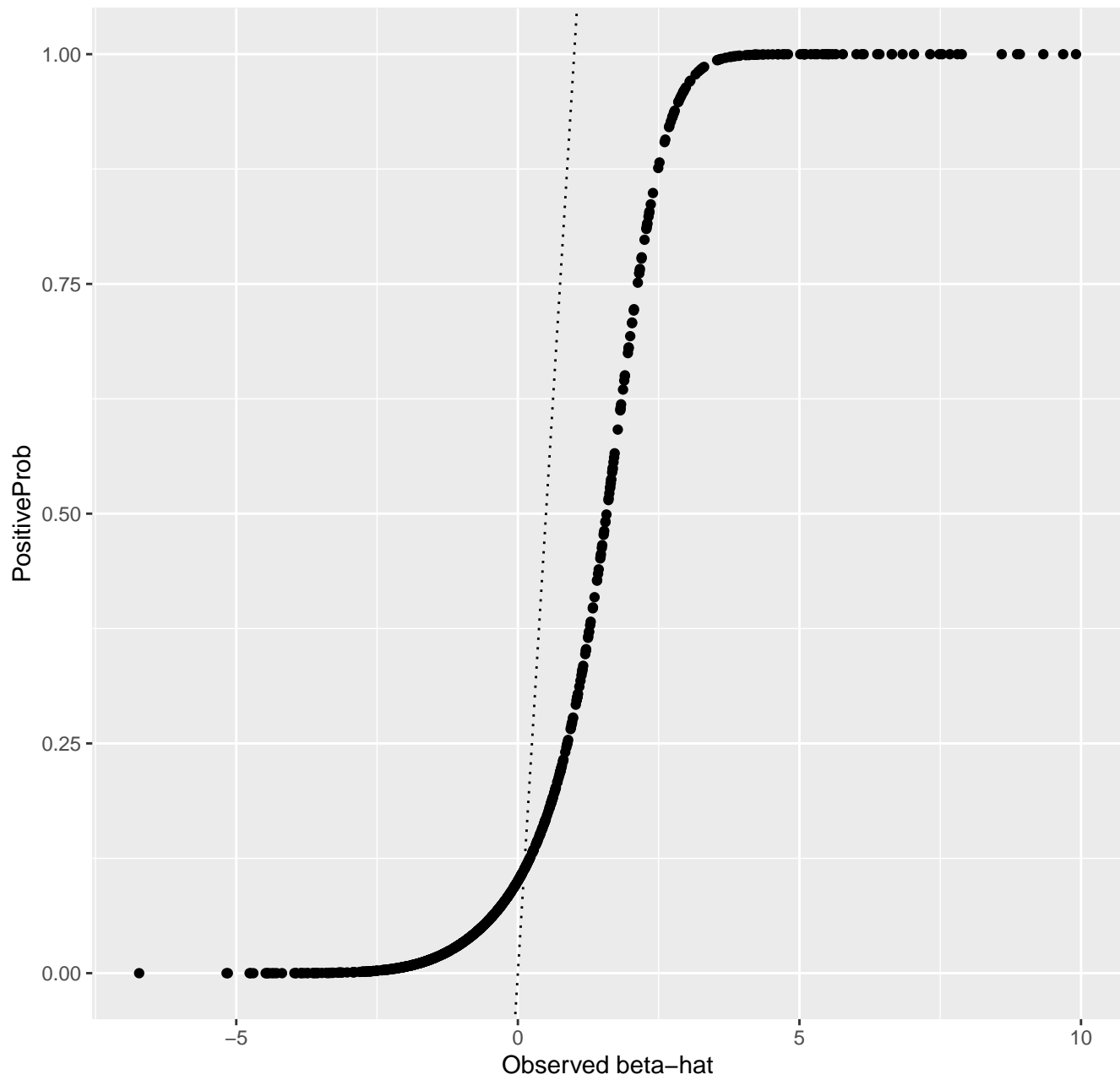
HNF4A_(3172) negative



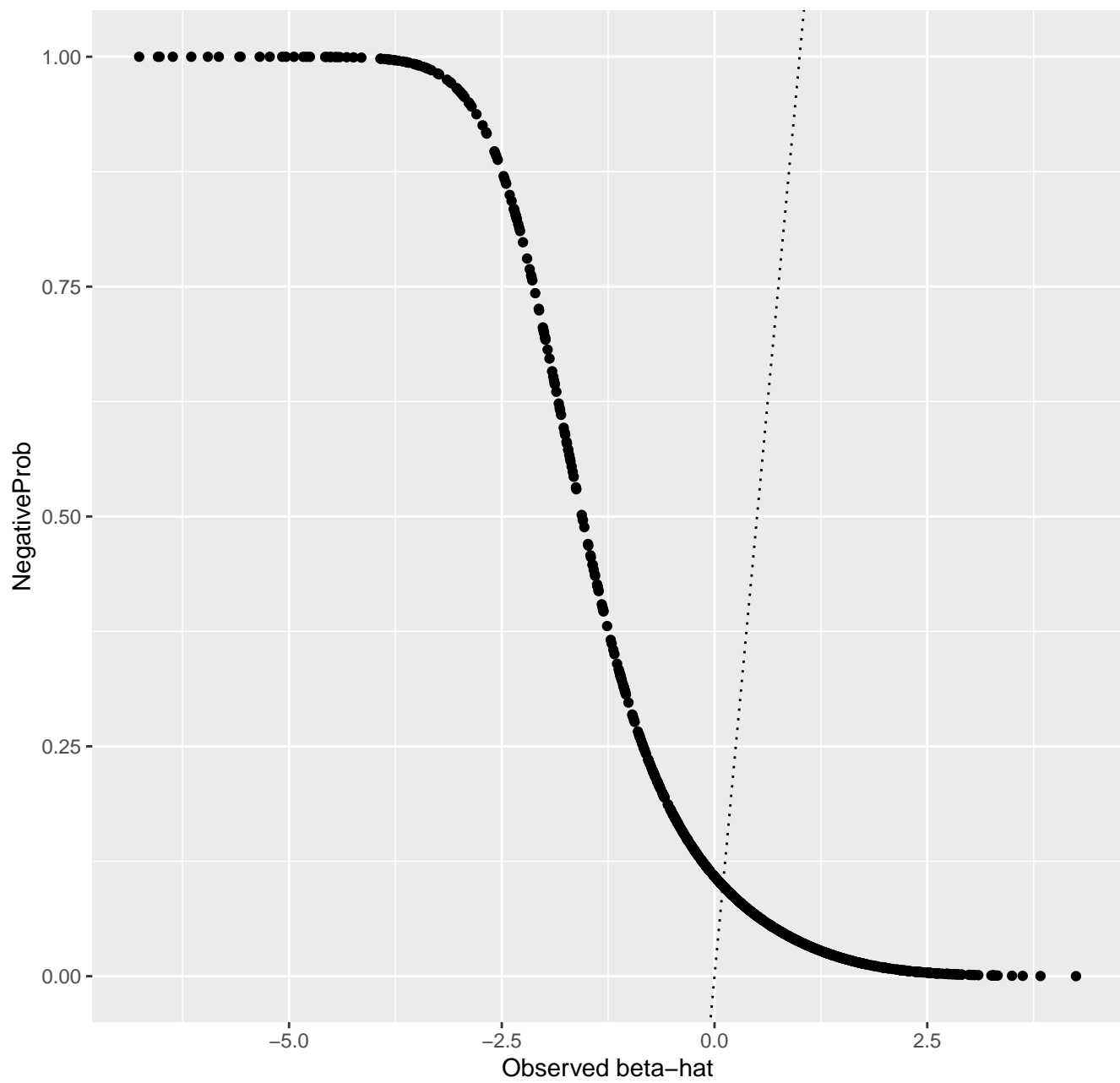
SLC16A1_(6566) negative



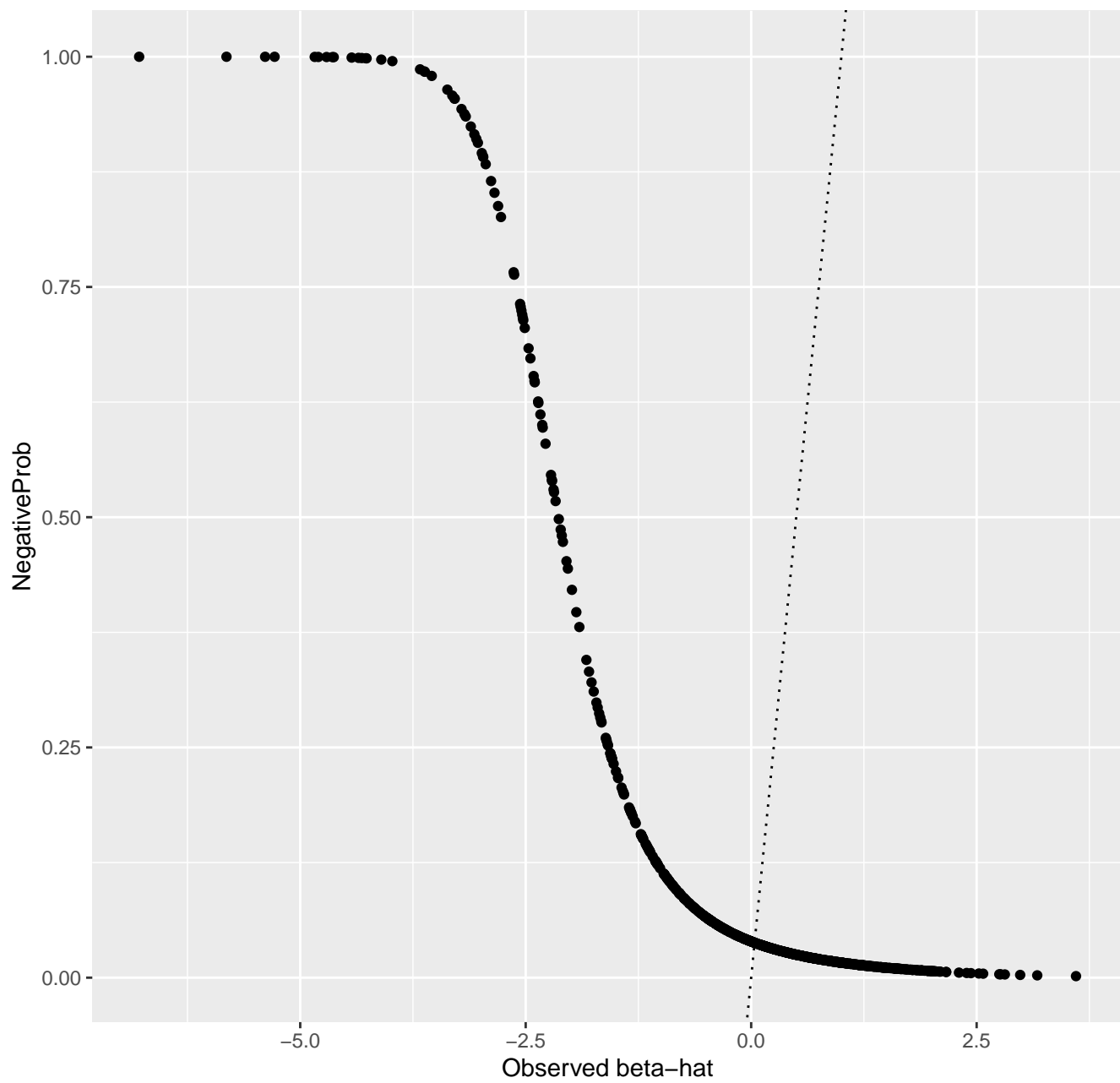
COPG1_(22820) positive



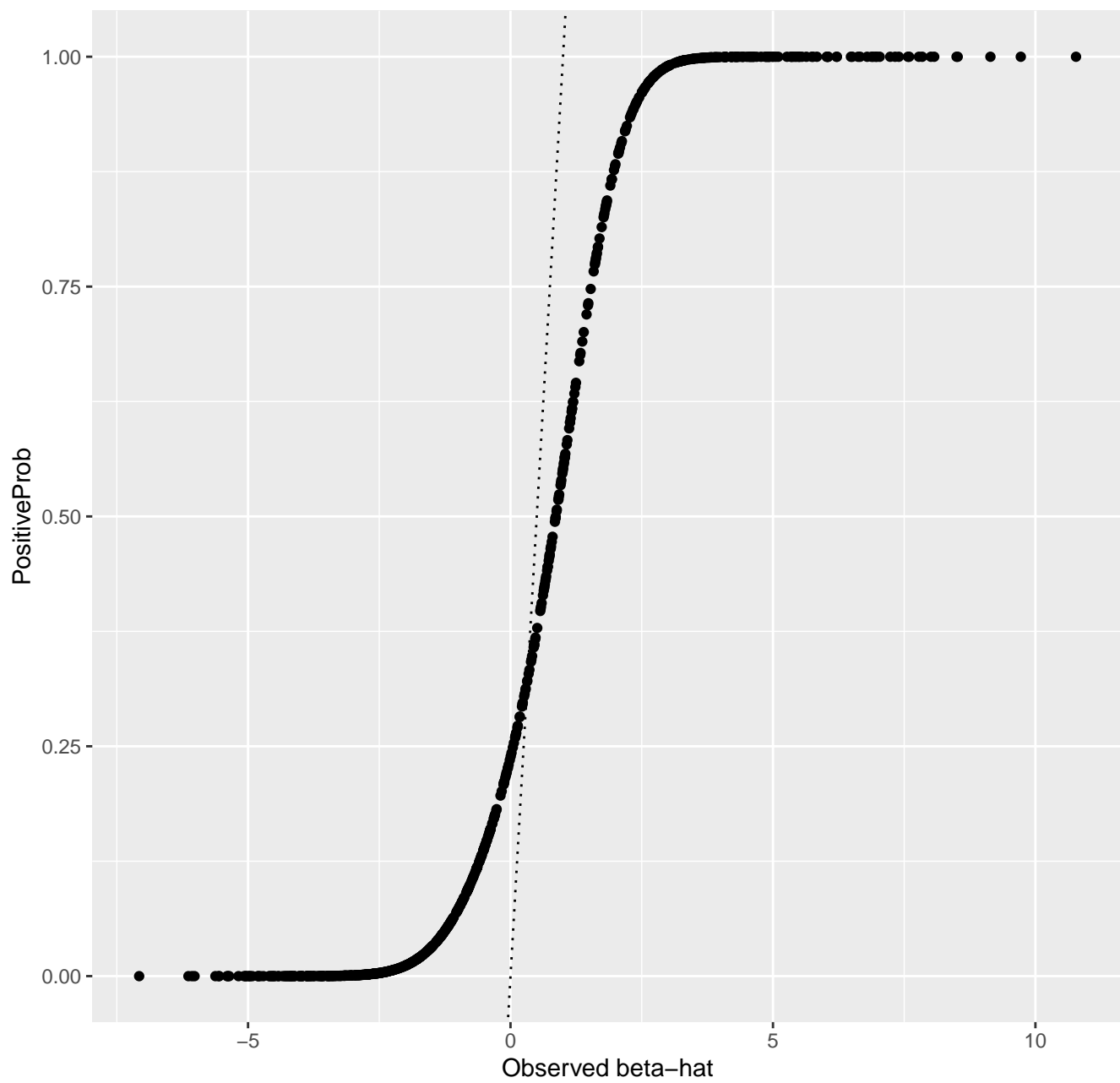
OXSM_(54995) negative



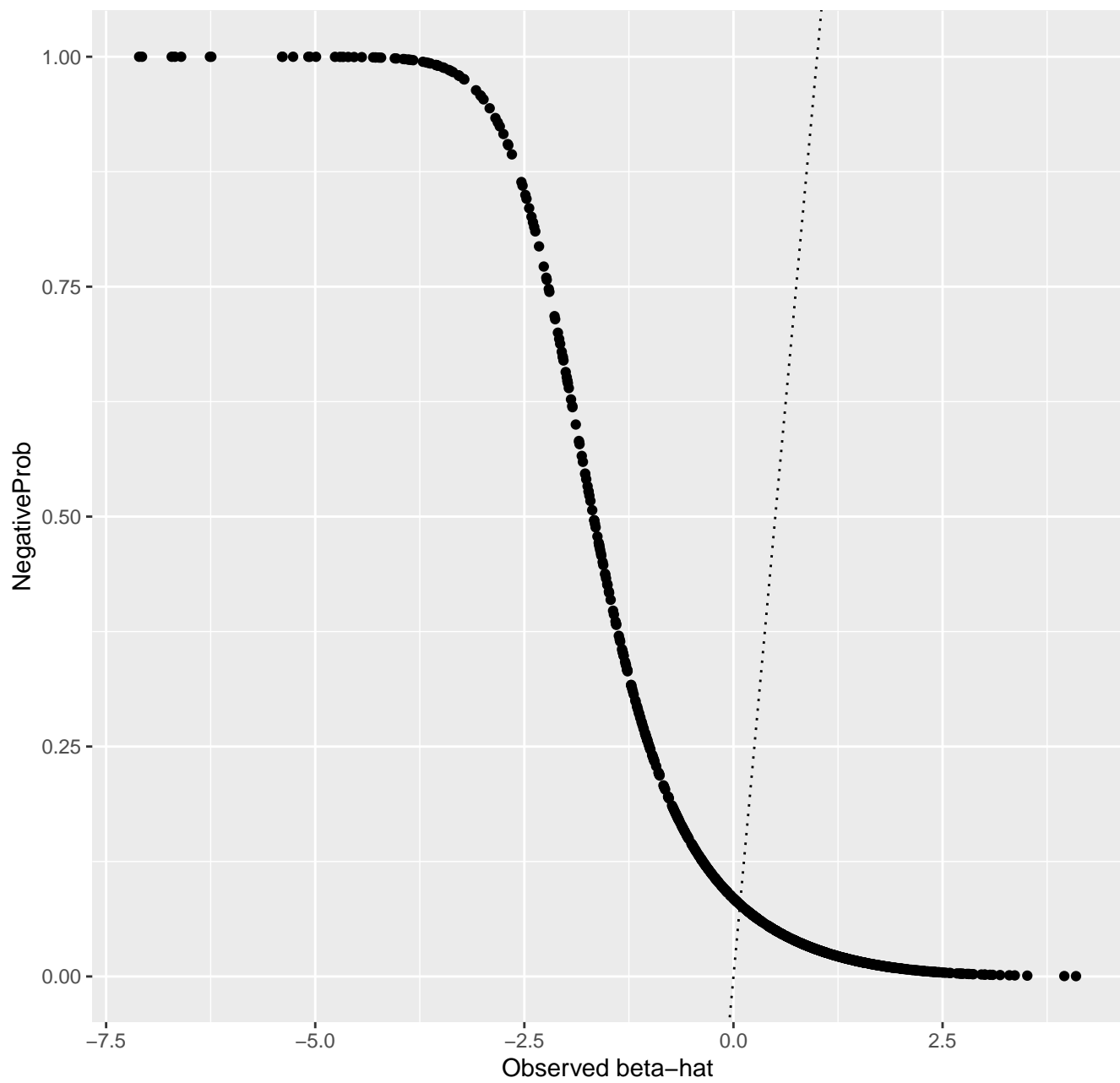
DMAC1_(90871) negative



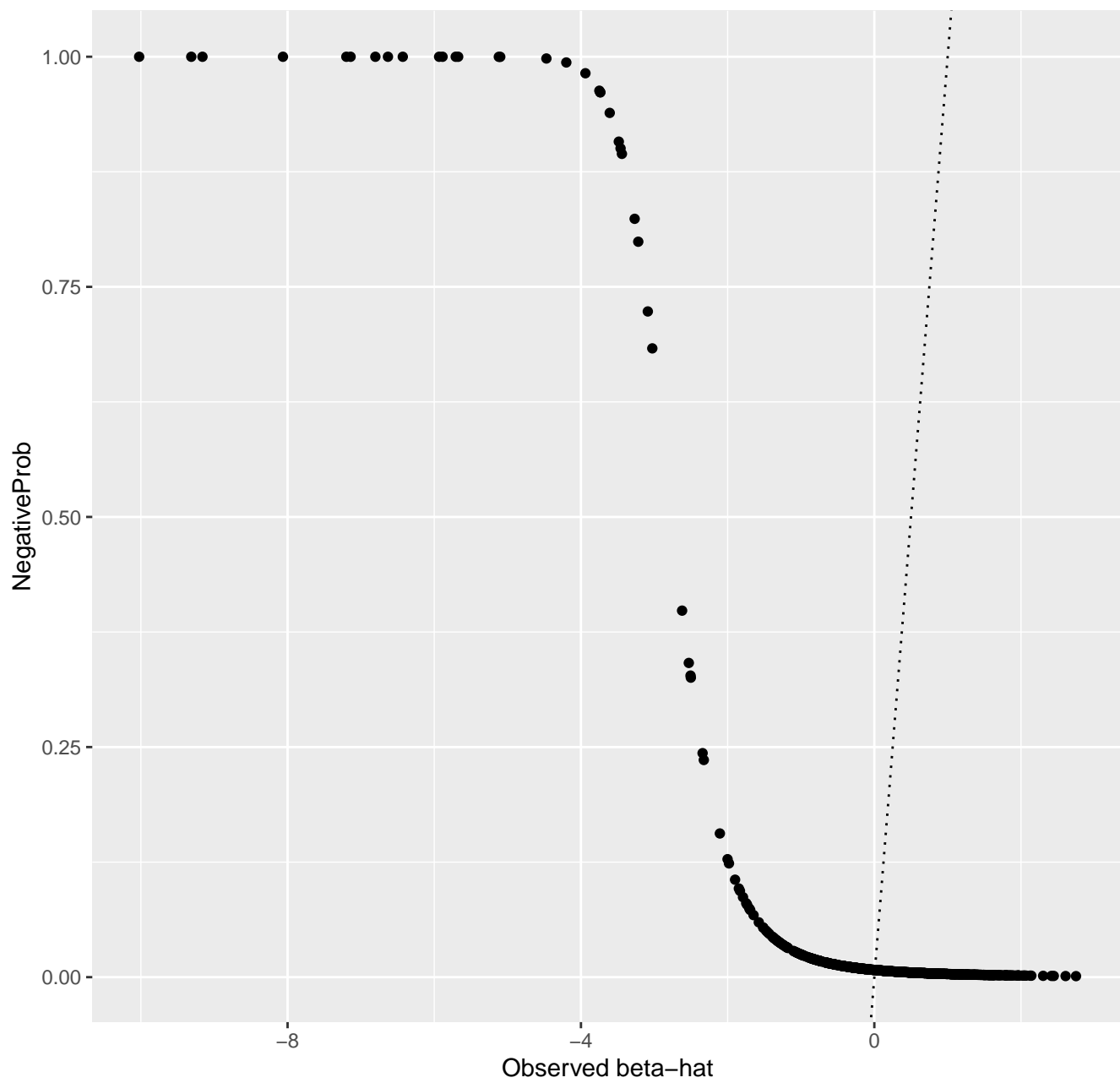
KIF18A_(81930) positive



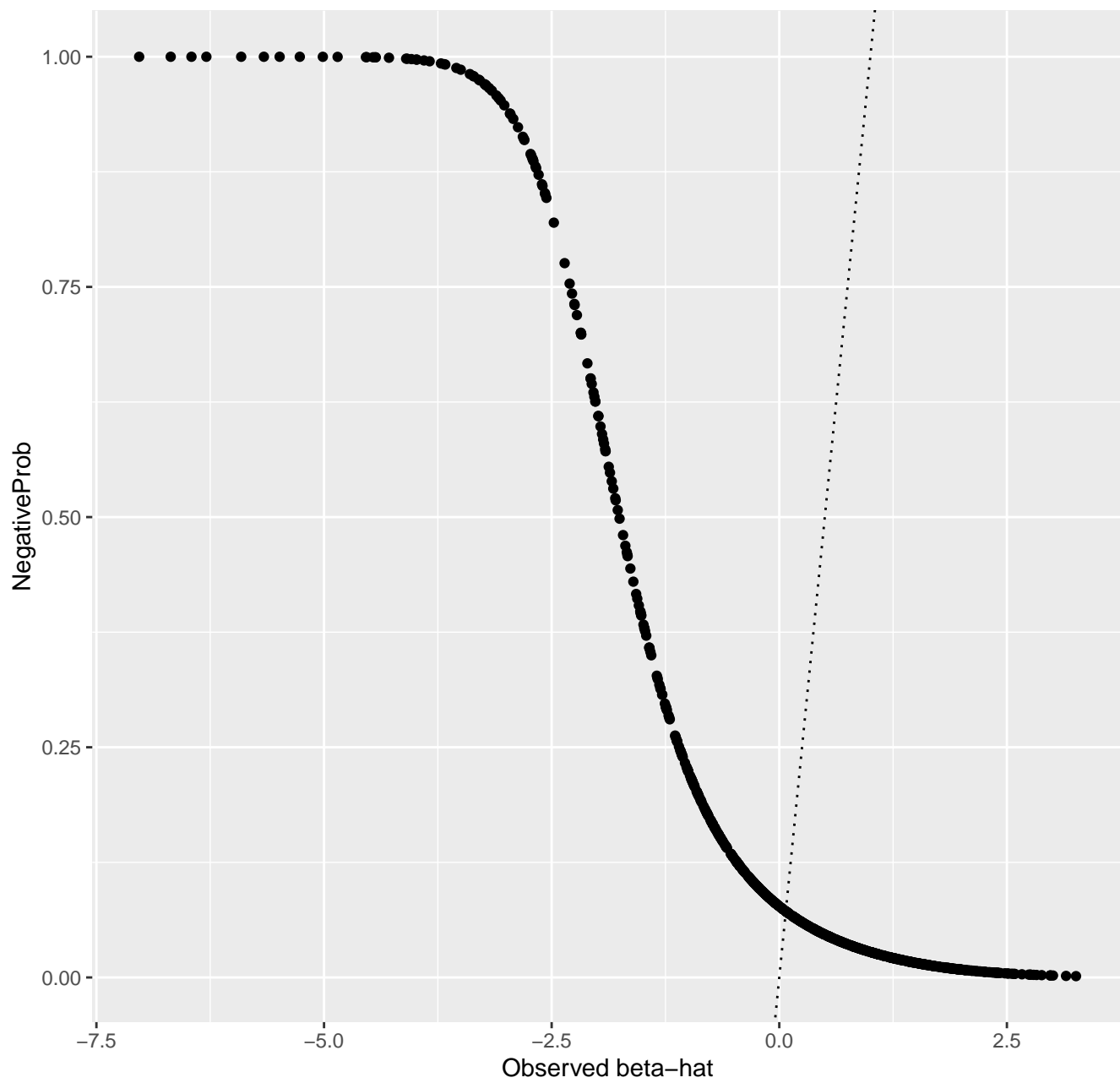
CRK_(1398) negative



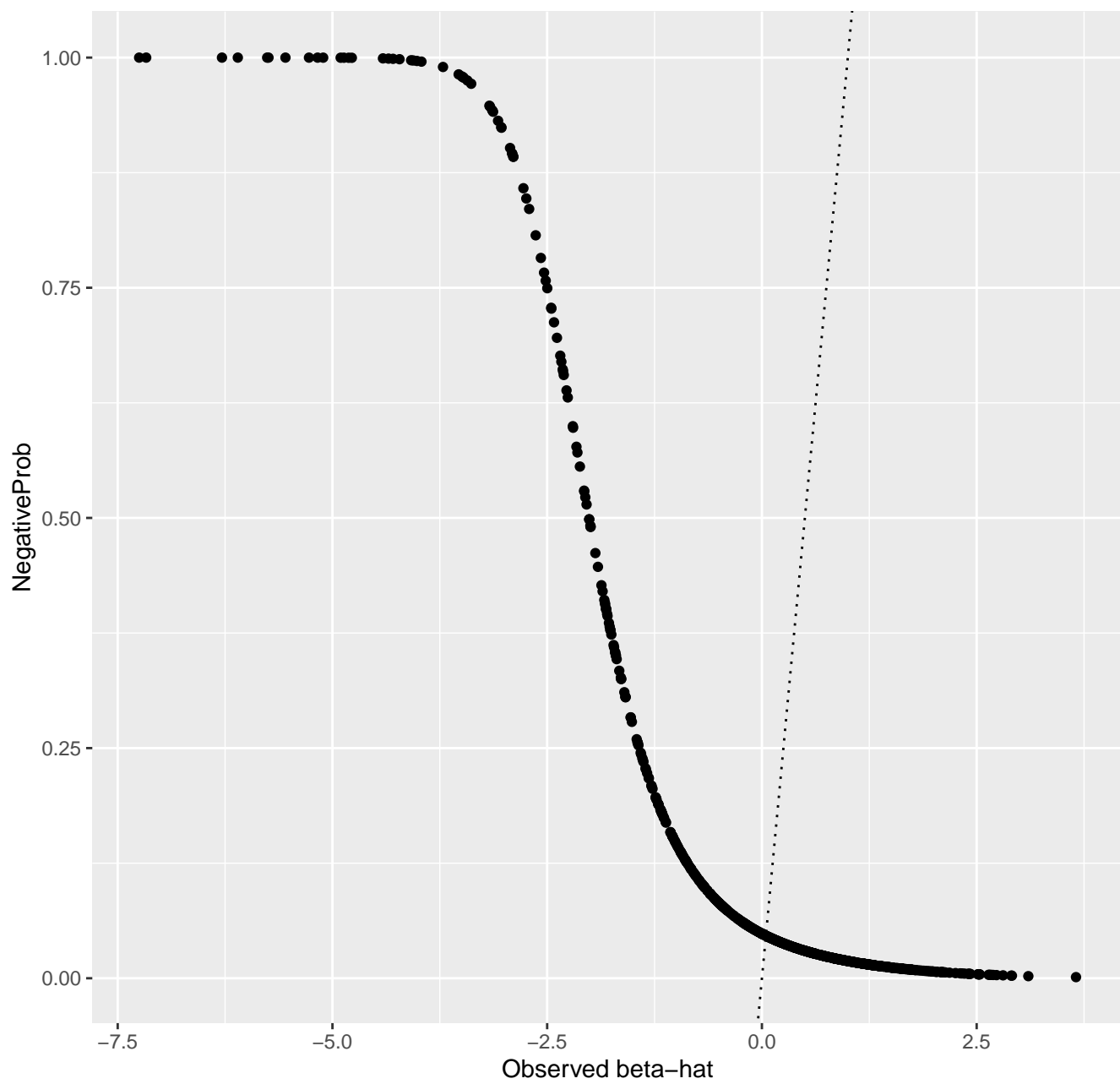
POU2AF1_(5450) negative



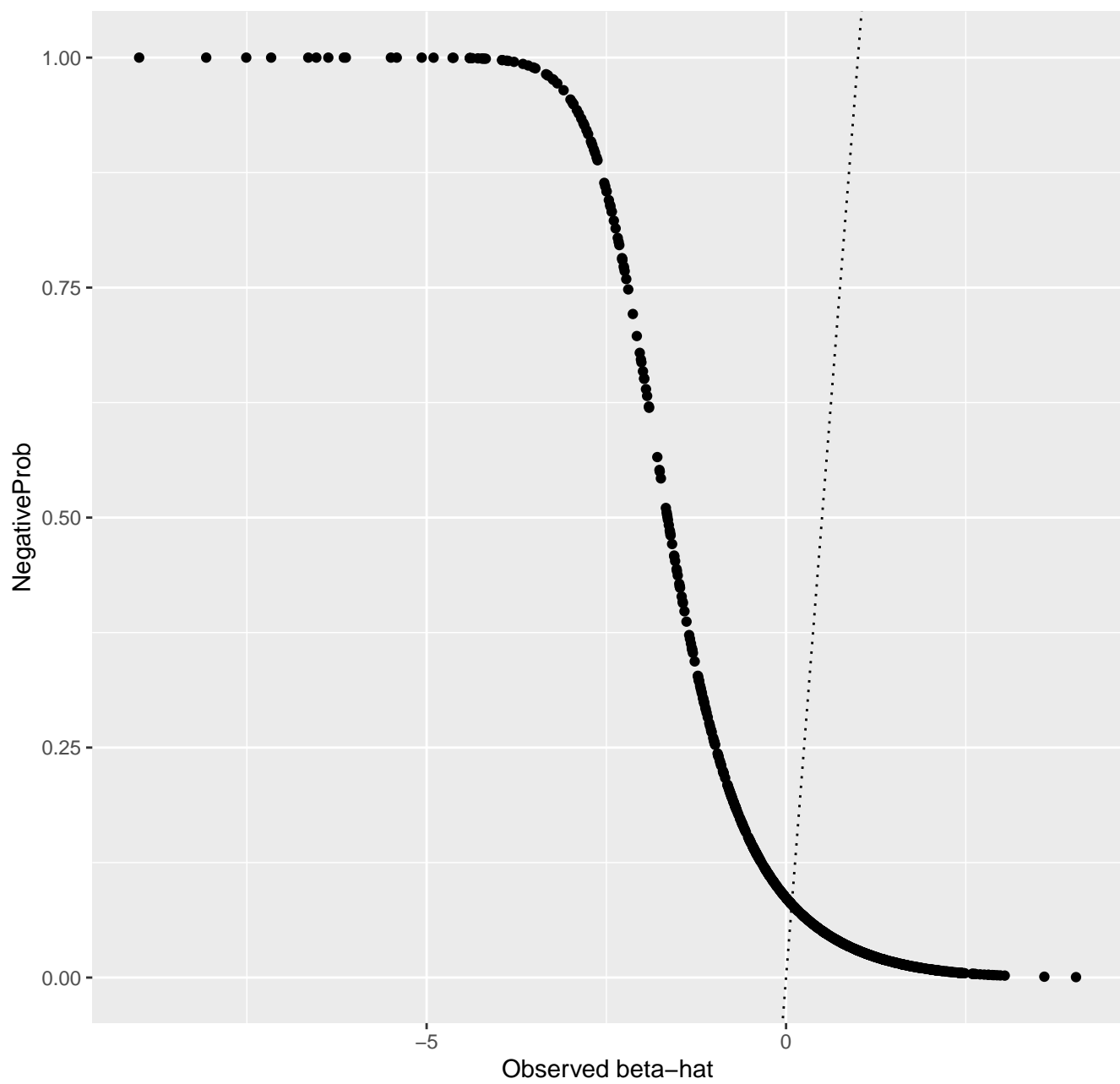
POC1A_(25886) negative



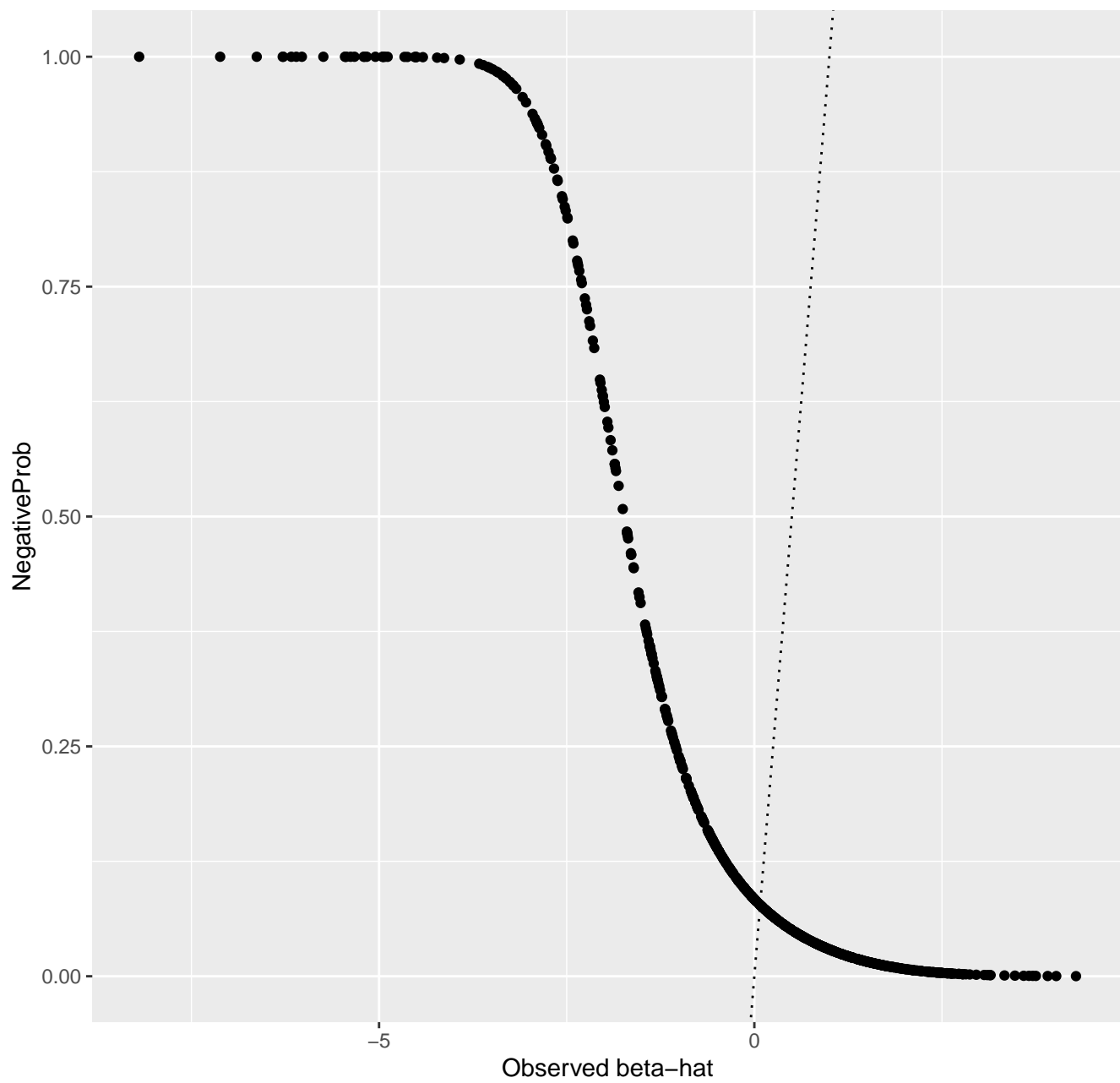
HTRA2_(27429) negative



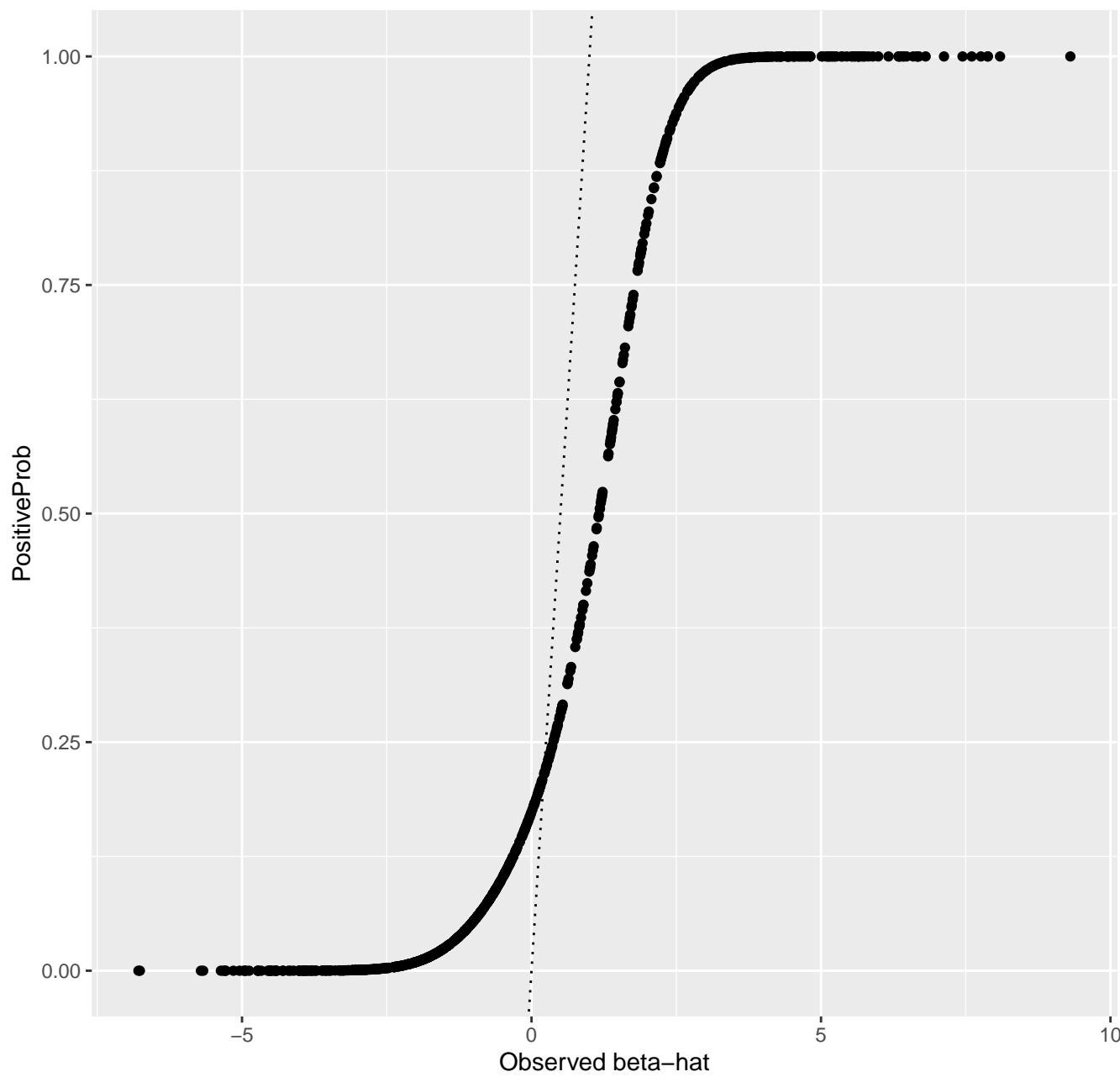
APEX2_(27301) negative



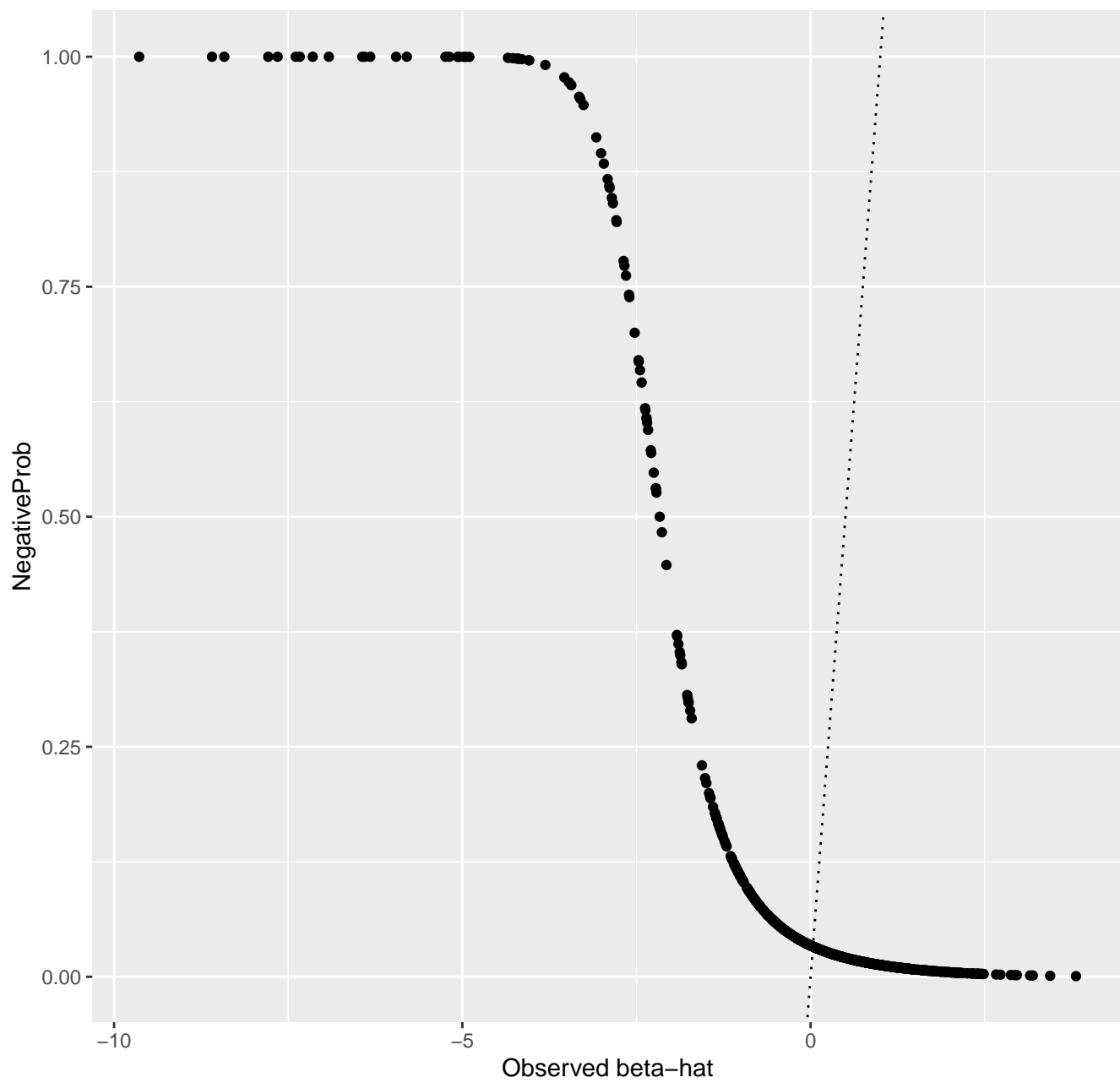
E2F3_(1871) negative



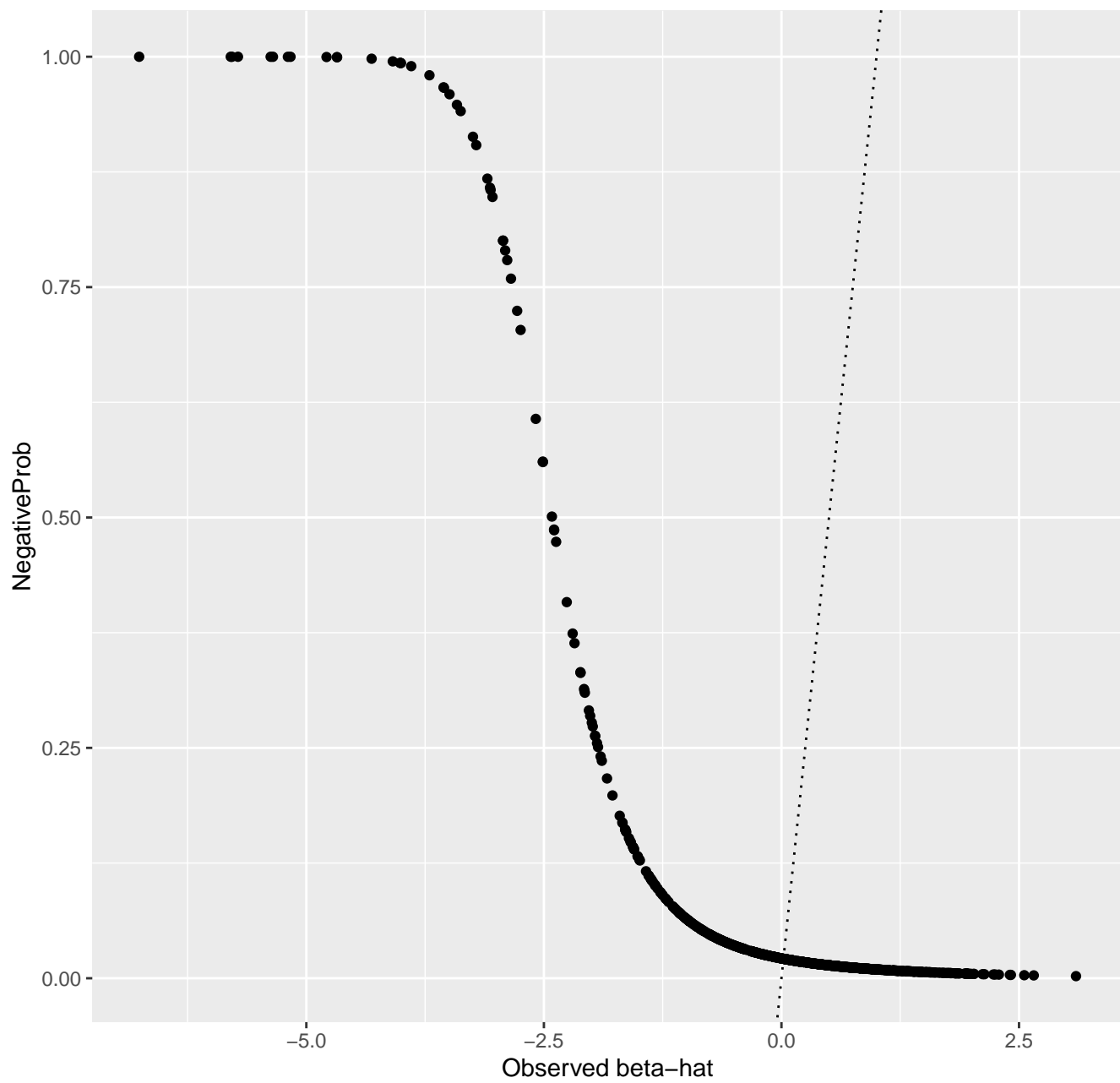
ATP1A1_(476) positive



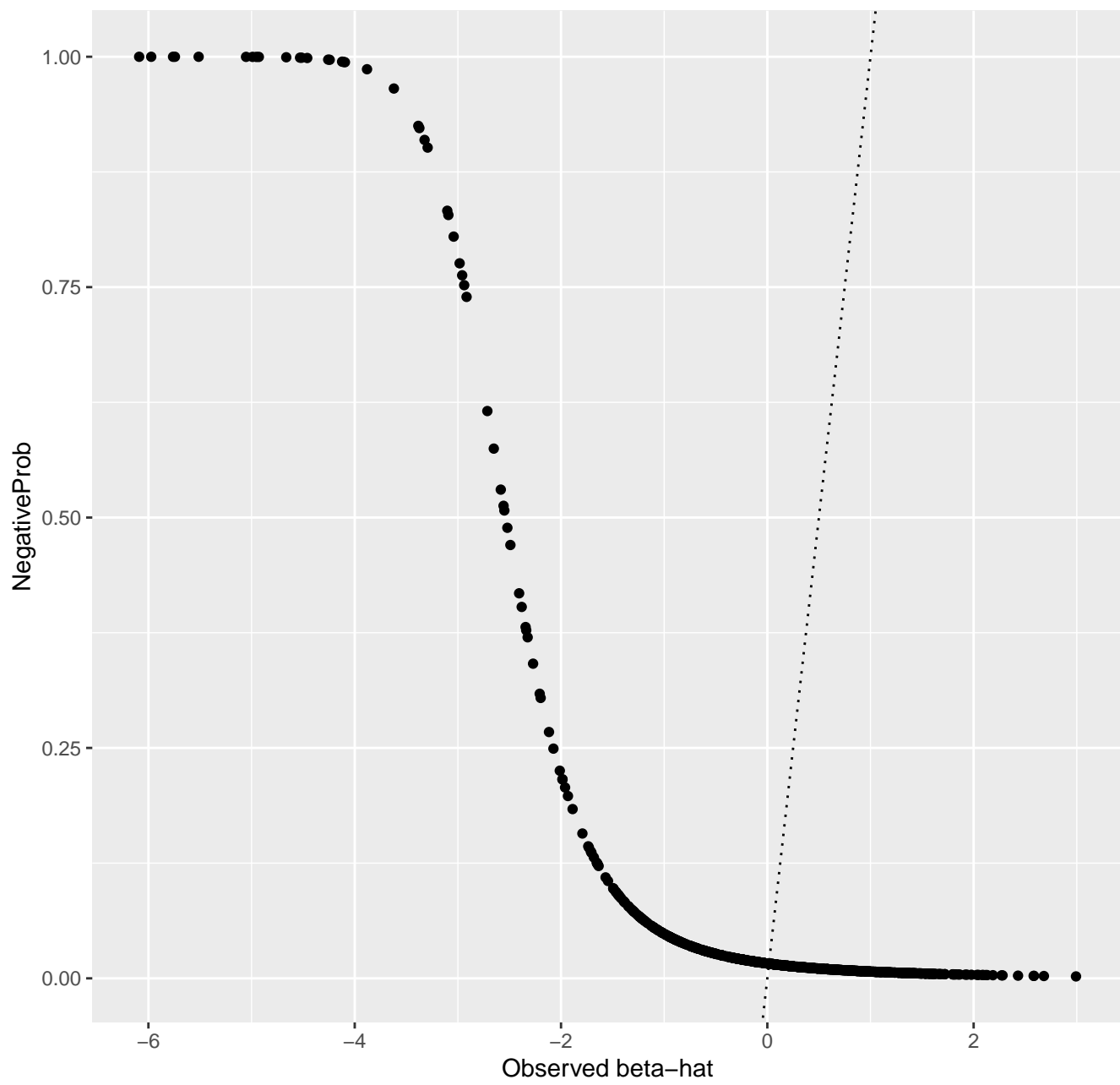
CCNE1_(898) negative



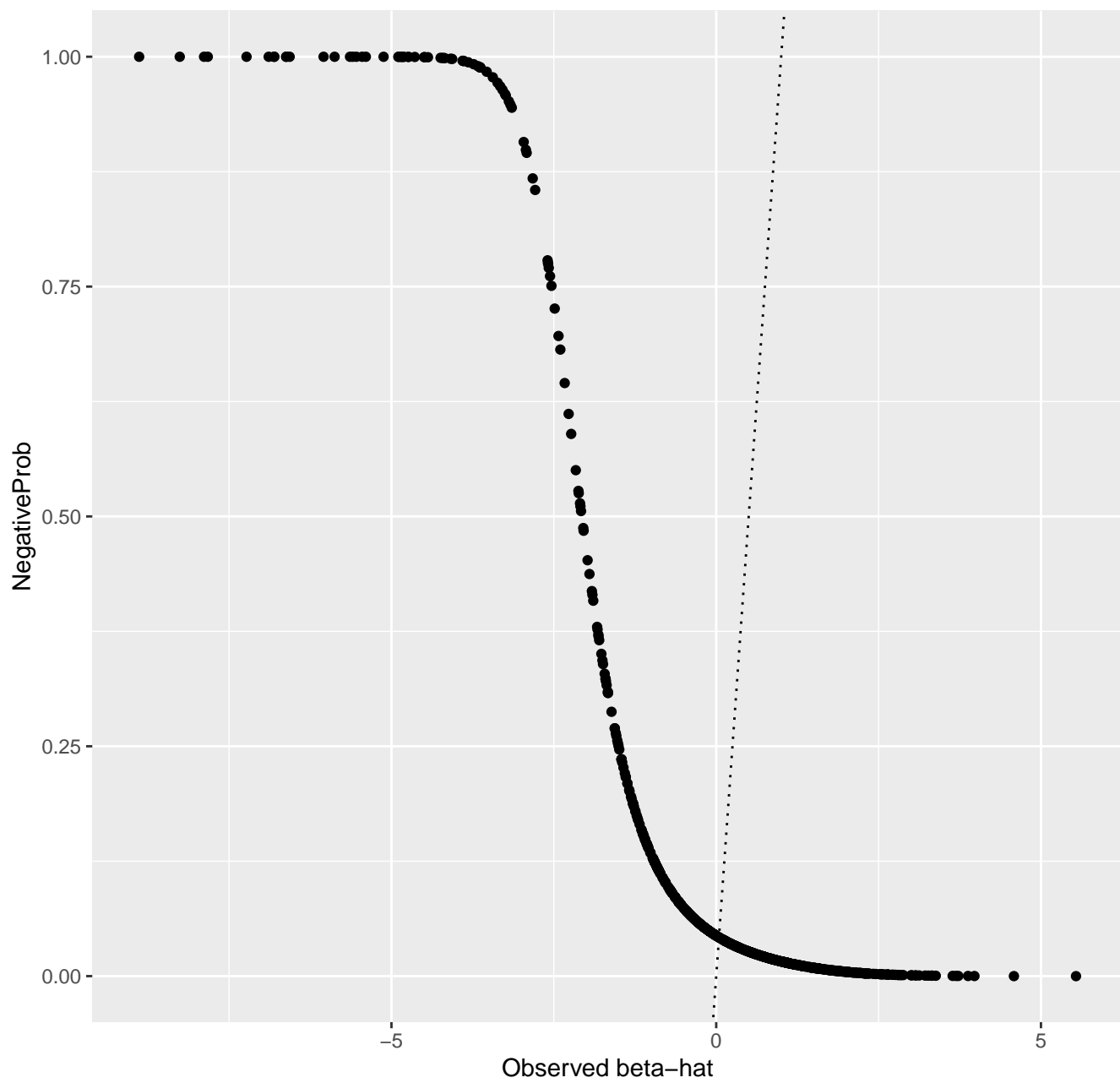
MTR_(4548) negative



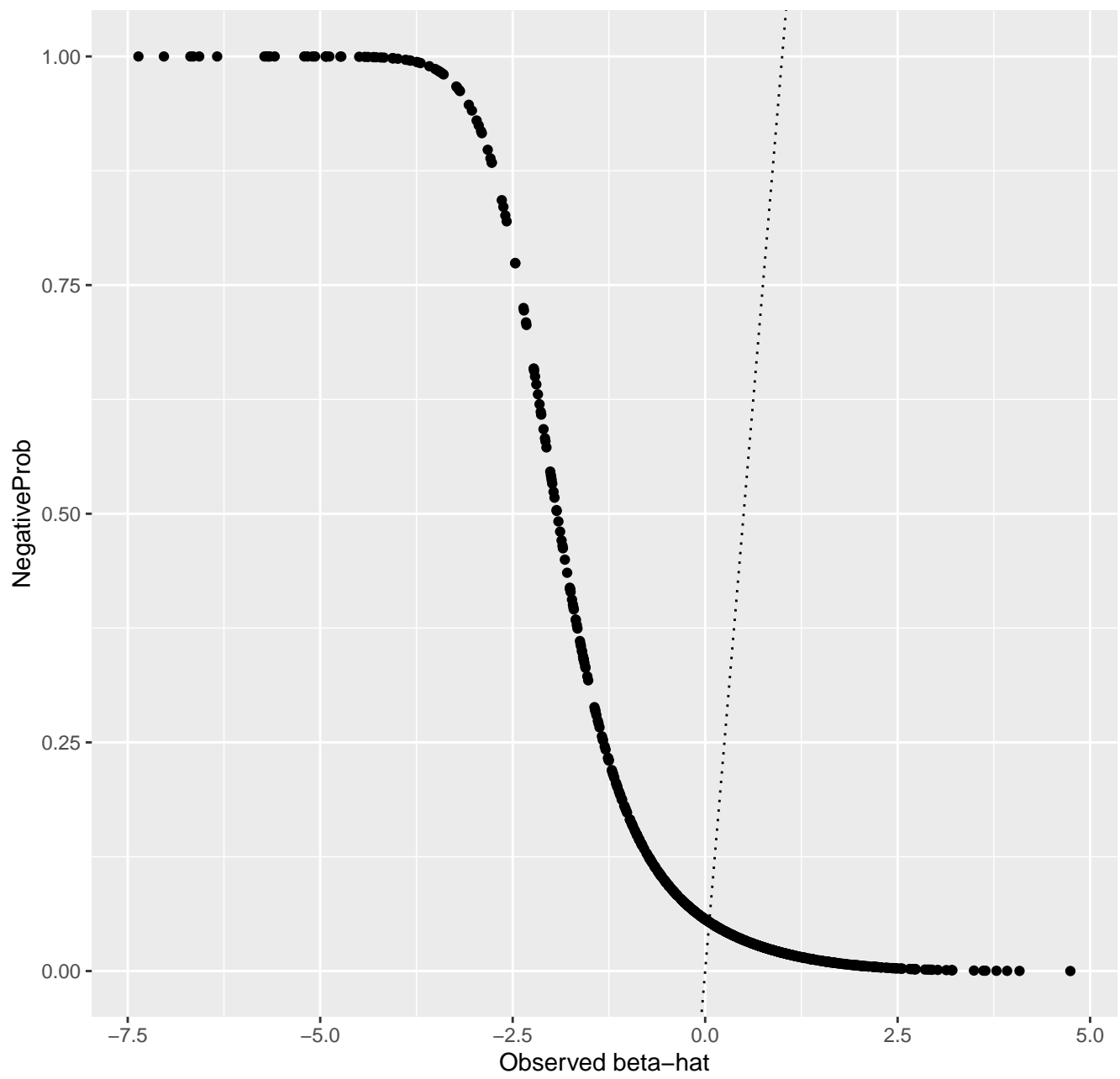
WRAP73_(49856) negative



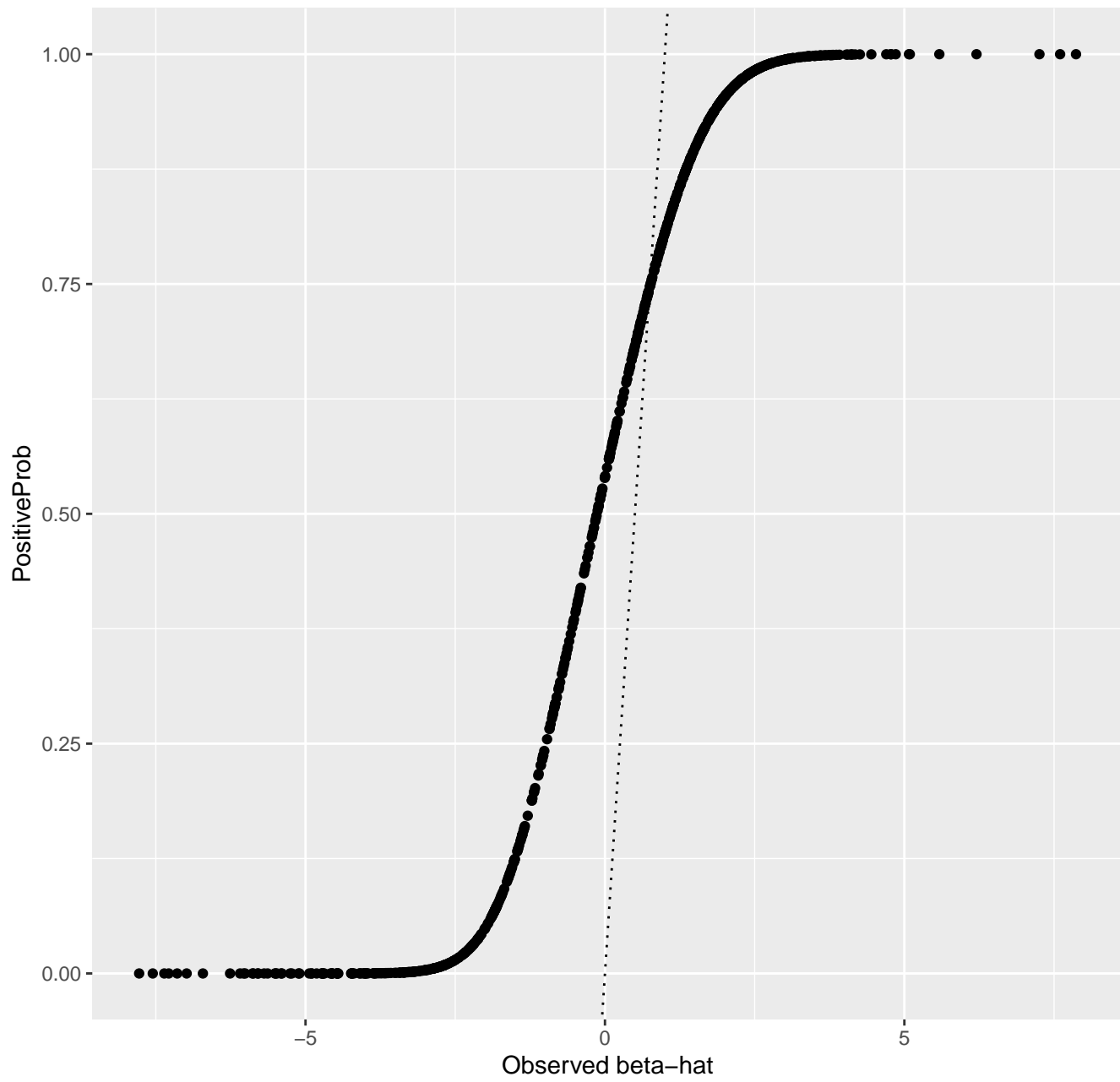
LDB1_(8861) negative



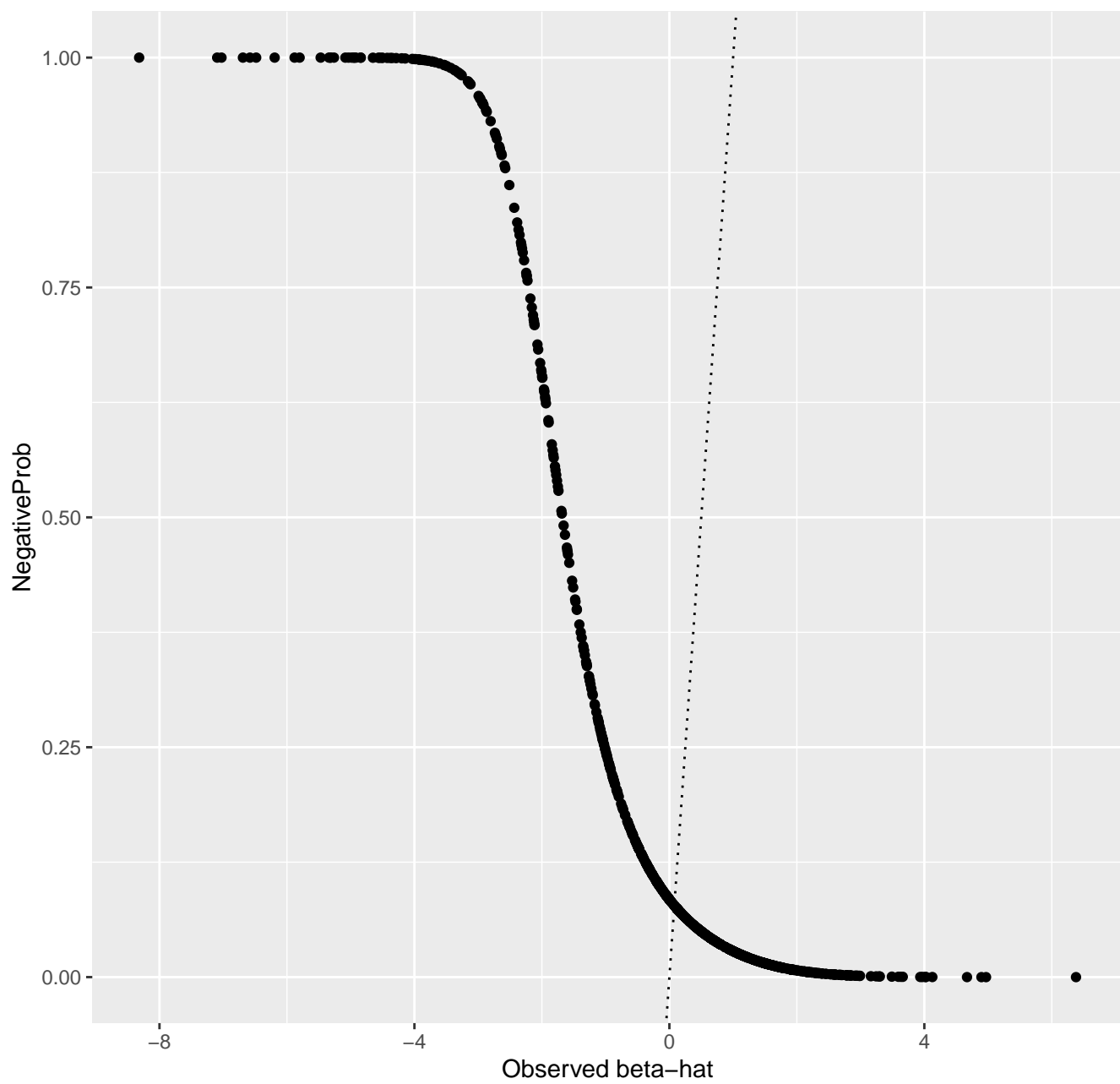
DNAJC13_(23317) negative



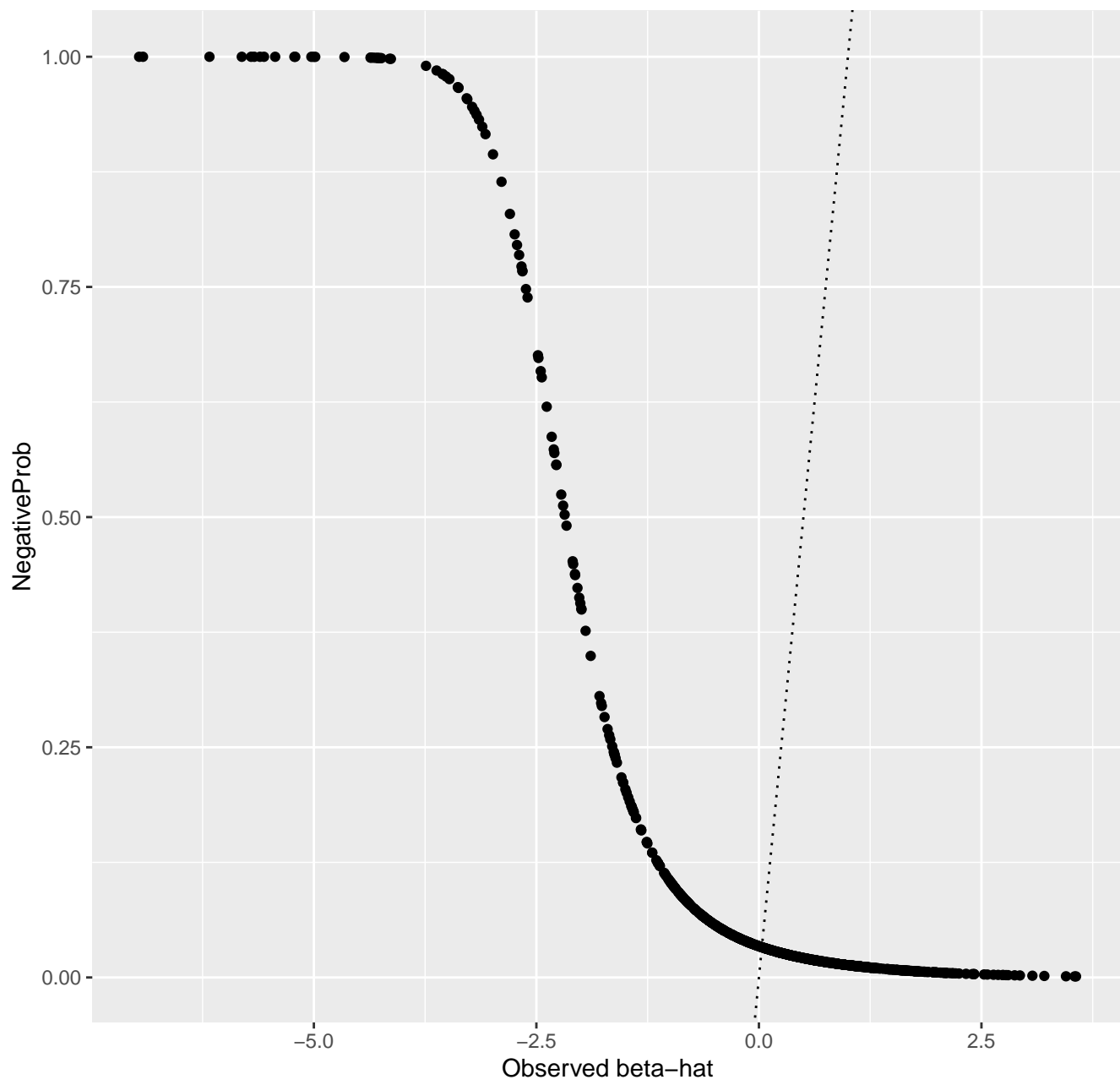
SLC2A1_(6513) positive



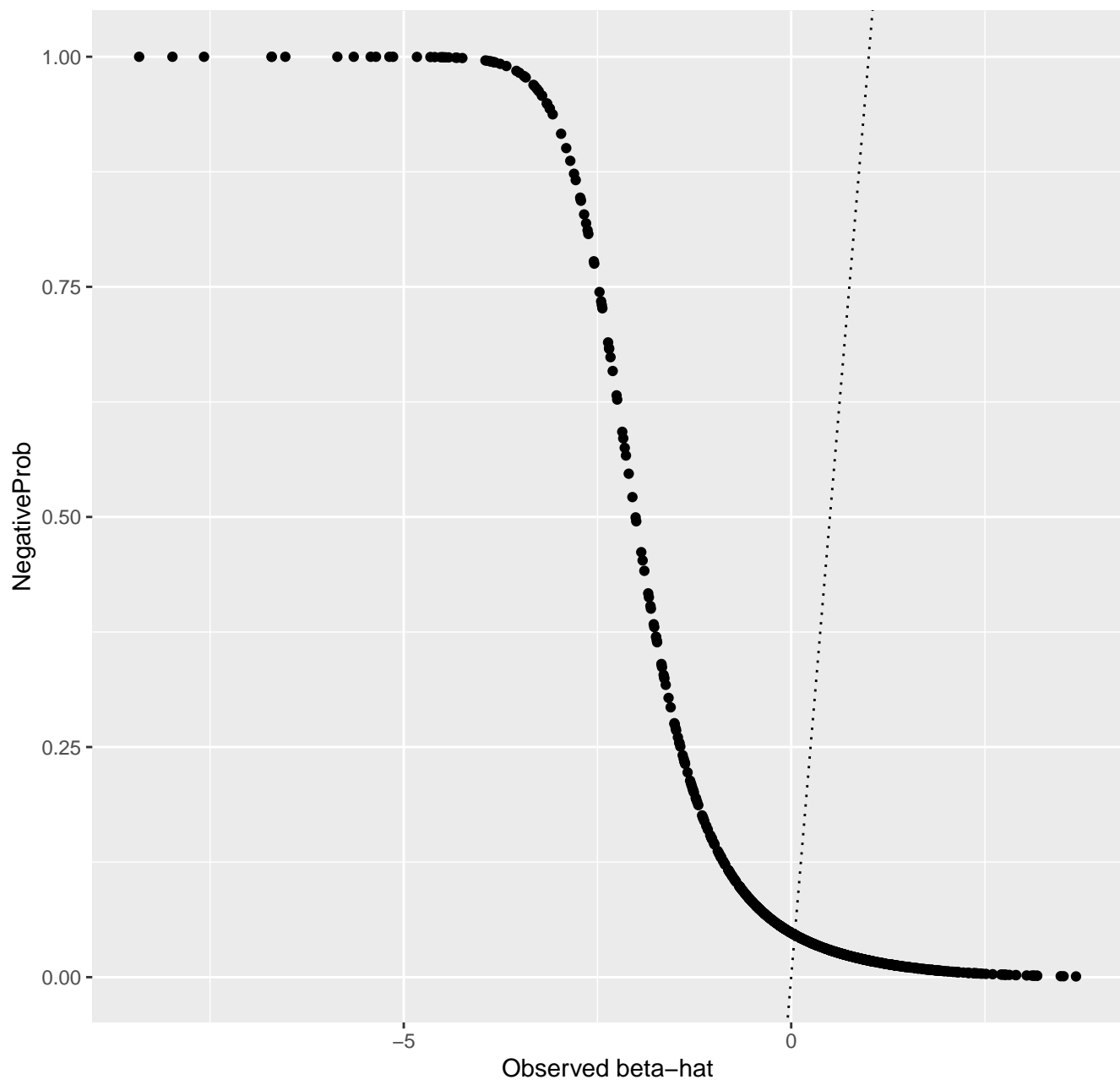
JUN_(3725) negative



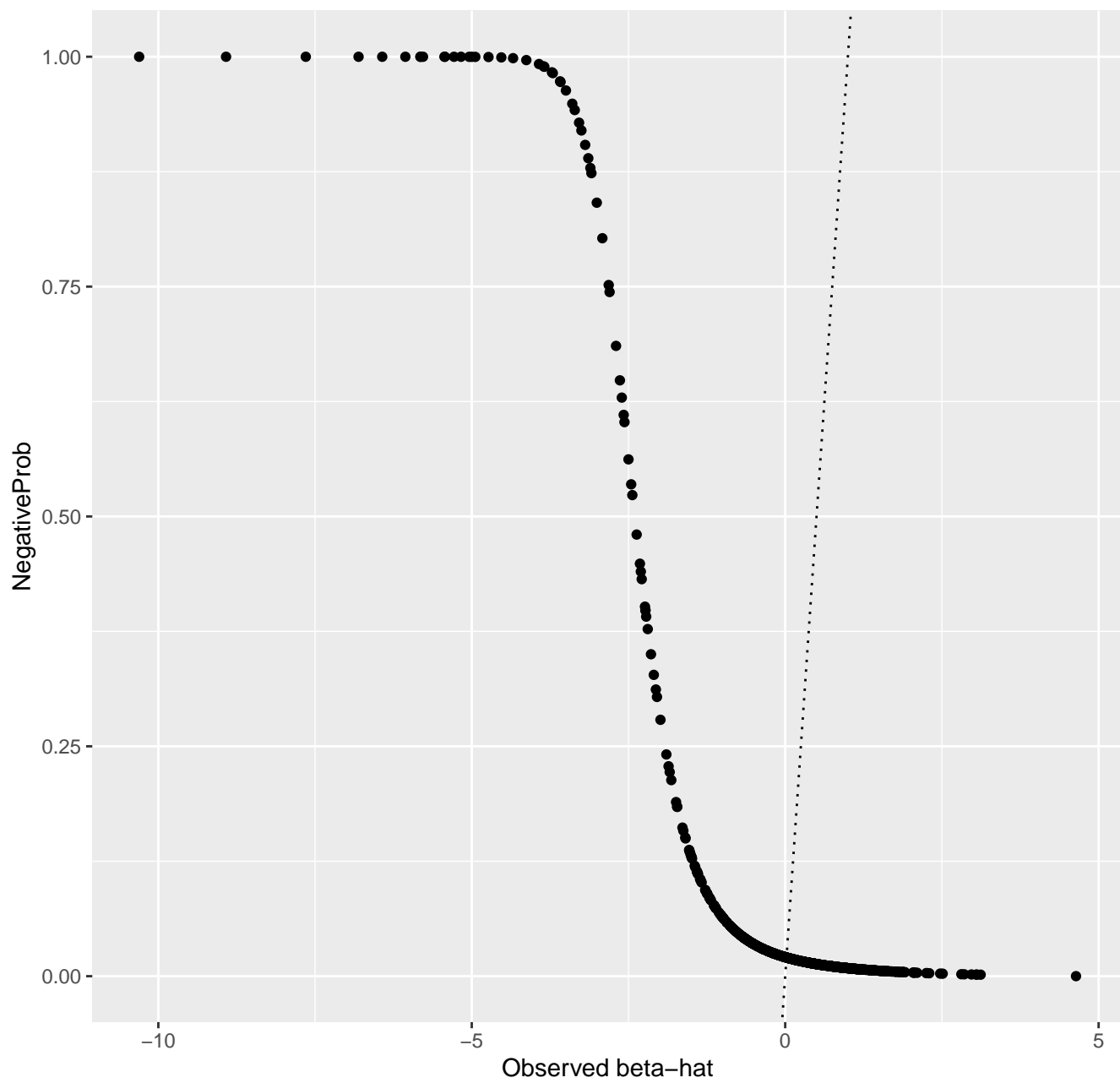
PARD3_(56288) negative



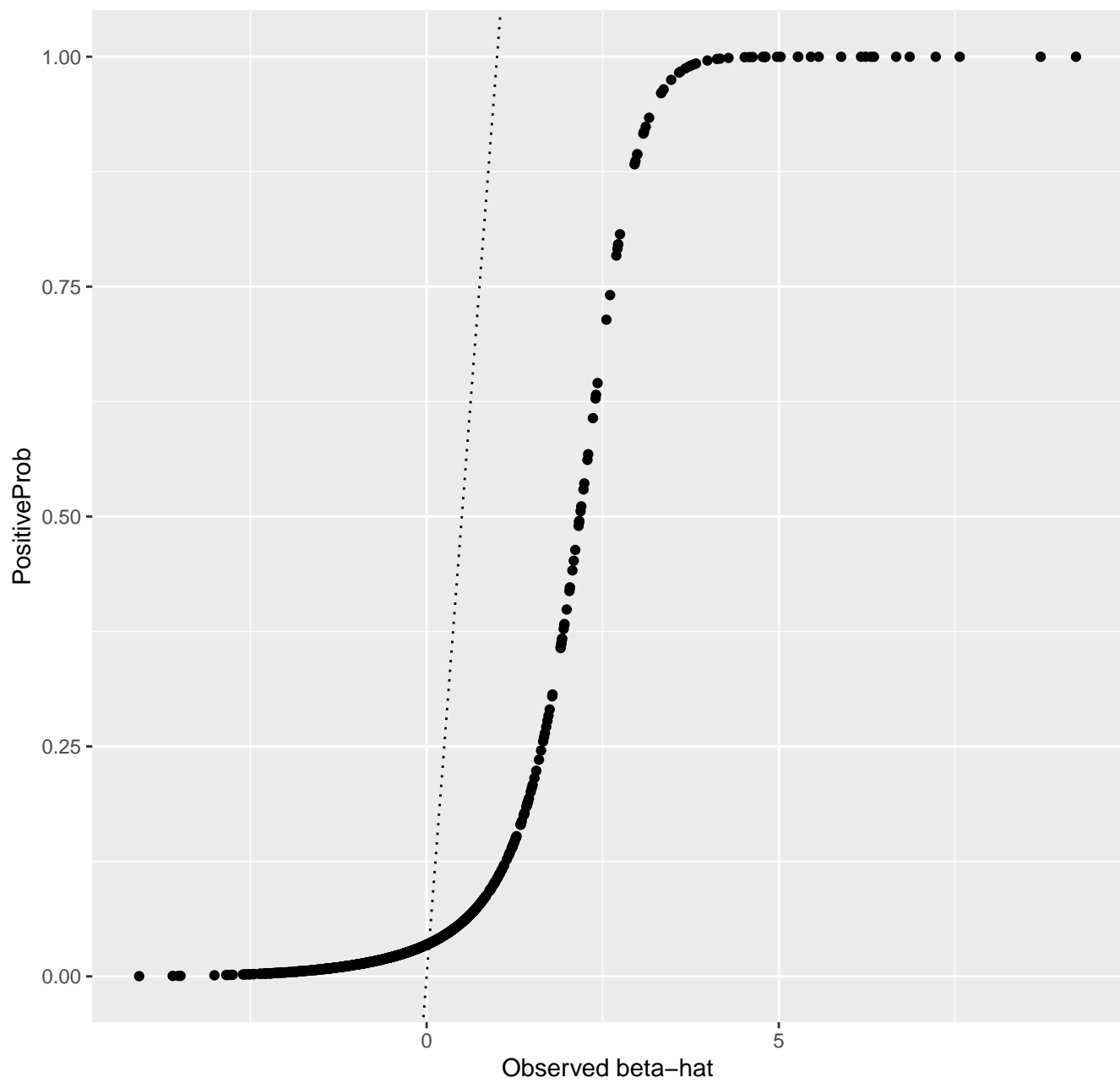
ITGA3_(3675) negative



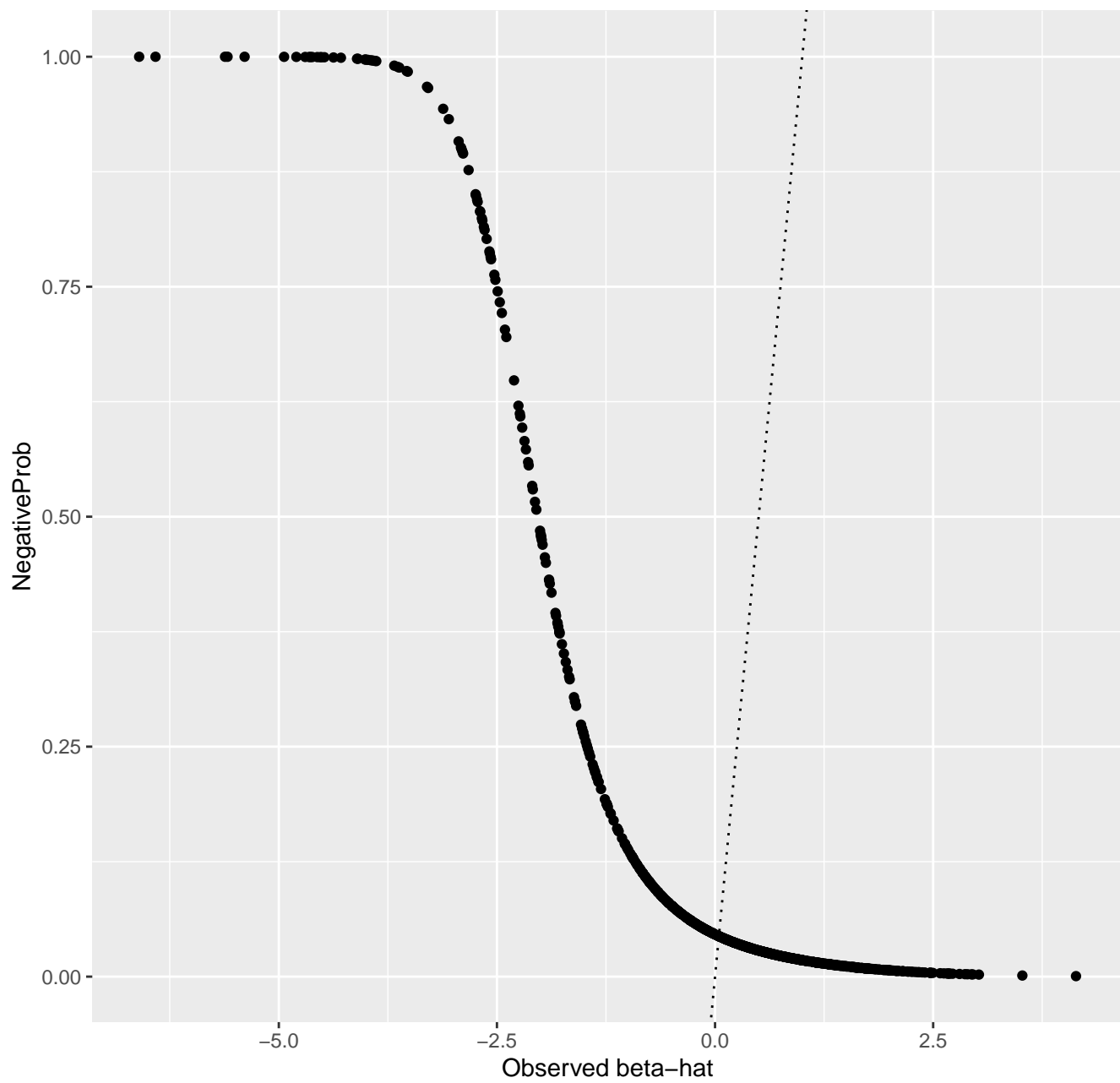
FERMT1_(55612) negative



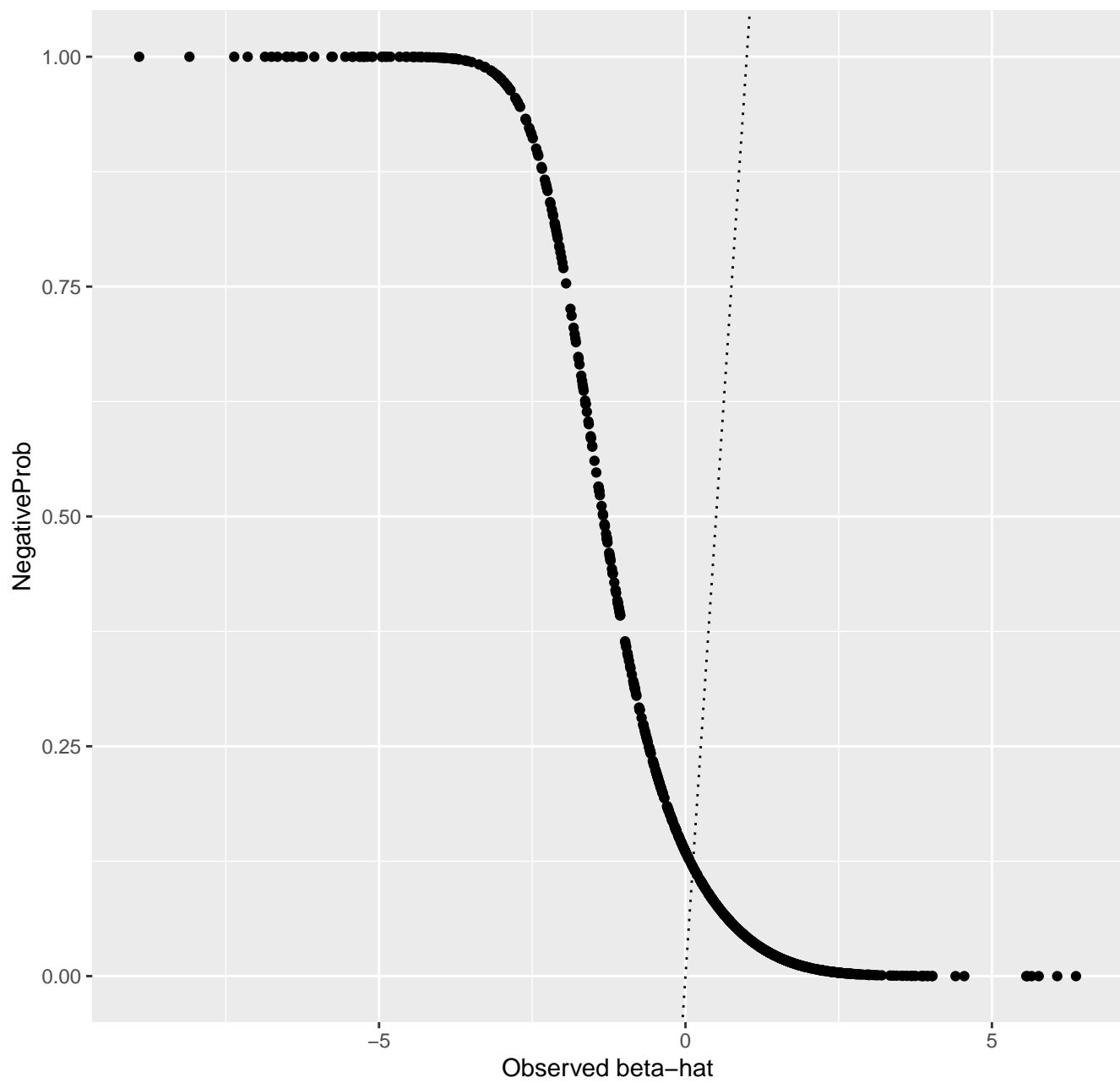
KIRREL1_(55243) positive



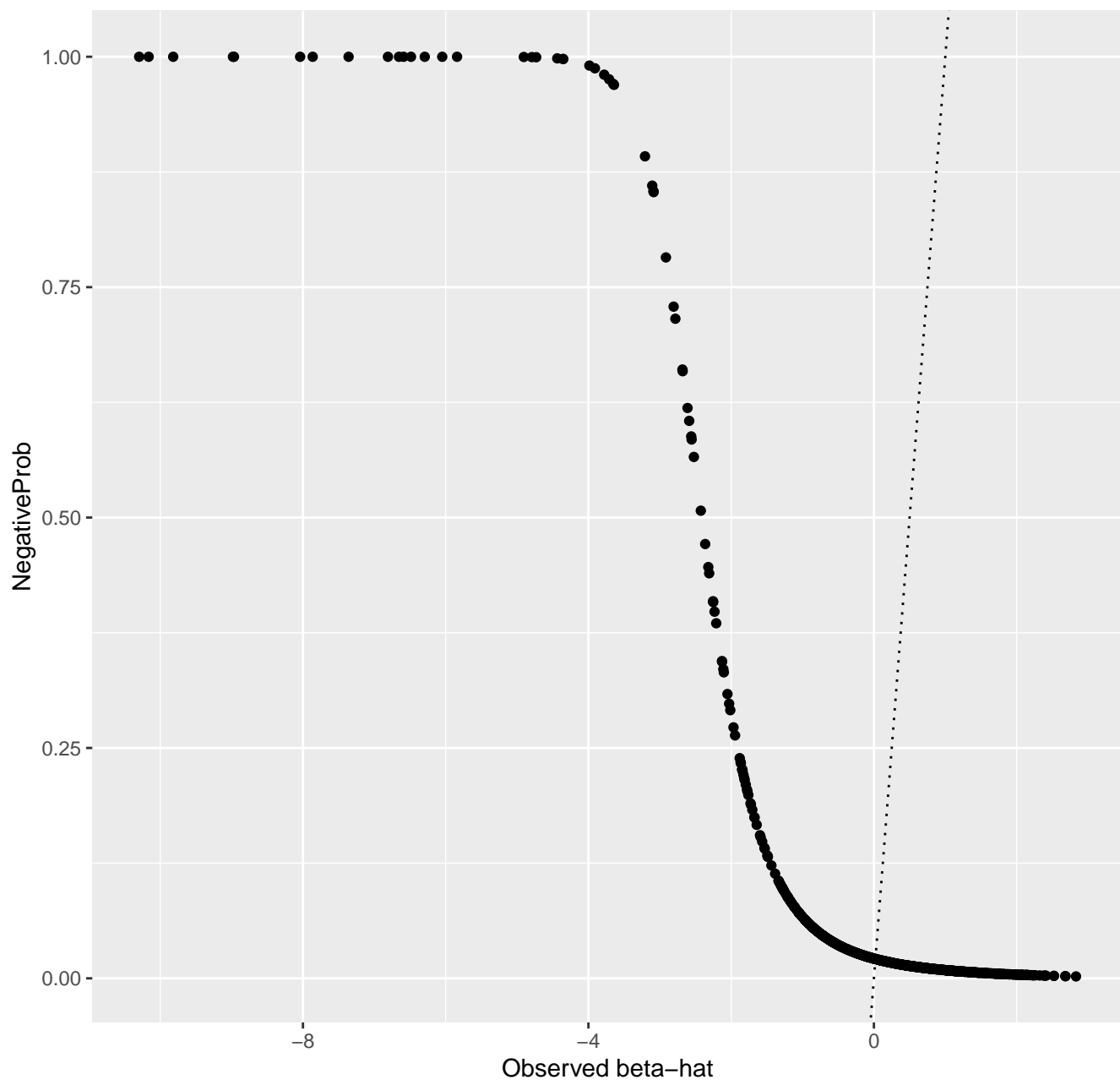
MDM4_(4194) negative



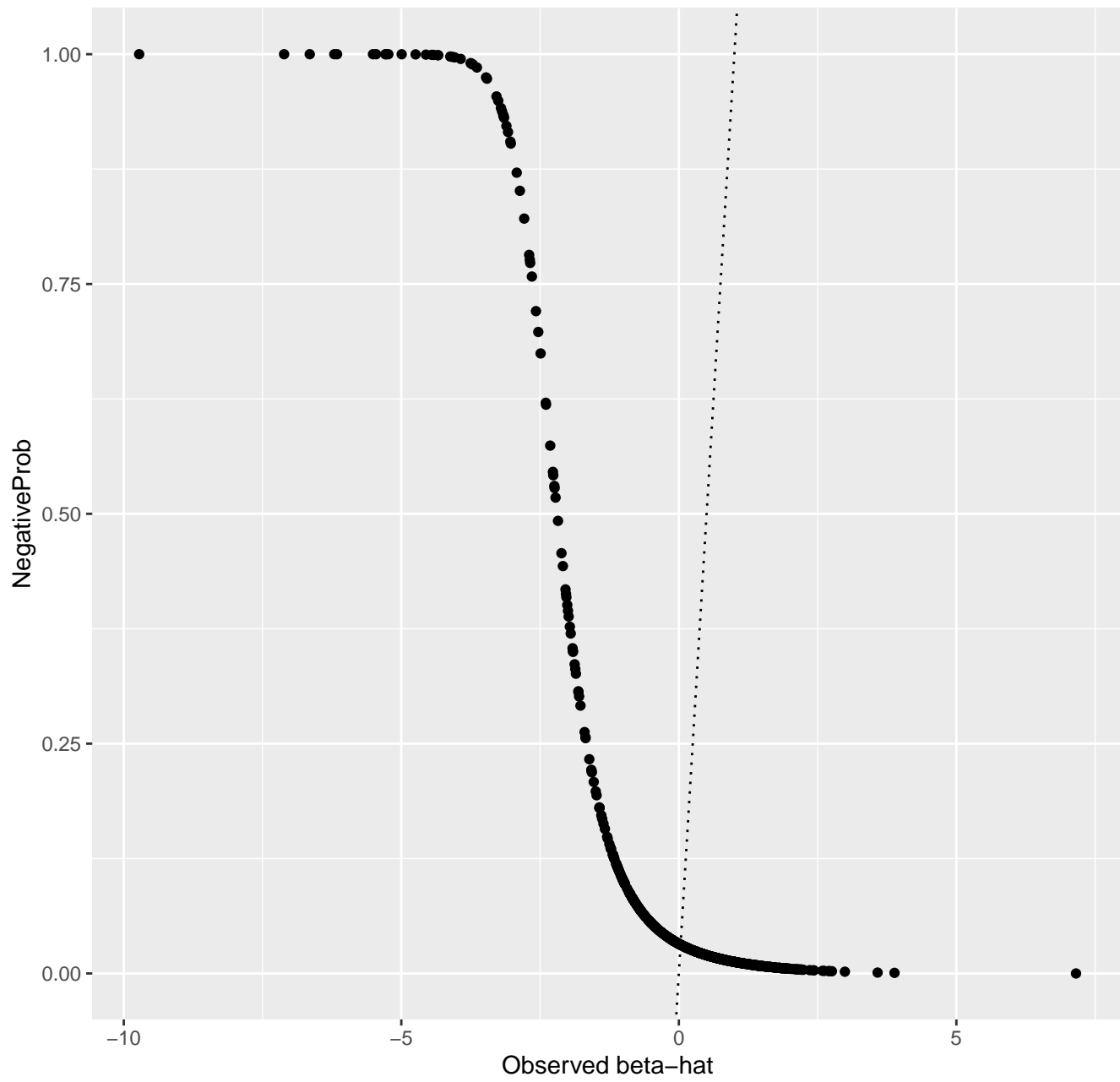
FBXO42_(54455) negative



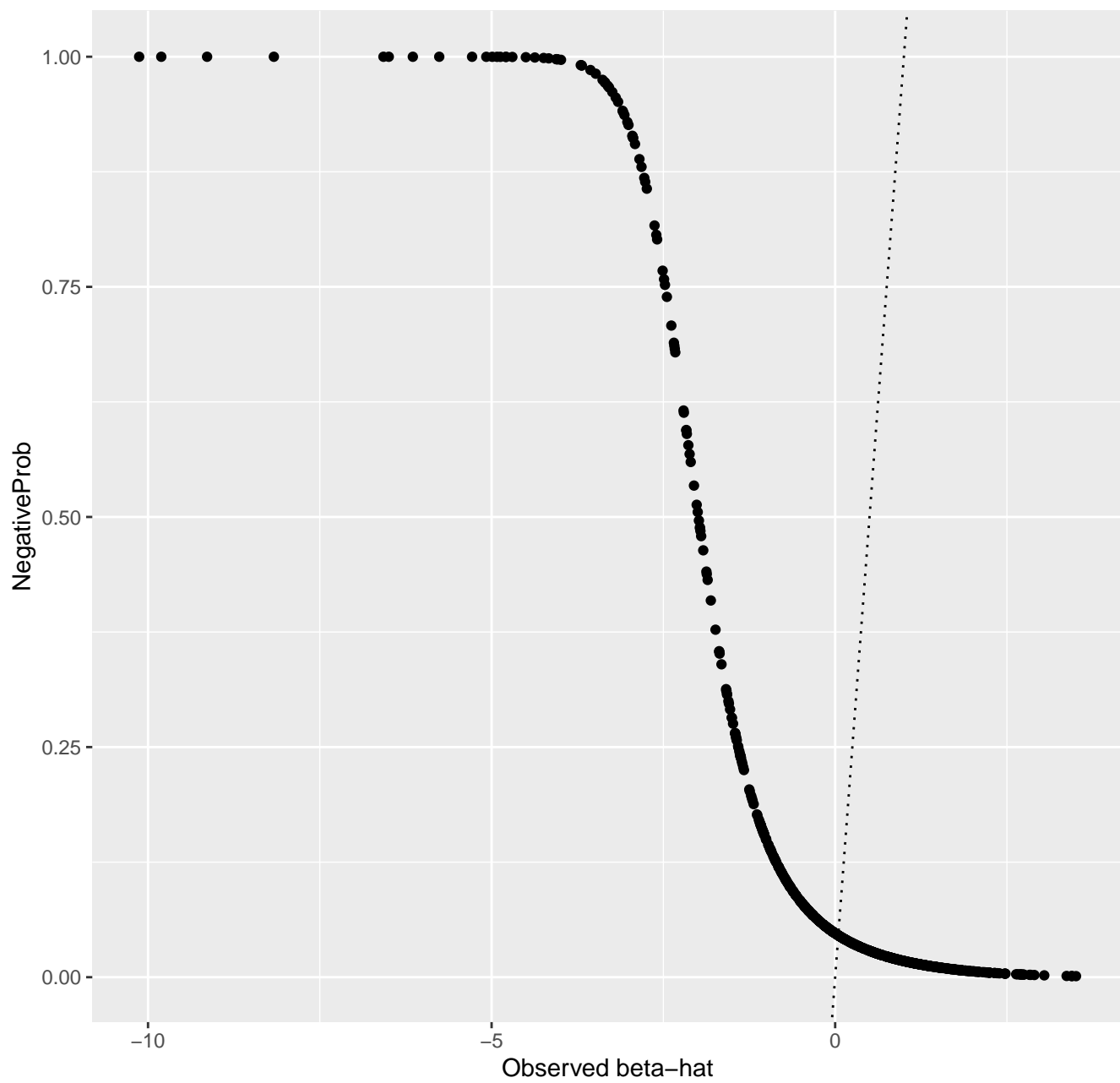
WRN_(7486) negative



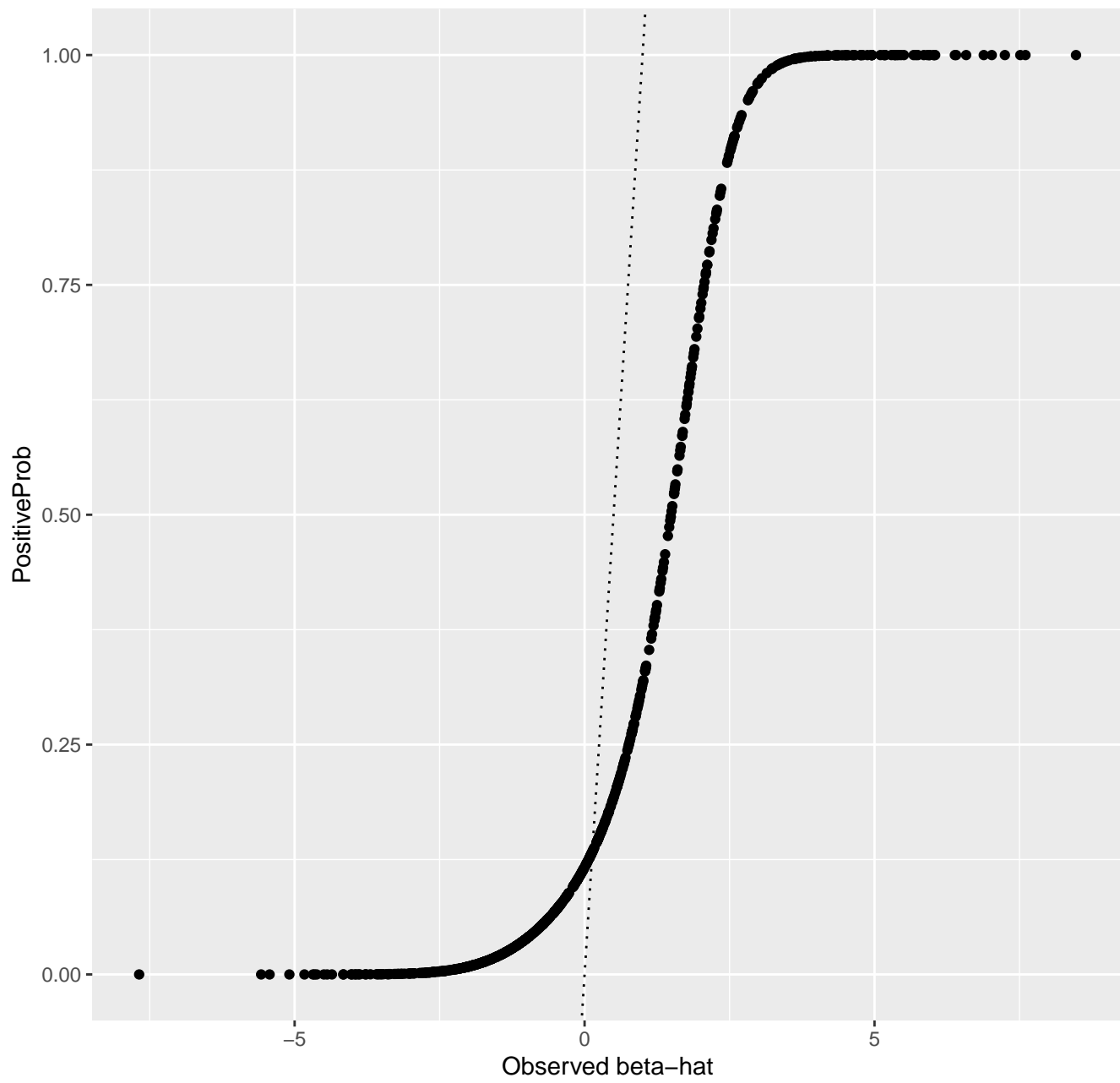
ACSL4_(2182) negative



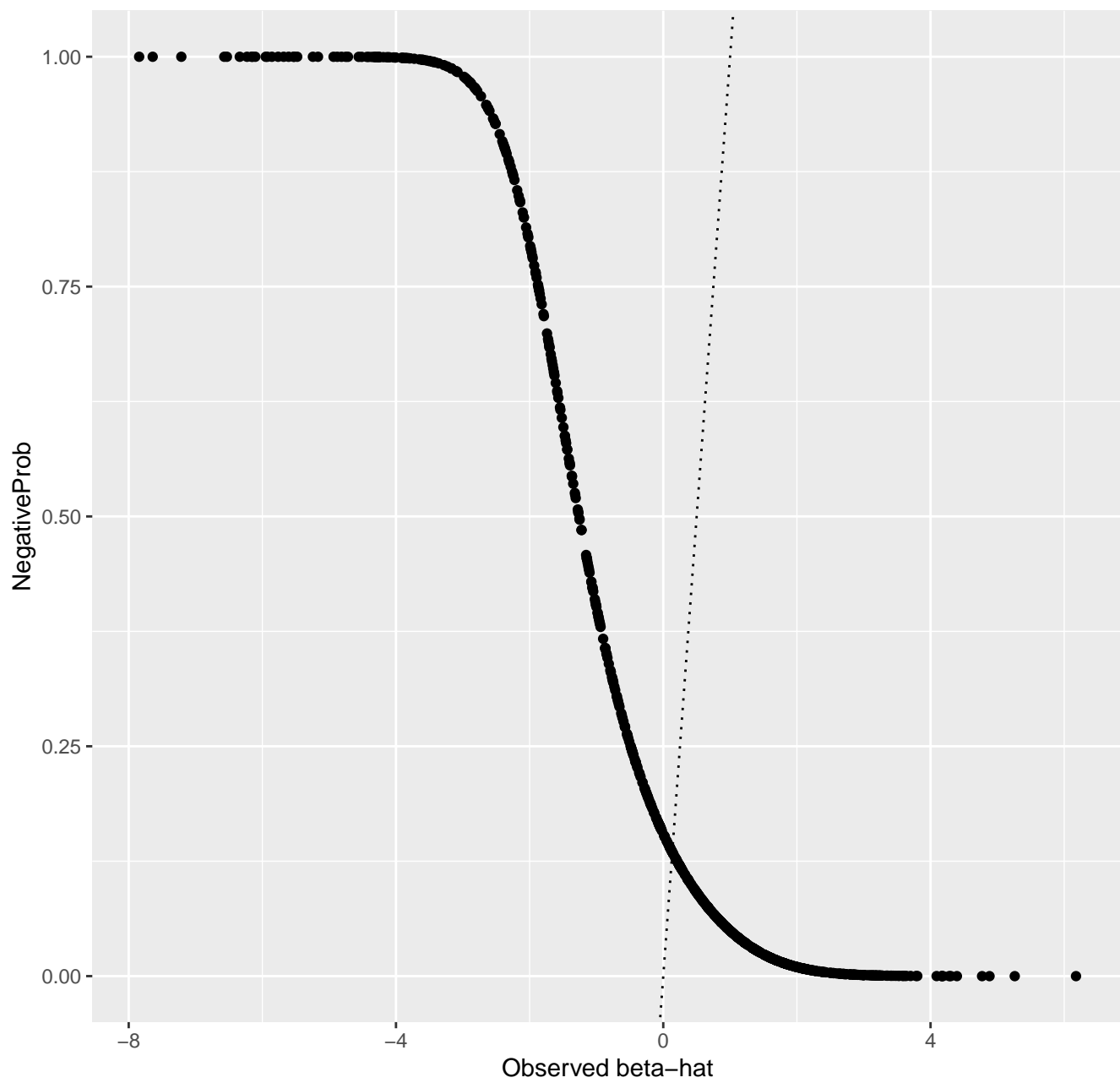
LMNA_(4000) negative



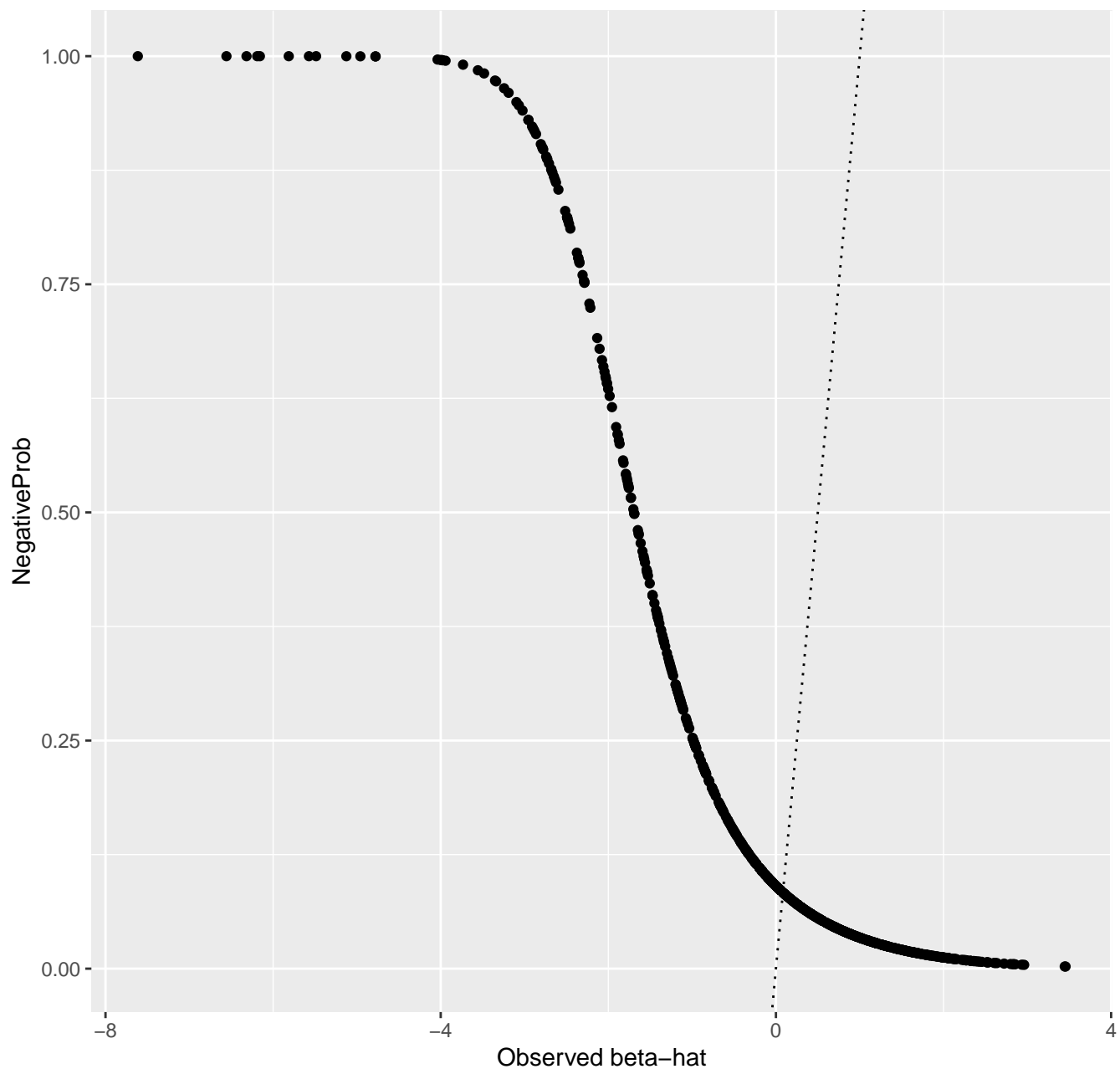
TUBB_(203068) positive



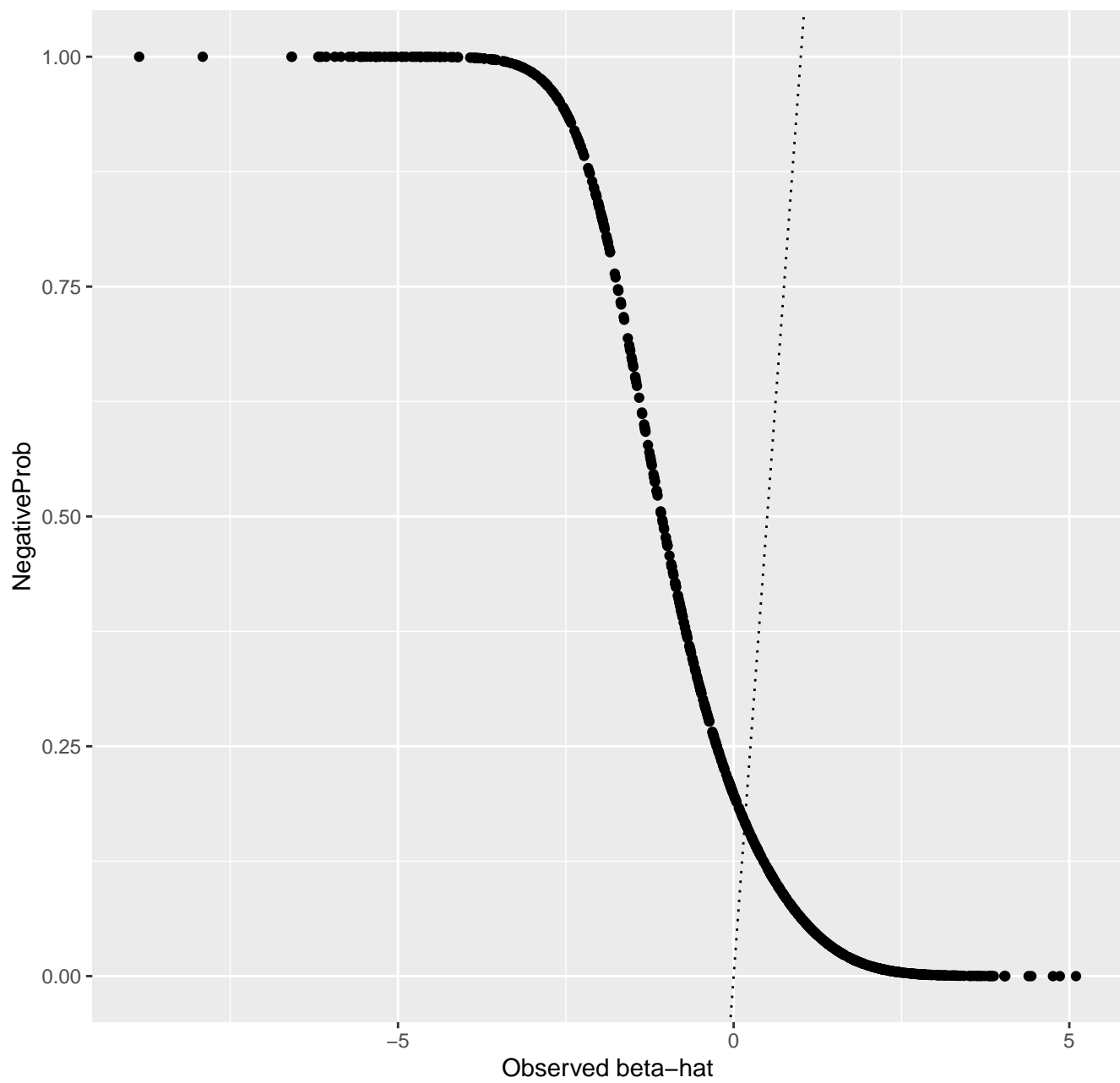
WWTR1_(25937) negative



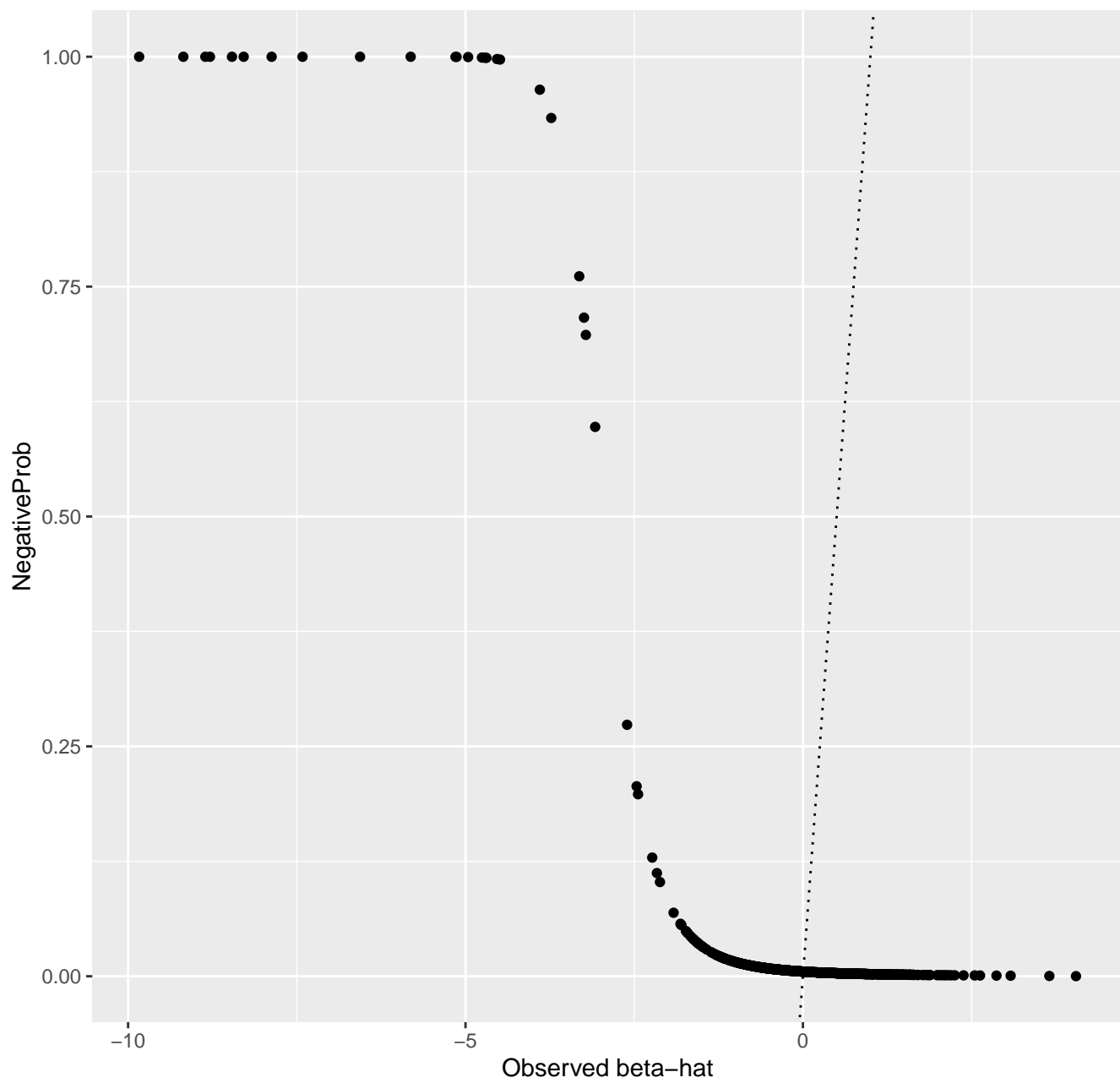
DPM2_(8818) negative



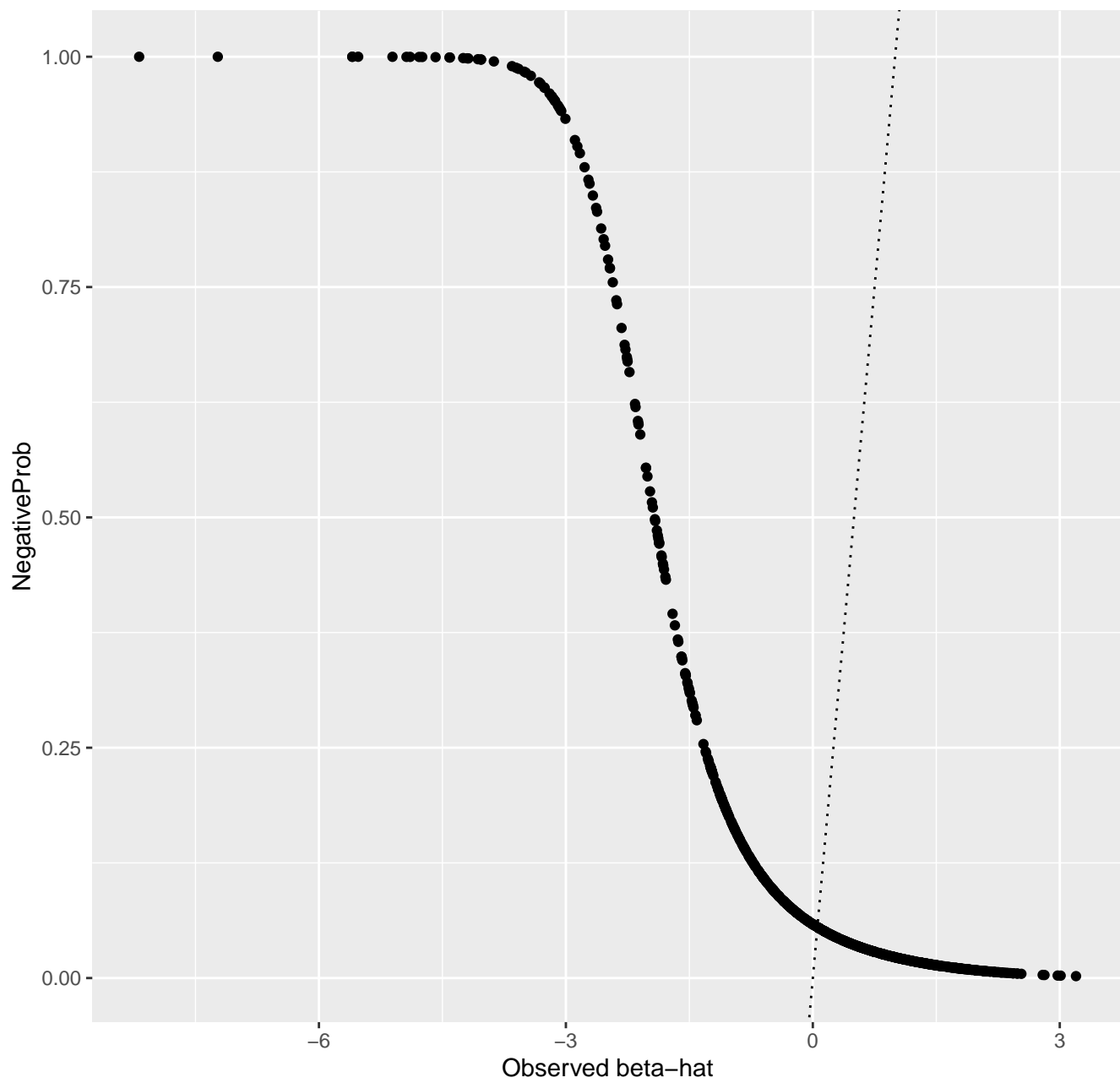
EAF1_(85403) negative



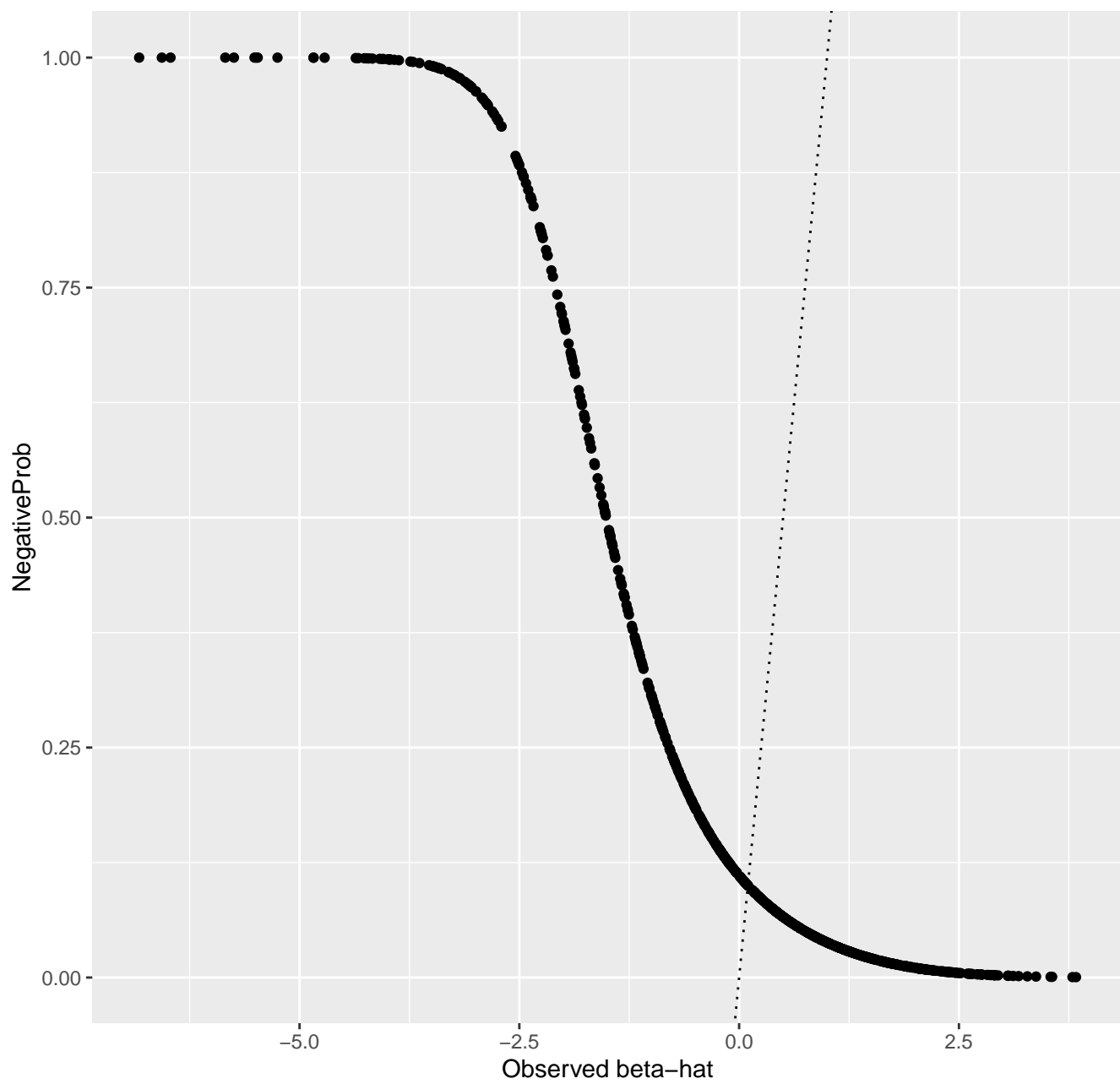
EBF1_(1879) negative



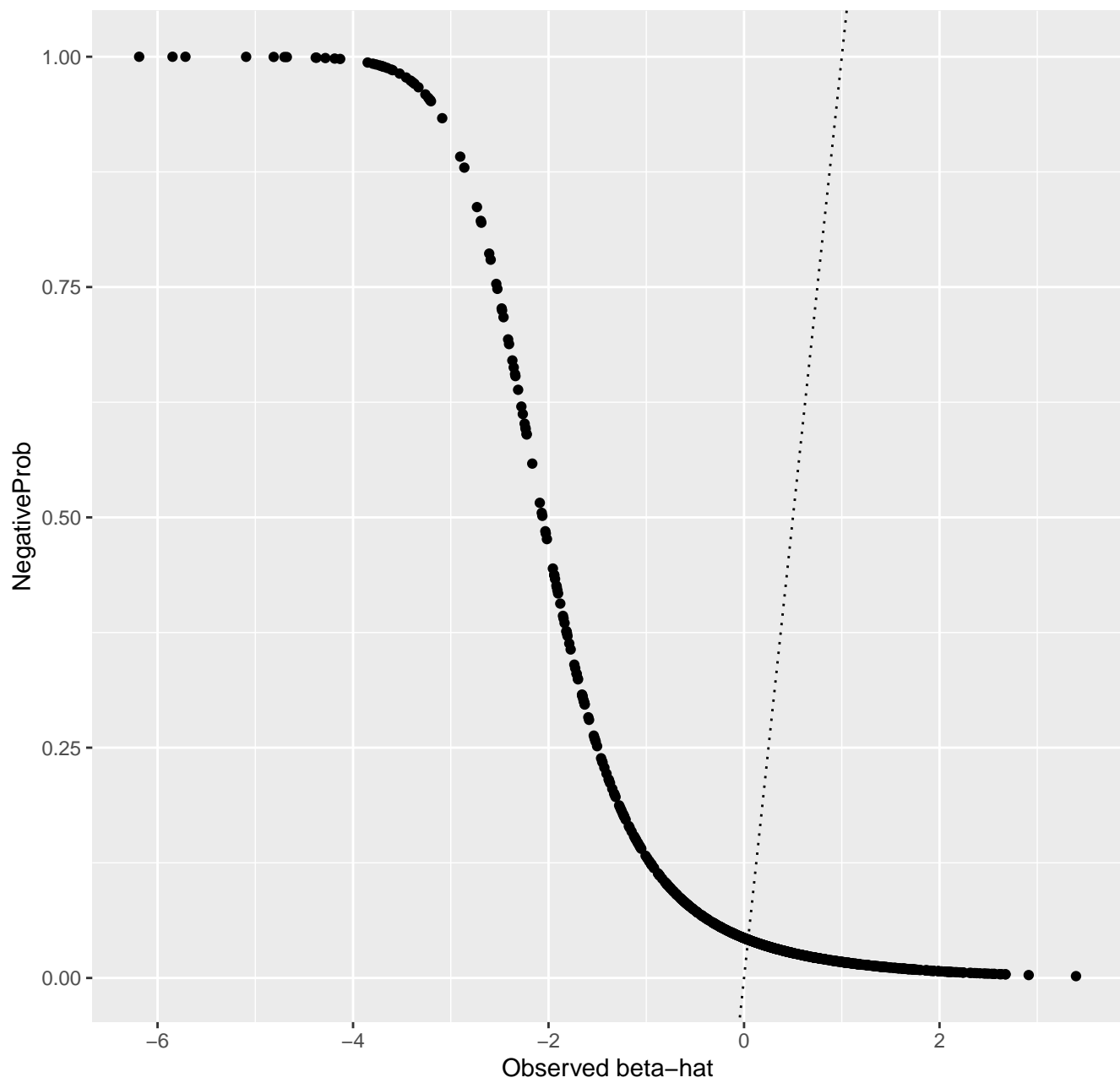
NME6_(10201) negative



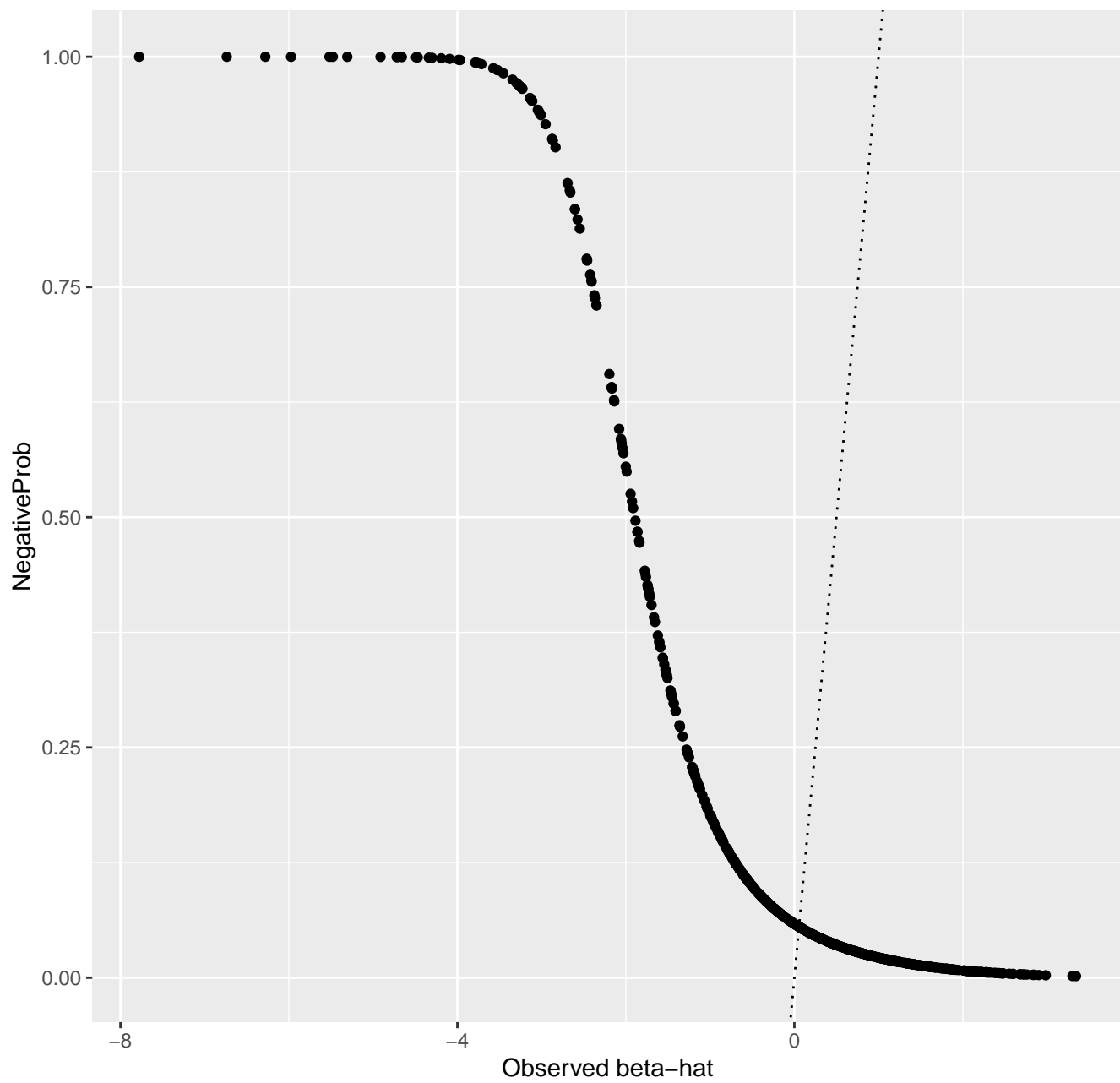
TRIP13_(9319) negative



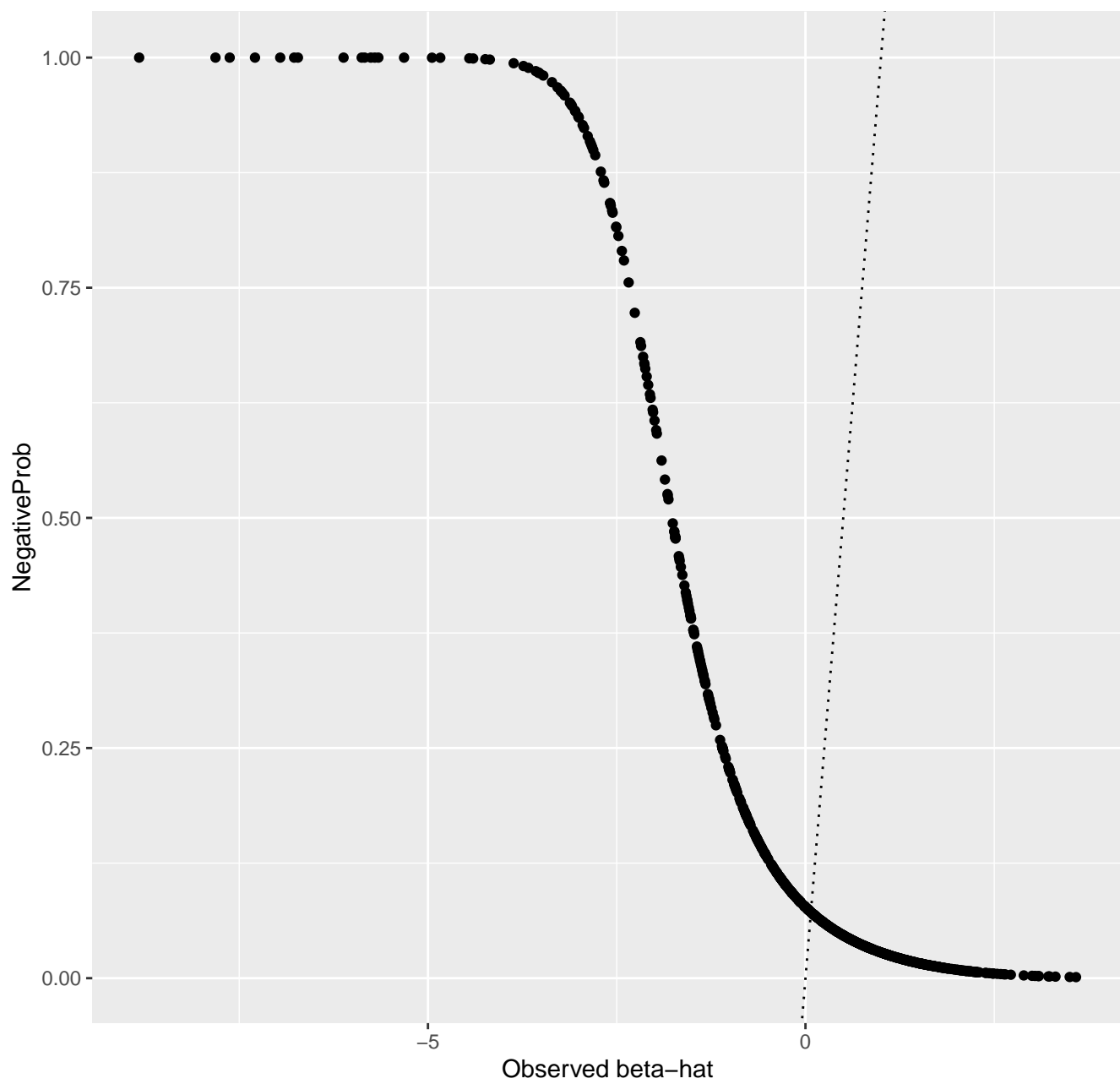
FANCI_(55215) negative



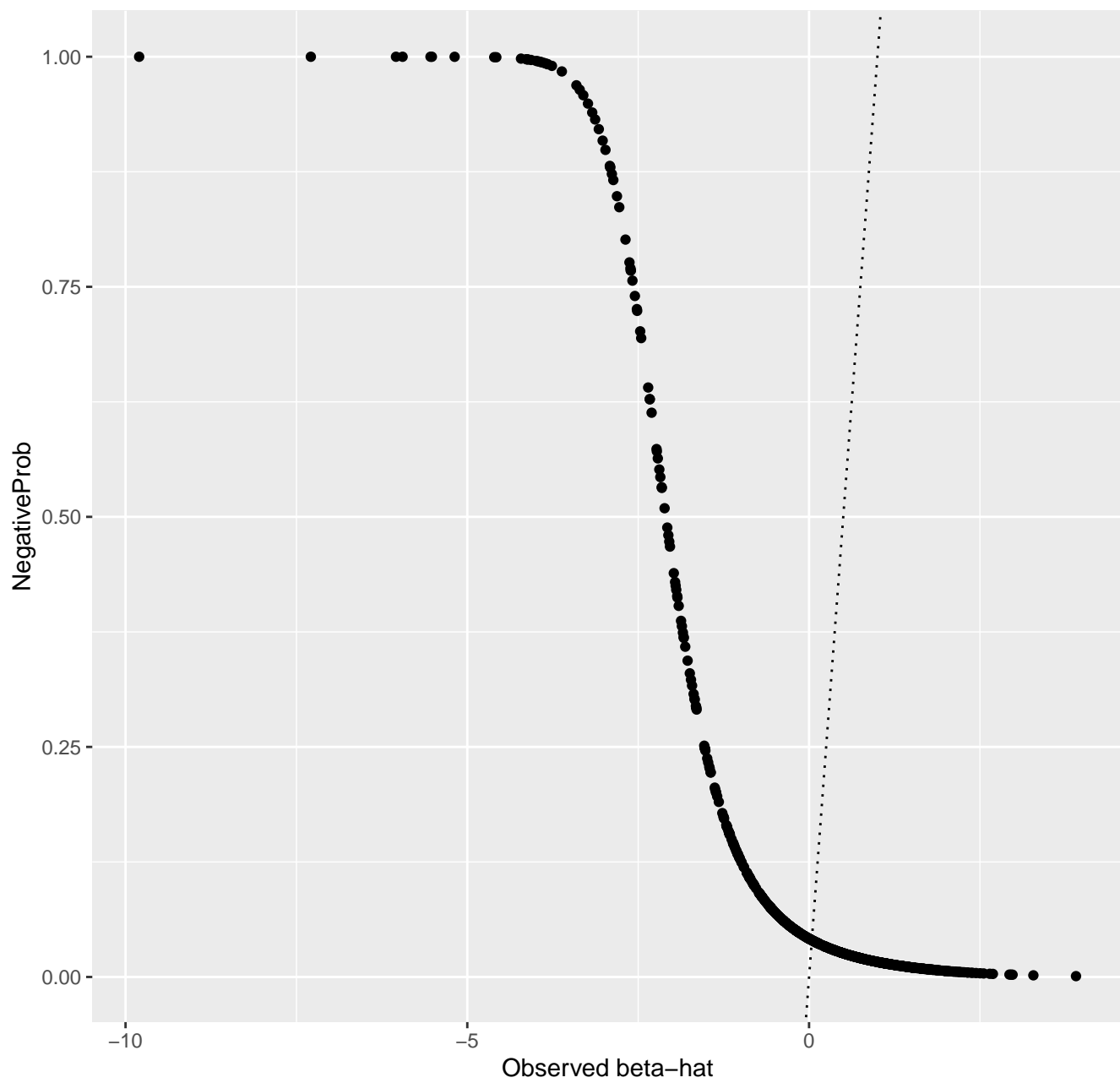
SURF4_(6836) negative



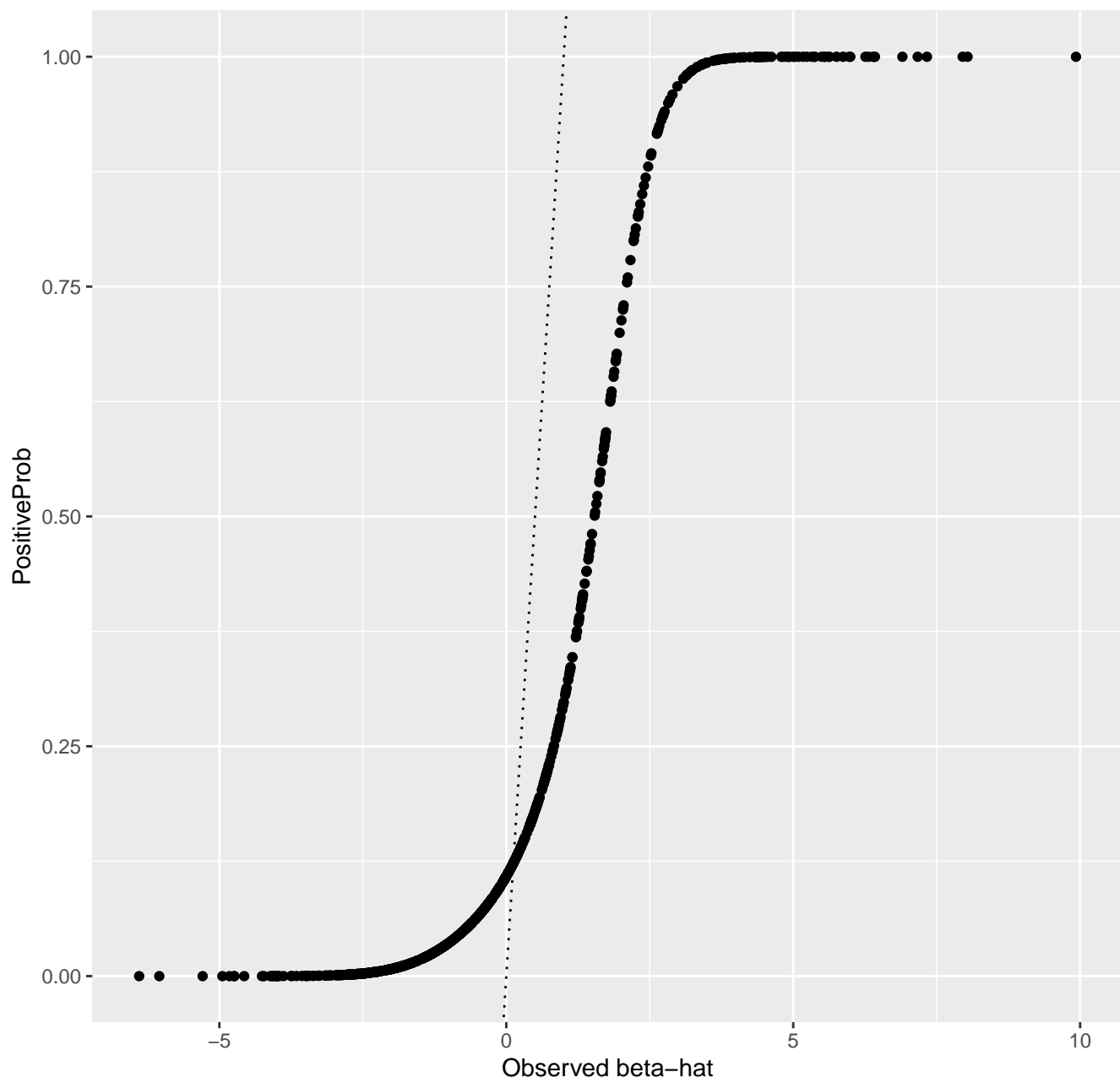
SOS1_(6654) negative



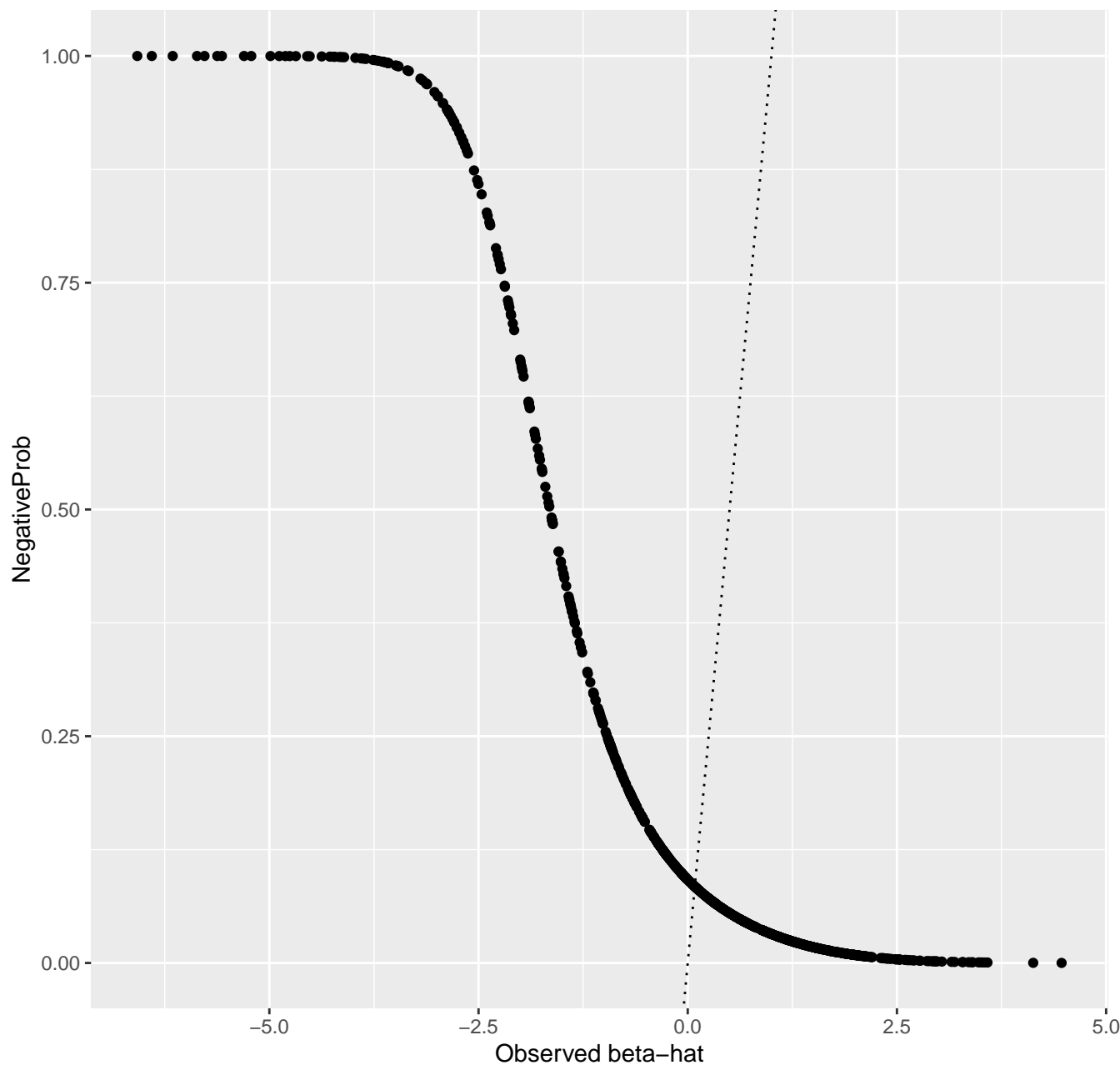
FBXO7_(25793) negative



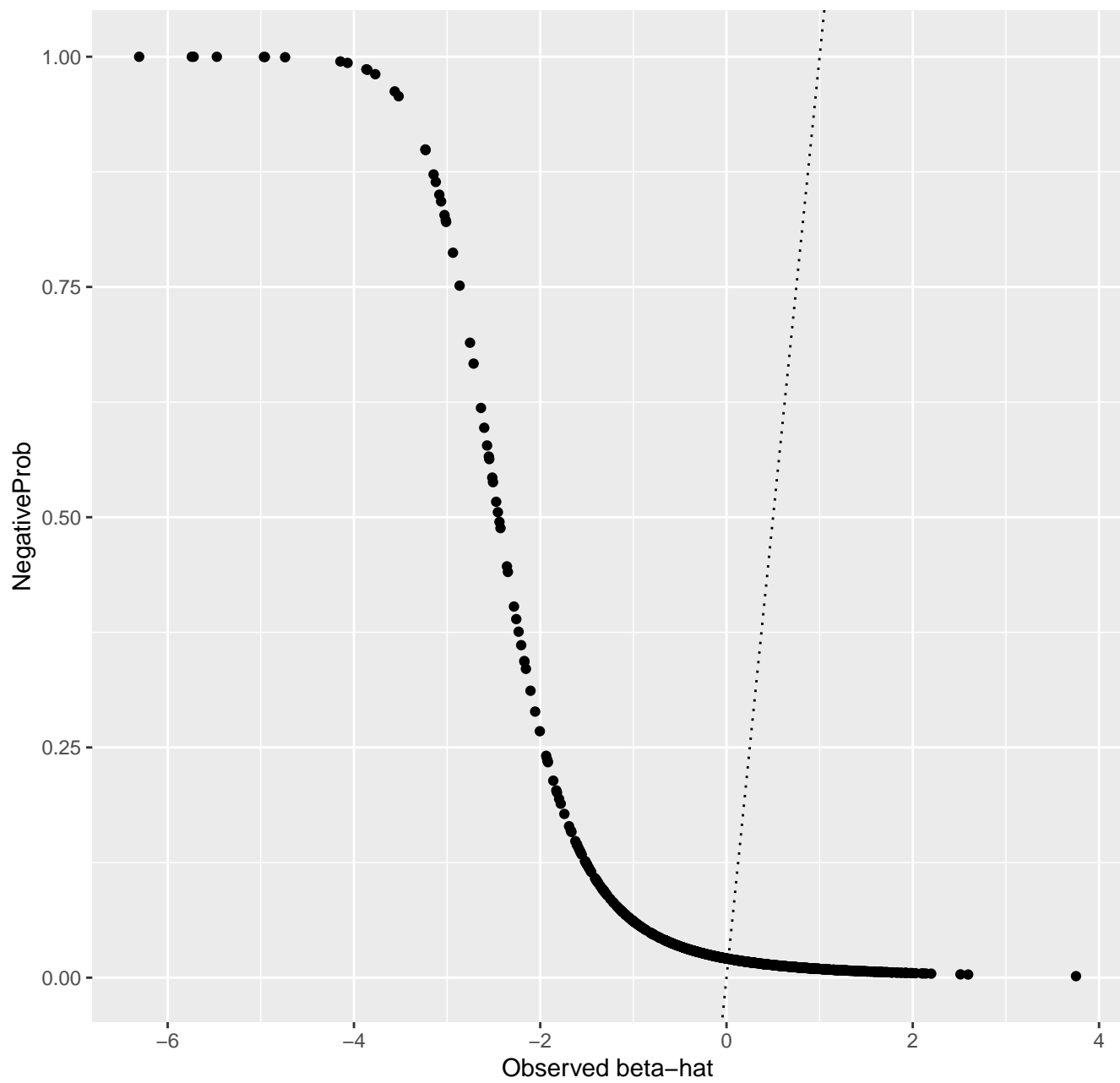
NCAPD3_(23310) positive



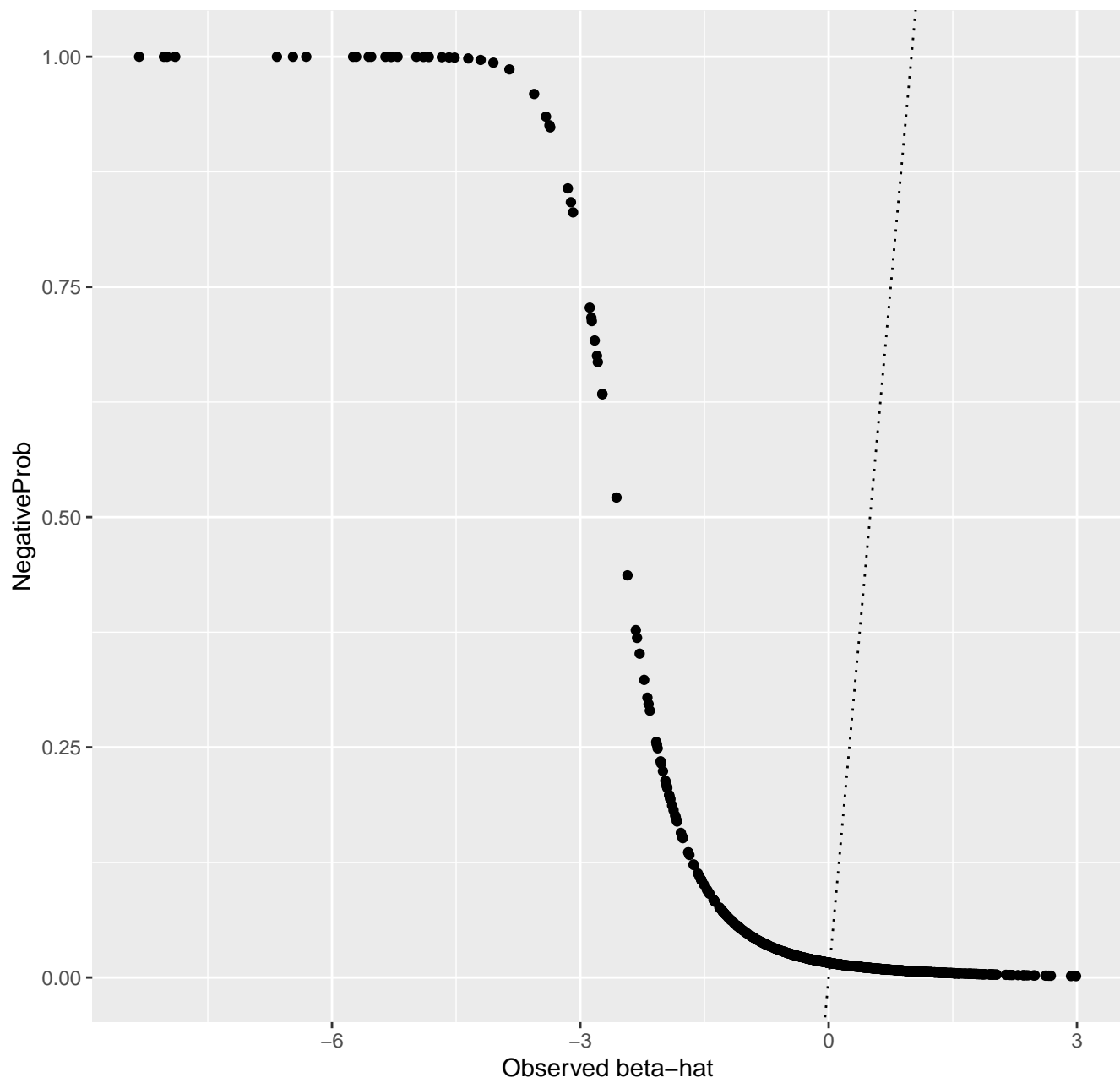
DDI2_(84301) negative



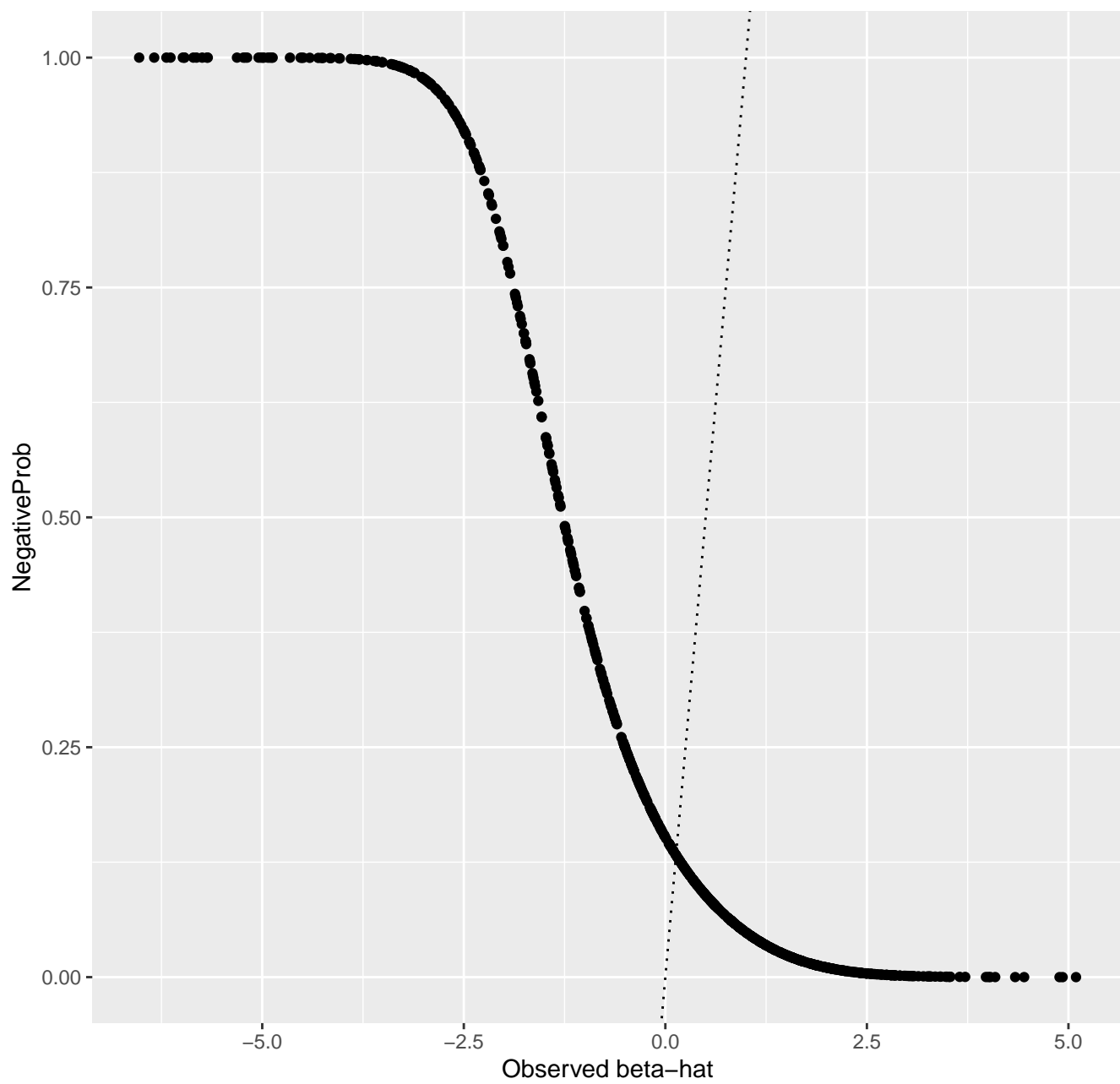
PIK3CB_(5291) negative



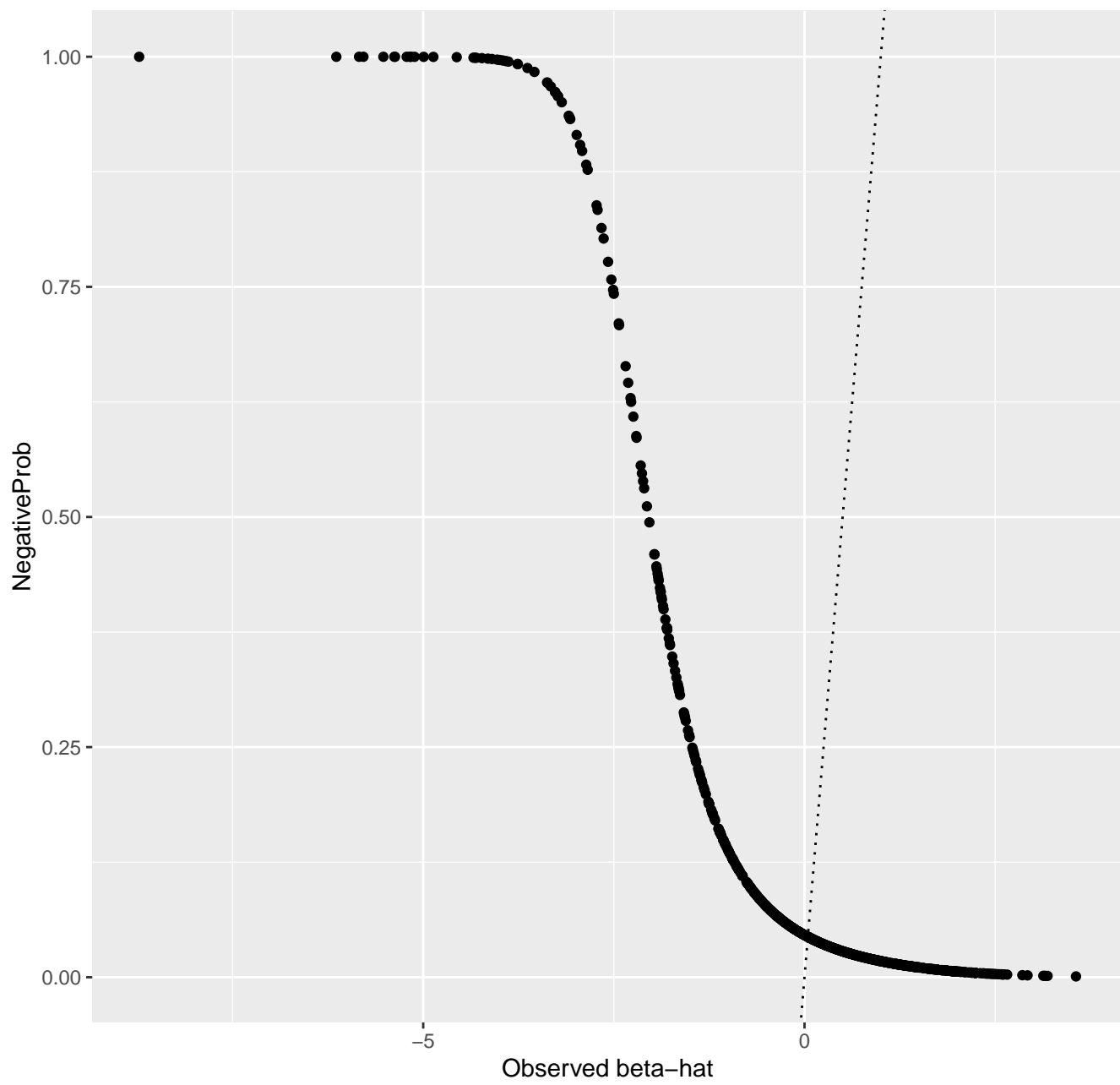
ACOX1_(51) negative



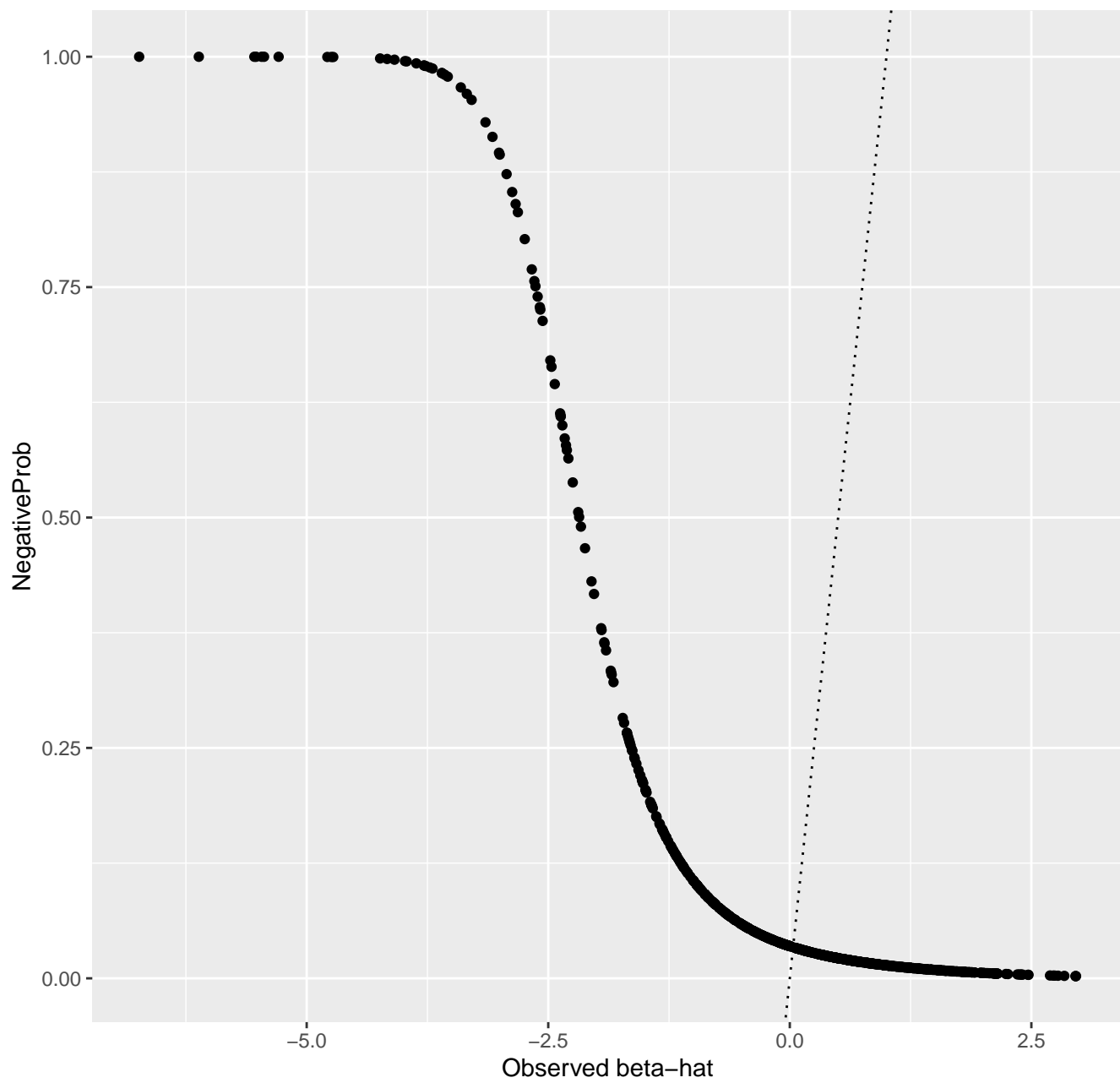
KDSR_(2531) negative



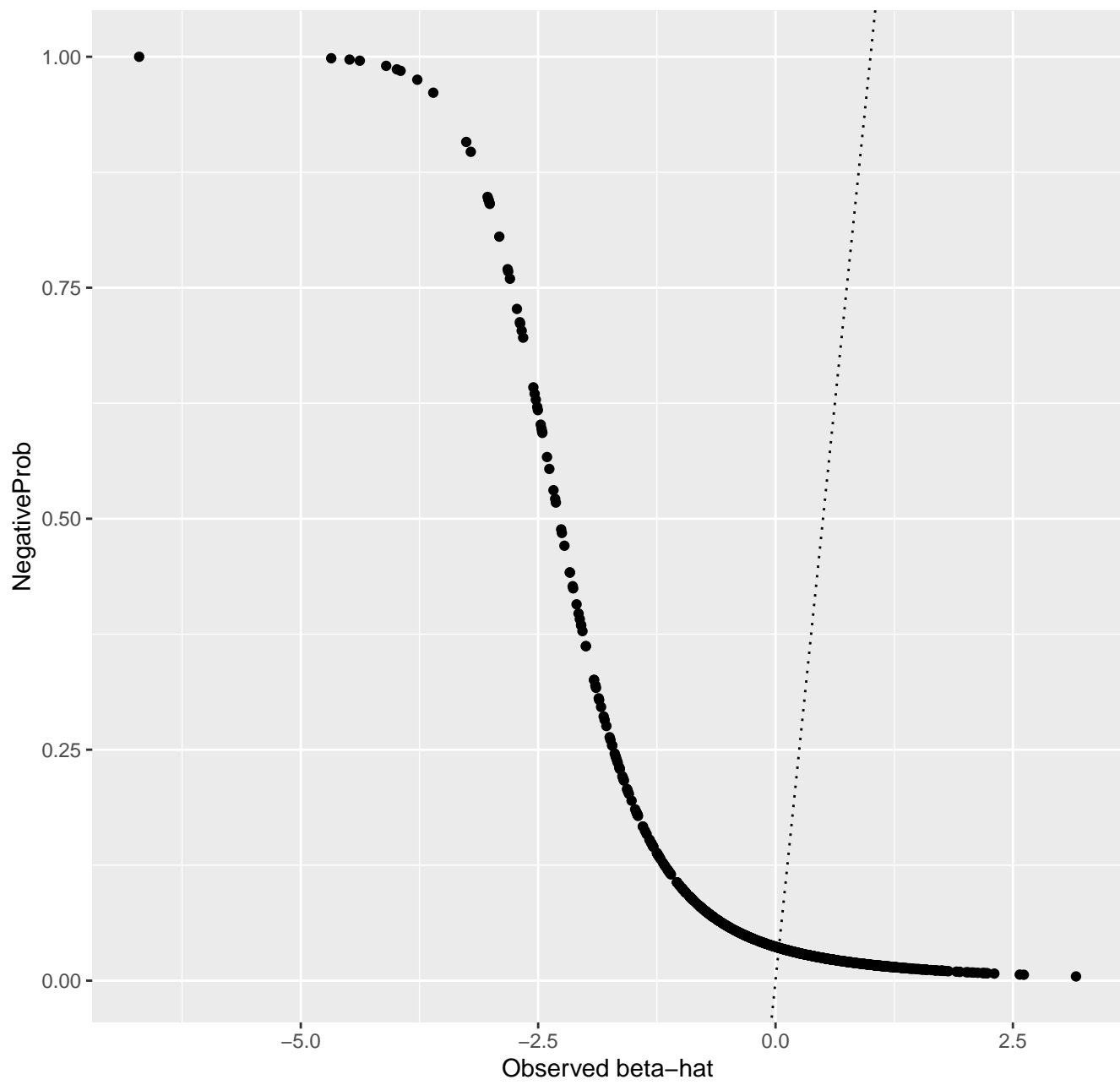
MTCH2_(23788) negative



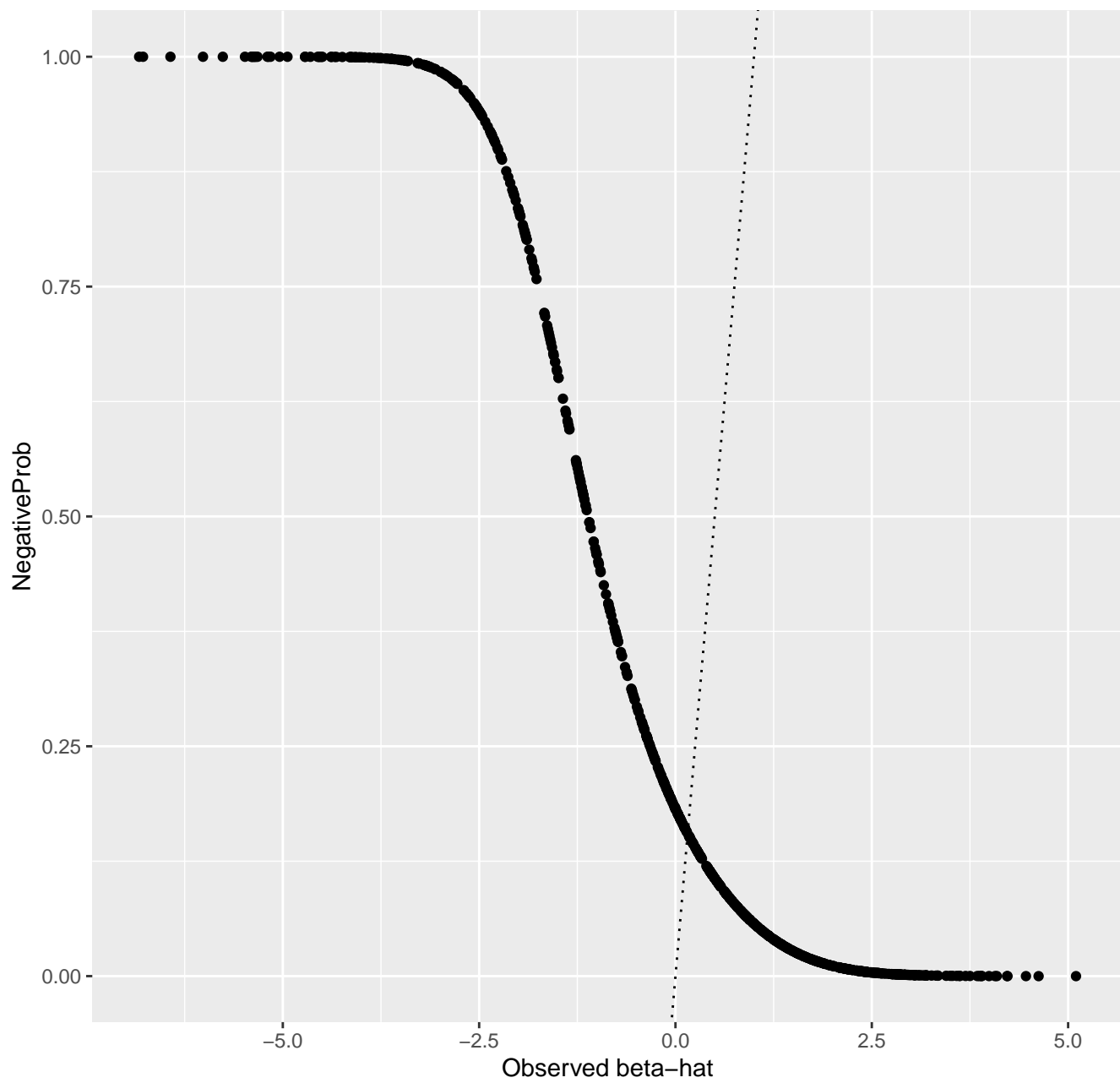
MPI_(4351) negative



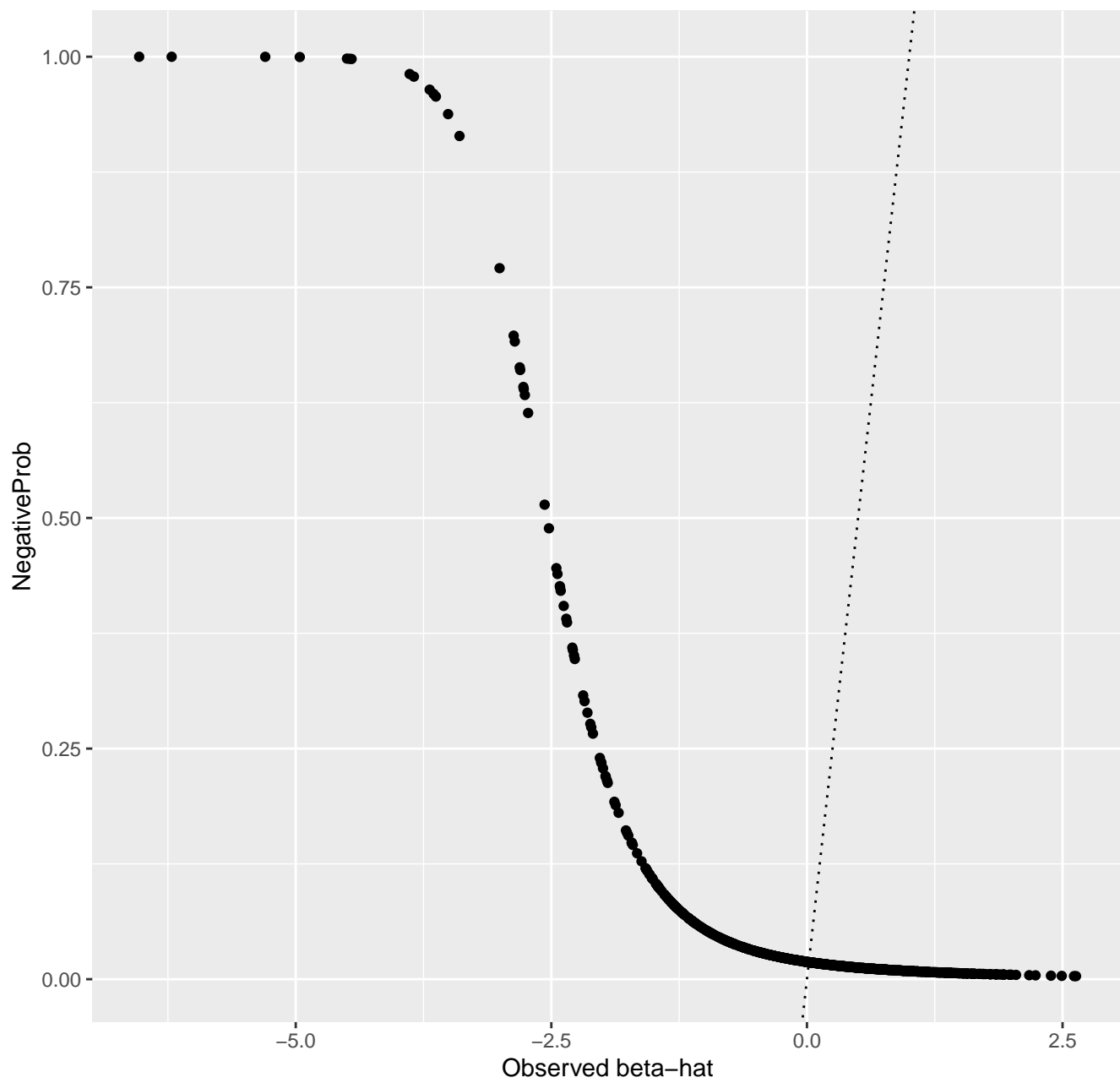
OSBPL9_(114883) negative



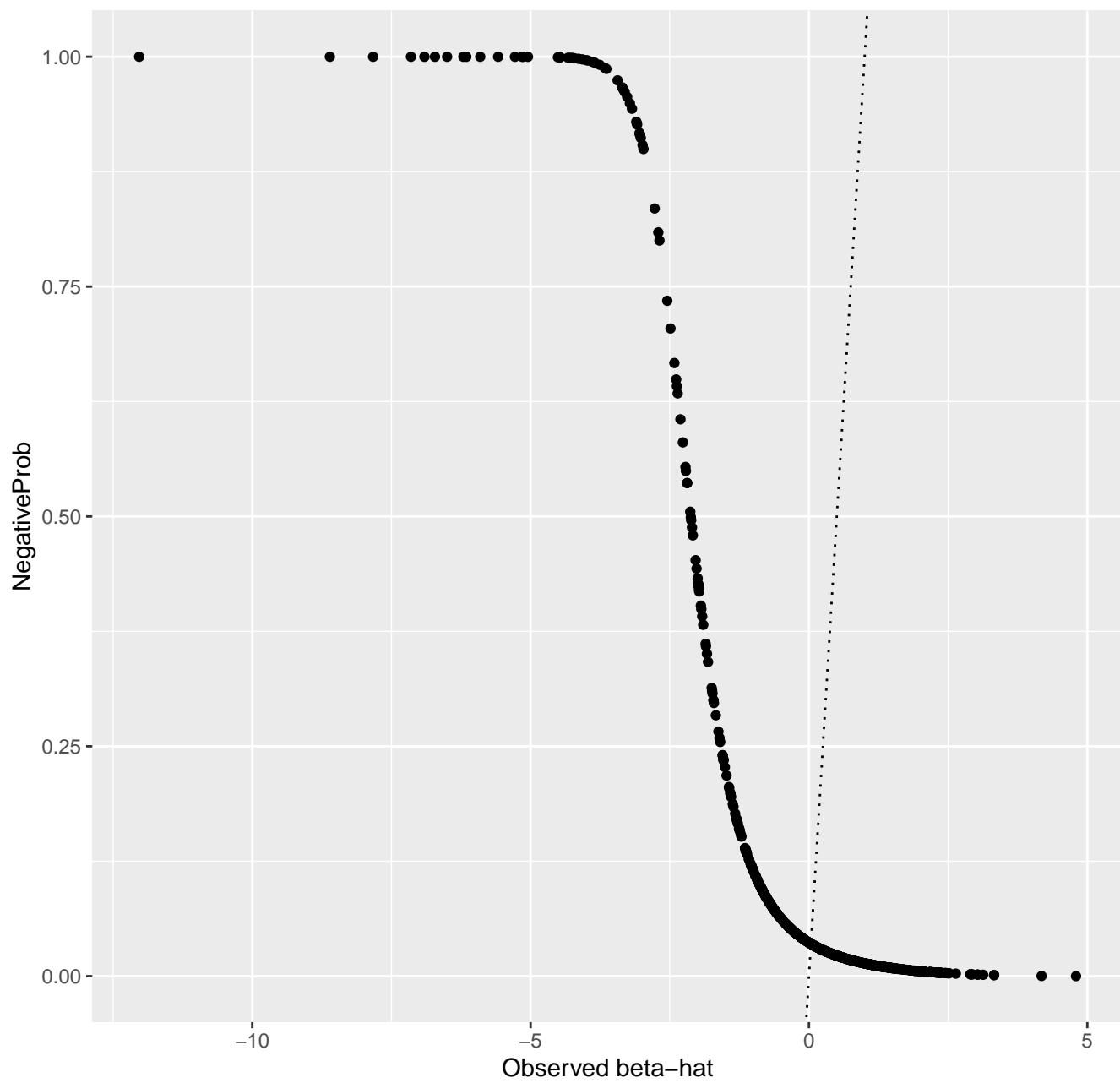
ARF4_(378) negative



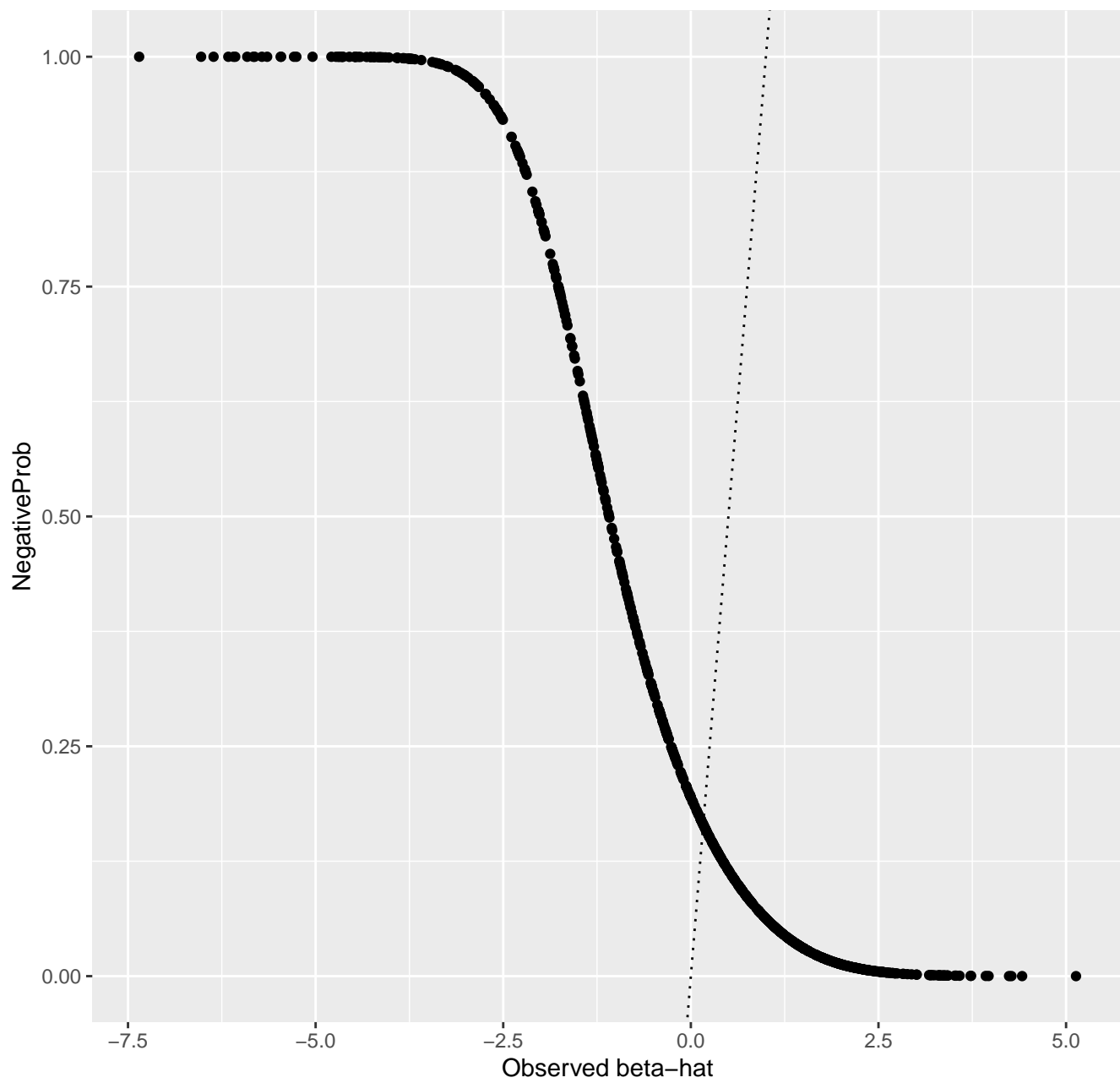
DYNC1LI1_(51143) negative



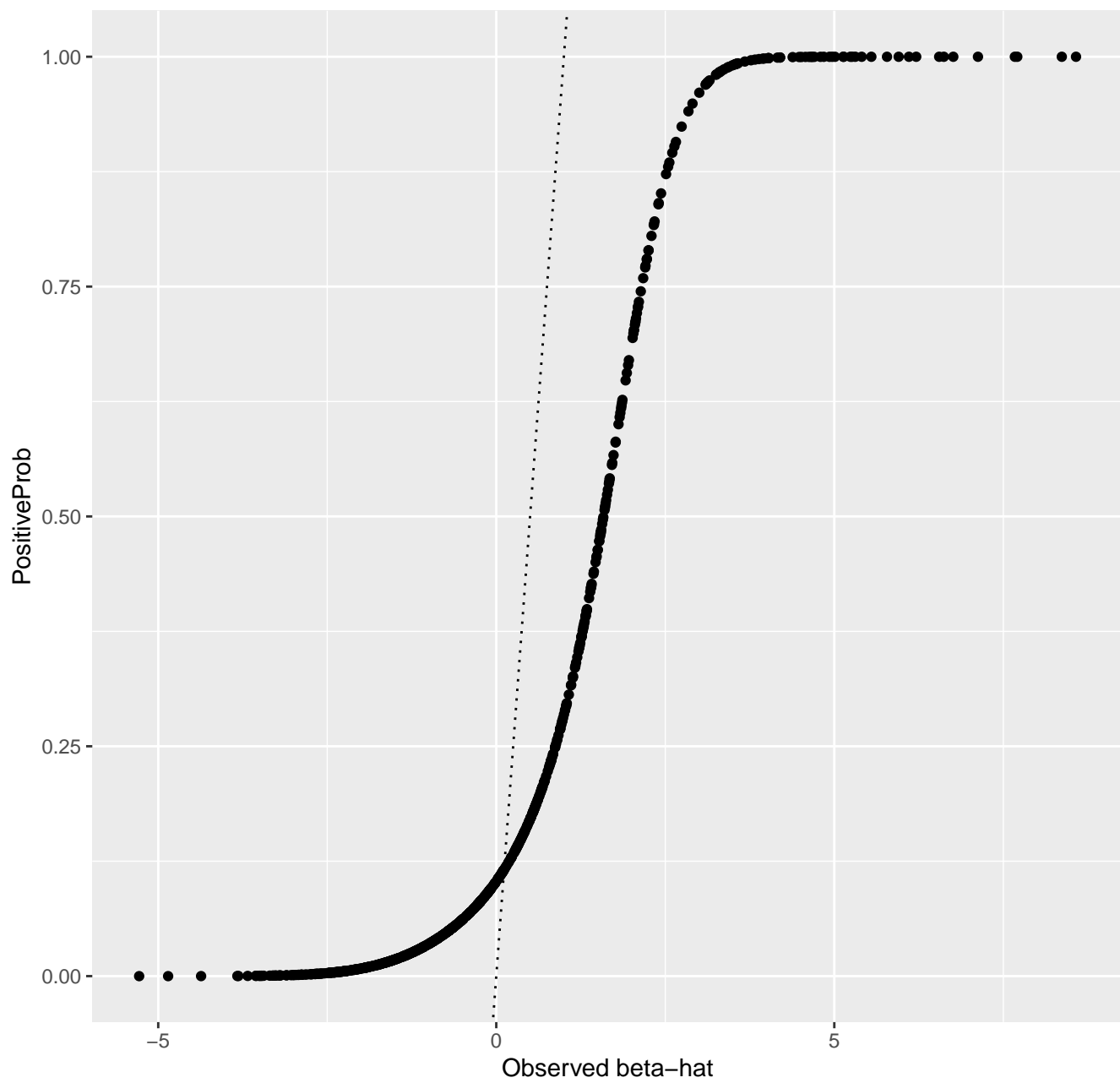
KIDINS220_(57498) negative



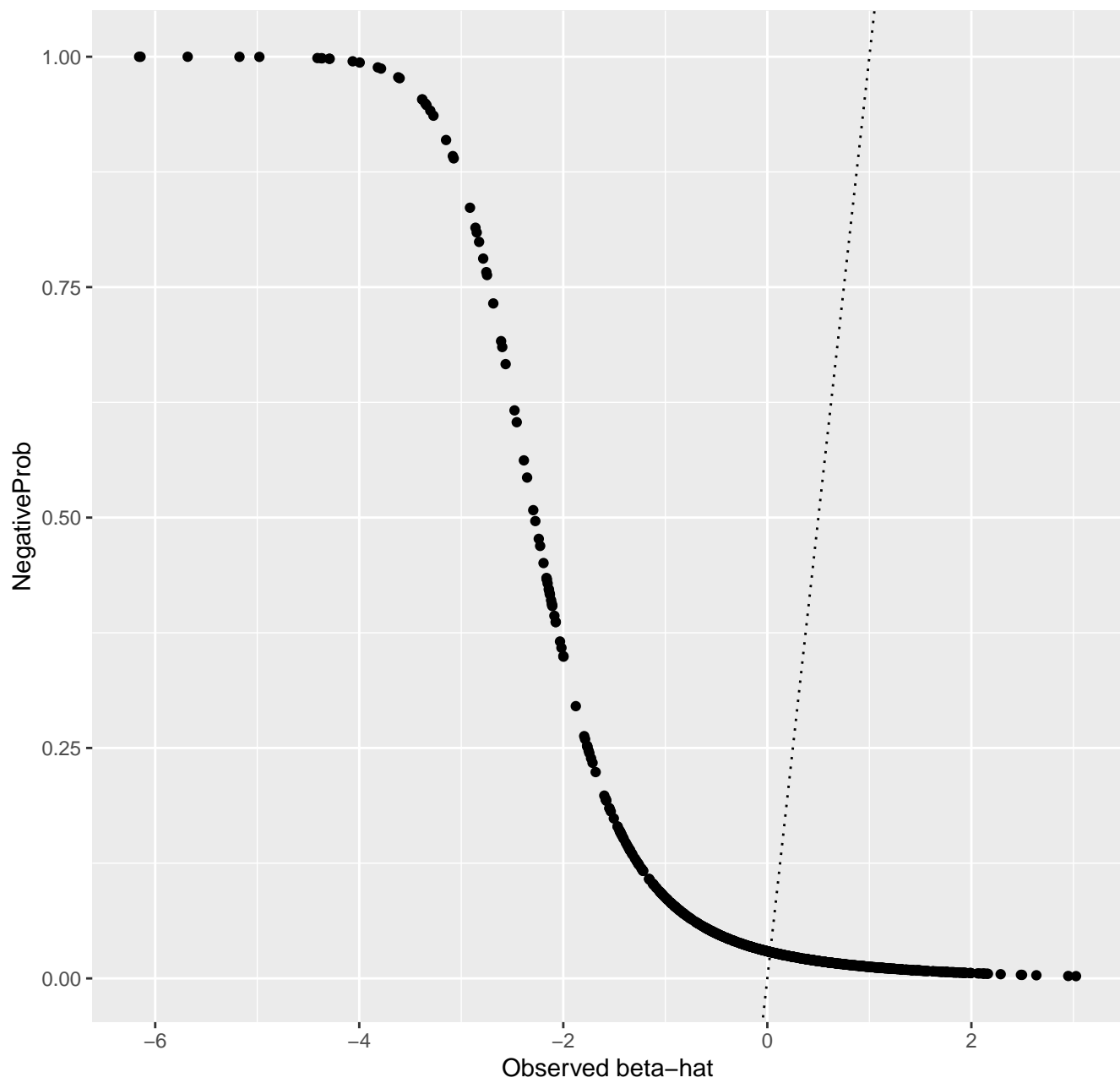
NDE1_(54820) negative



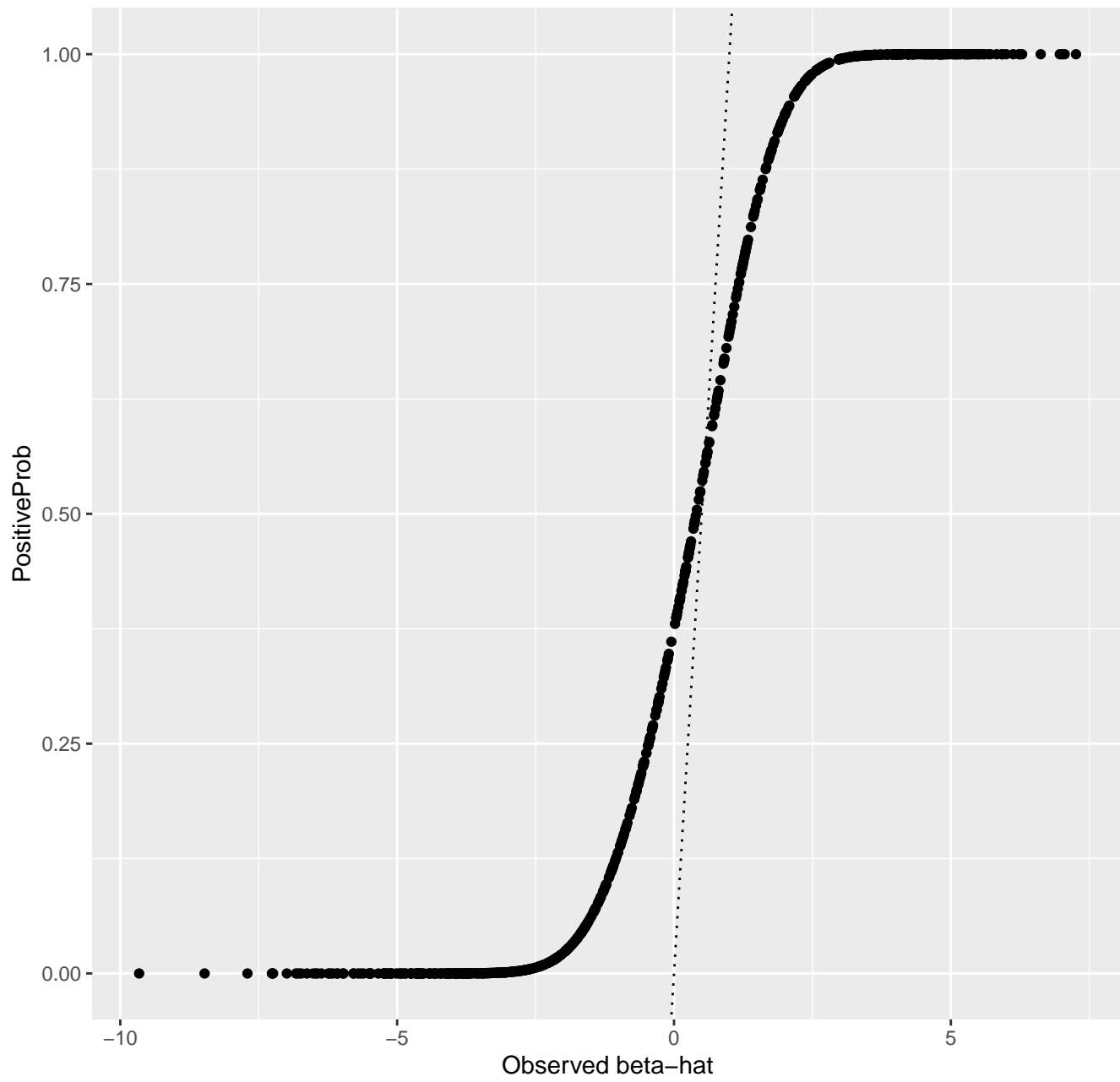
TAOK1_(57551) positive



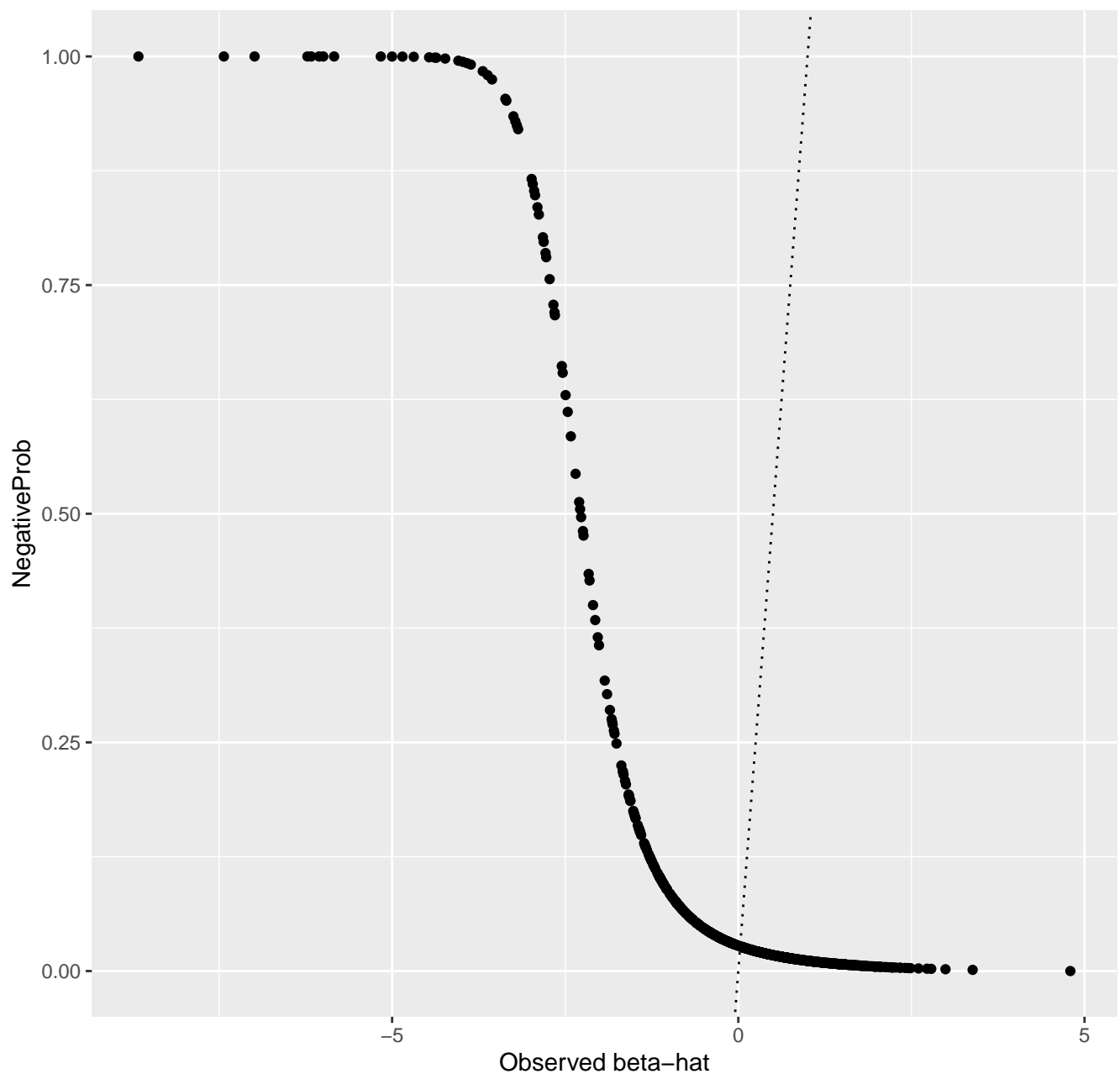
NFE2L1_(4779) negative



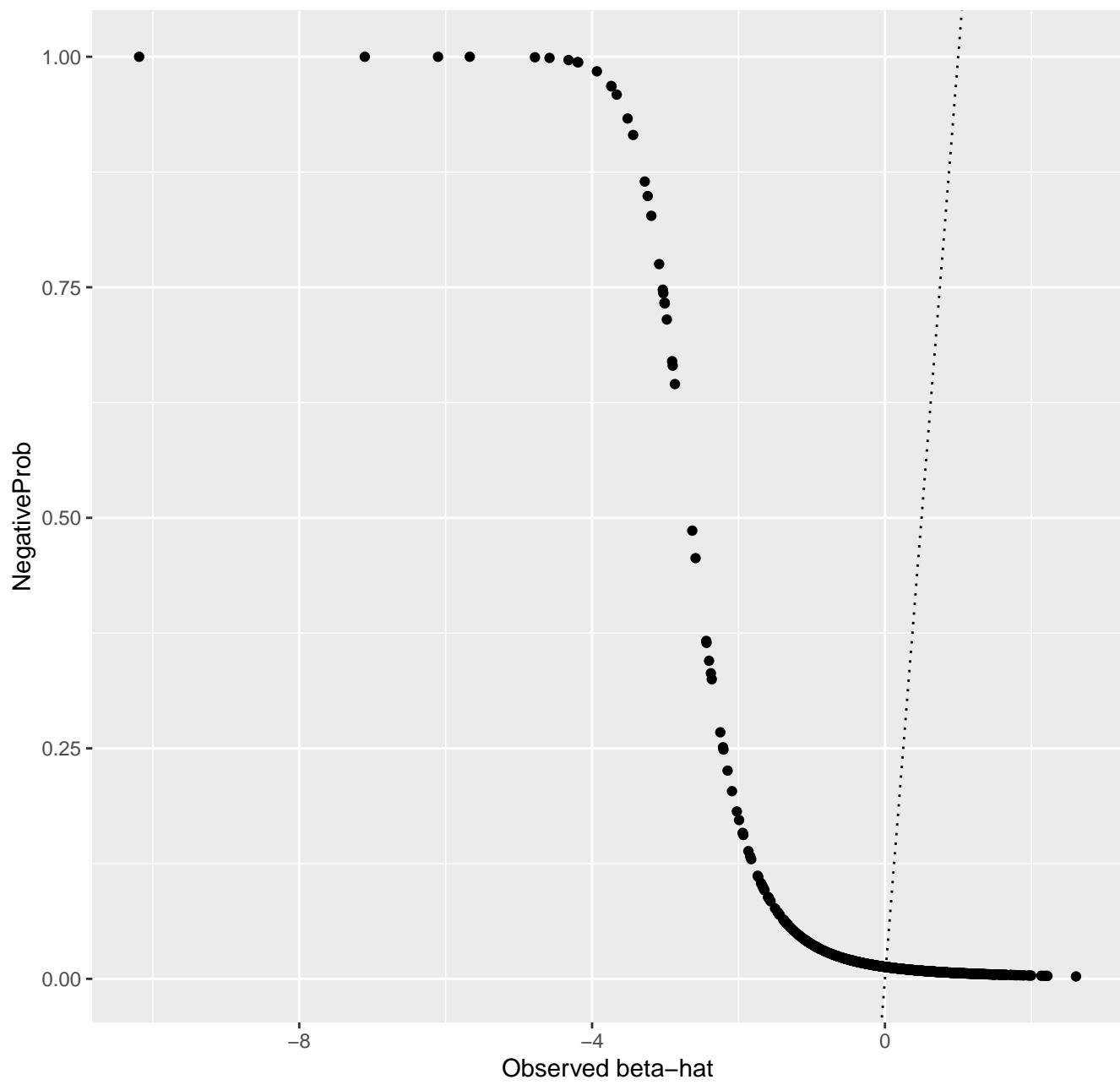
SNAP23_(8773) positive



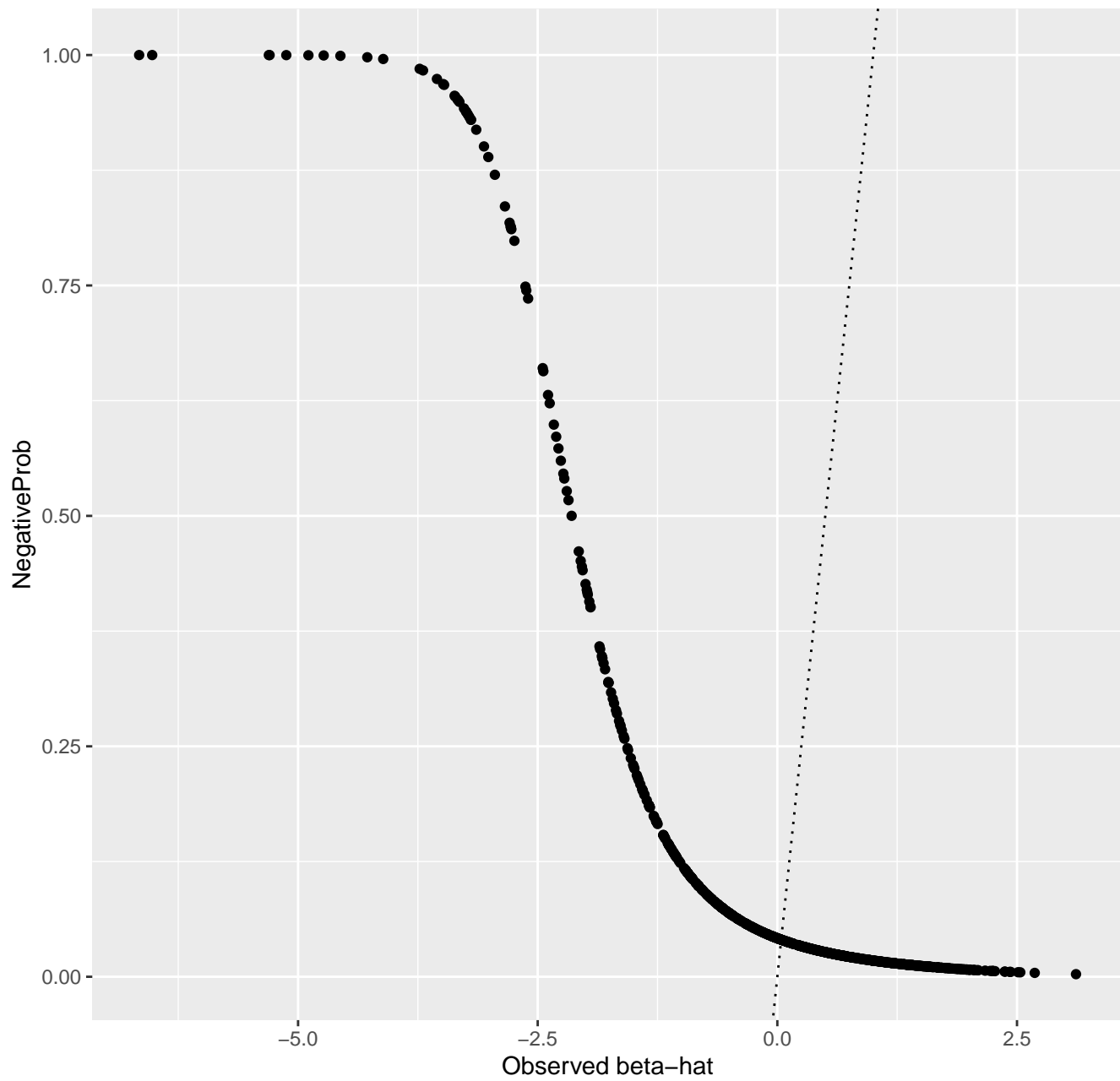
SMAD7_(4092) negative



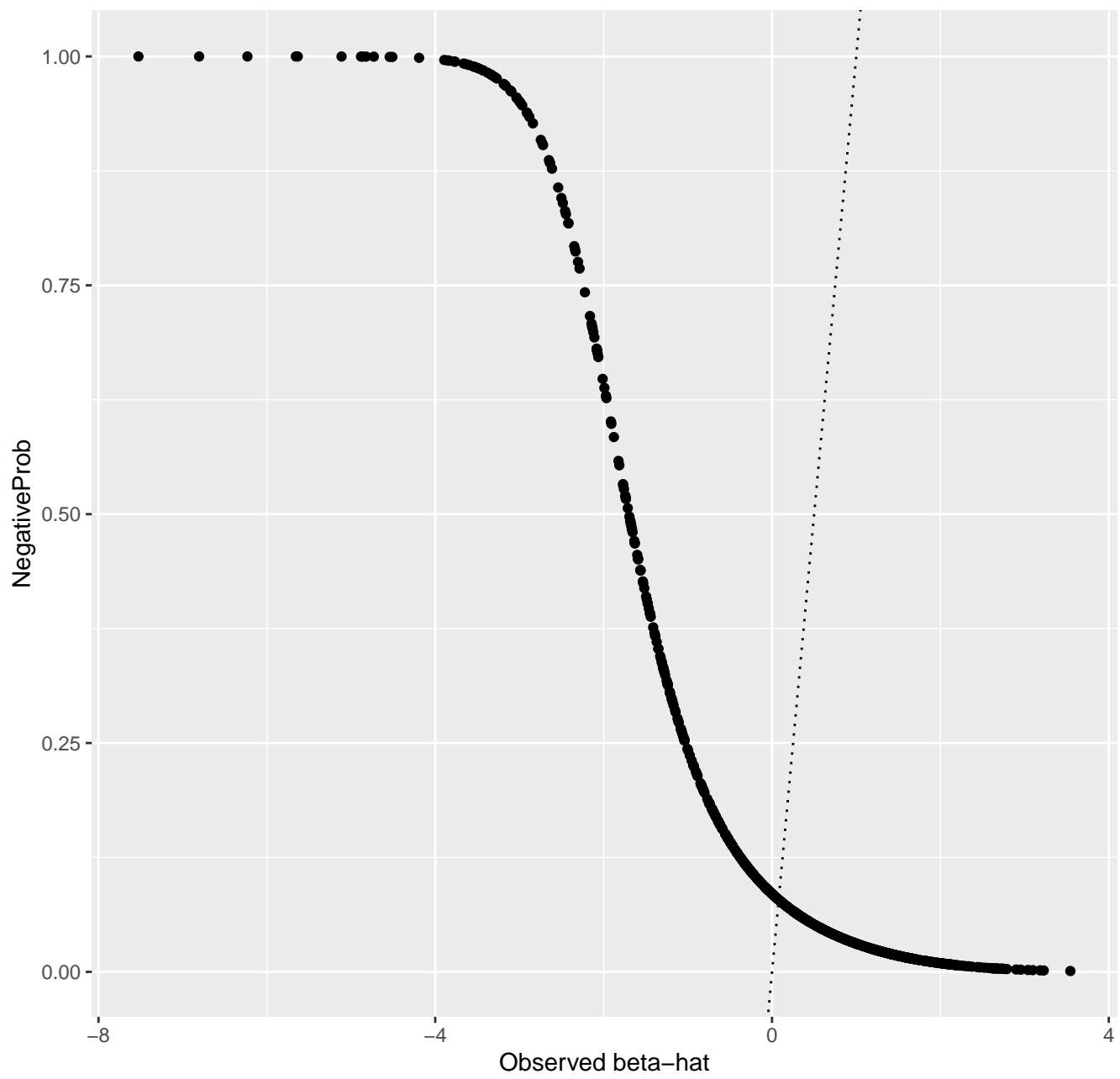
ITGB3_(3690) negative



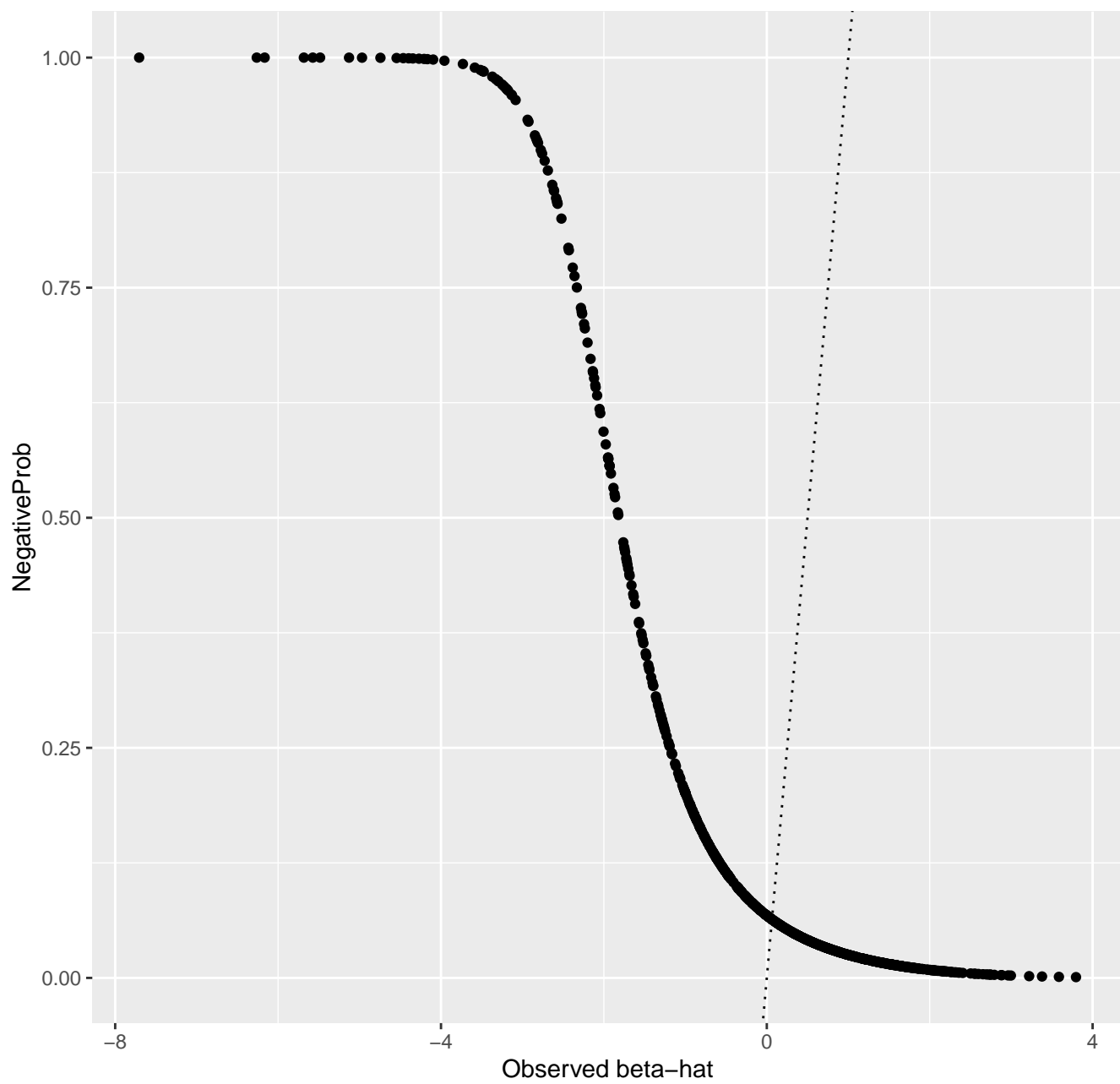
MECR_(51102) negative



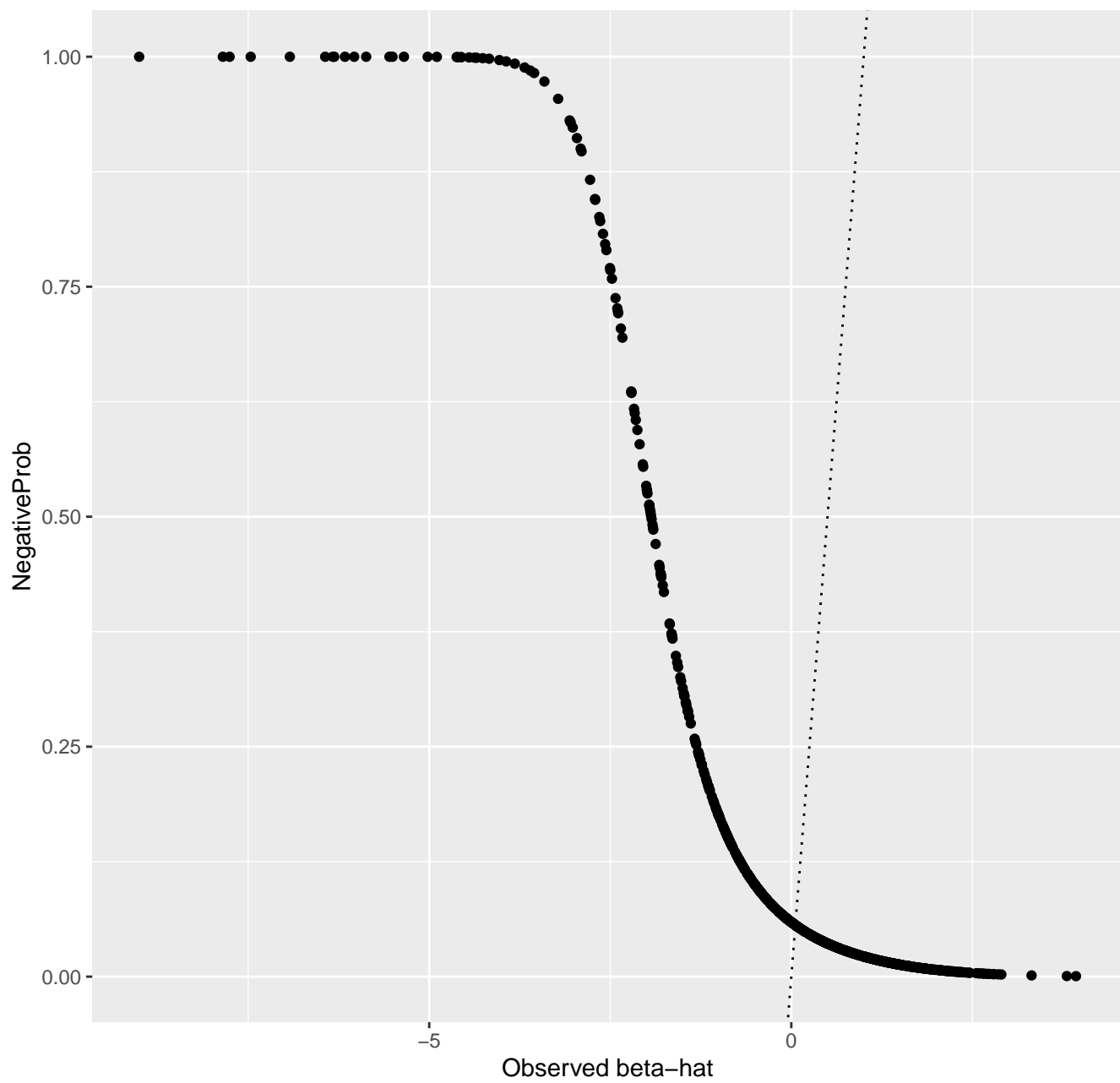
POLE4_(56655) negative



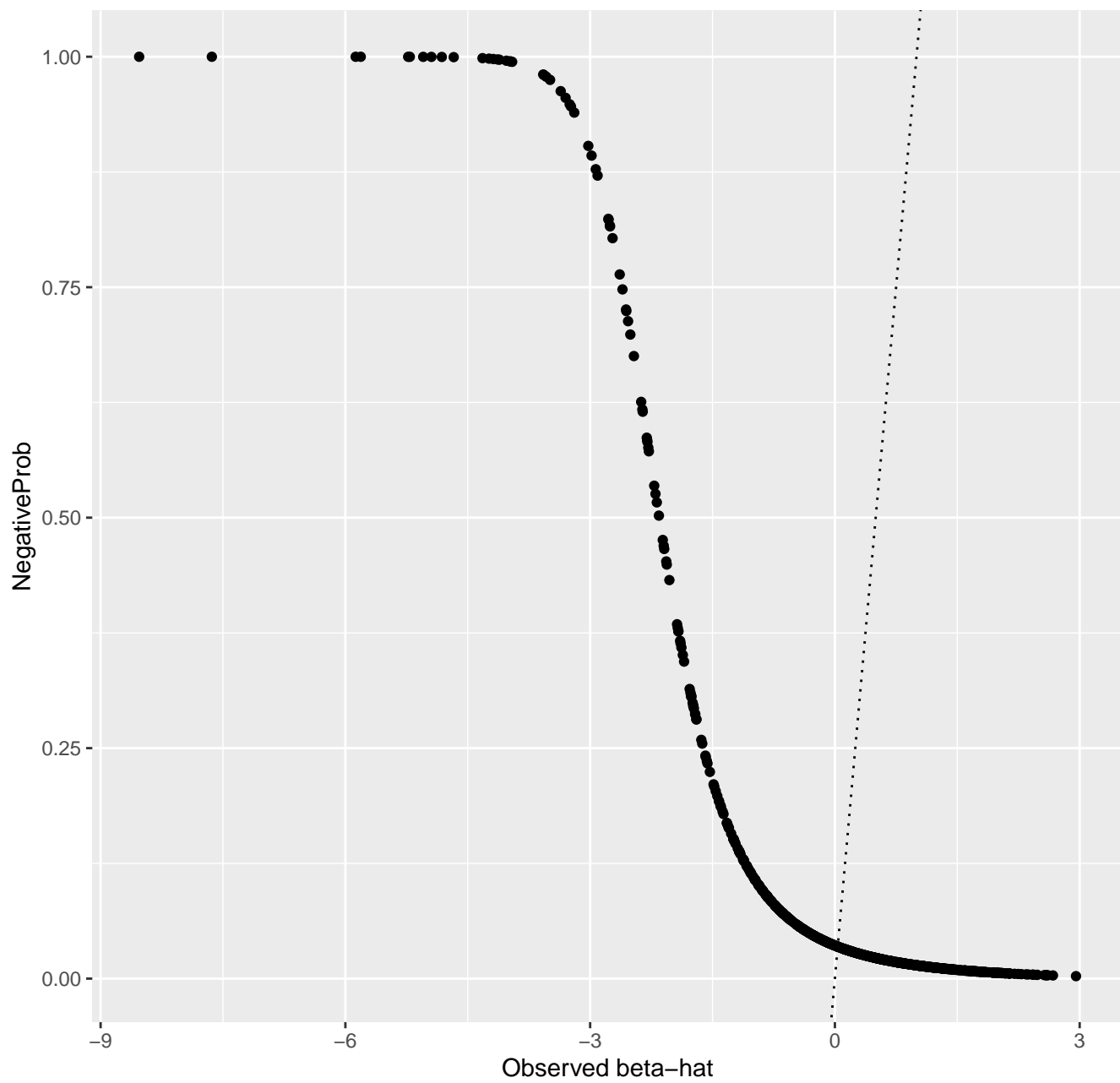
PAK2_(5062) negative



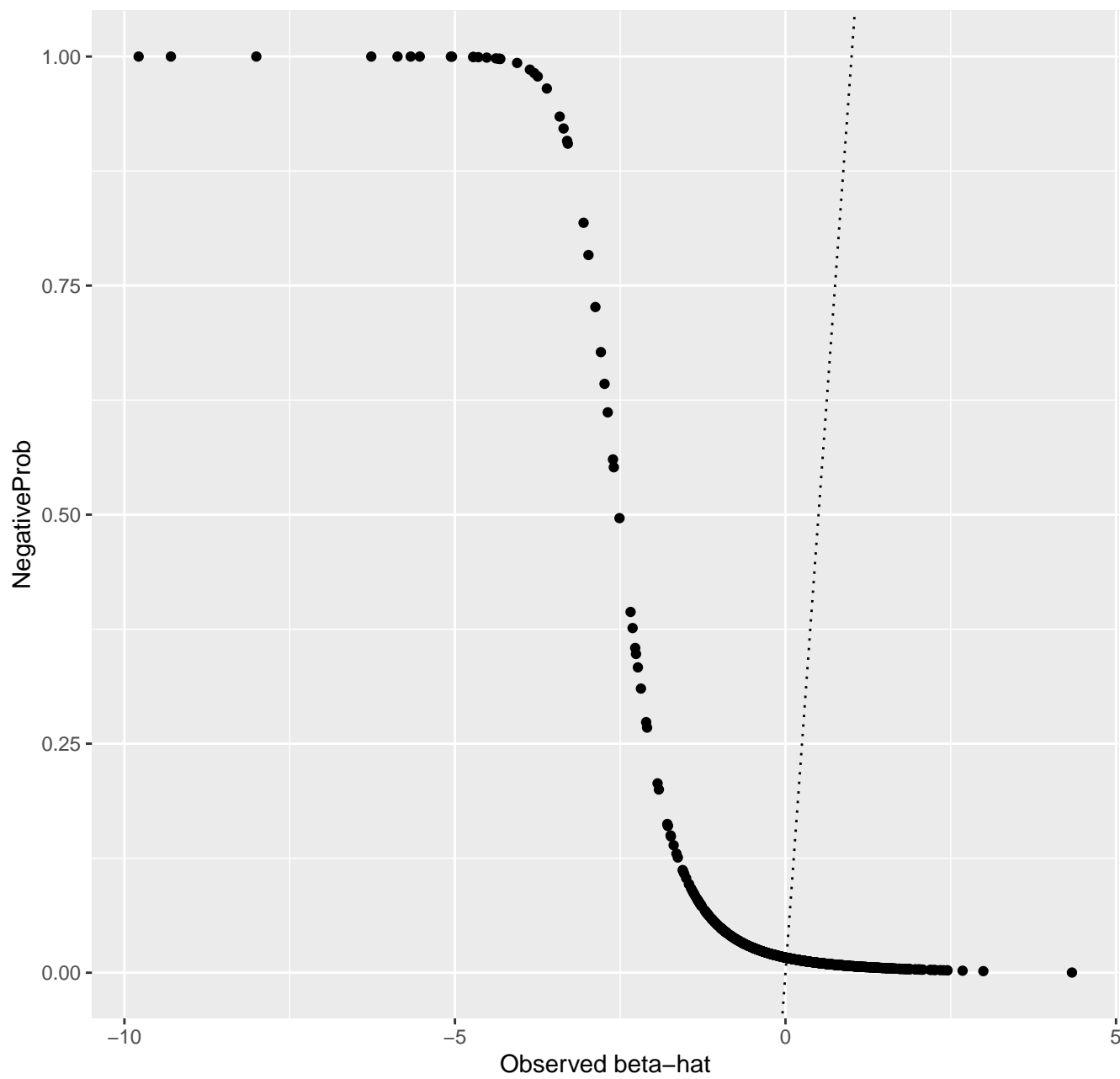
RBBP7_(5931) negative



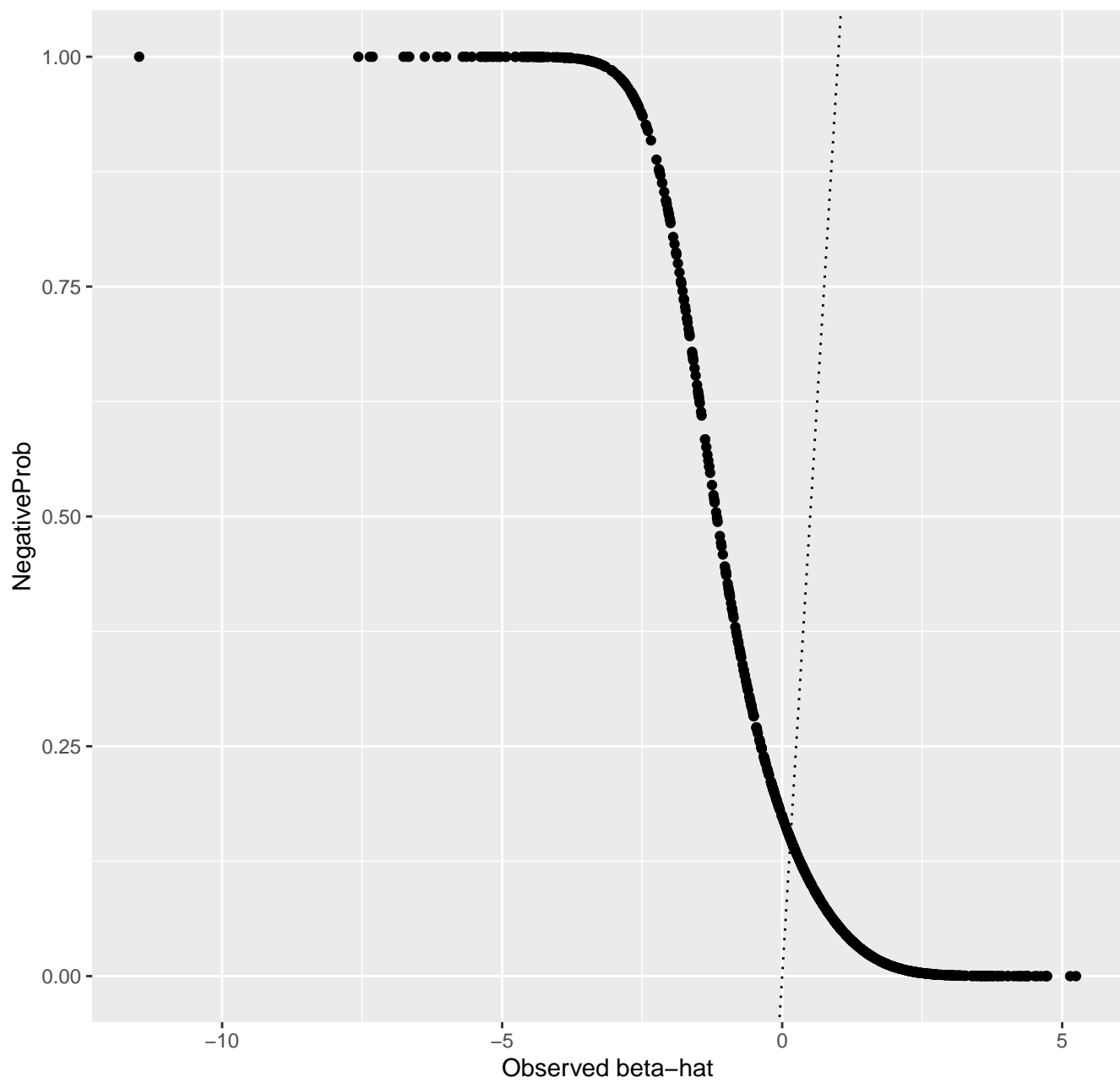
TGFBRAP1_(9392) negative



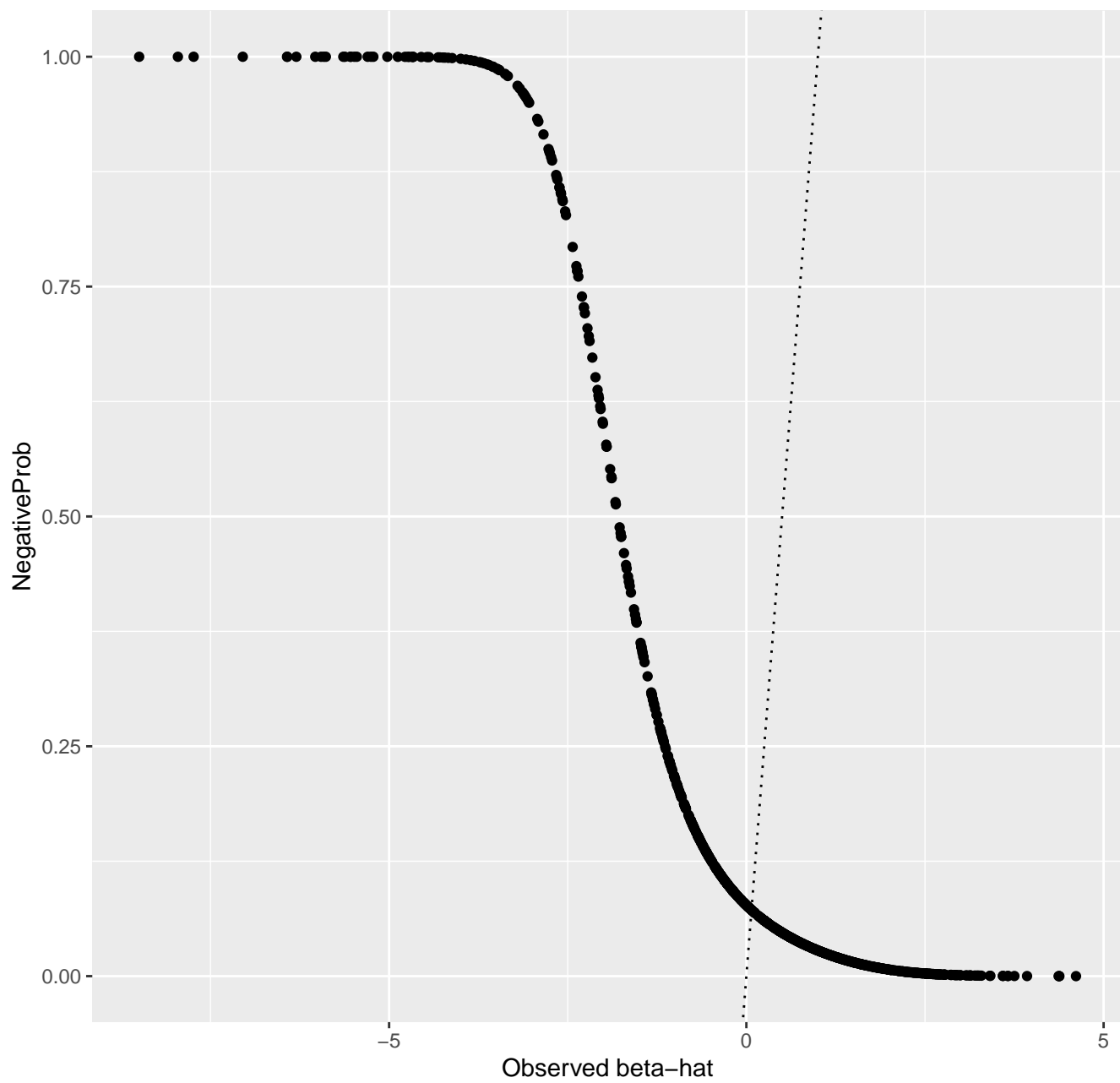
SEC23B_(10483) negative



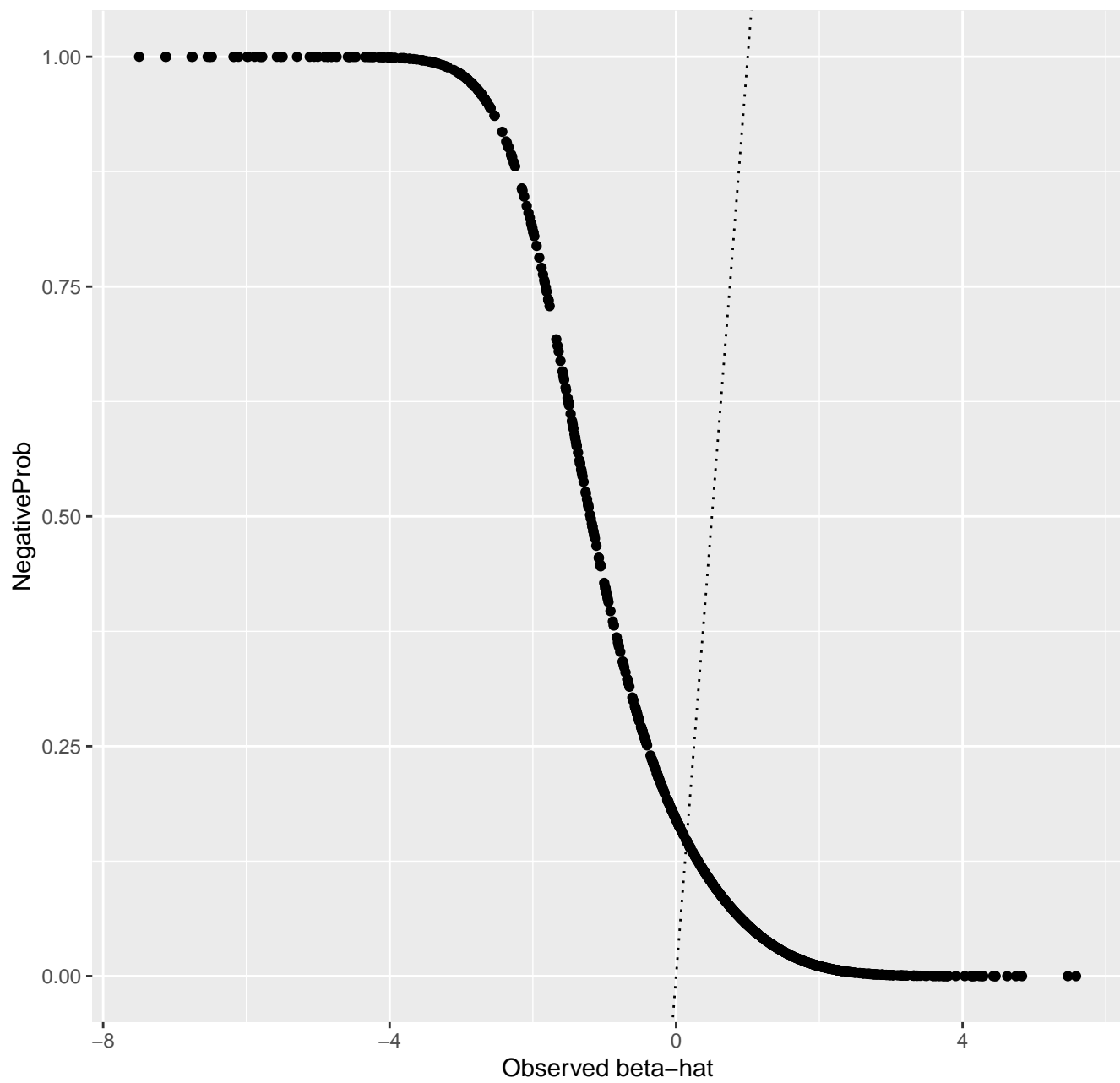
UMPS_(7372) negative



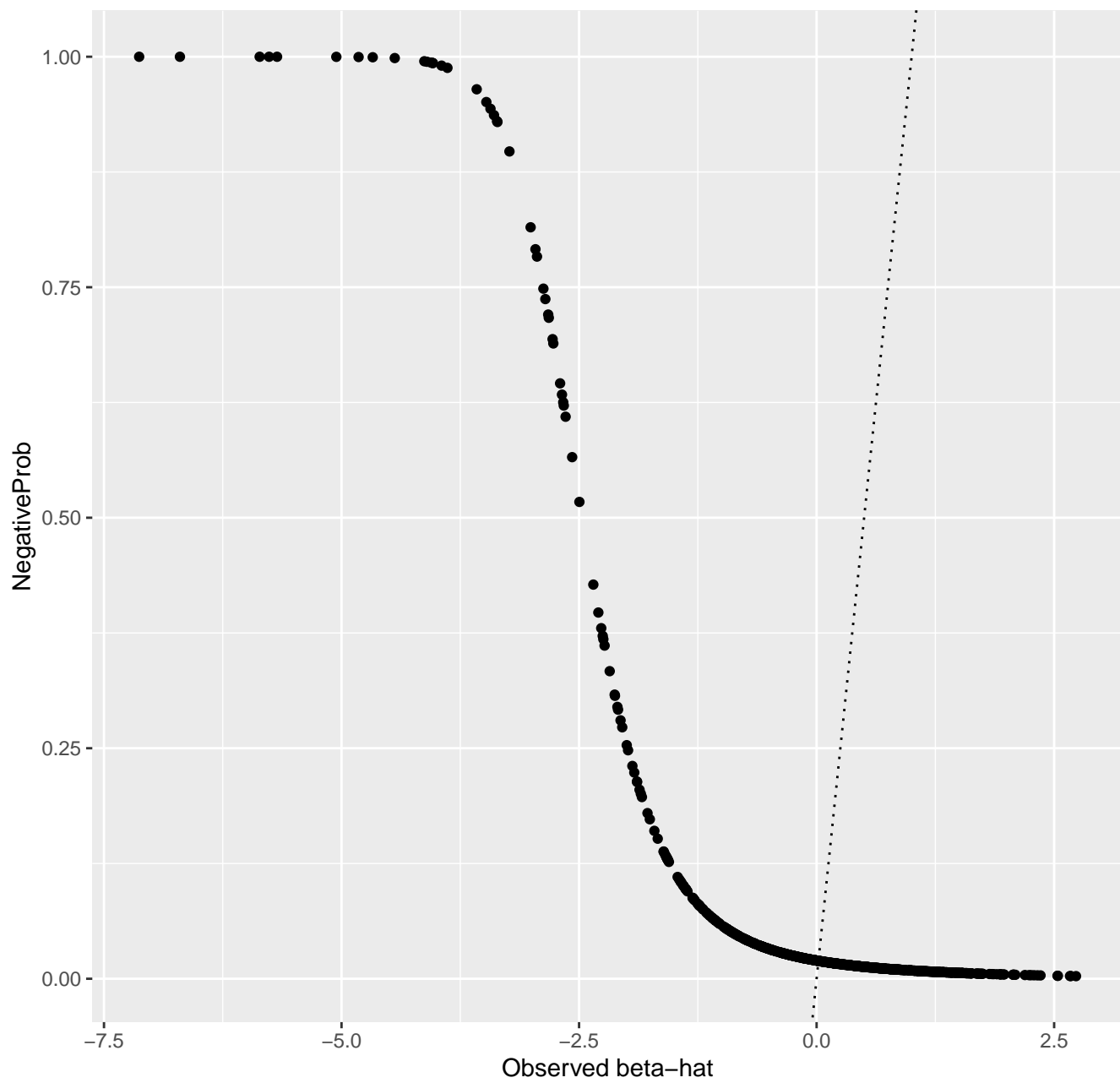
DHODH_(1723) negative



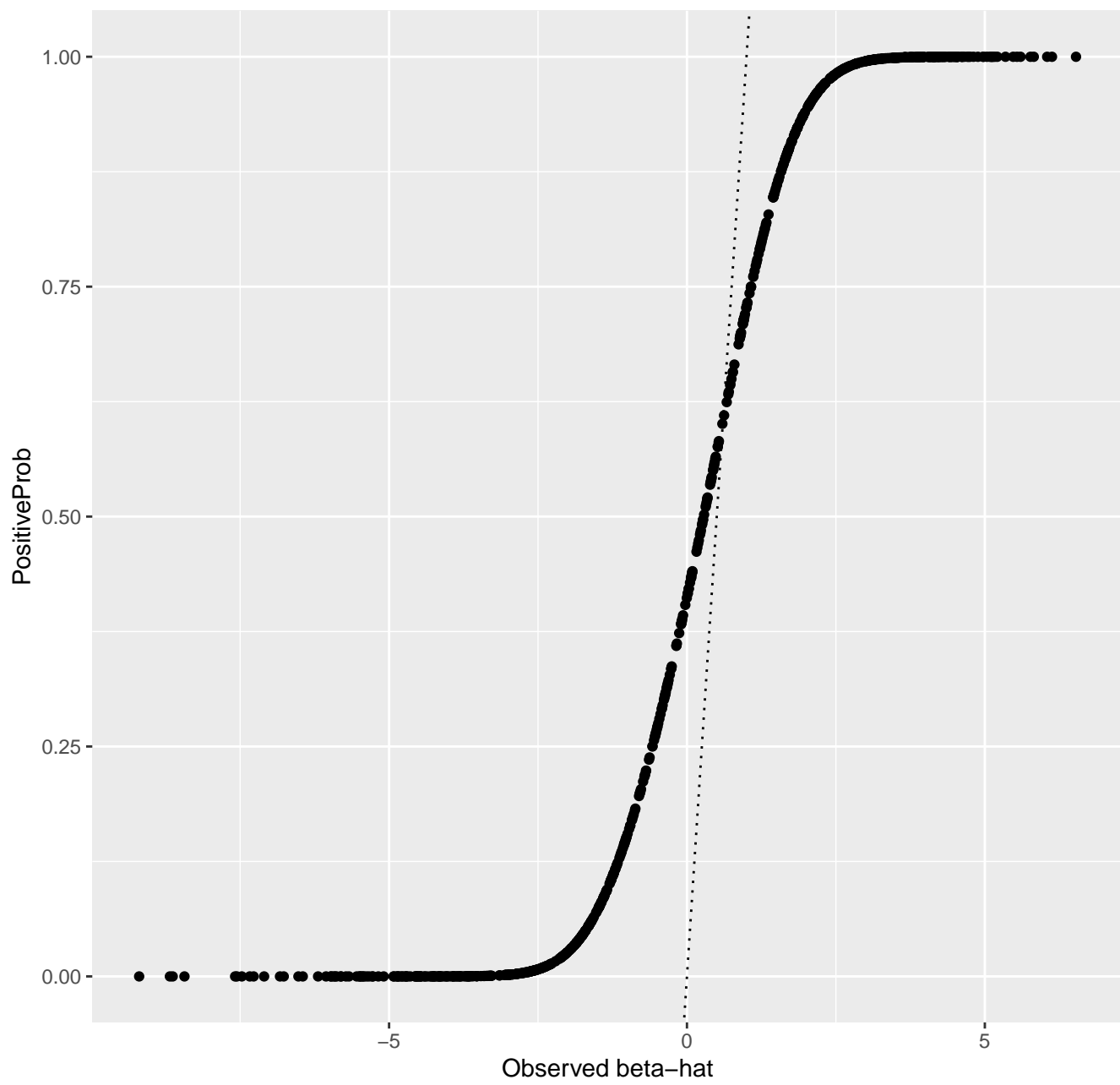
FOSL1_(8061) negative



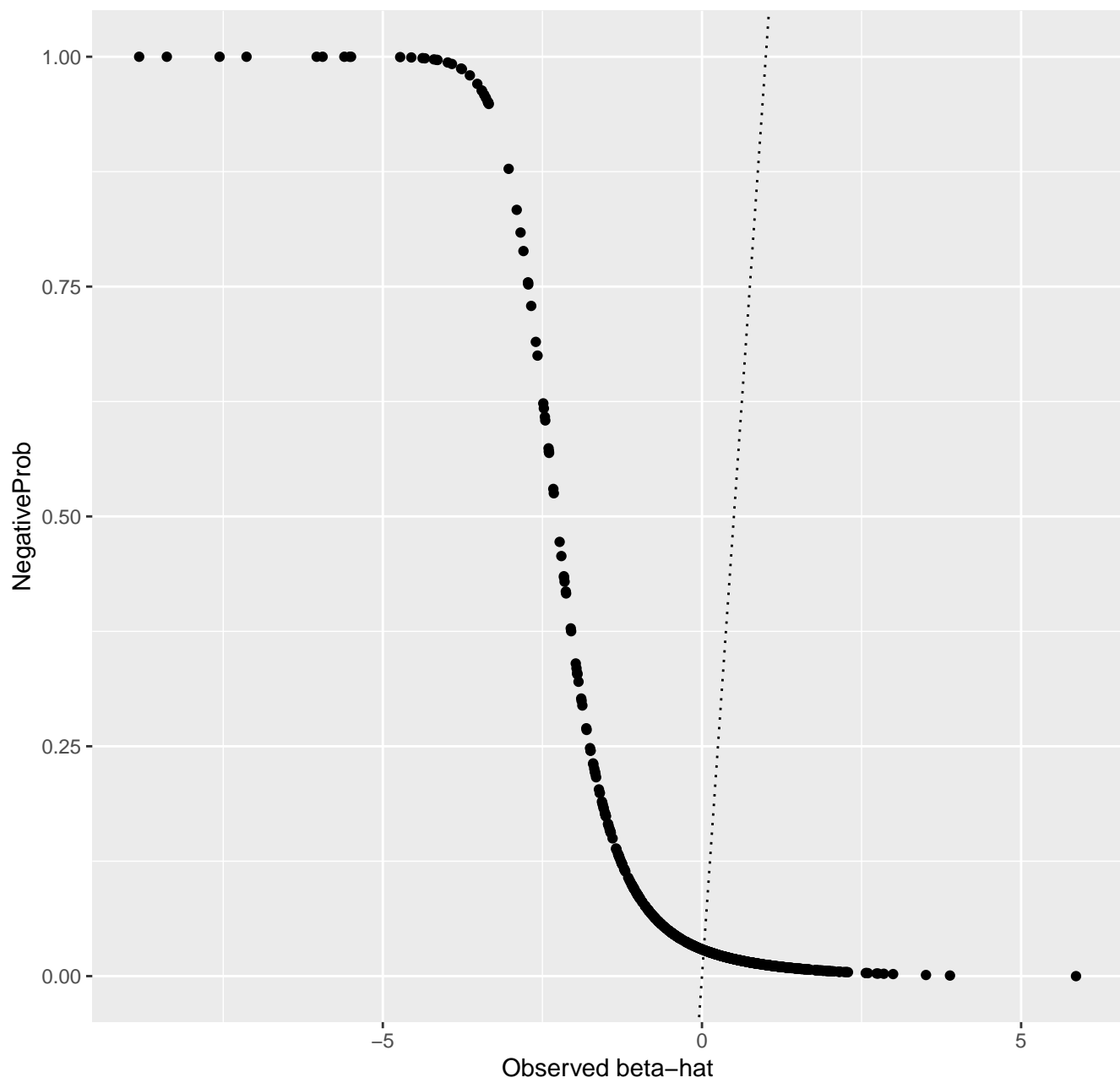
PPP1CC_(5501) negative



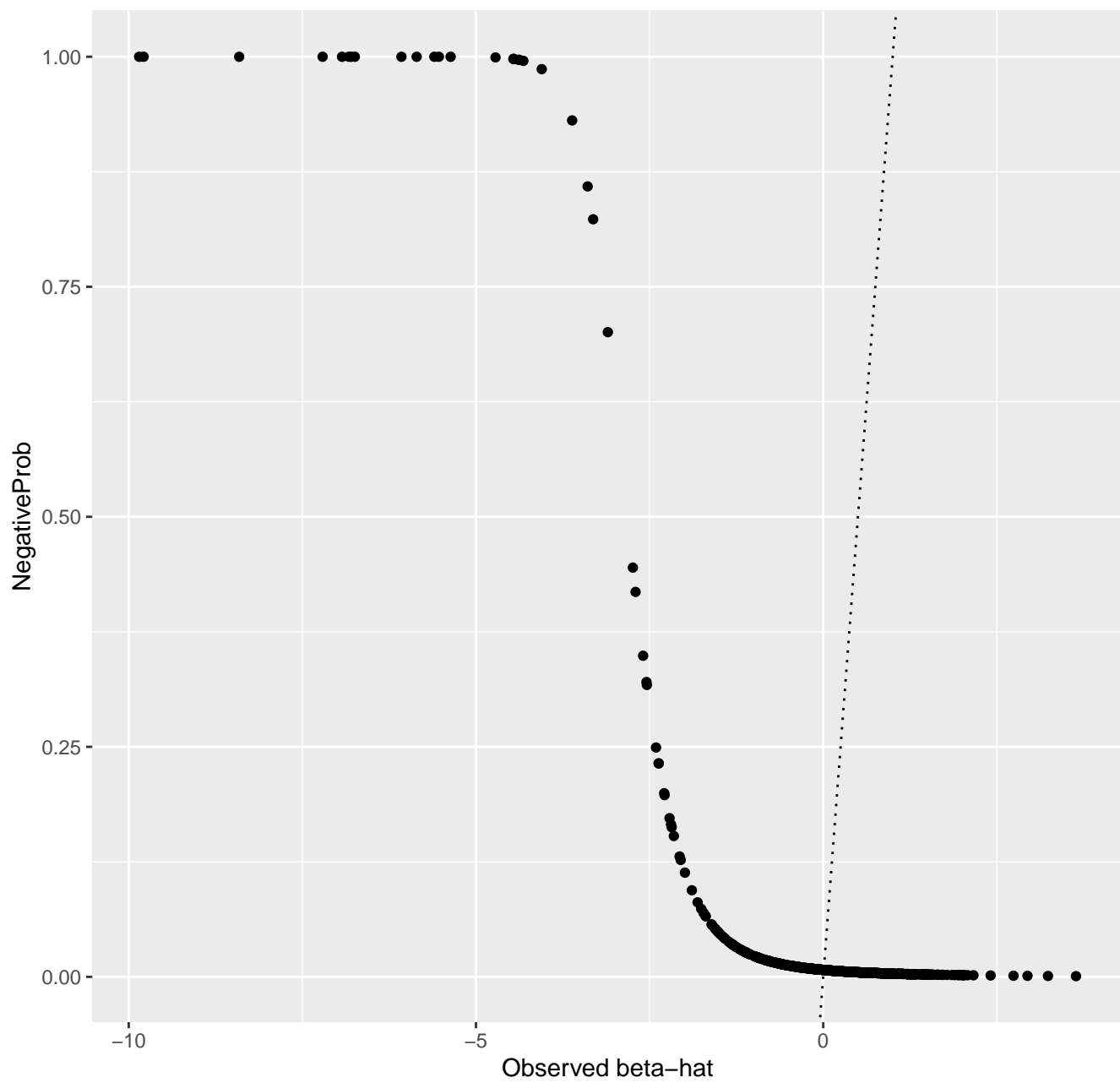
CDK6_(1021) positive



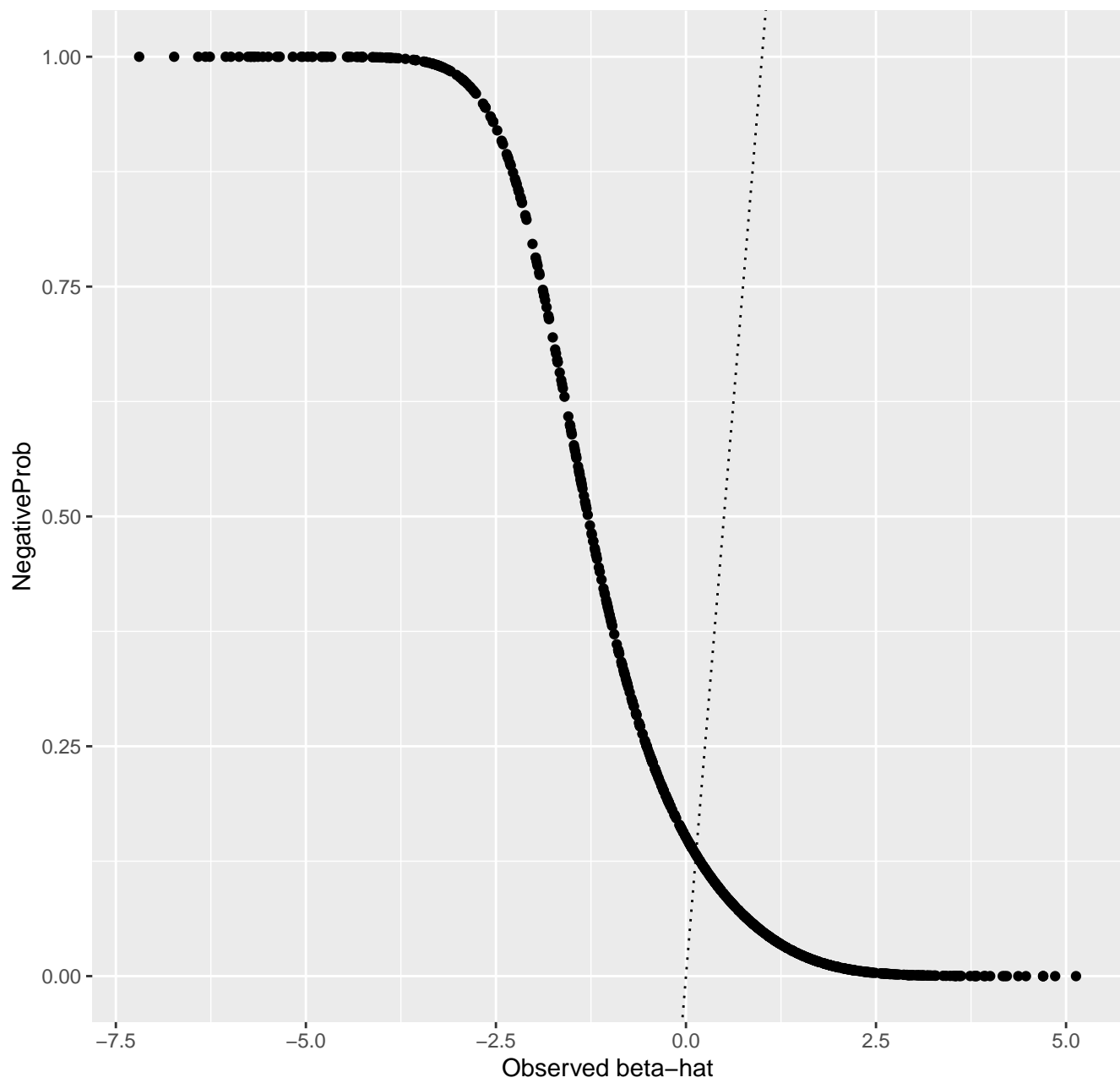
SLC25A37_(51312) negative



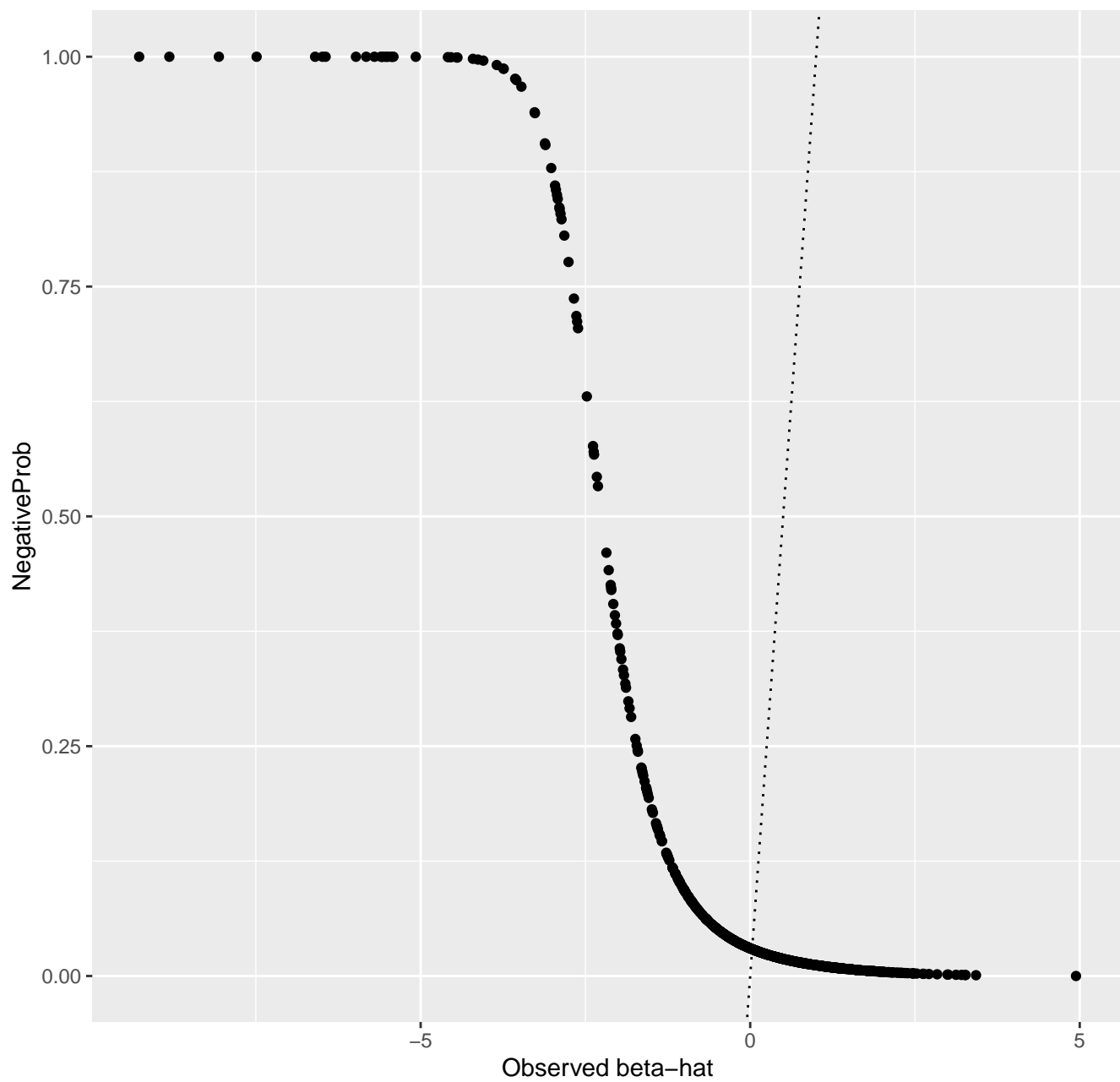
SLC5A3_(6526) negative



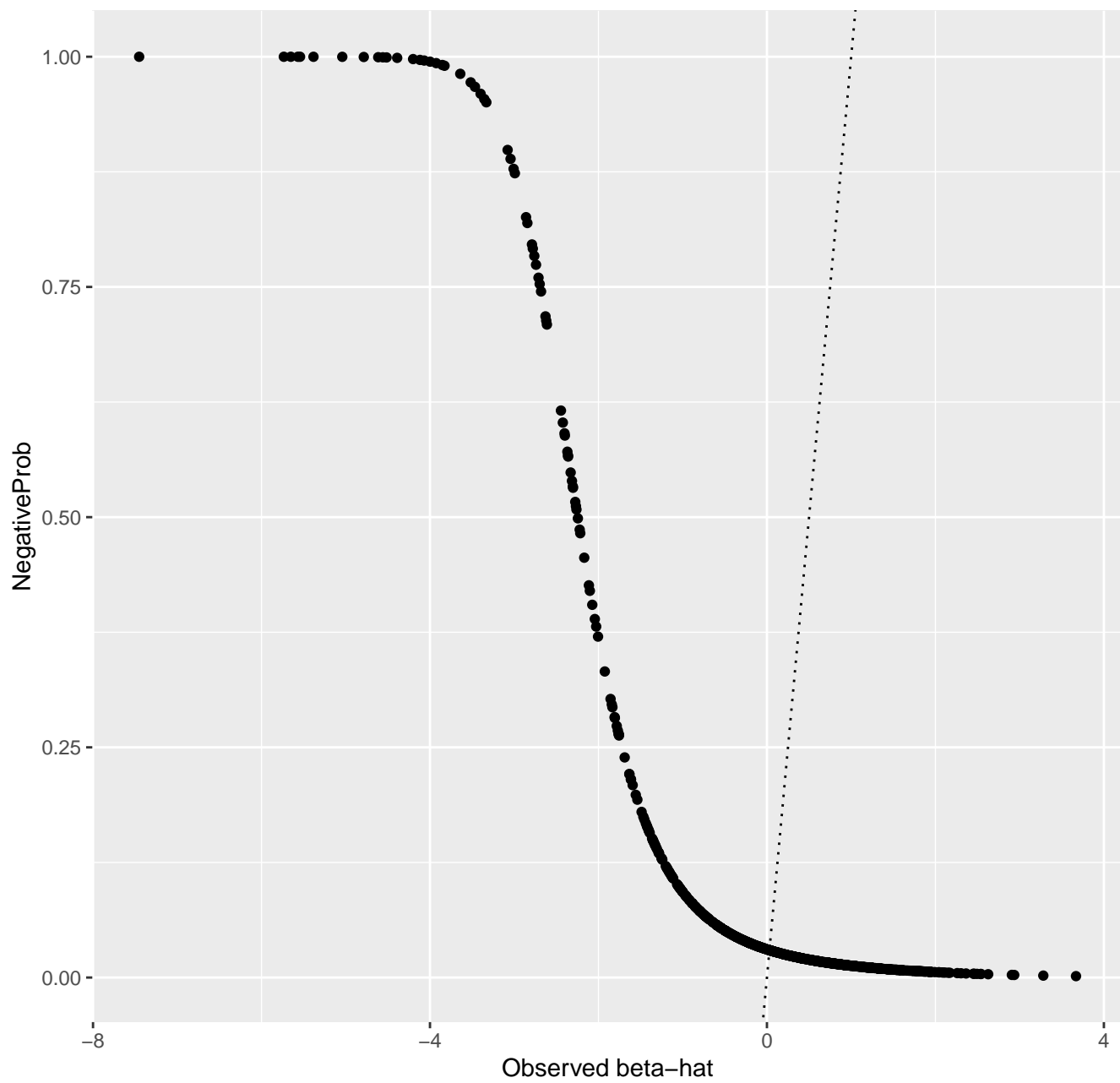
PPAT_(5471) negative



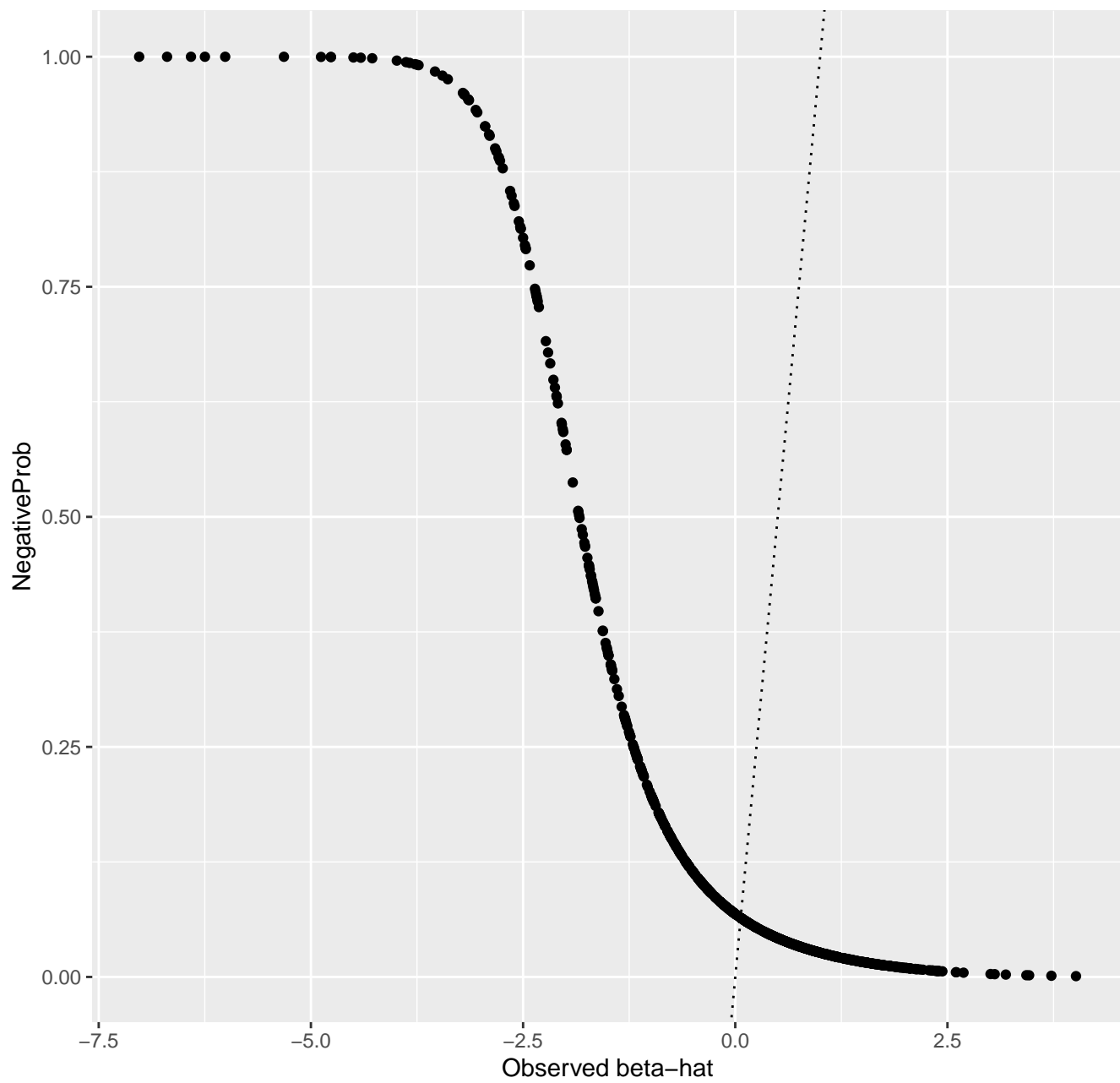
PEX6_(5190) negative



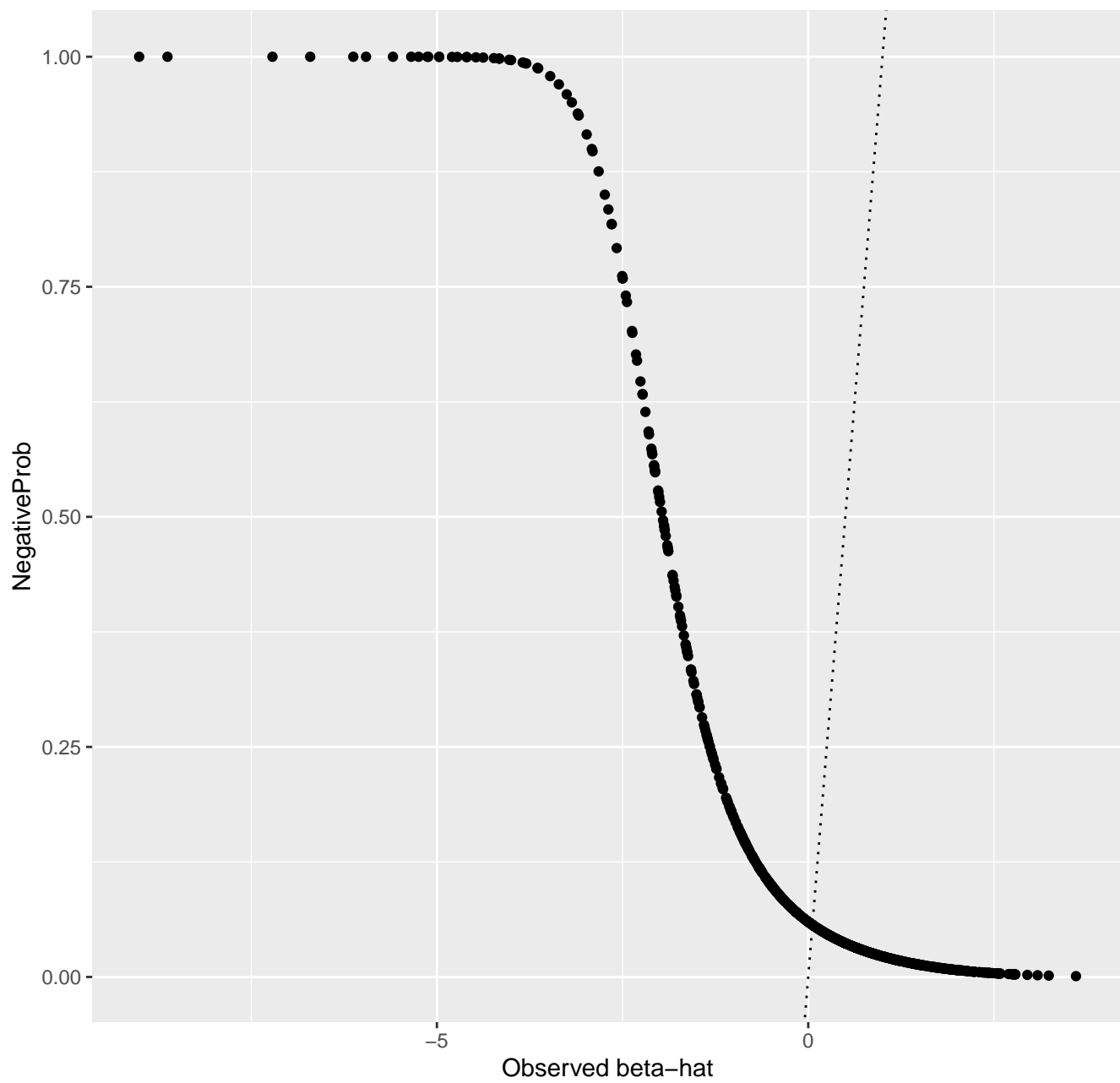
DDX19A_(55308) negative



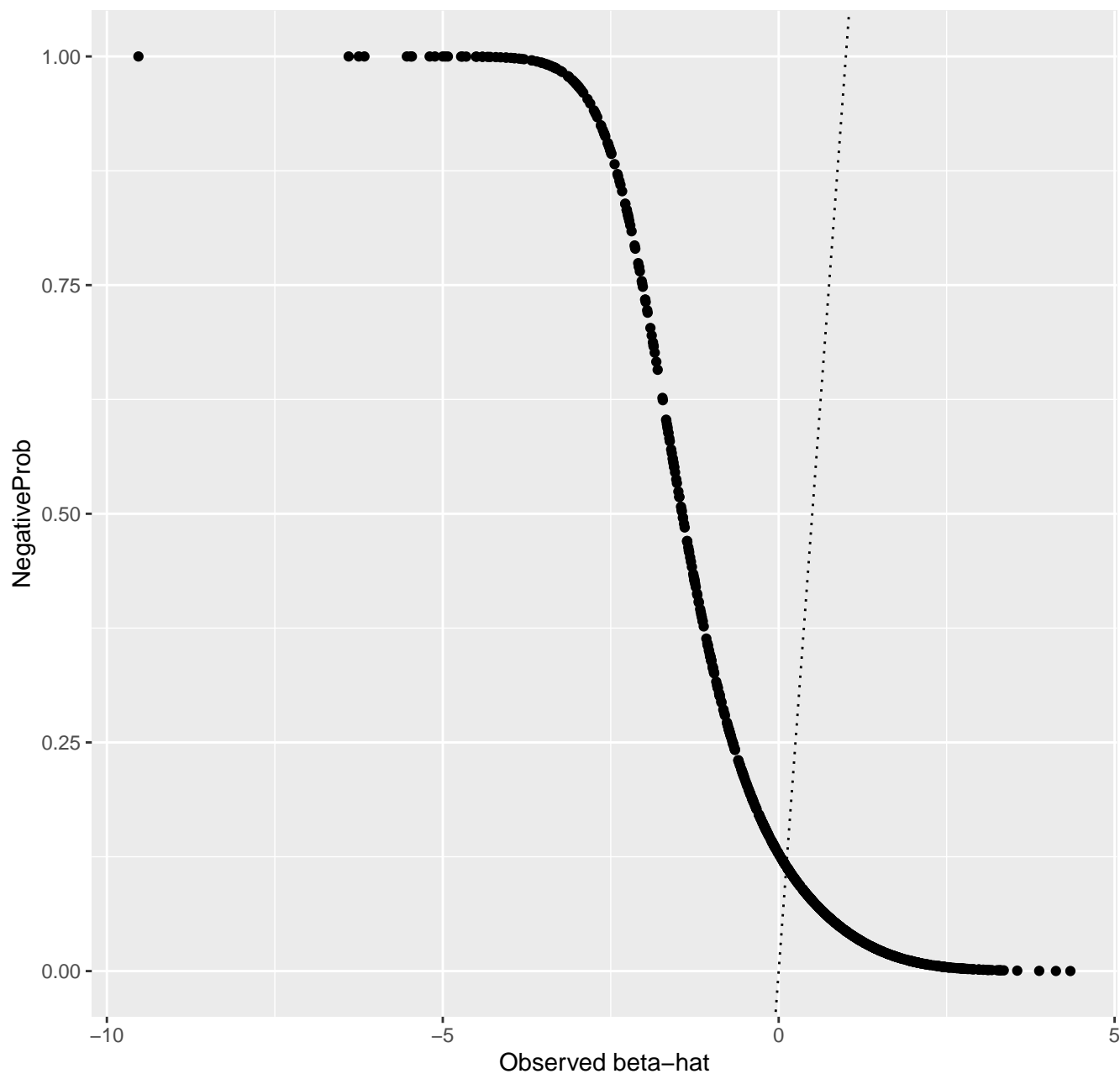
GRK2_(156) negative



NCOA4_(8031) negative



EXT1_(2131) negative



SLC39A6_(25800) negative

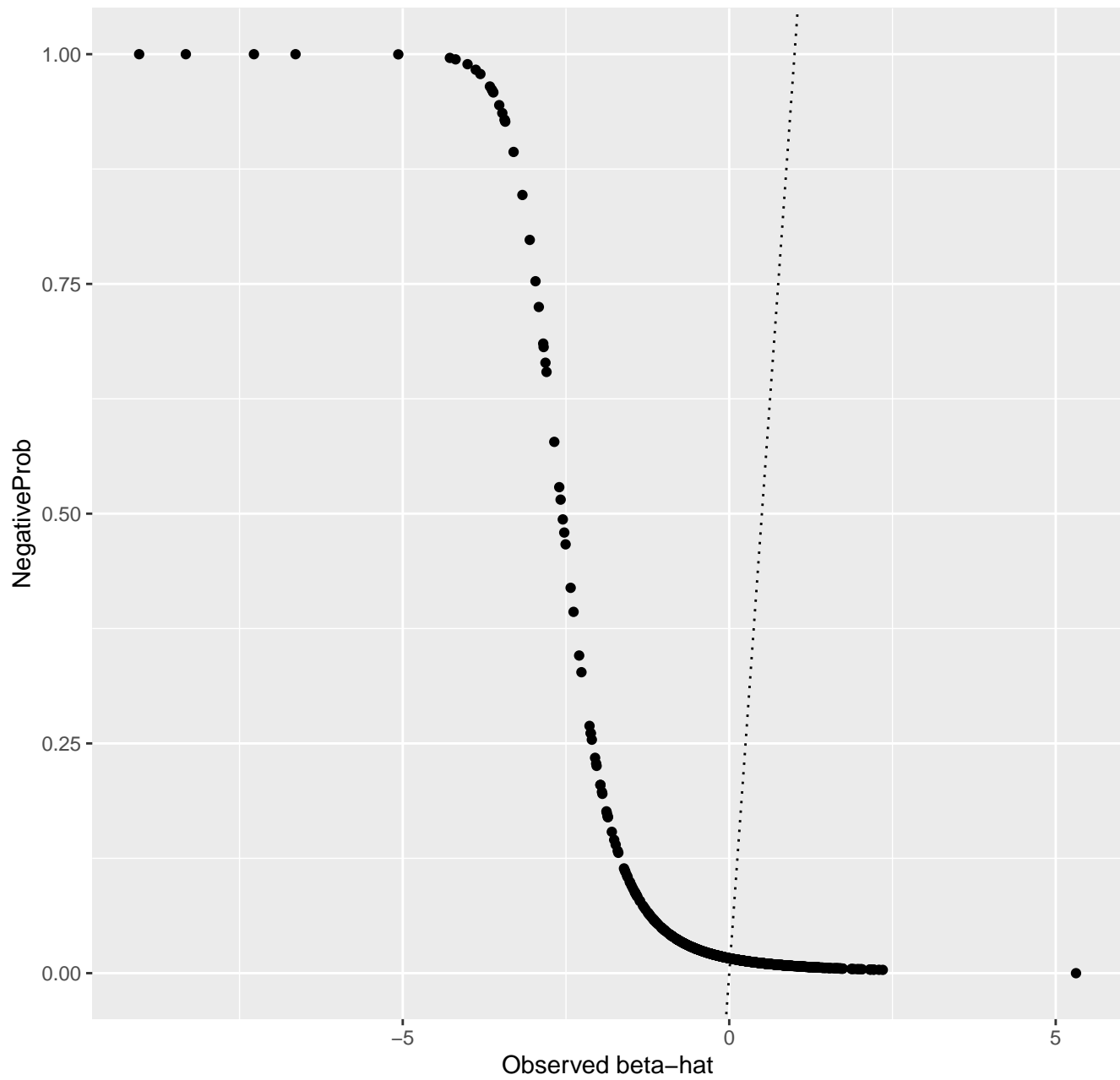
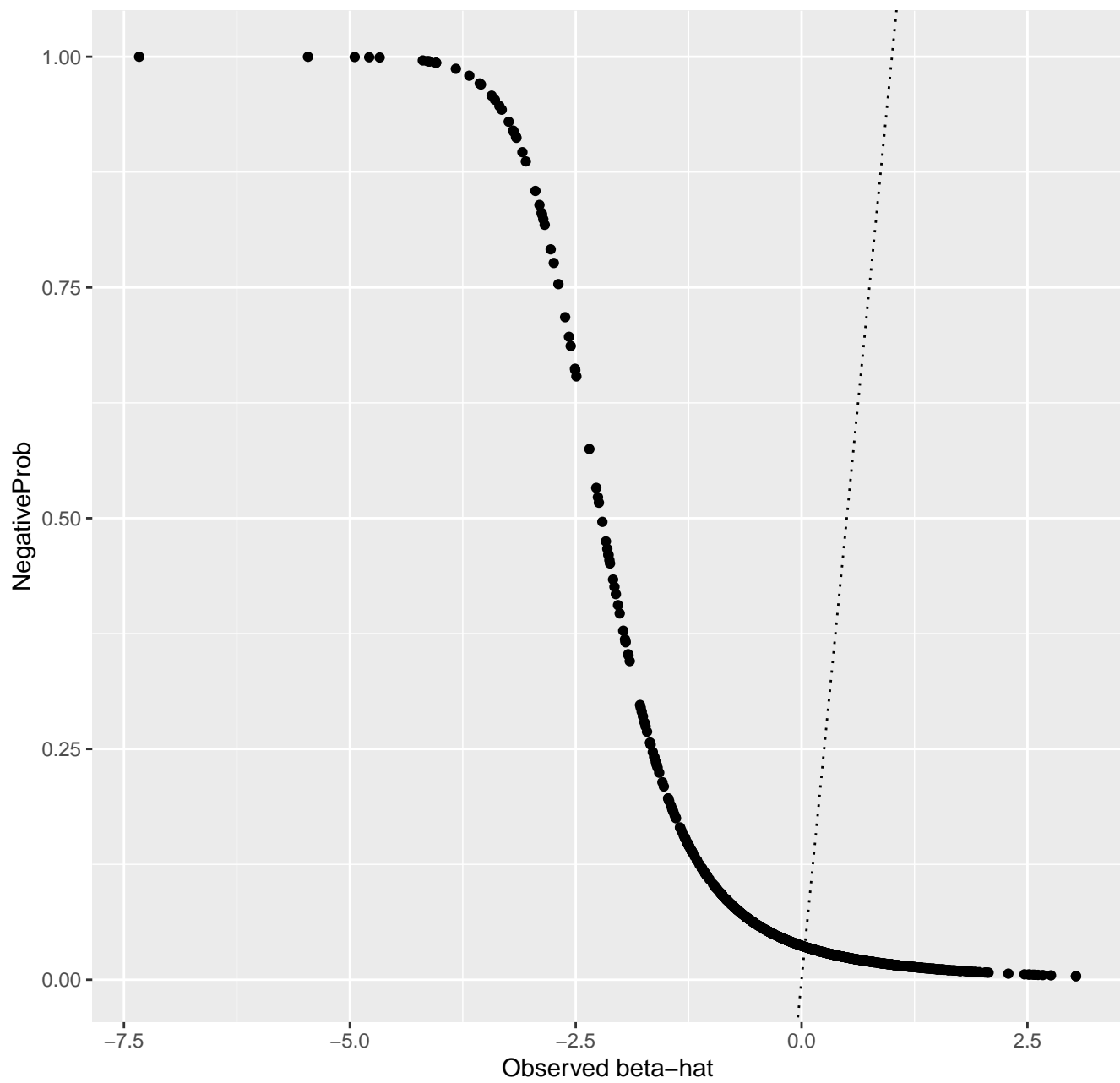
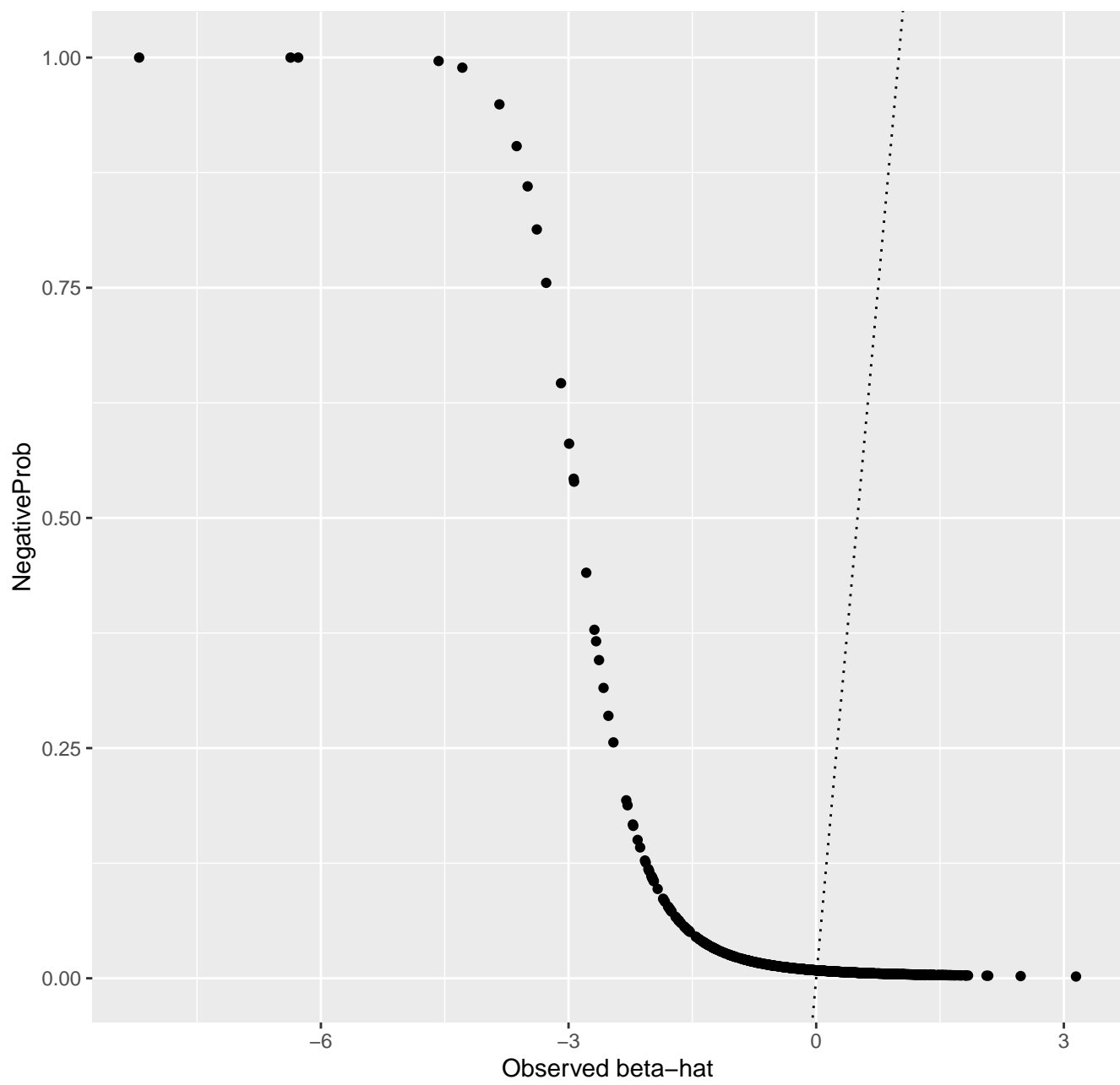


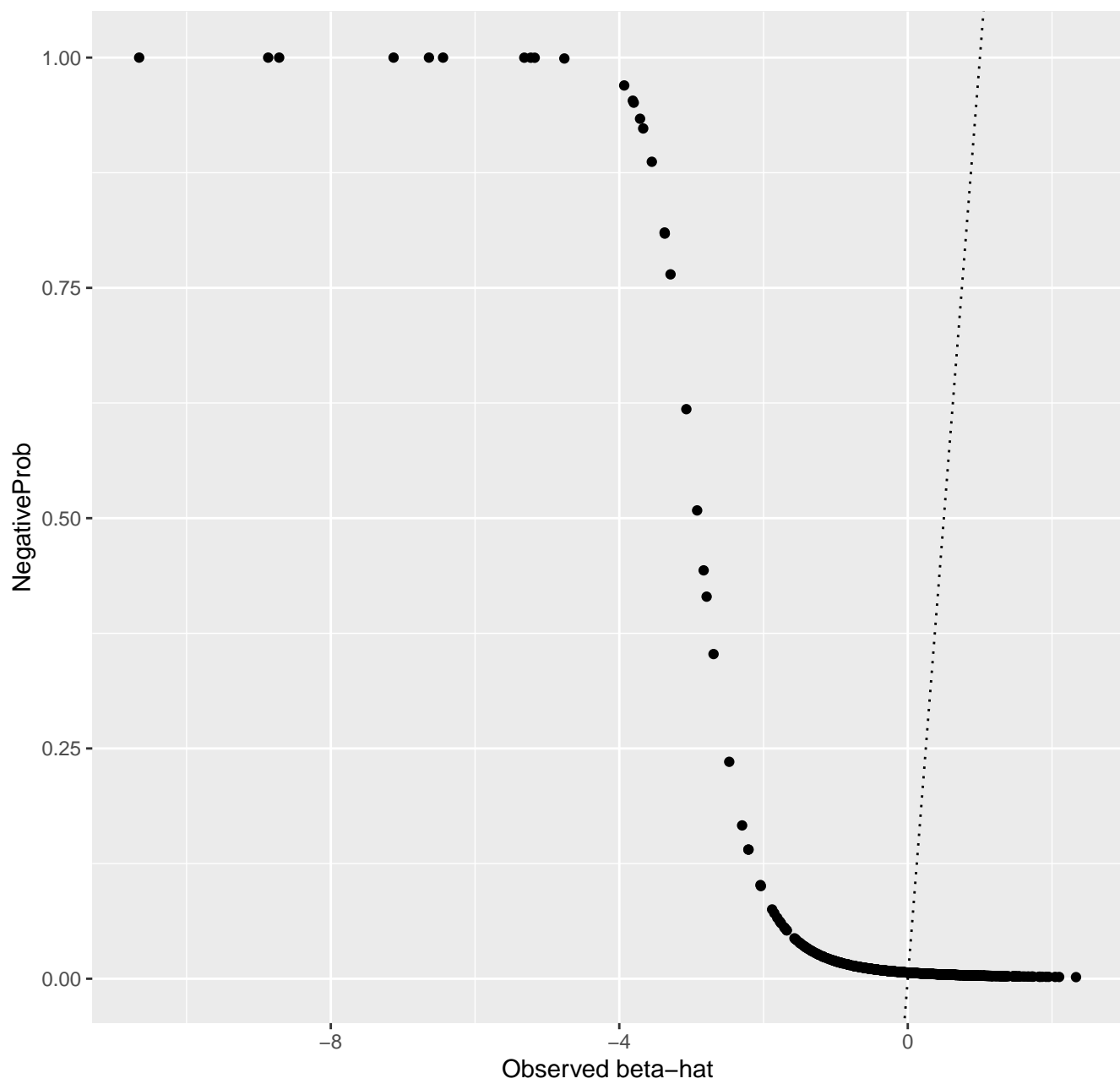
FIG4_(9896) negative



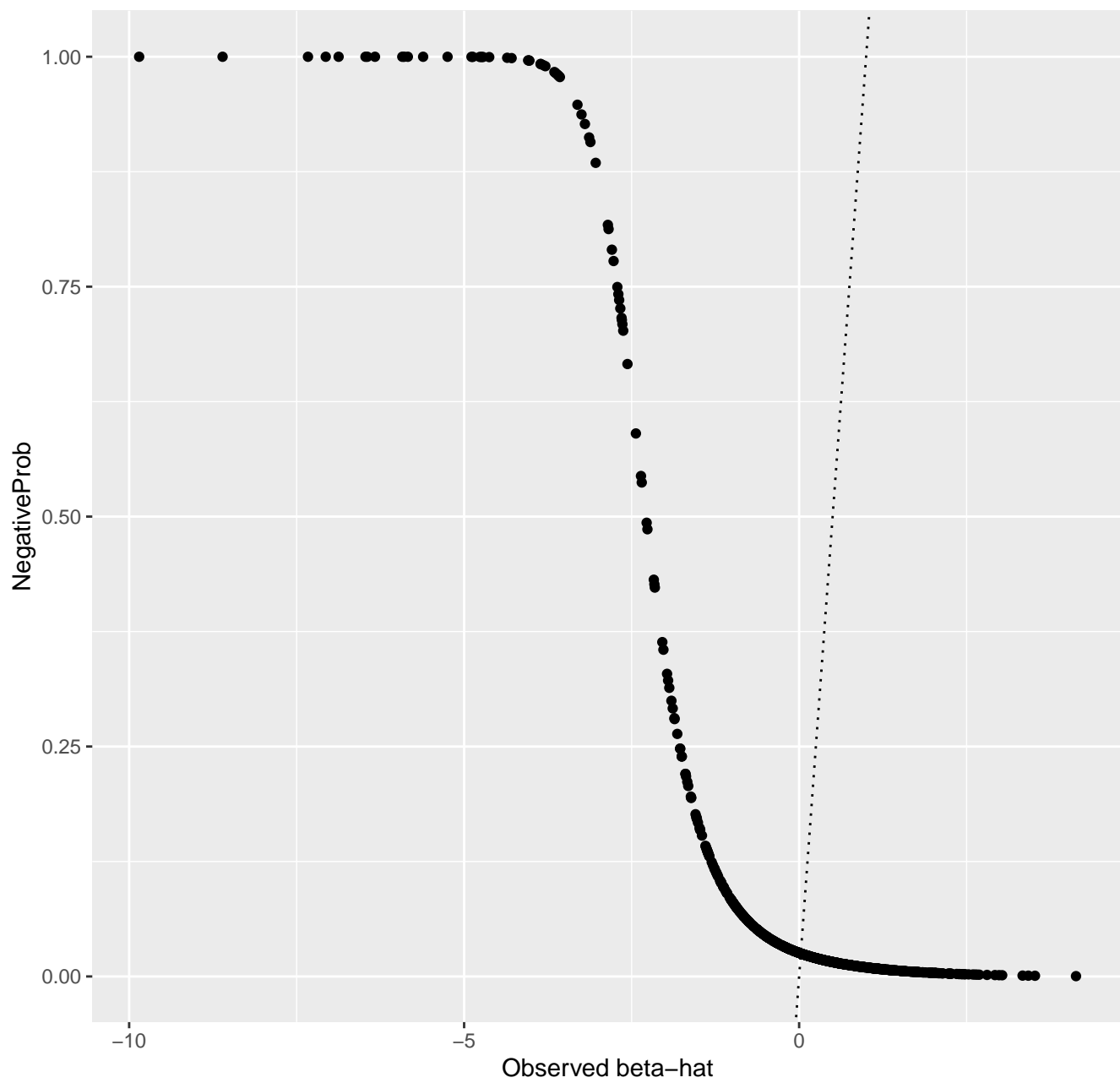
DENND4C_(55667) negative



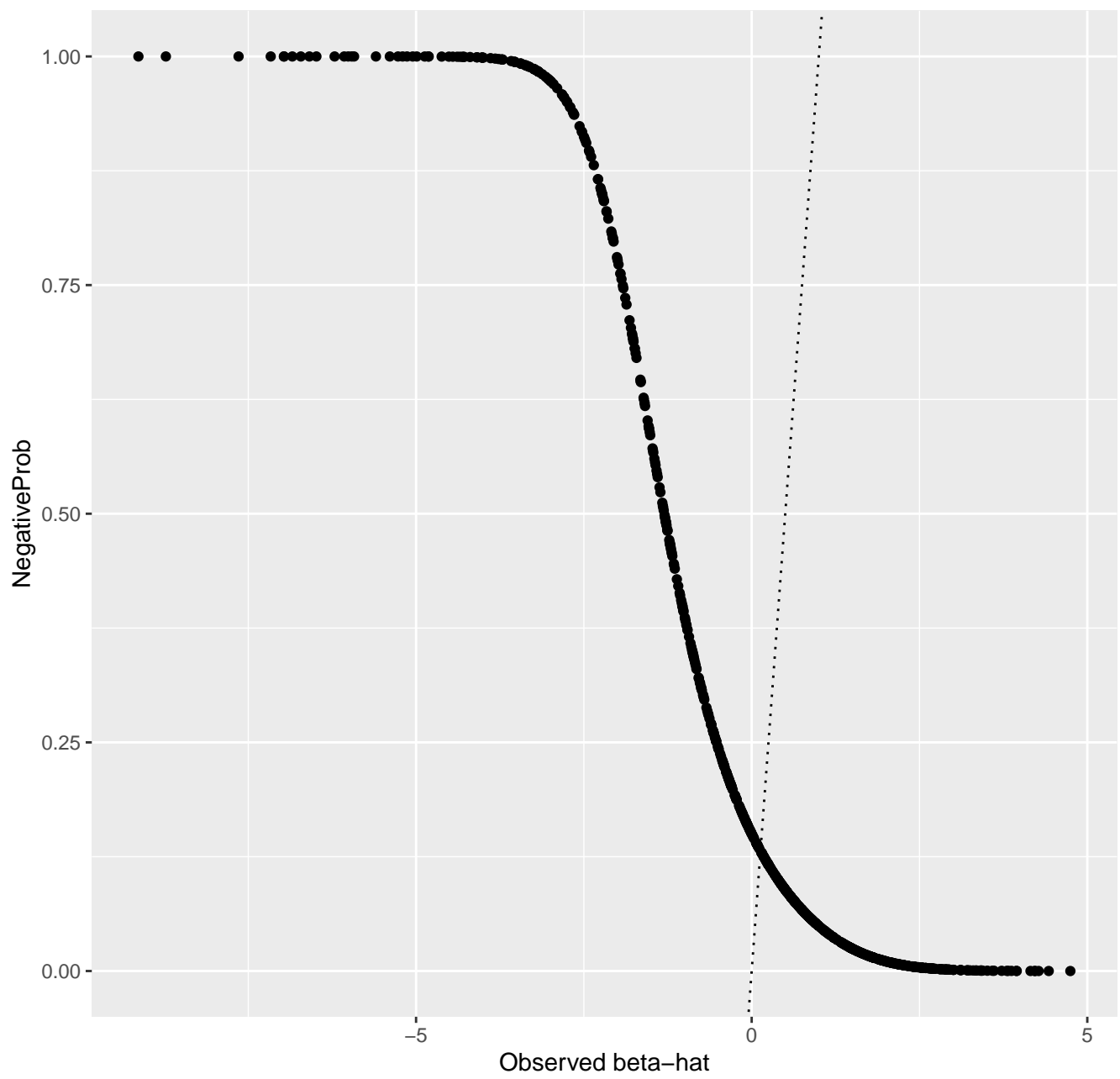
SMARCAL1_(50485) negative



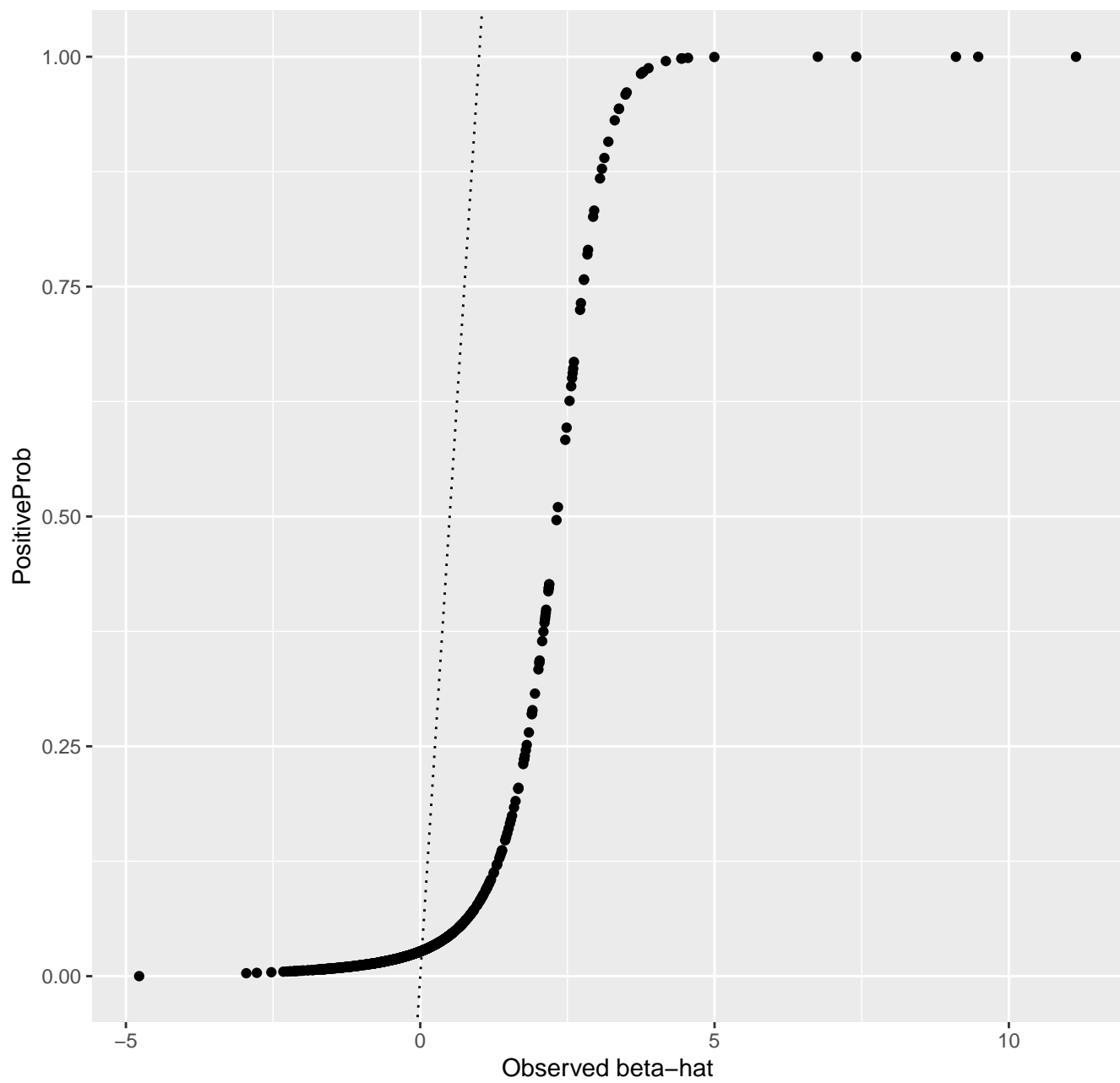
TFAP2A_(7020) negative



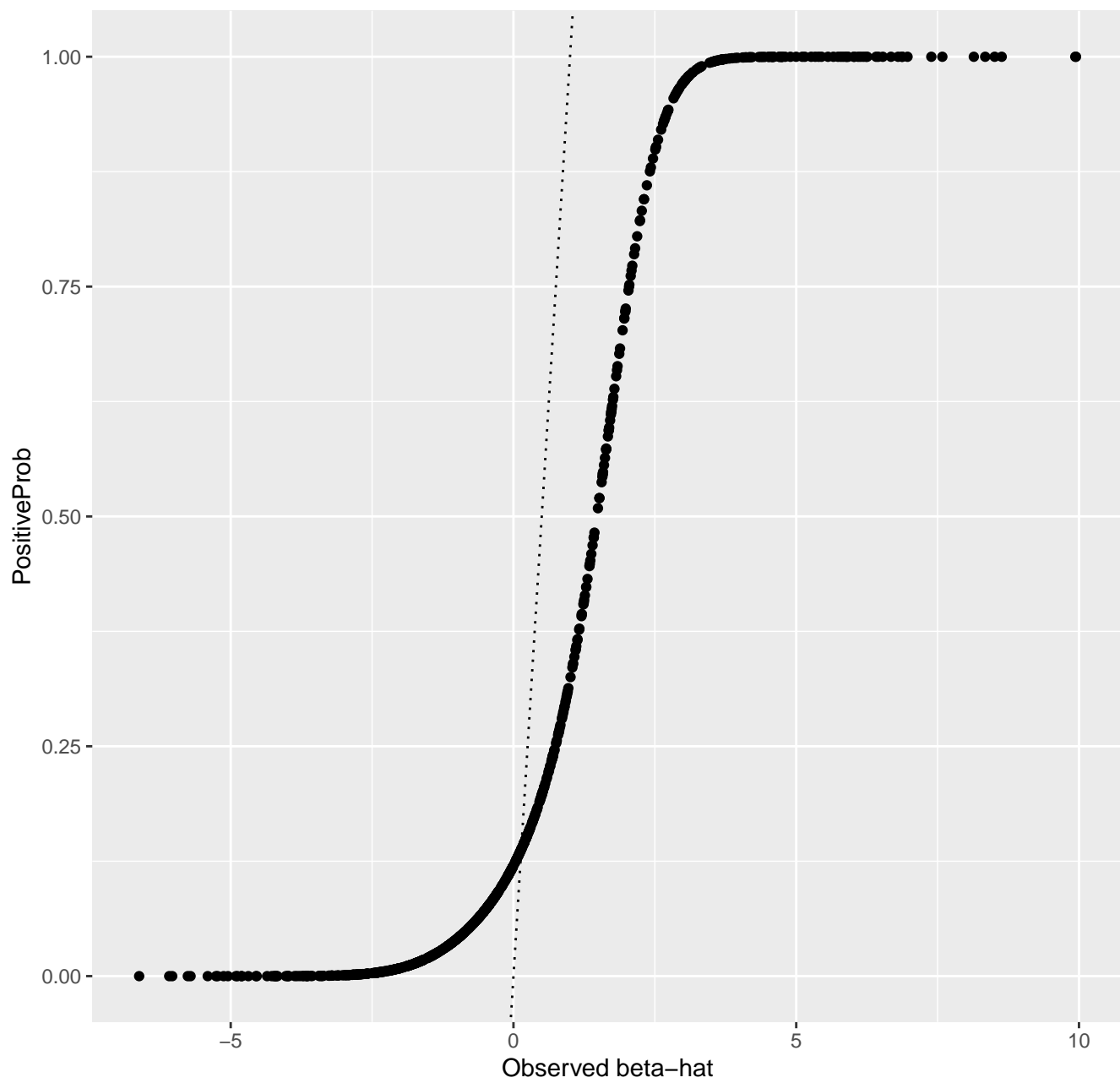
POLE3_(54107) negative



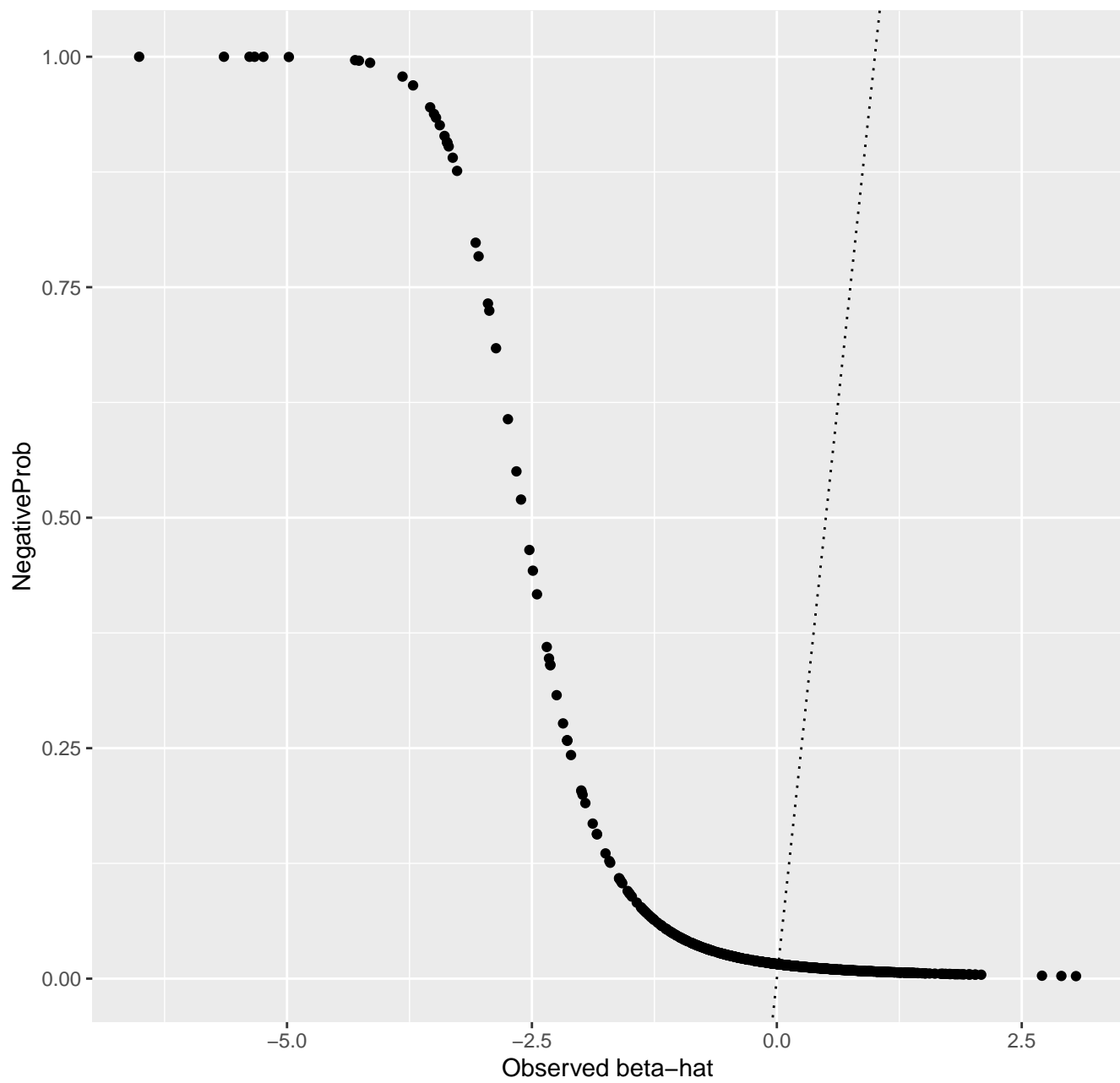
NRP1_(8829) positive



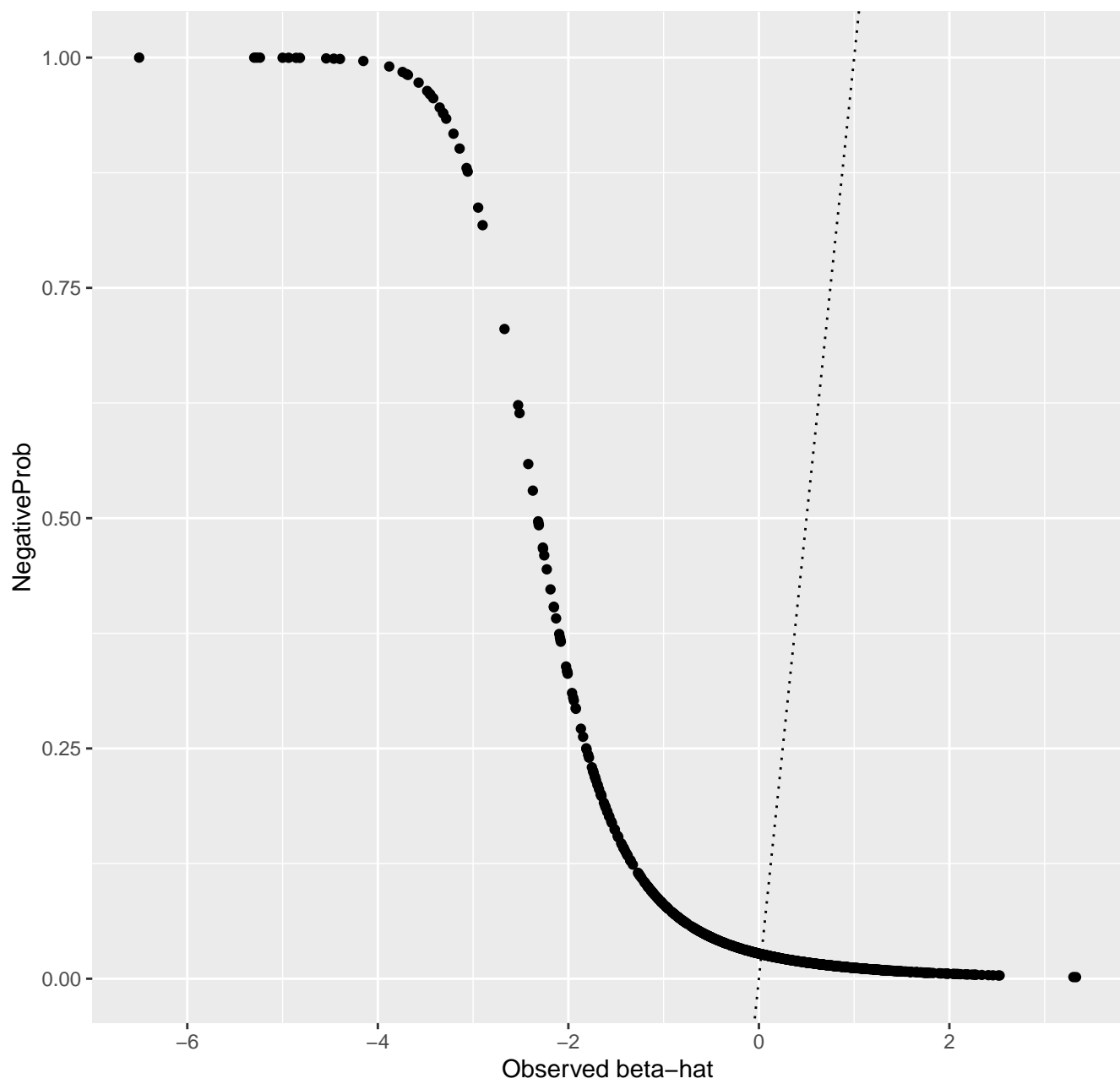
DNM2_(1785) positive



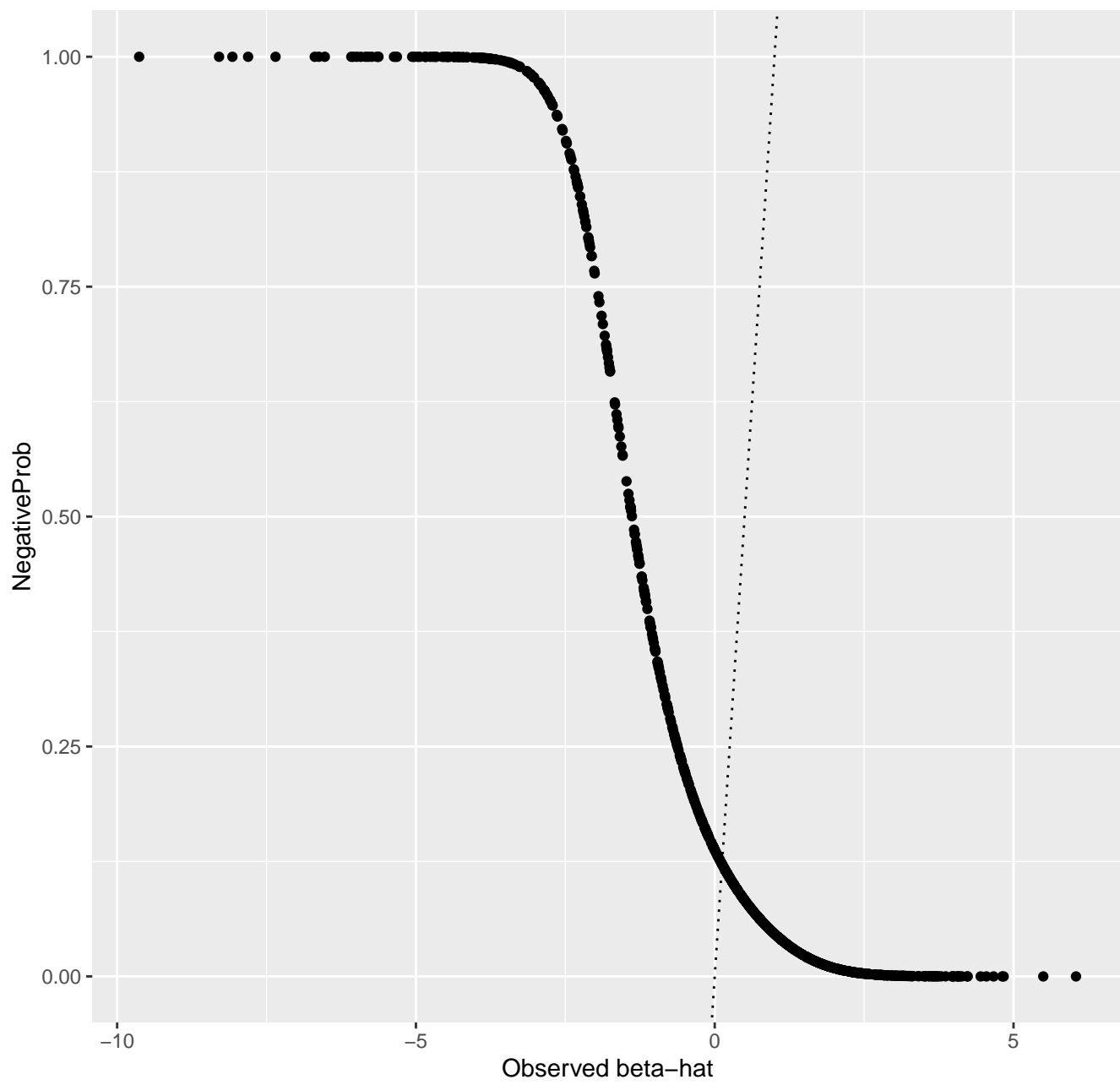
LPAR1_(1902) negative



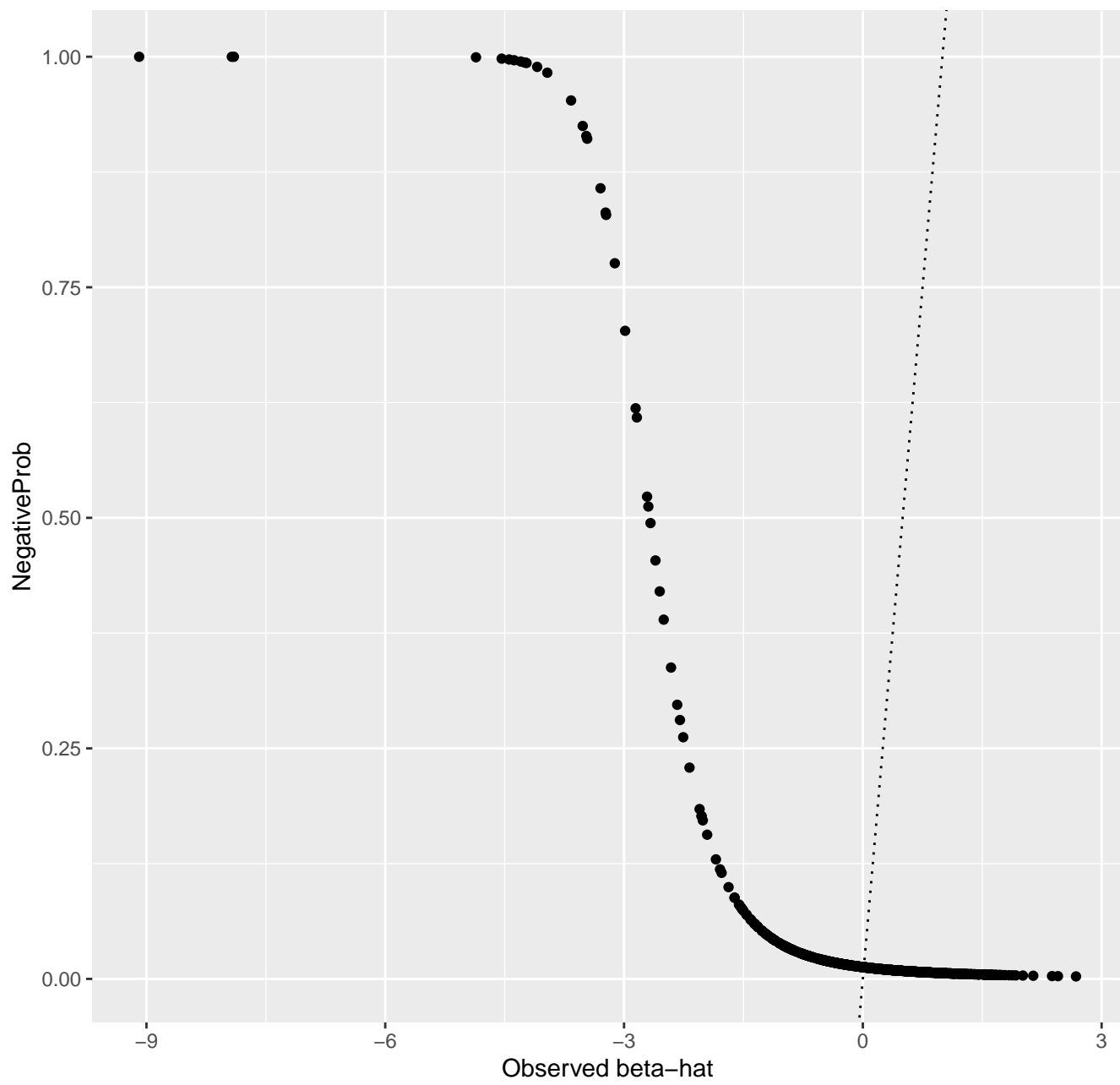
SEPT2_(4735) negative



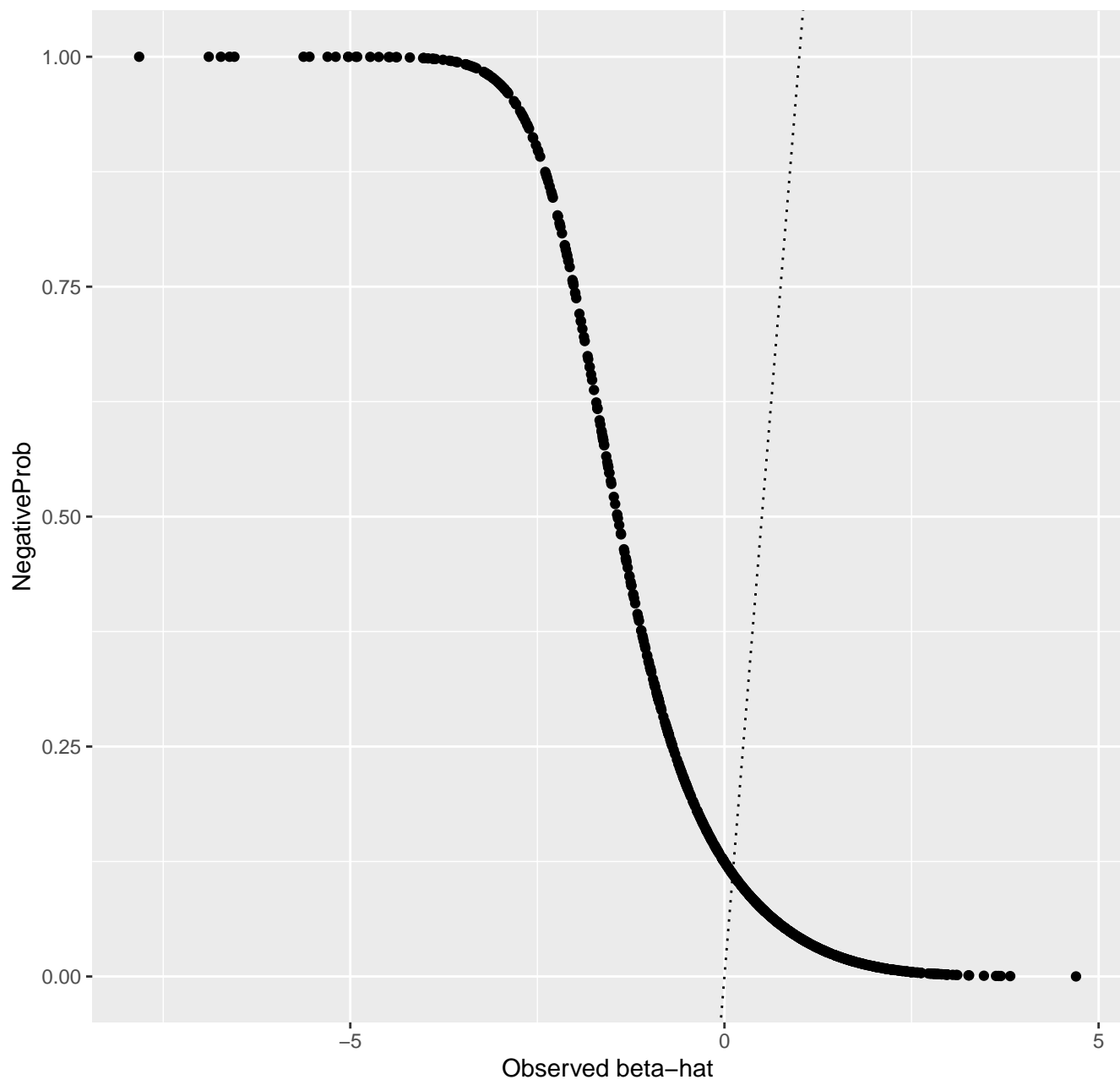
CAD_(790) negative



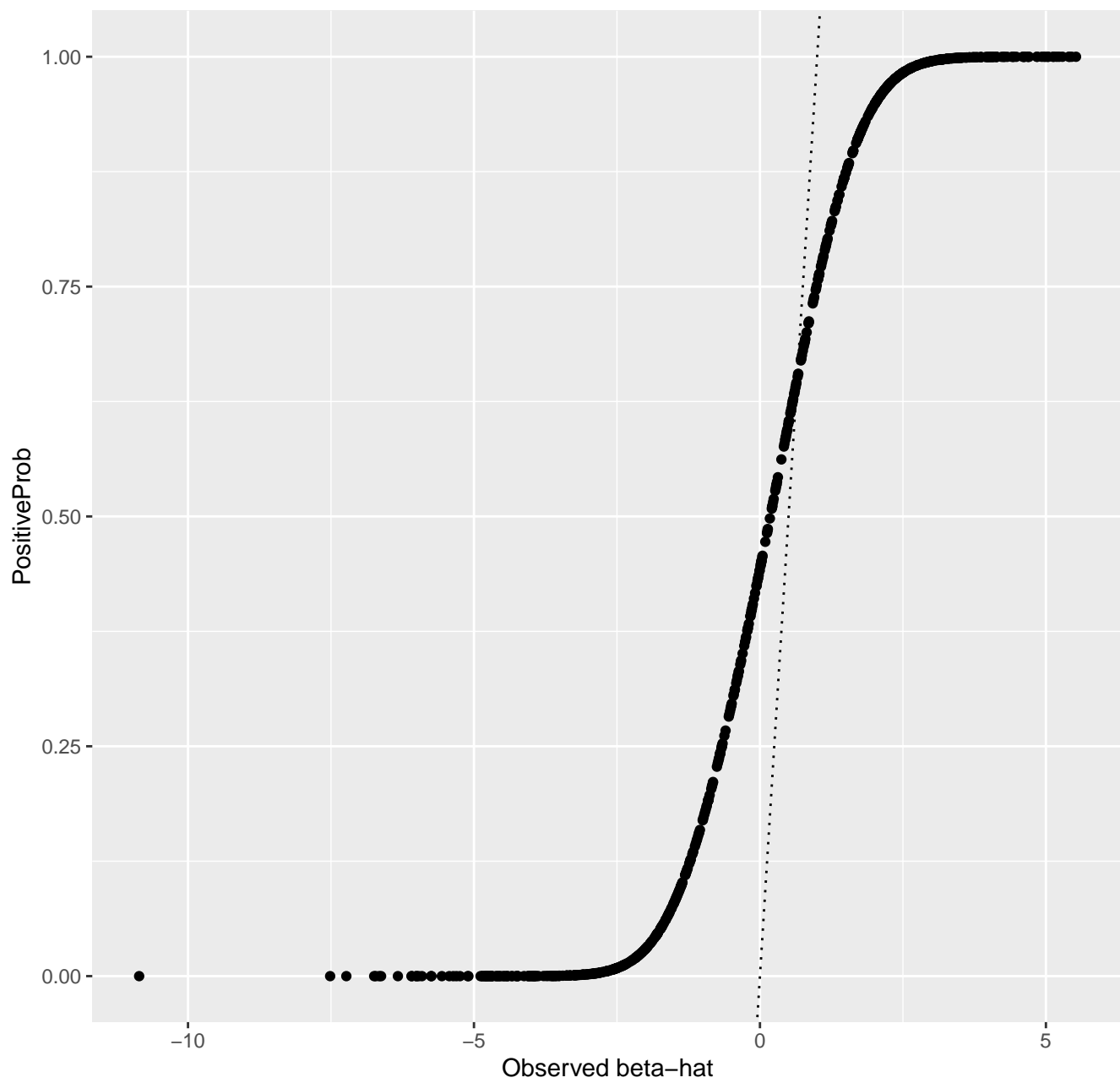
LDHA_(3939) negative



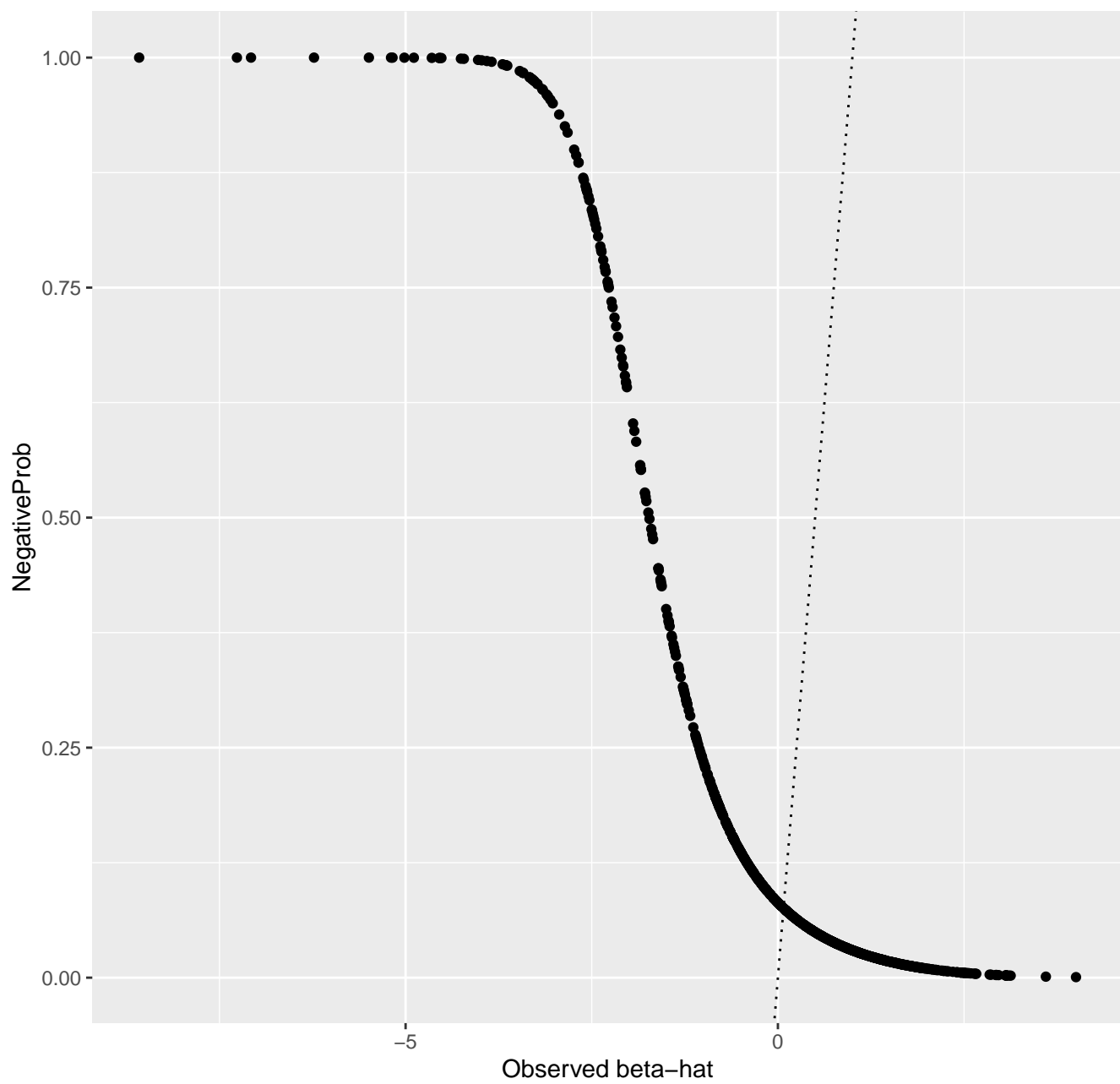
PITRM1_(10531) negative



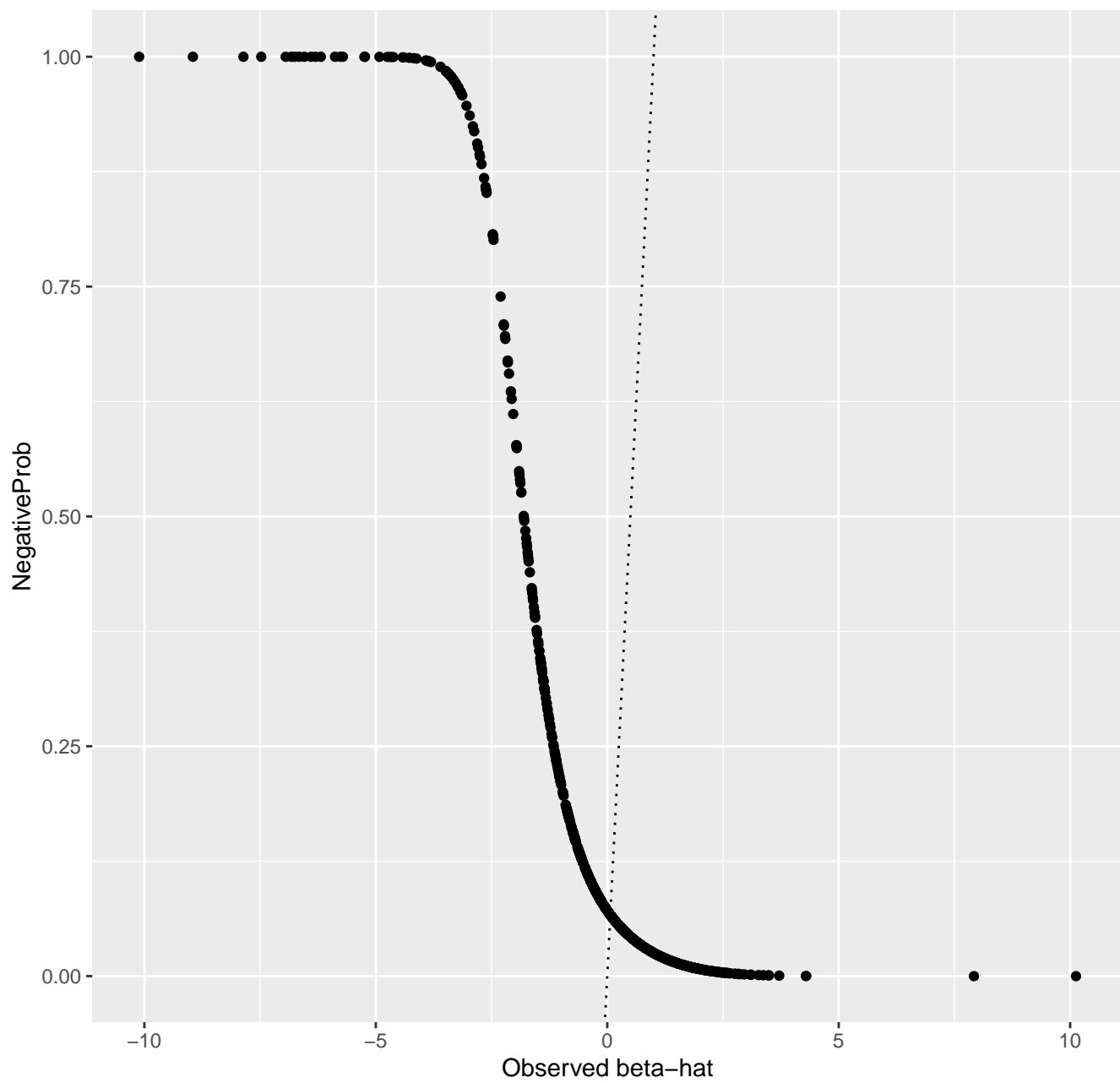
RAB6A_(5870) positive



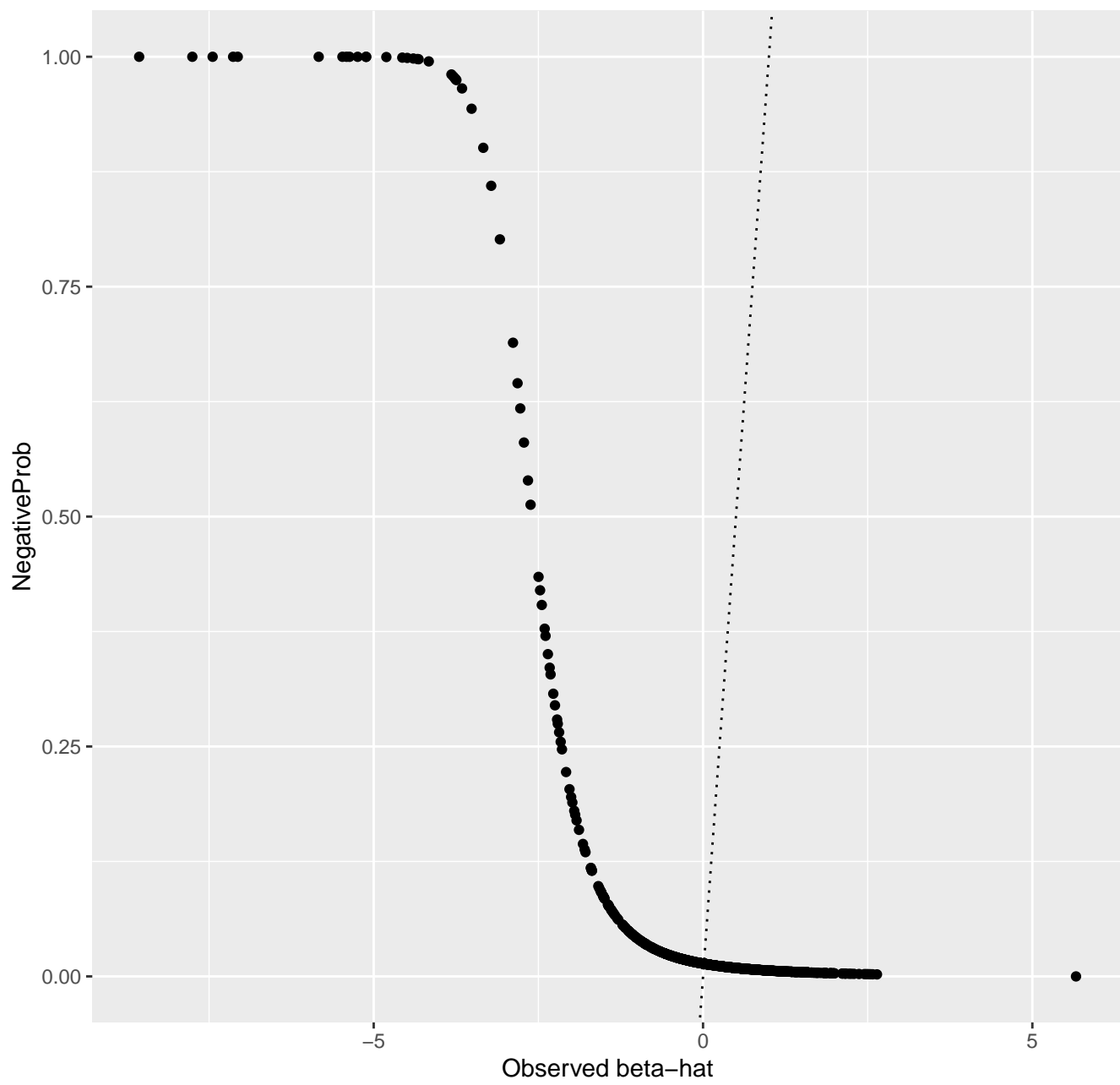
CLPB_(81570) negative



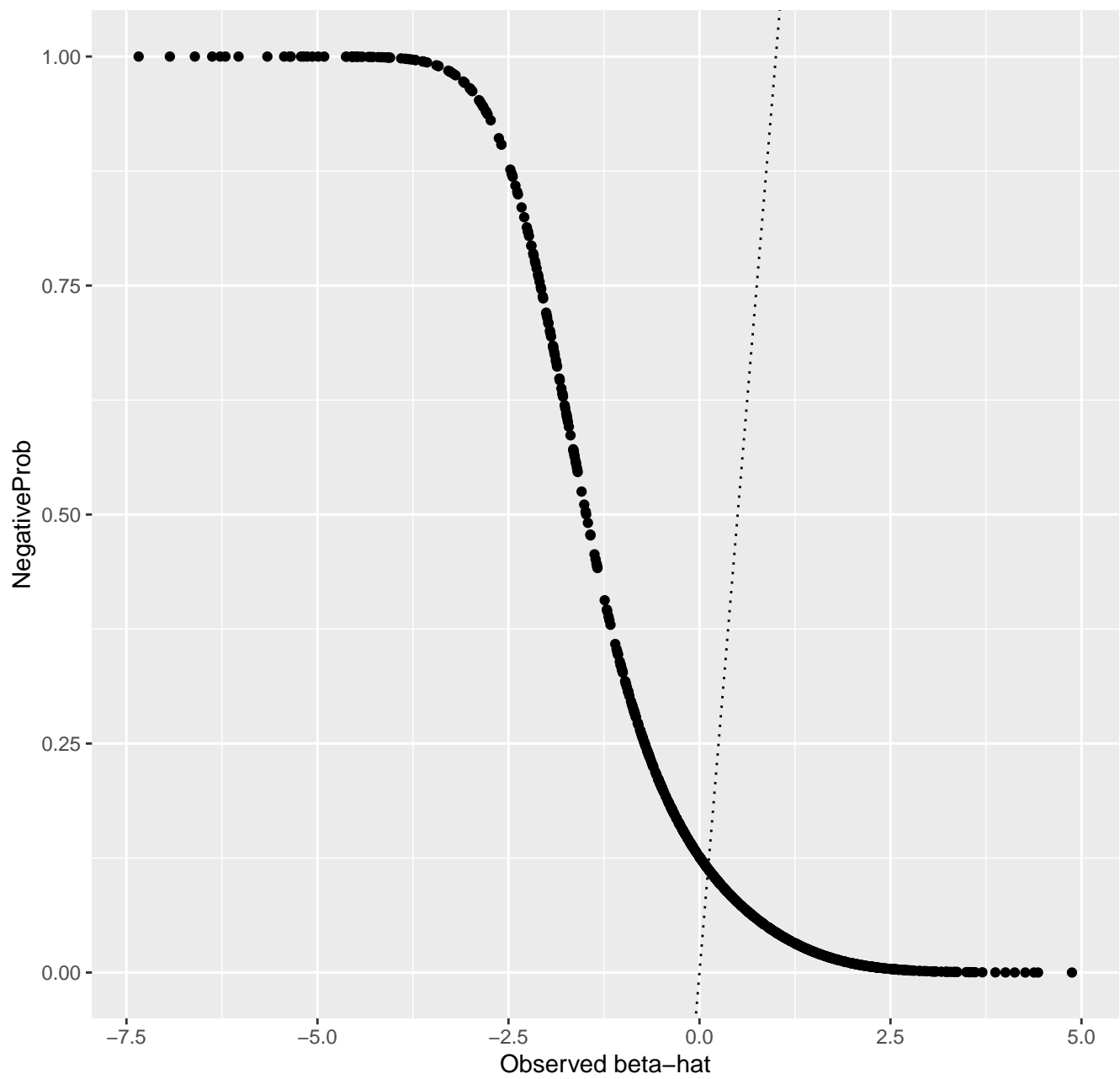
FADD_(8772) negative



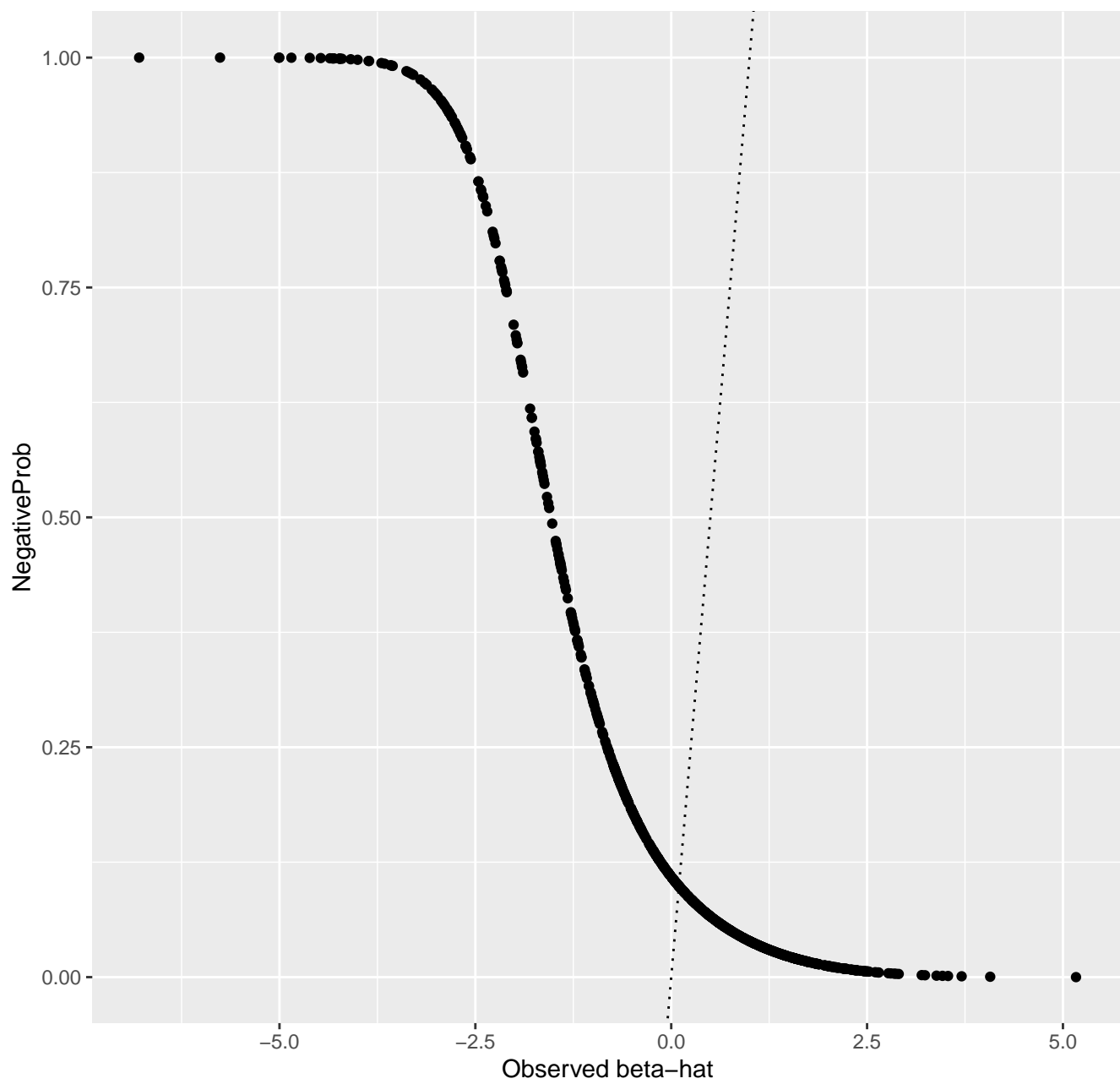
FOXA2_(3170) negative



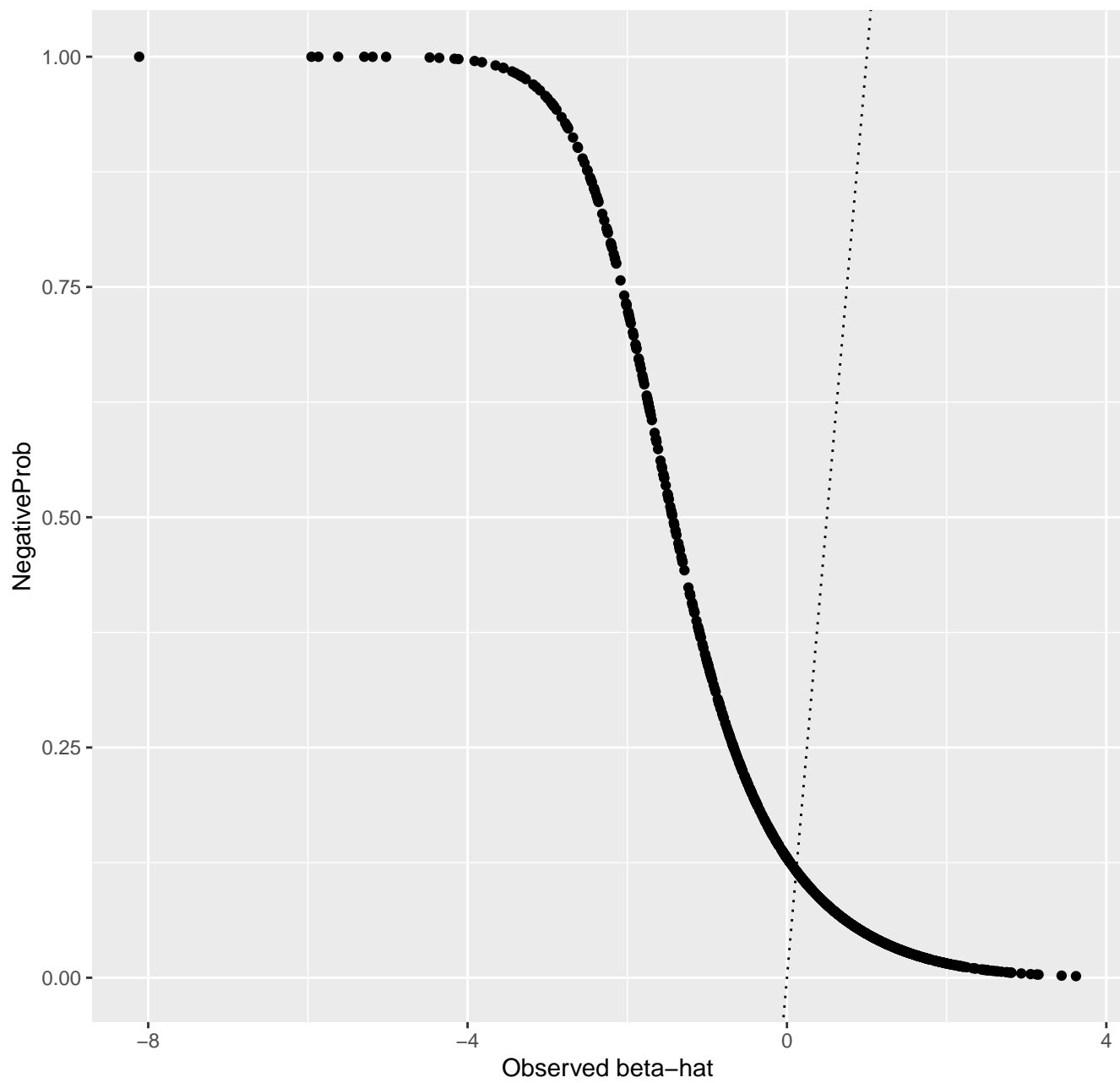
ARMC5_(79798) negative



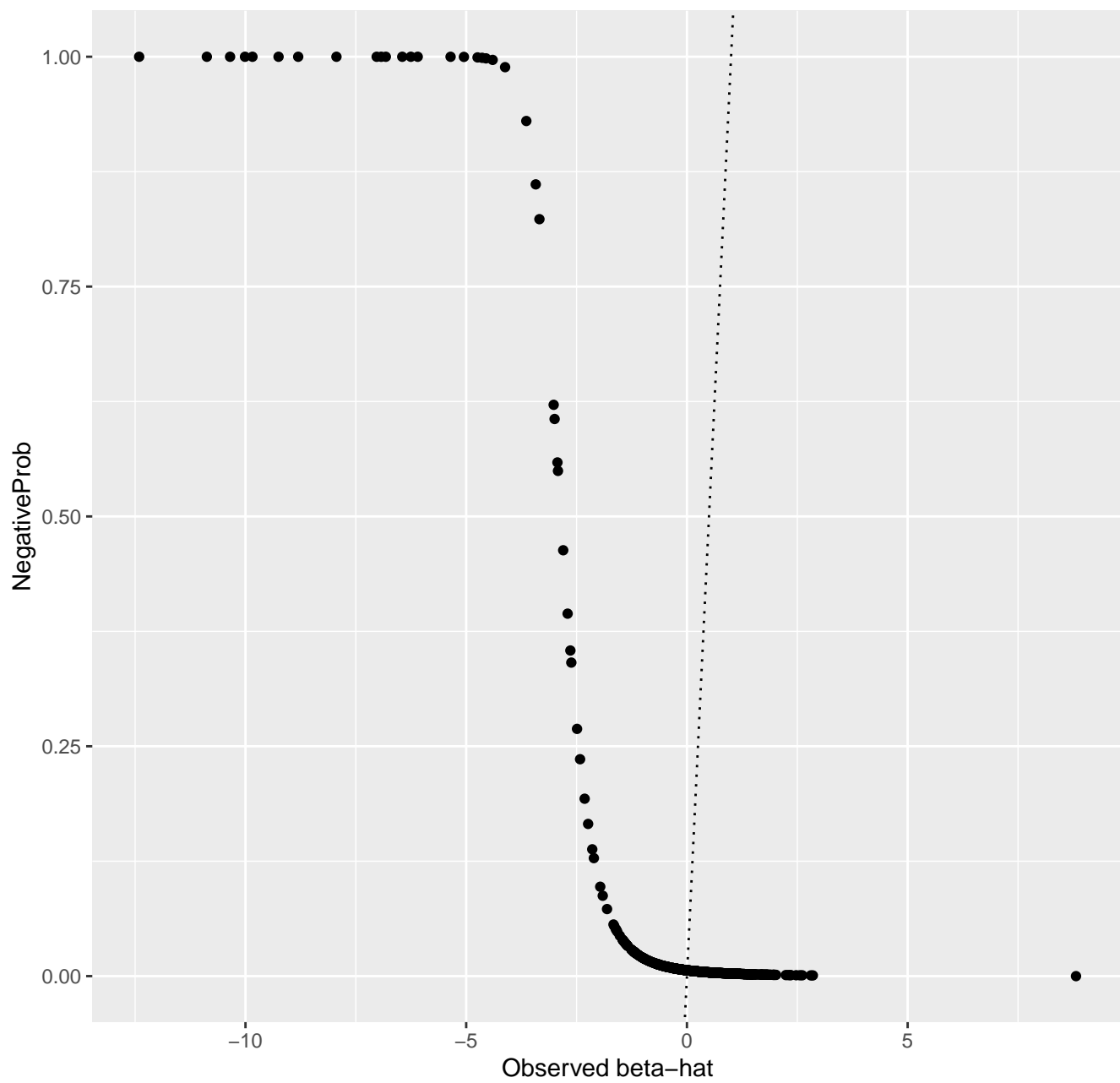
SLC38A2_(54407) negative



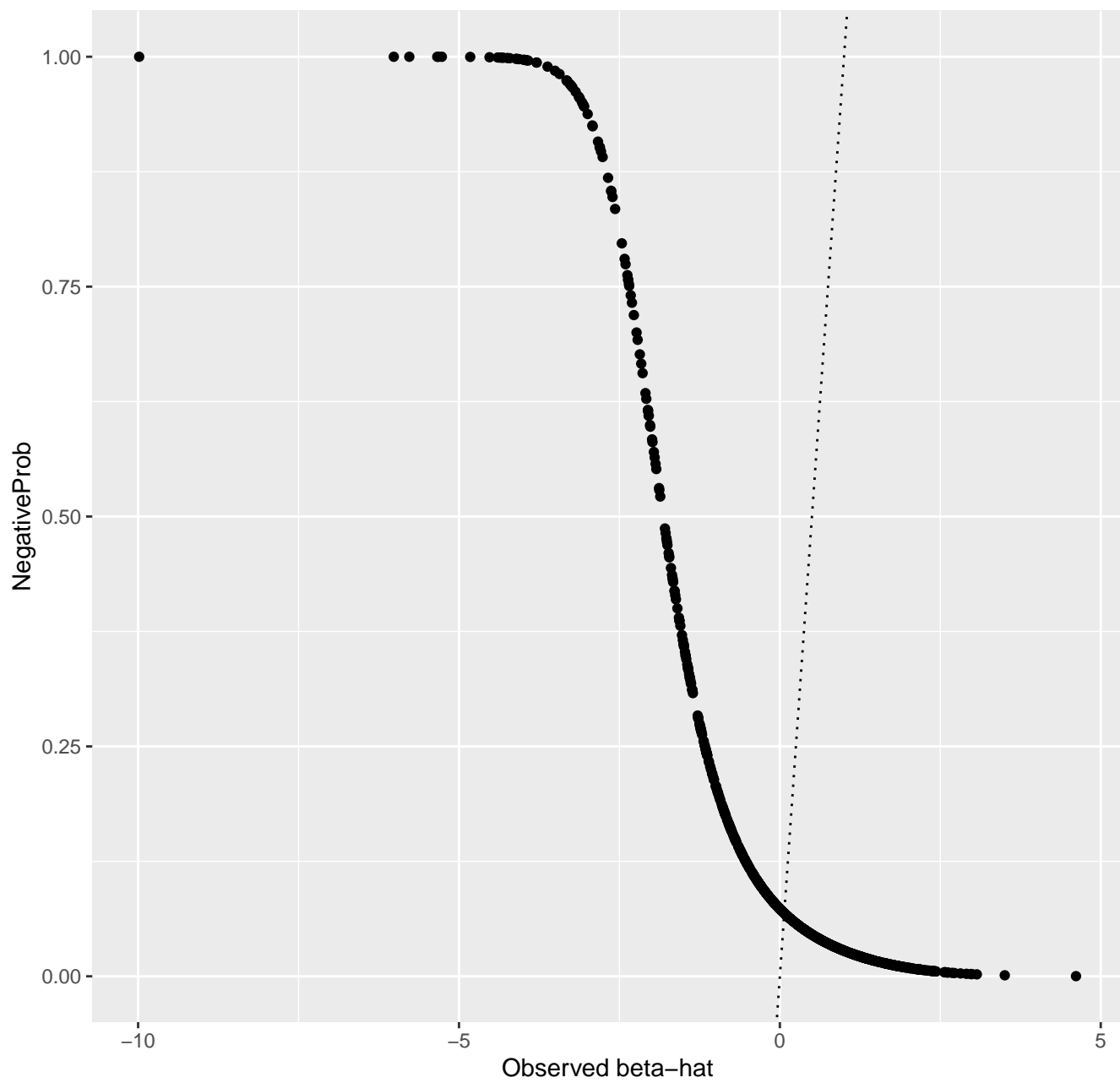
IREB2_(3658) negative



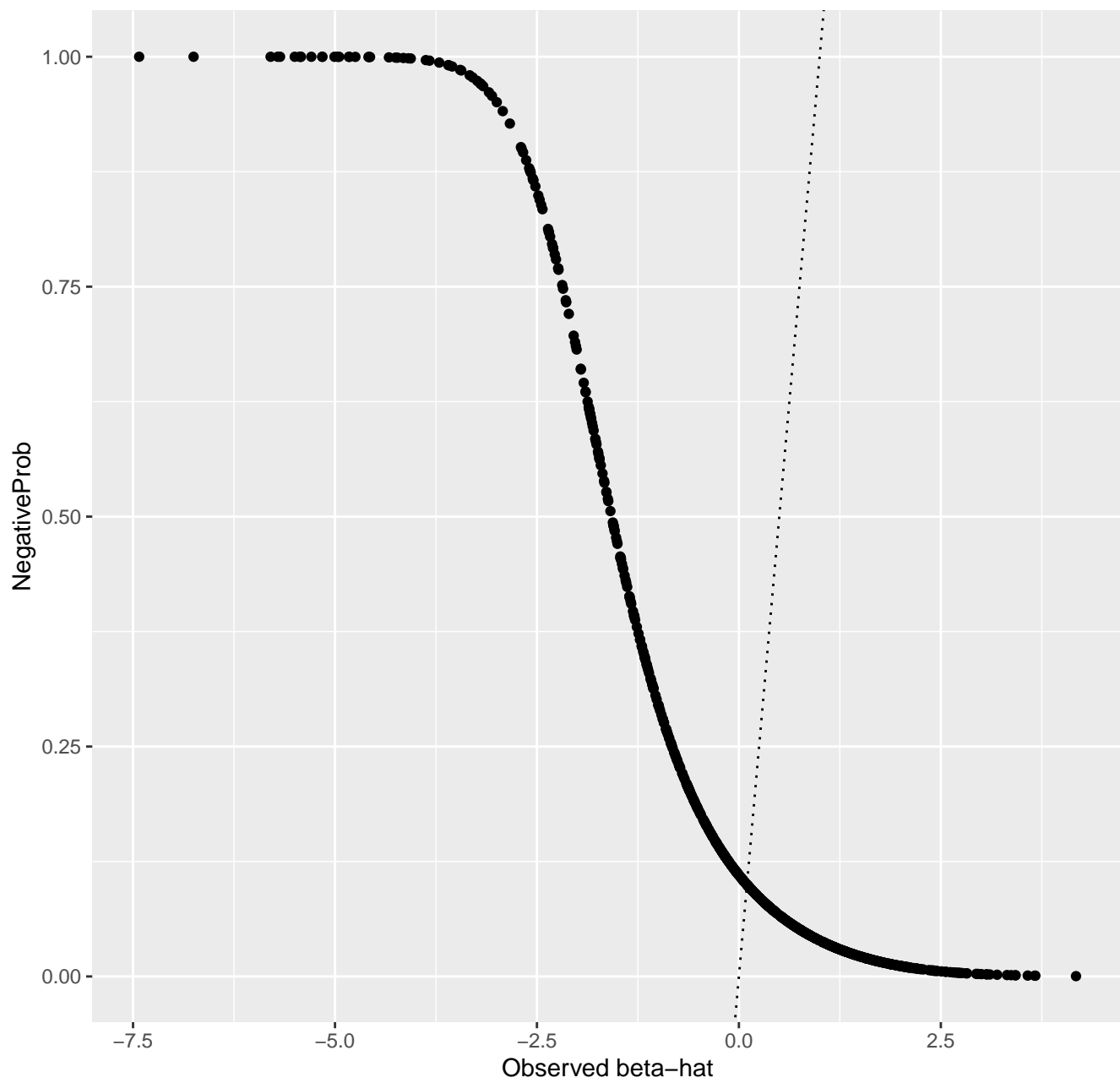
MEF2C_(4208) negative



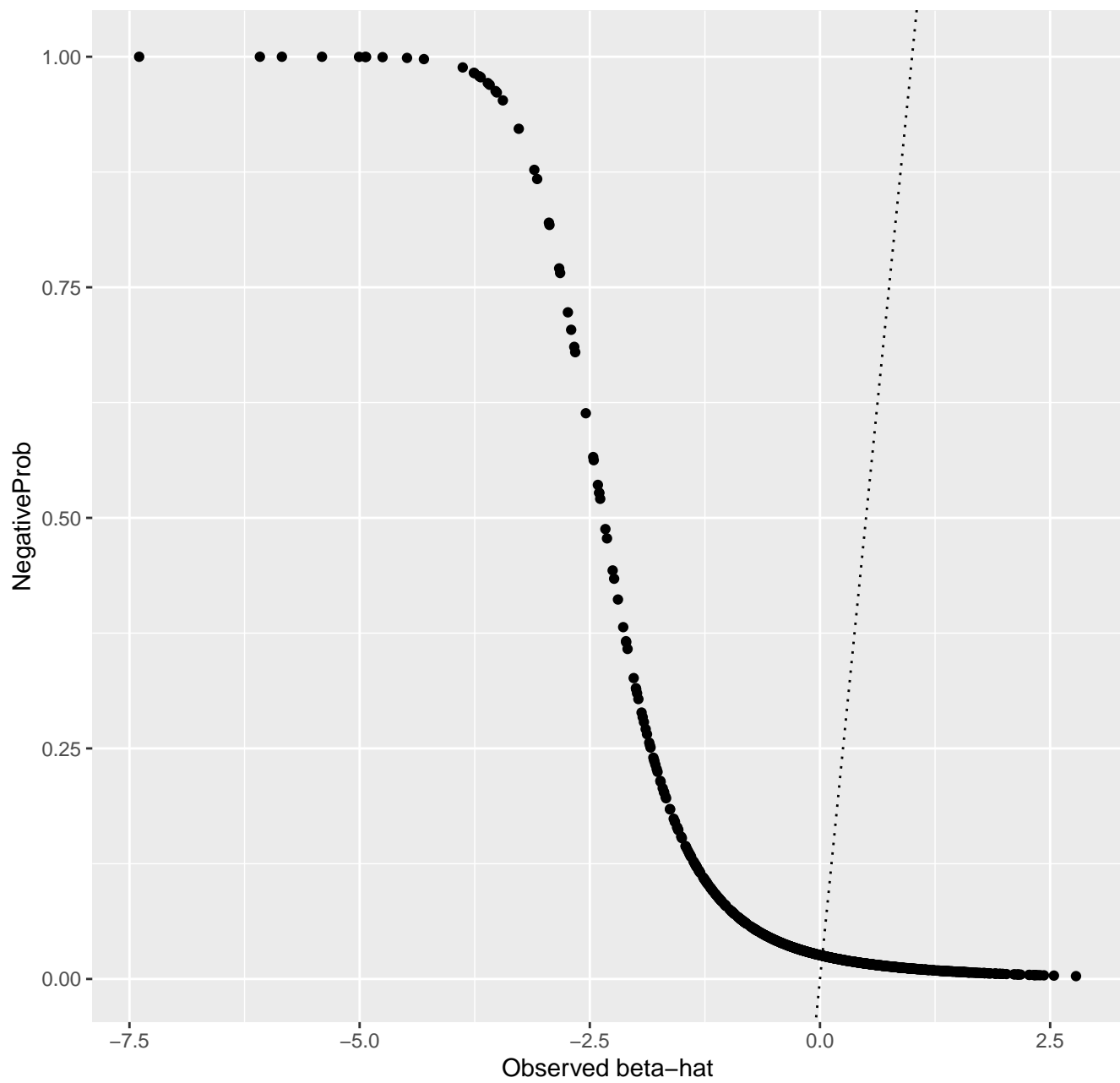
UBE3D_(90025) negative



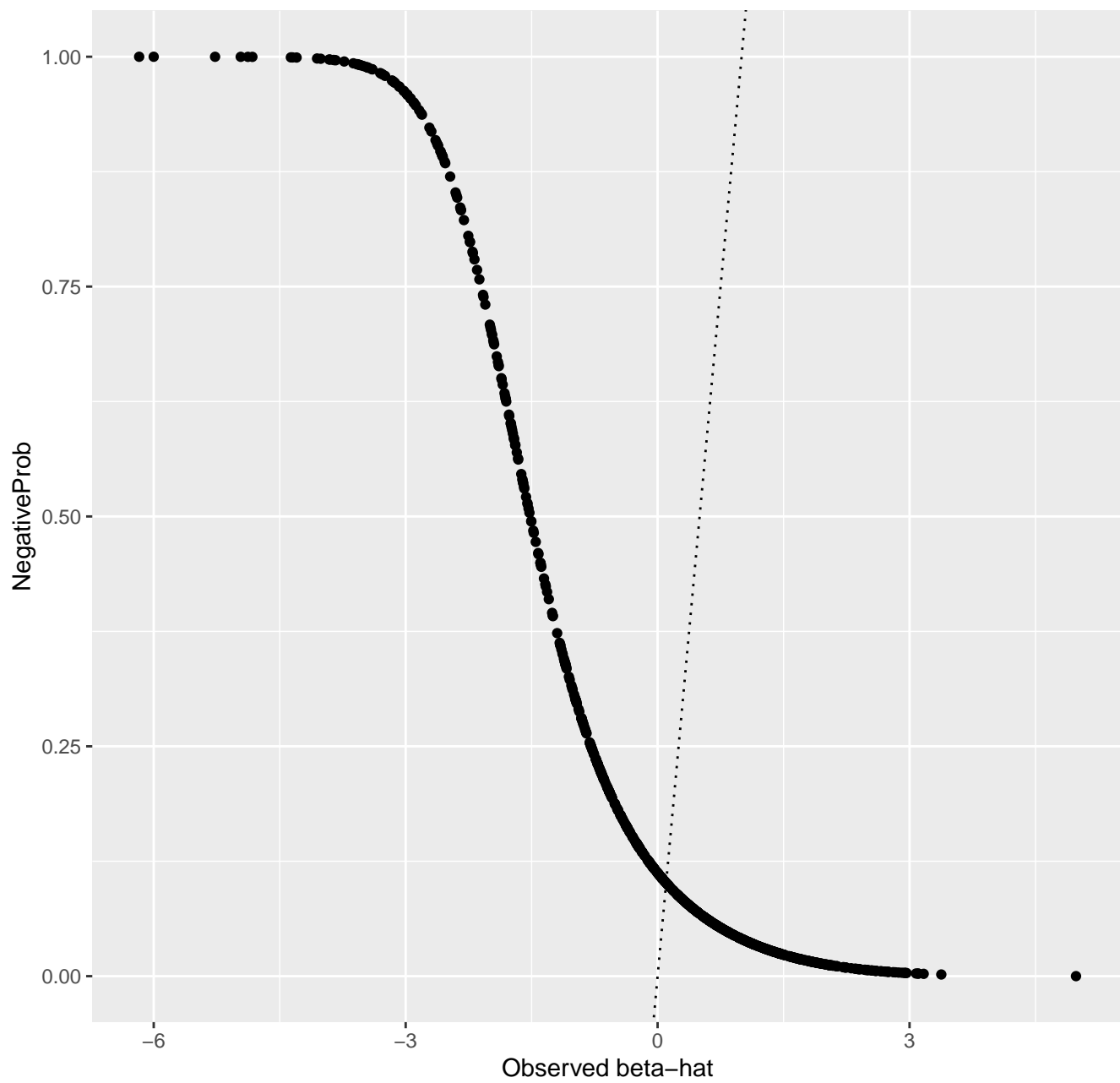
TRAPPC13_(80006) negative



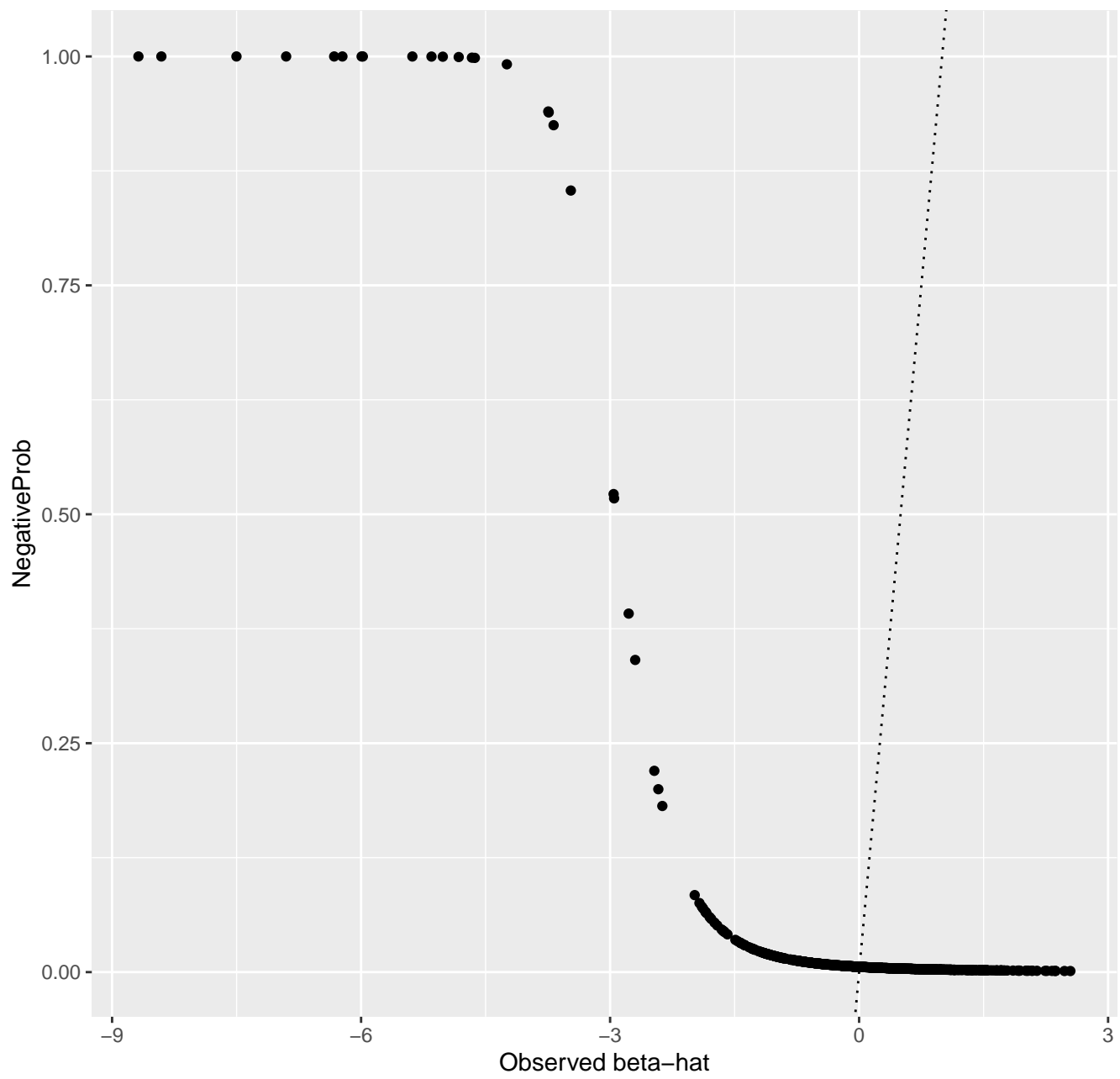
DERL1_(79139) negative



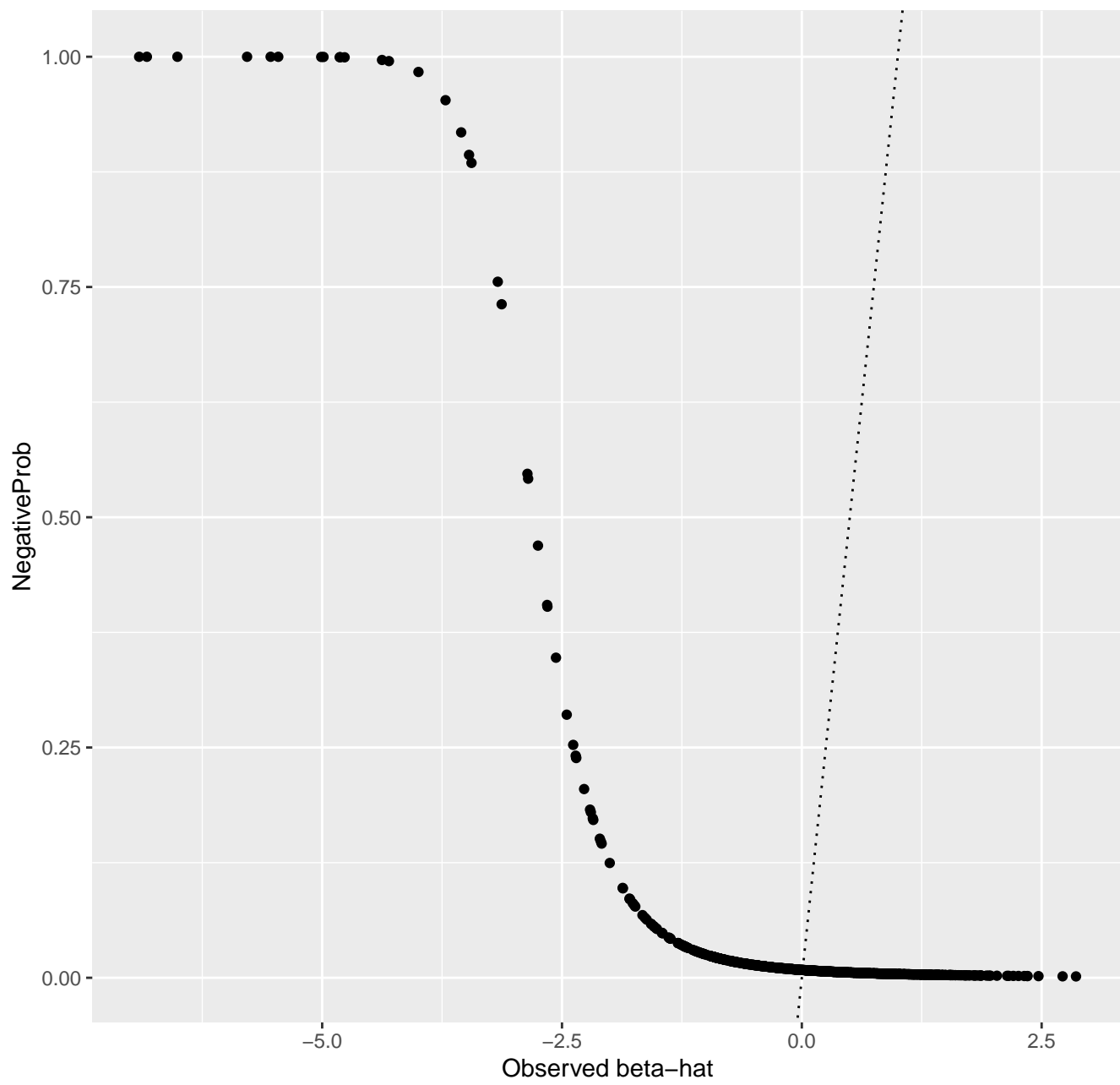
EXT2_(2132) negative



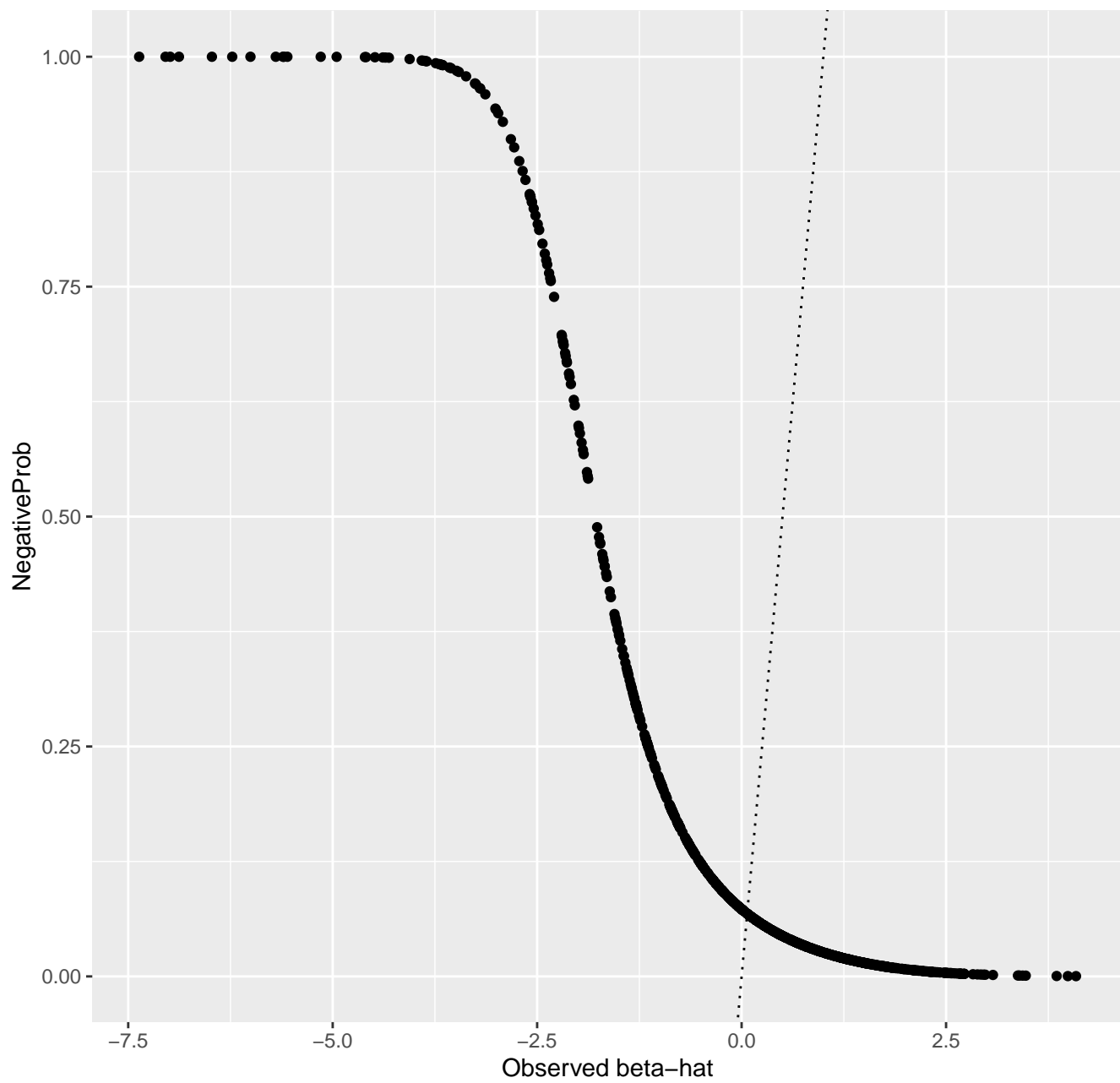
LMO2_(4005) negative



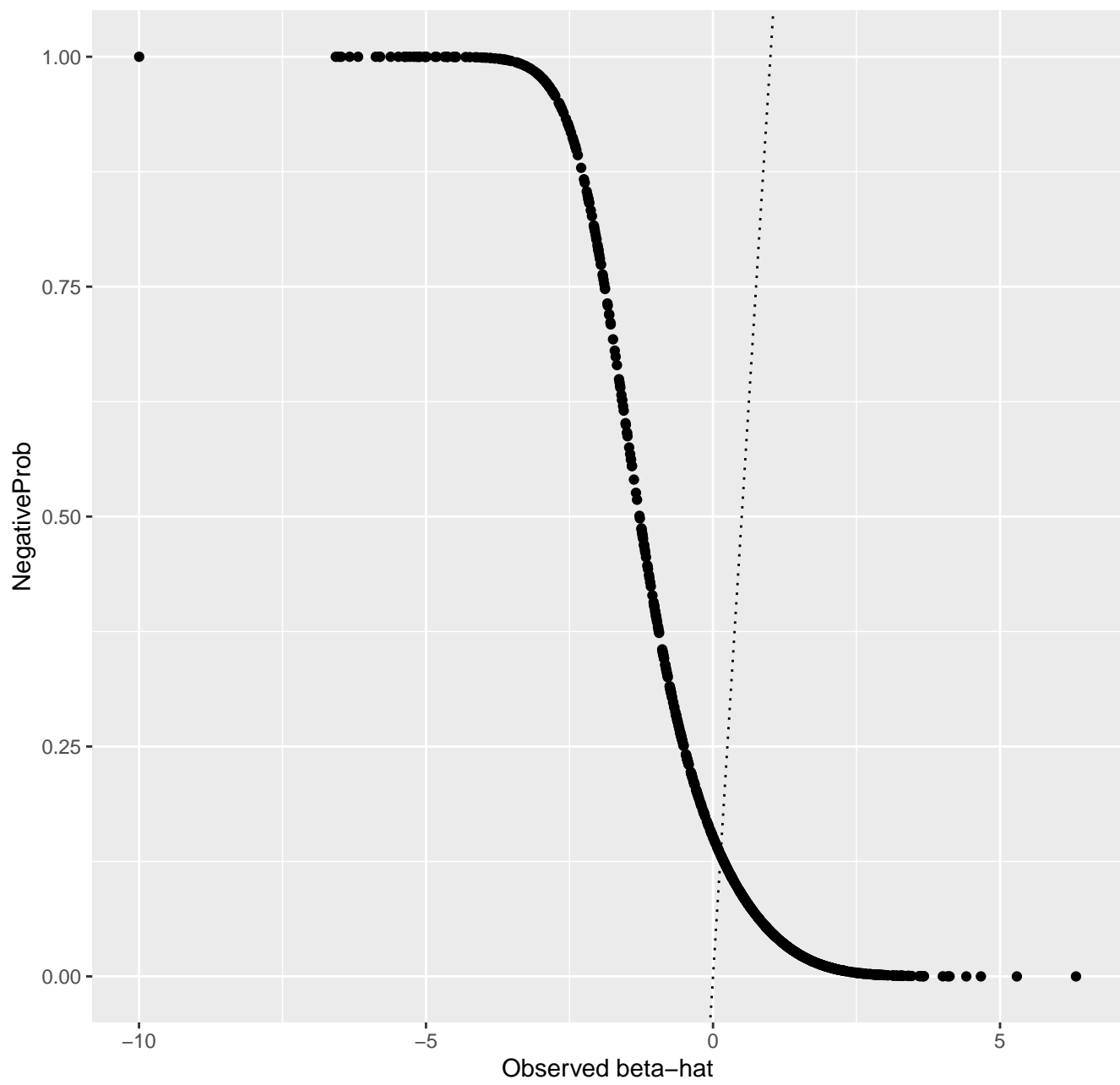
LRP8_(7804) negative



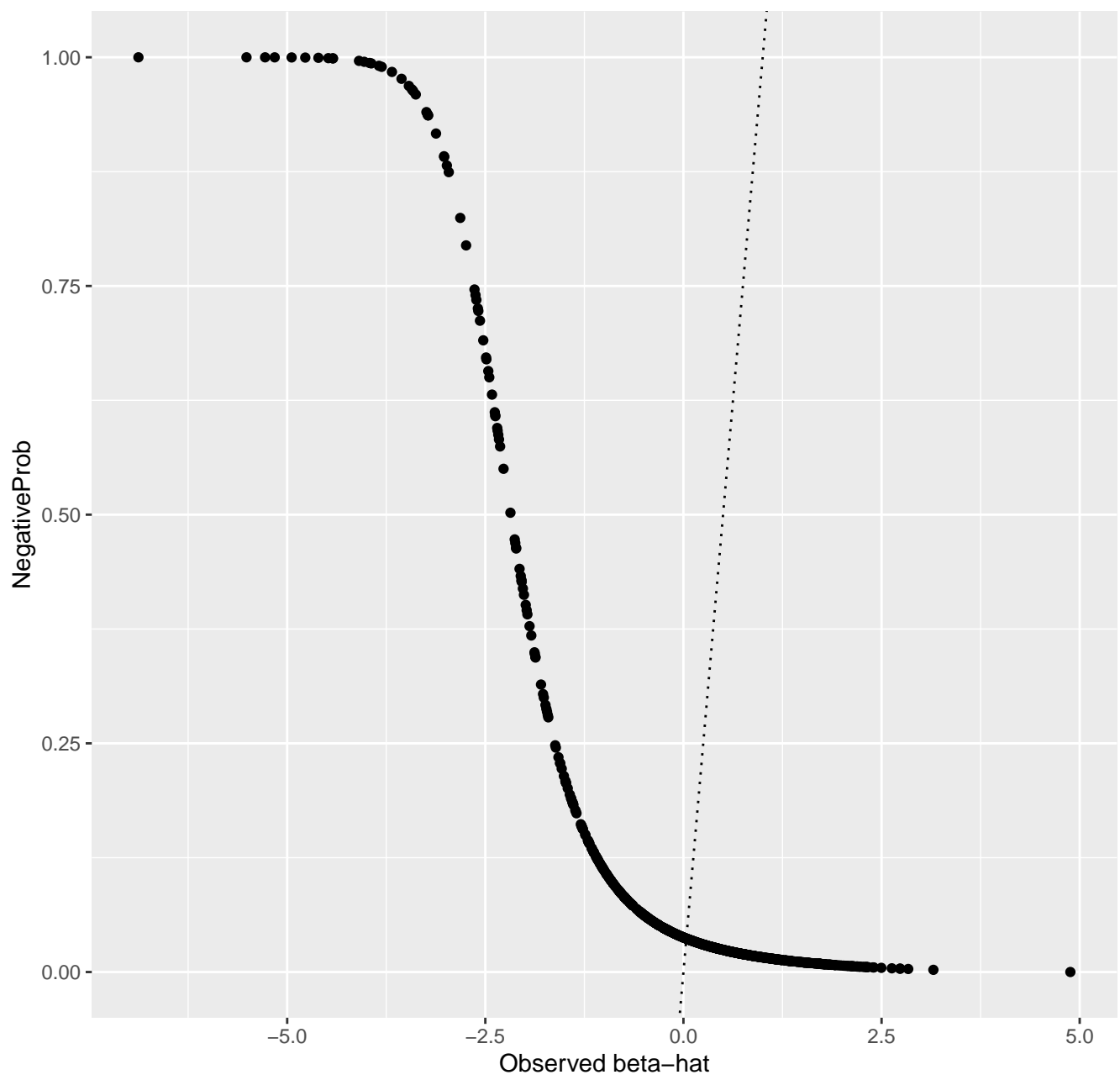
MTF1_(4520) negative



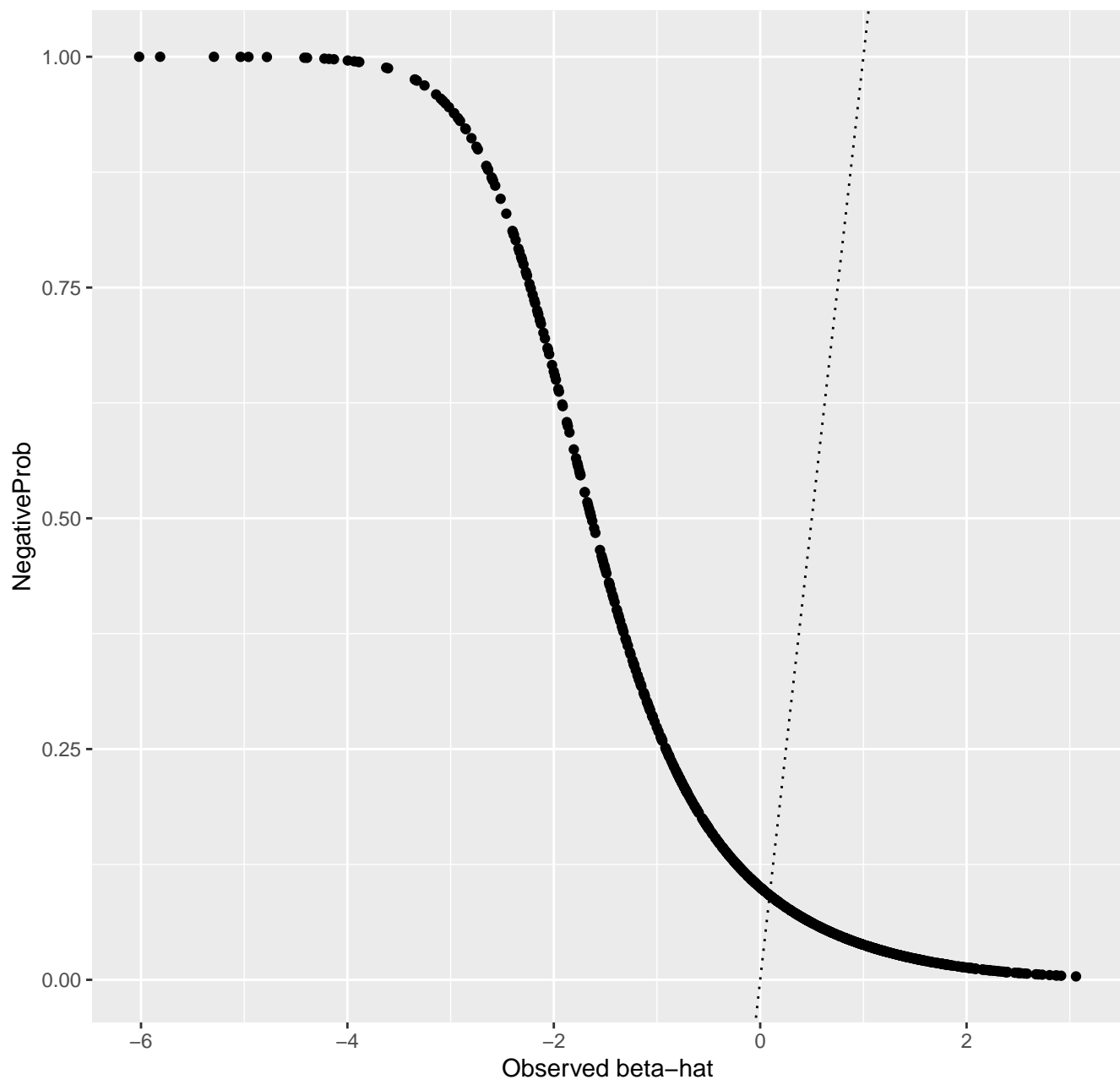
TEAD1_(7003) negative



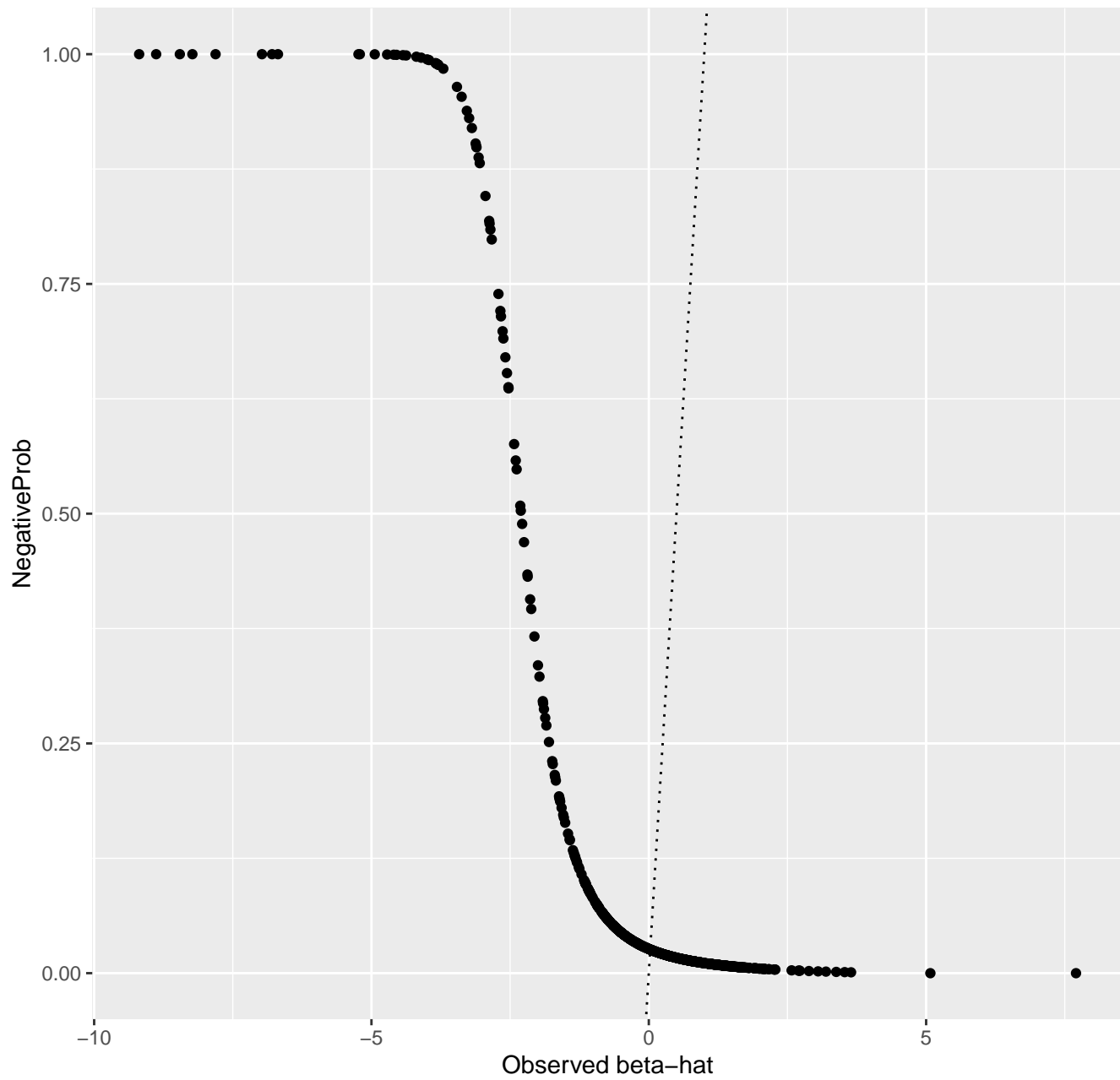
KIF2A_(3796) negative



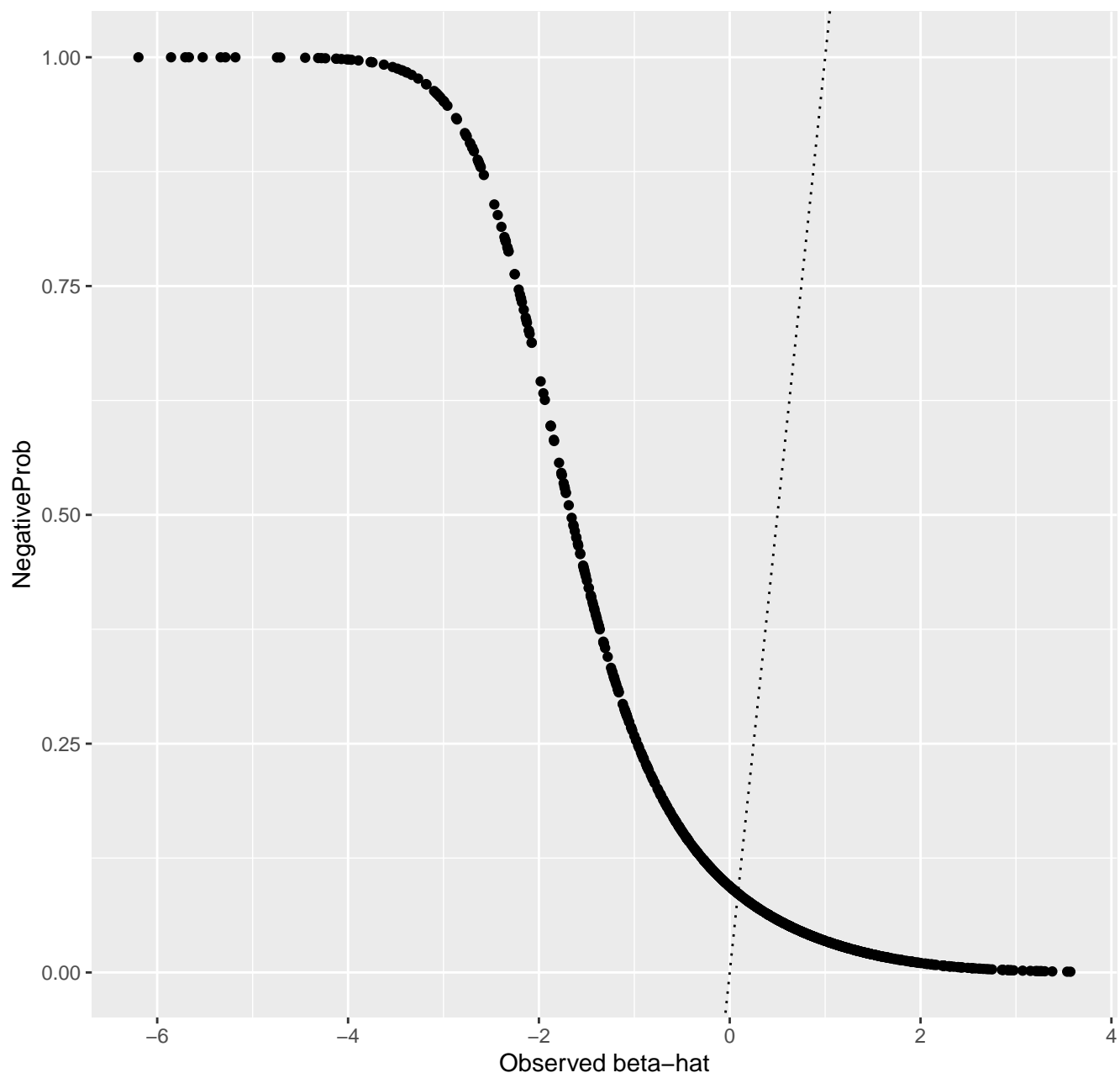
ZNF511_(118472) negative



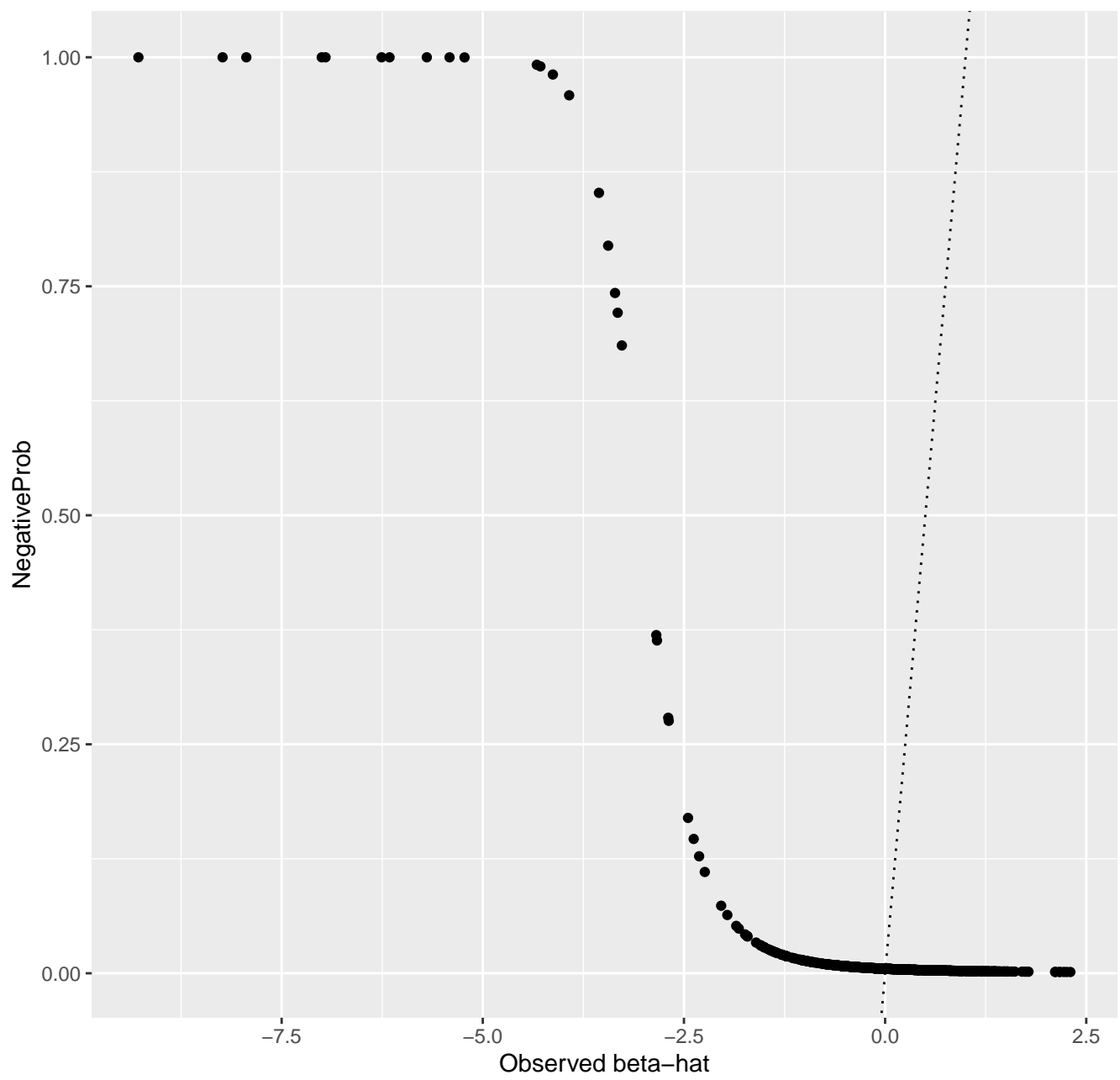
SMCHD1_(23347) negative



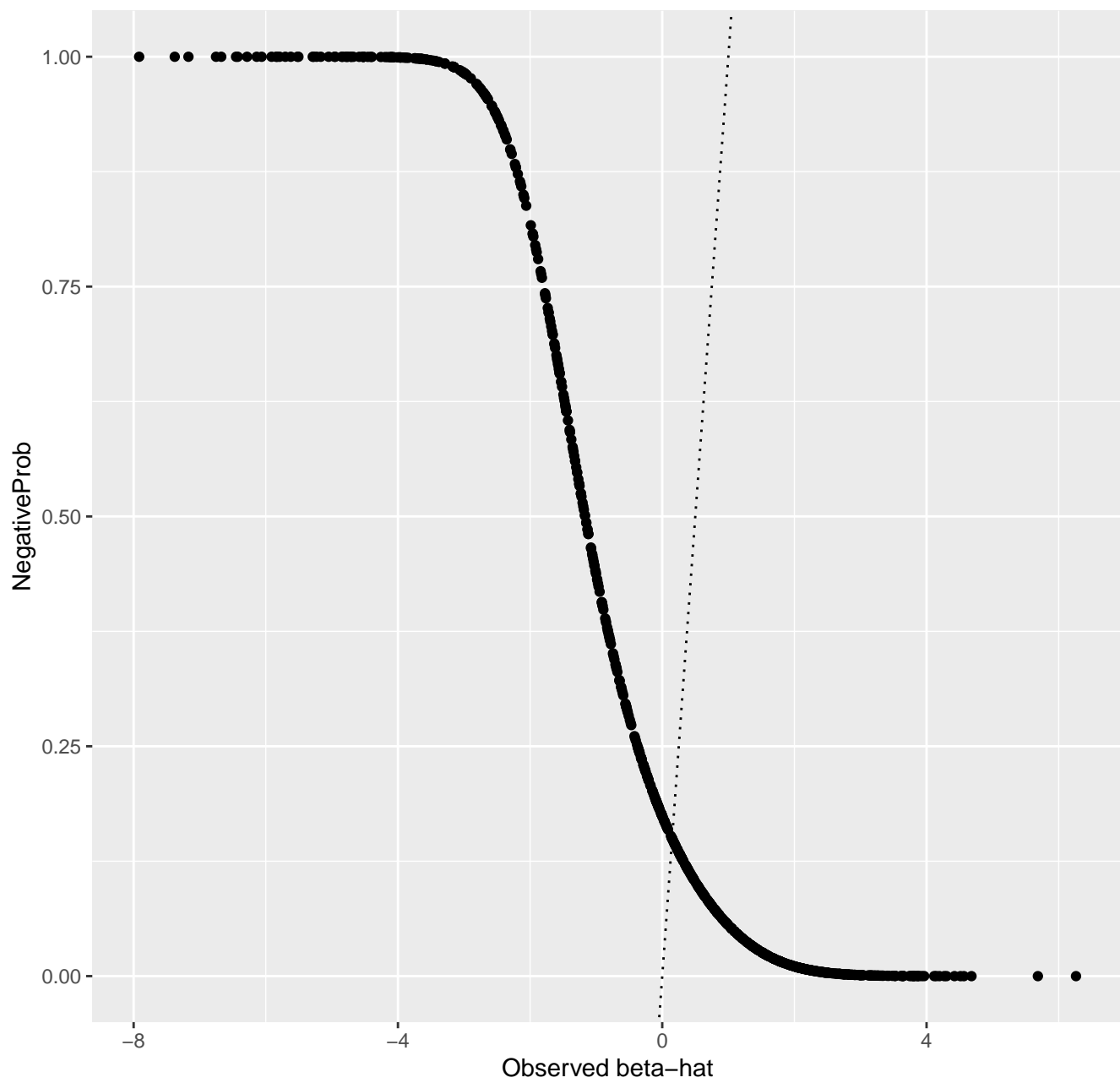
PPM1D_(8493) negative



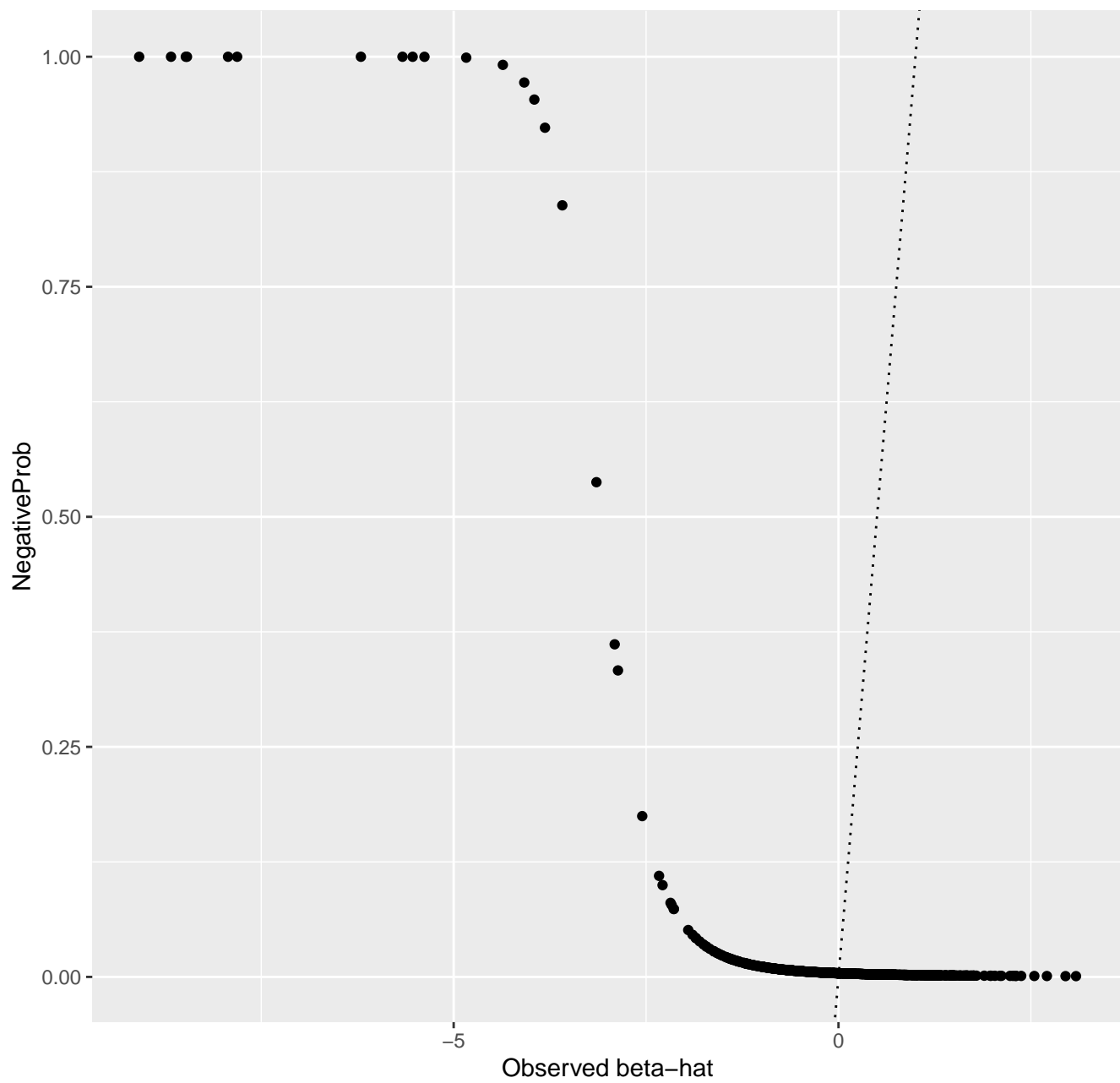
FAM126B_(285172) negative



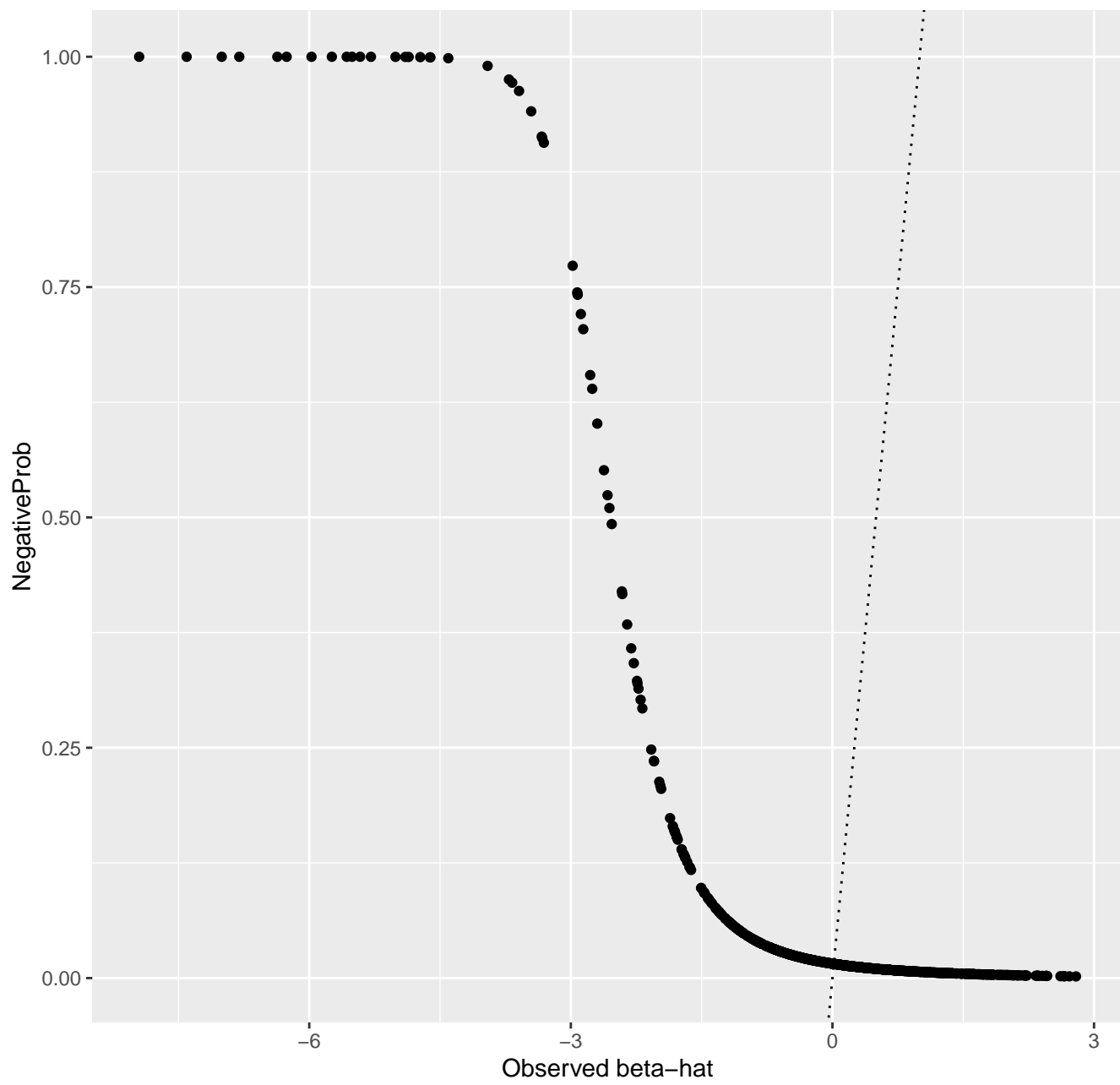
ARHGEF7_(8874) negative



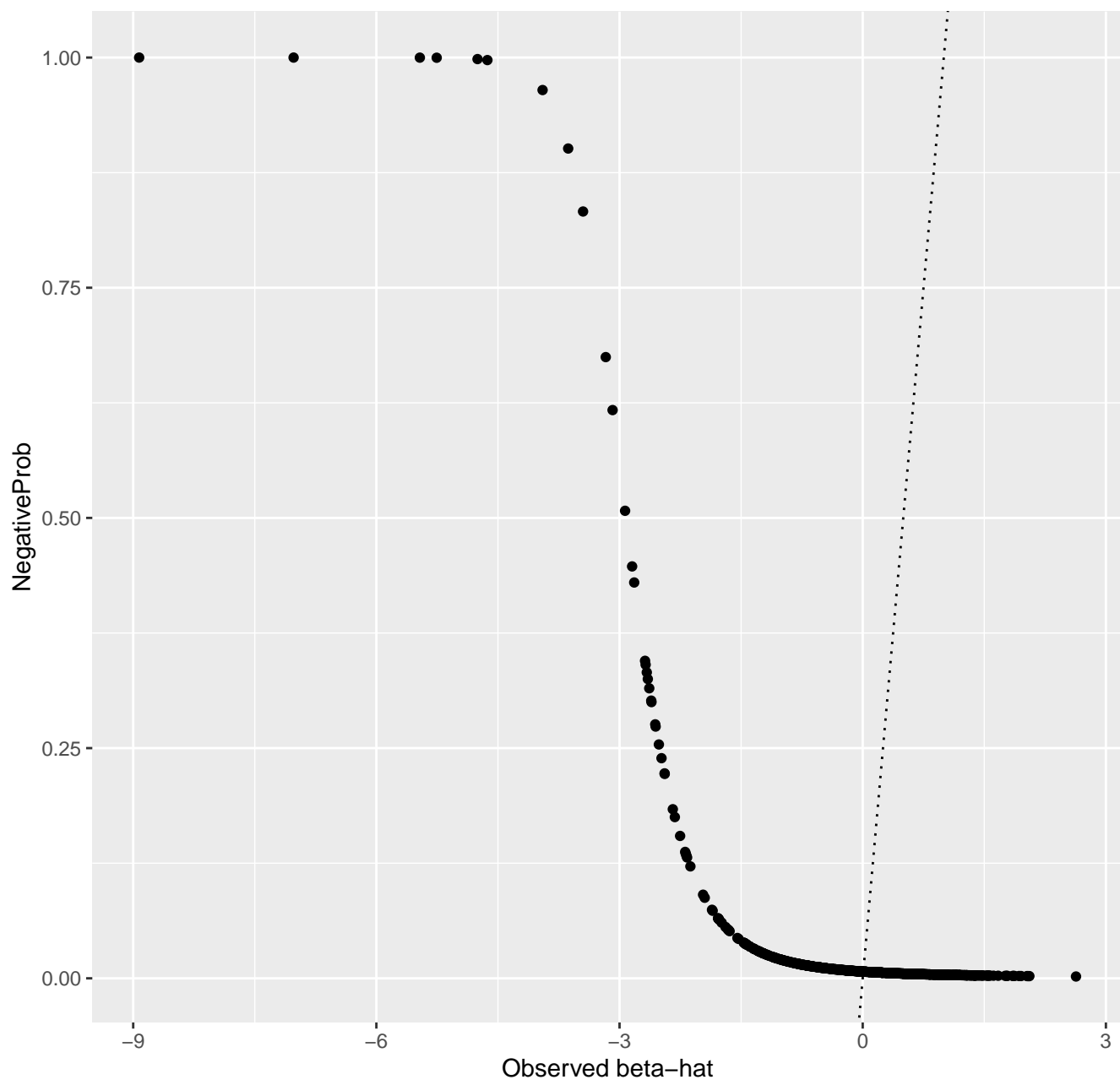
ISL1_(3670) negative



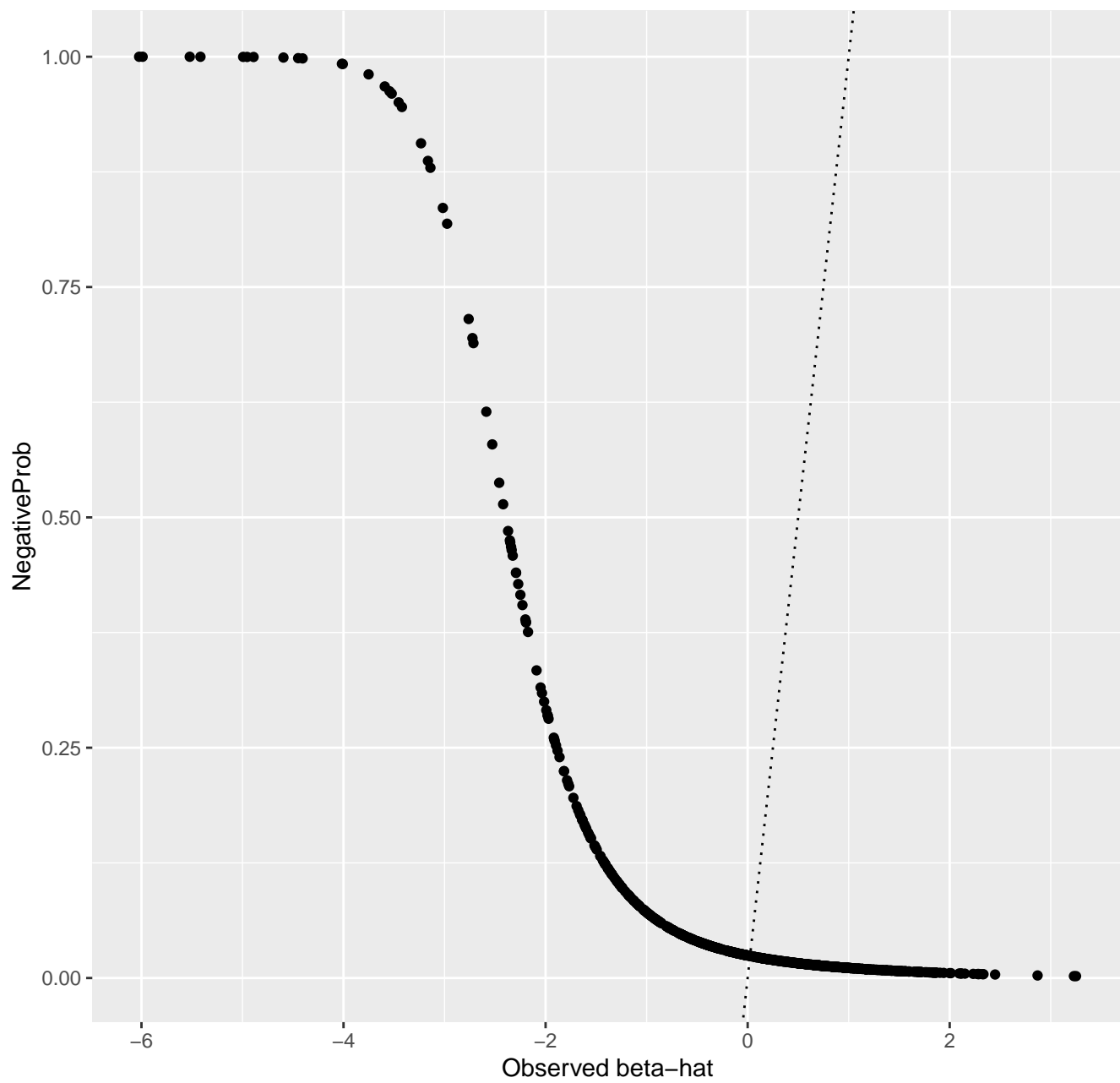
GATA3_(2625) negative



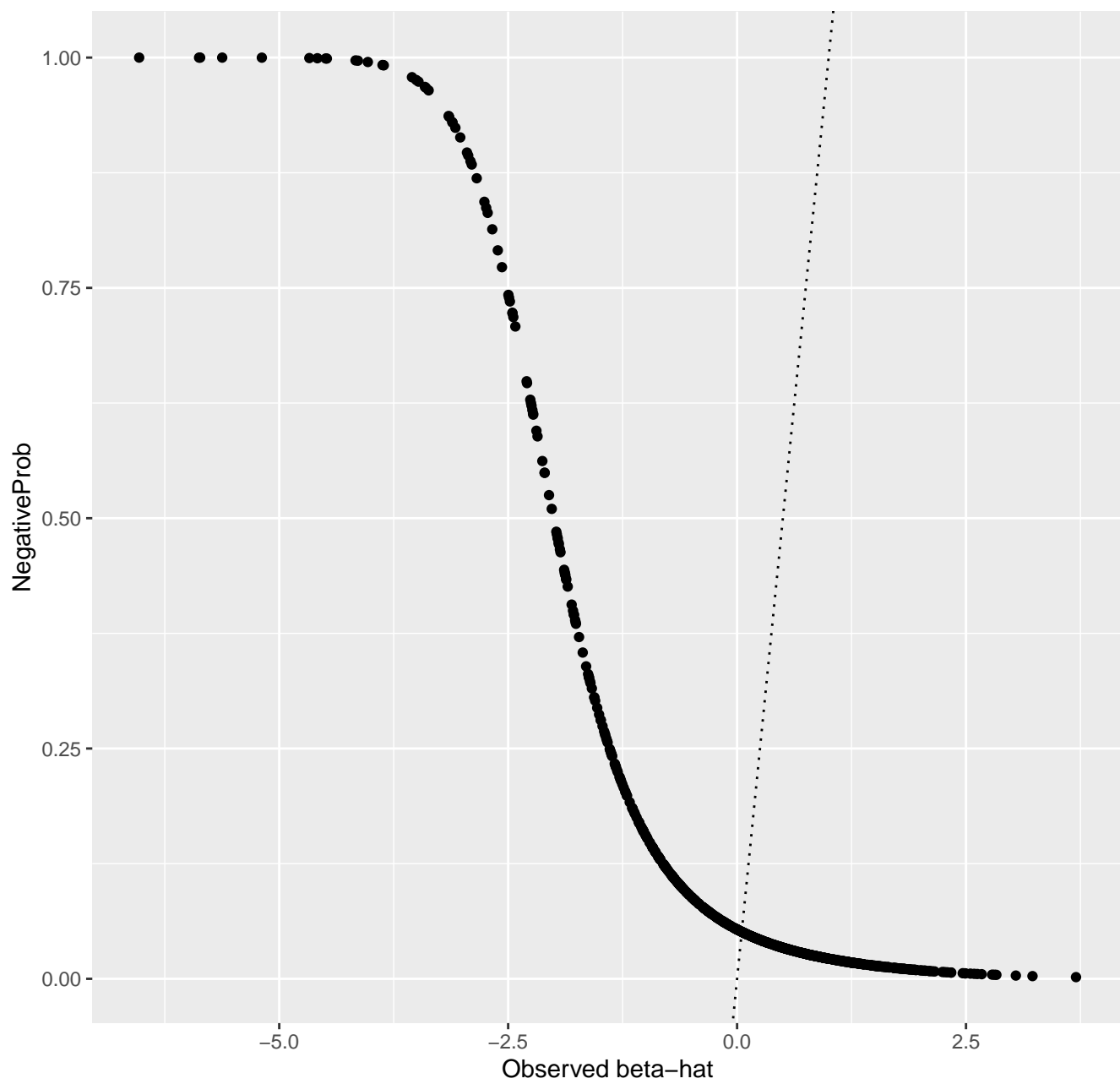
FAM83D_(81610) negative



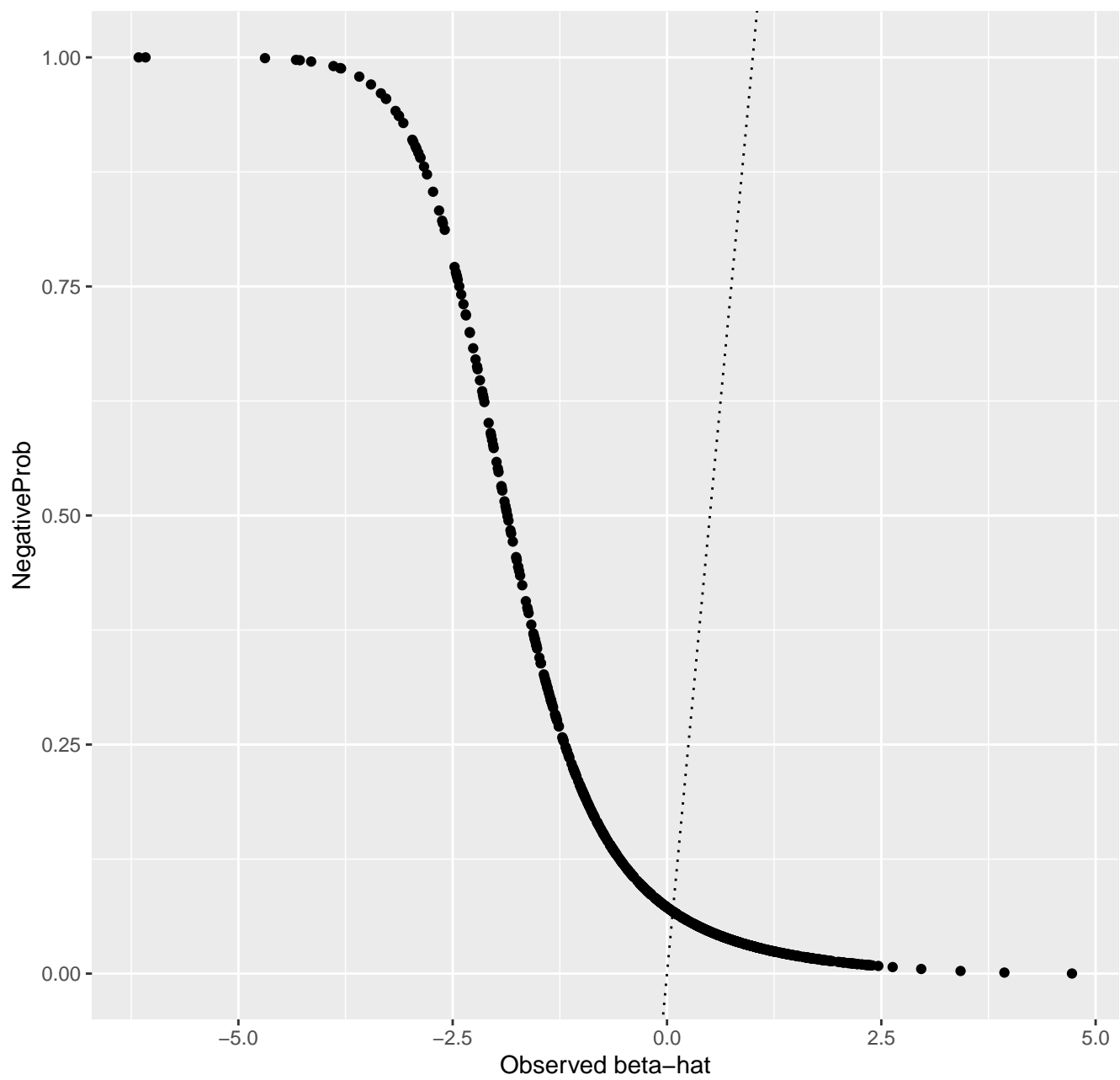
BCAP31_(10134) negative



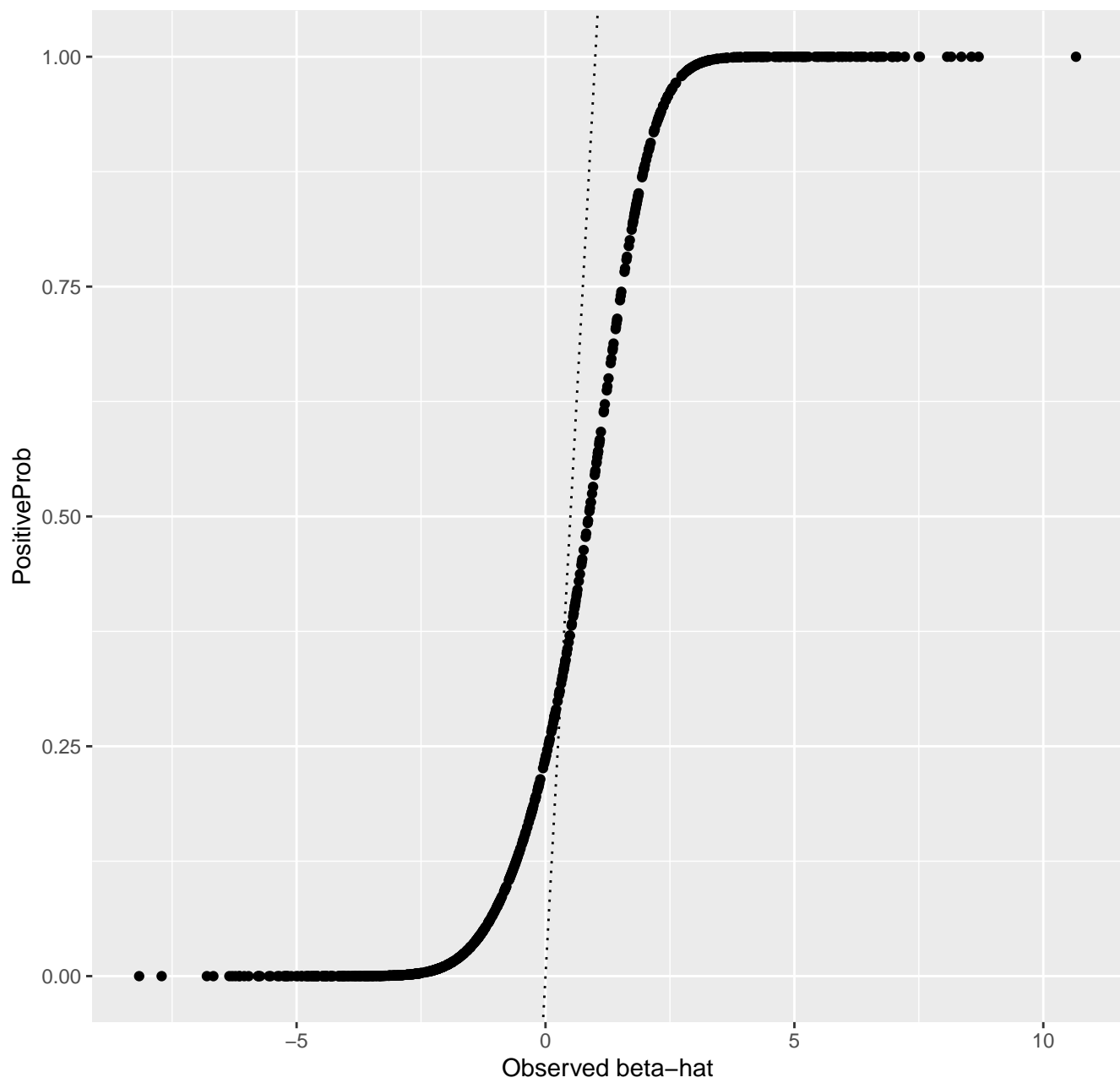
ASF1A_(25842) negative



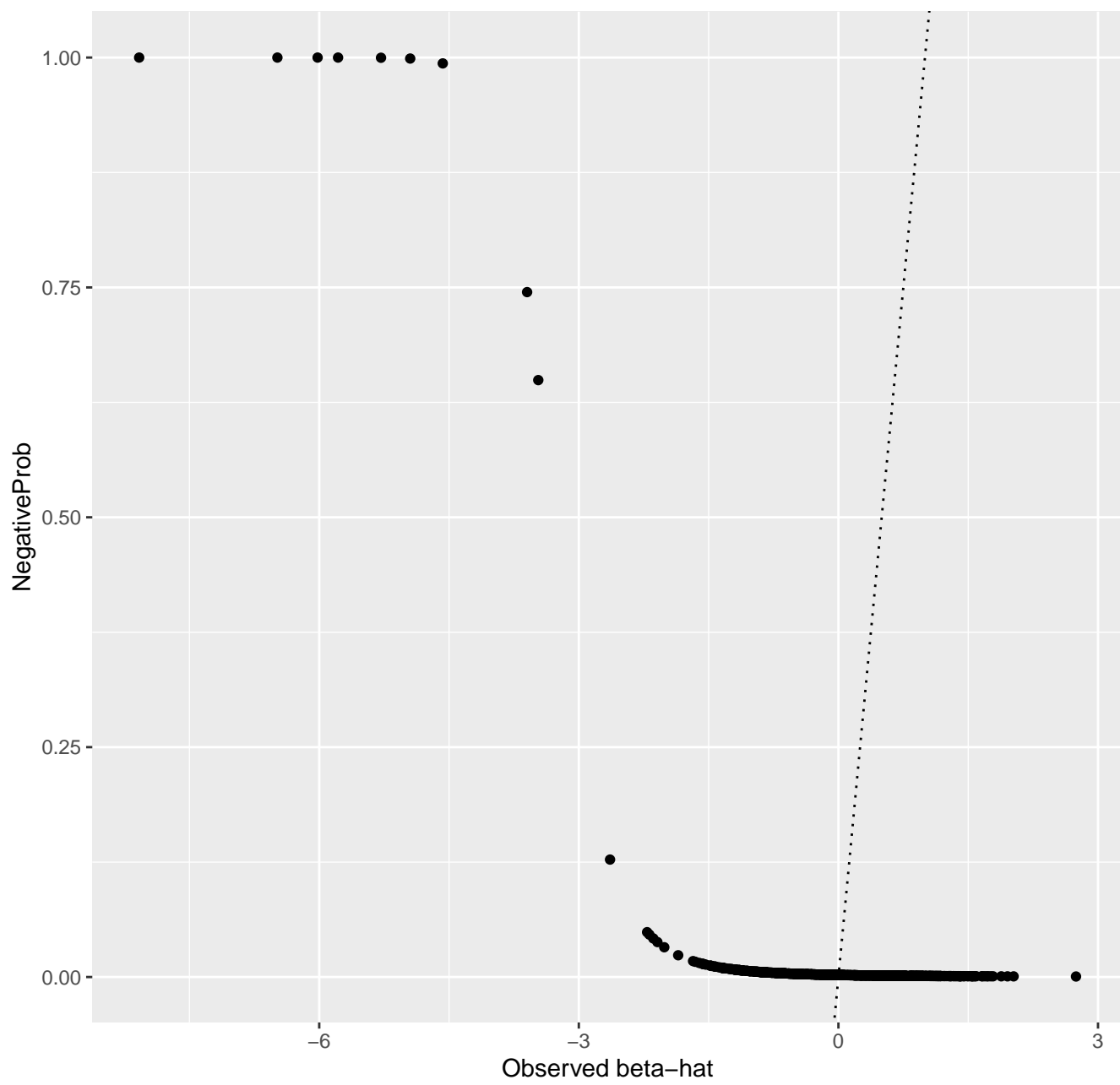
AP1M1_(8907) negative



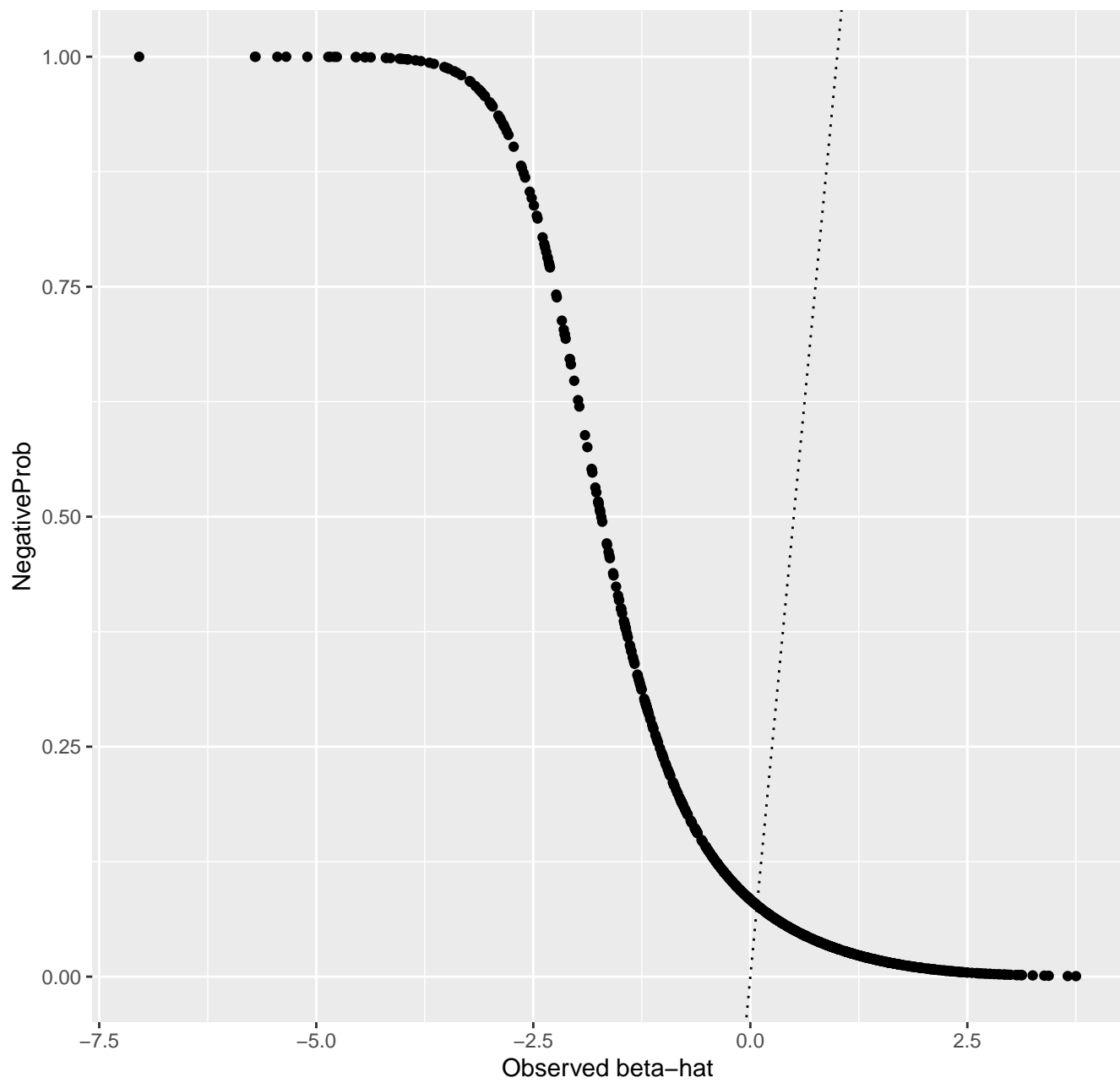
GRB2_(2885) positive



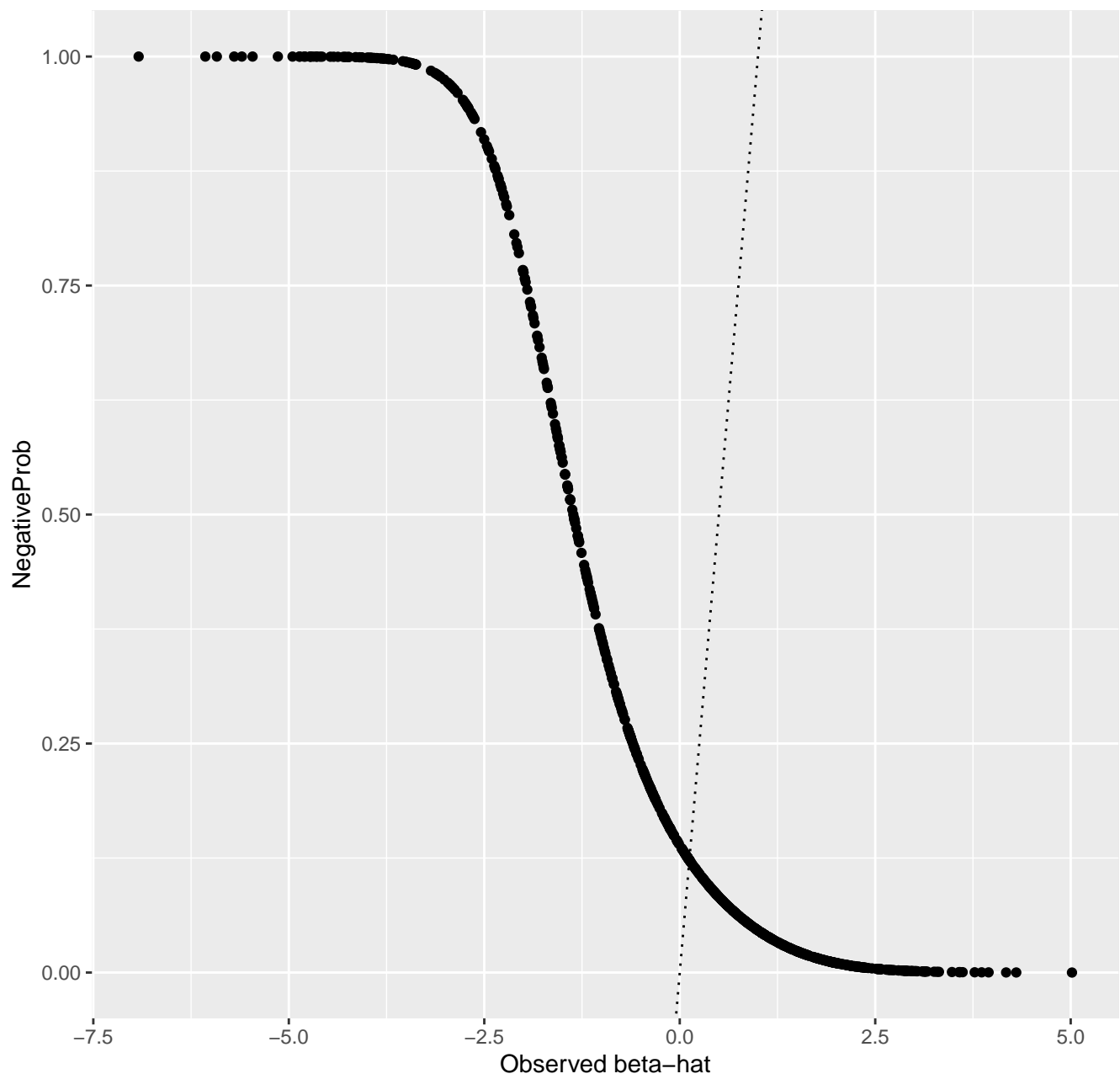
STAT5B_(6777) negative



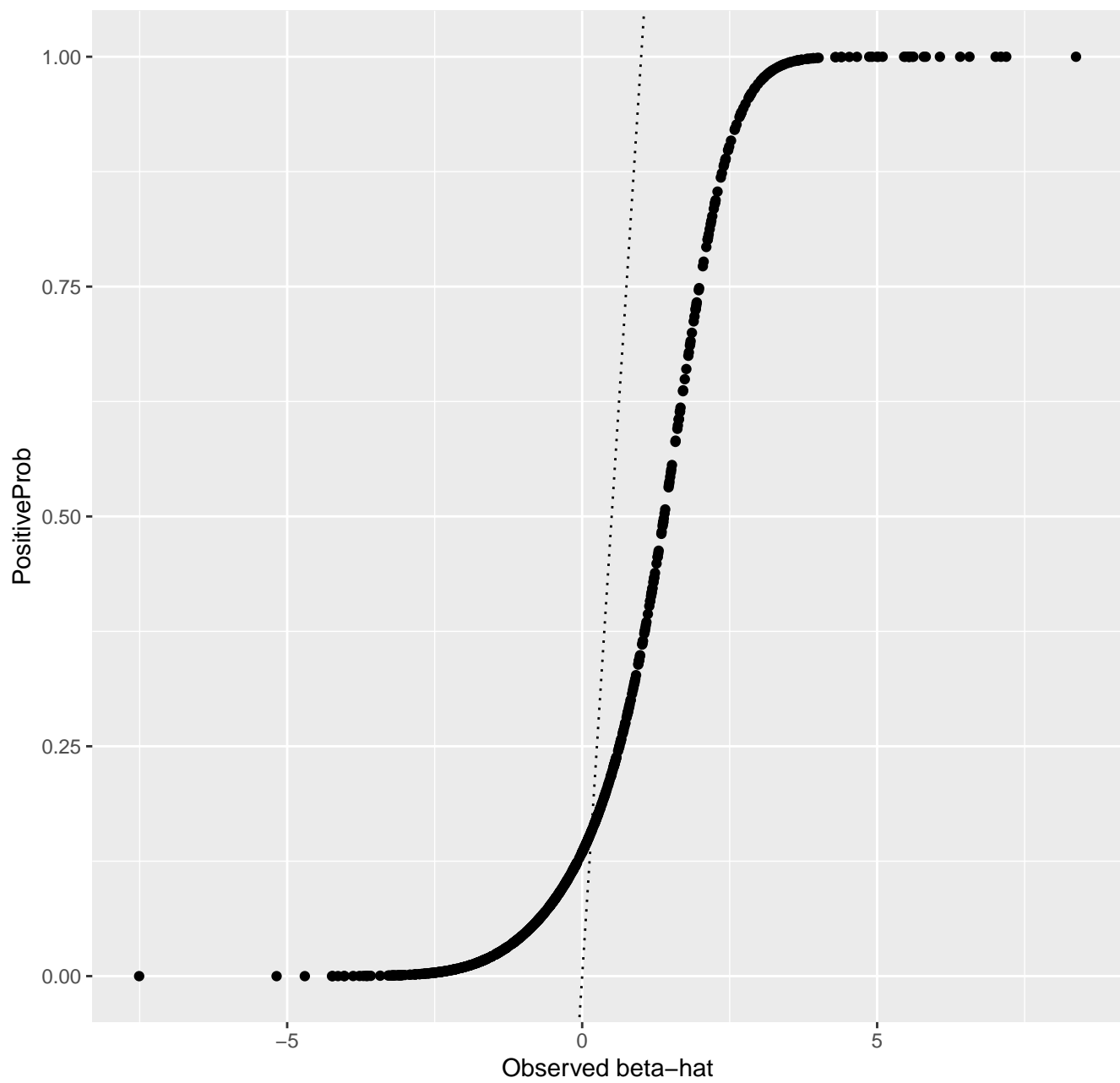
ATP2B1_(490) negative



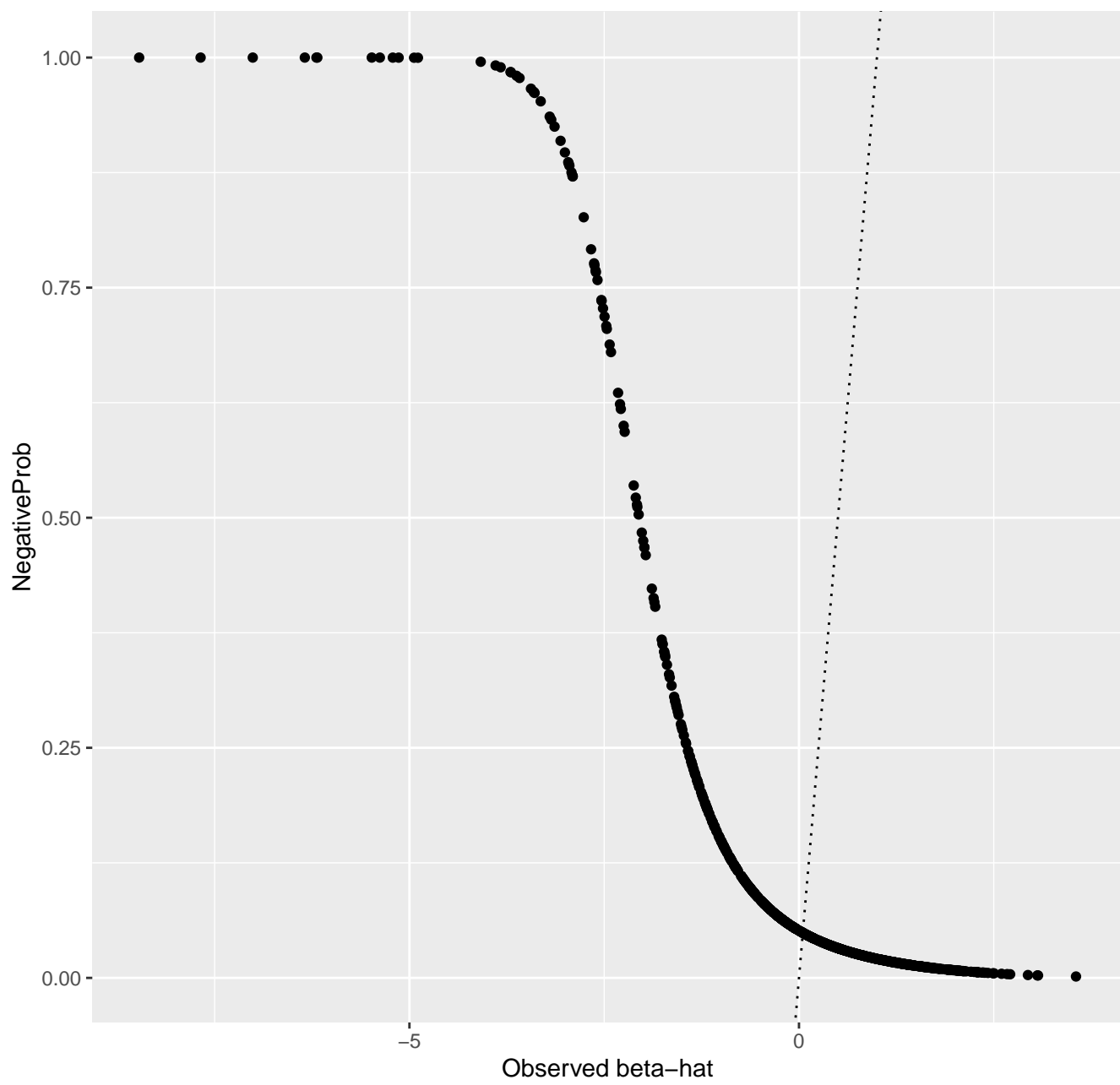
IDH3A_(3419) negative



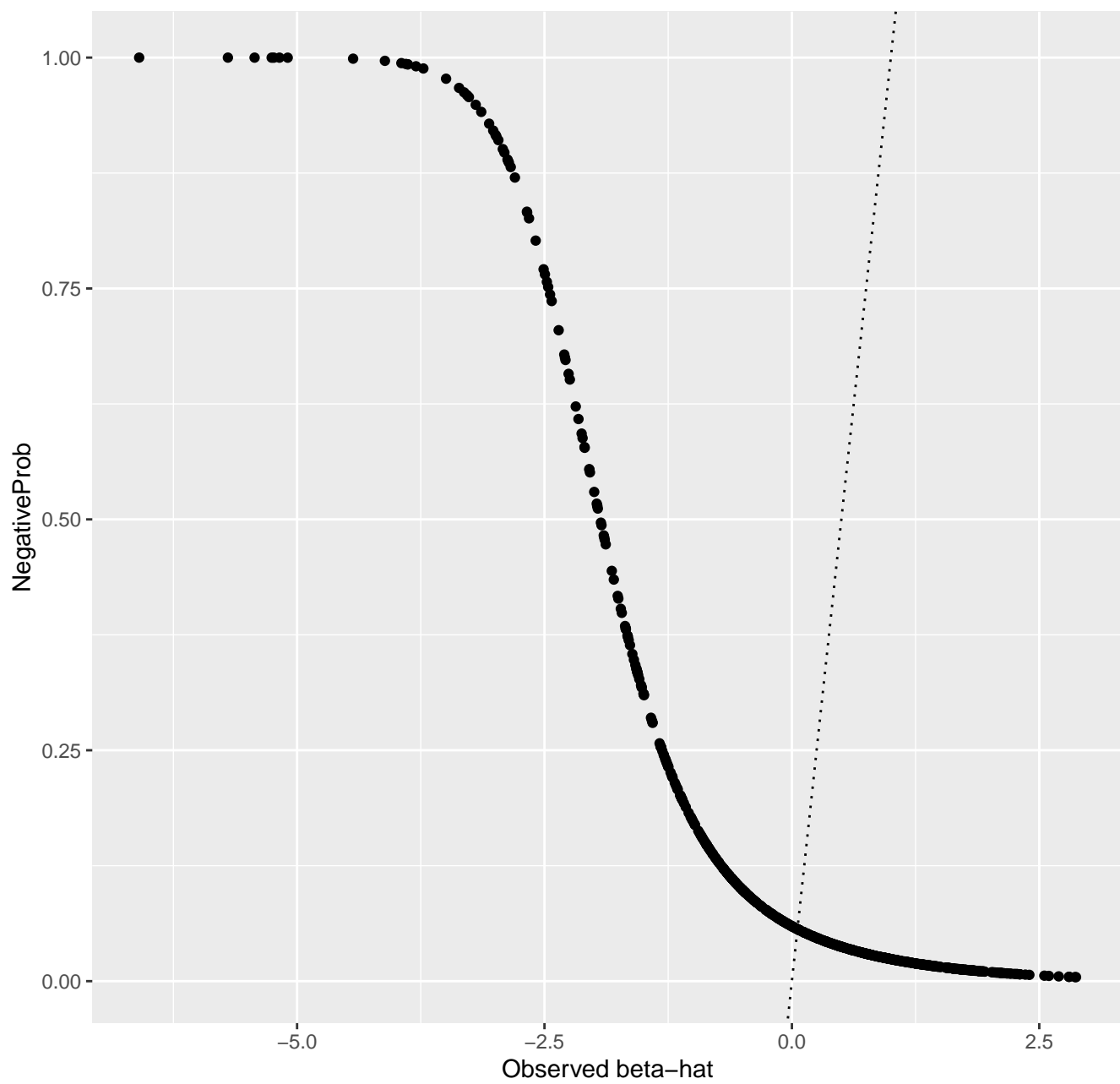
PYROXD1_(79912) positive



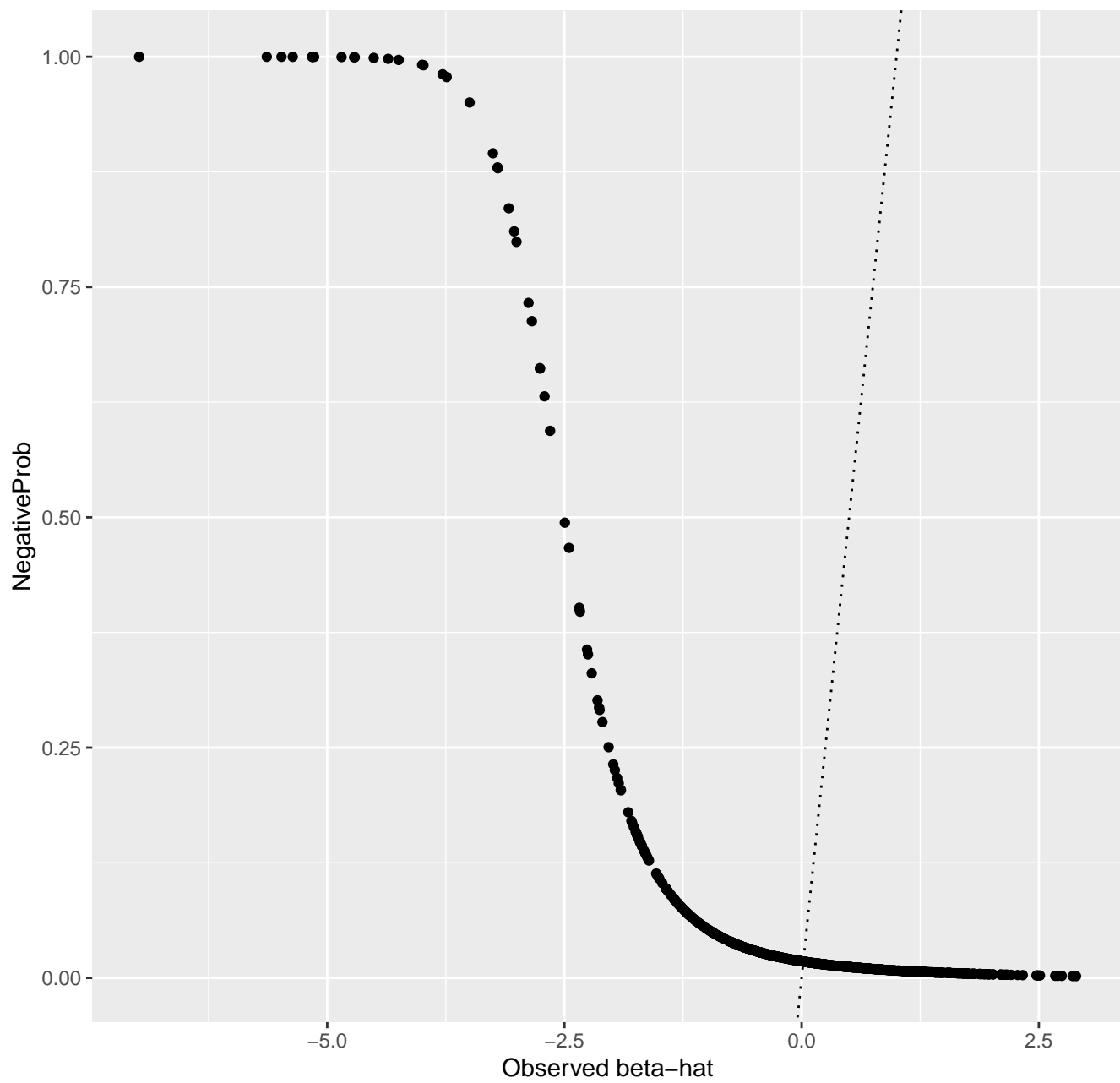
TBC1D20_(128637) negative



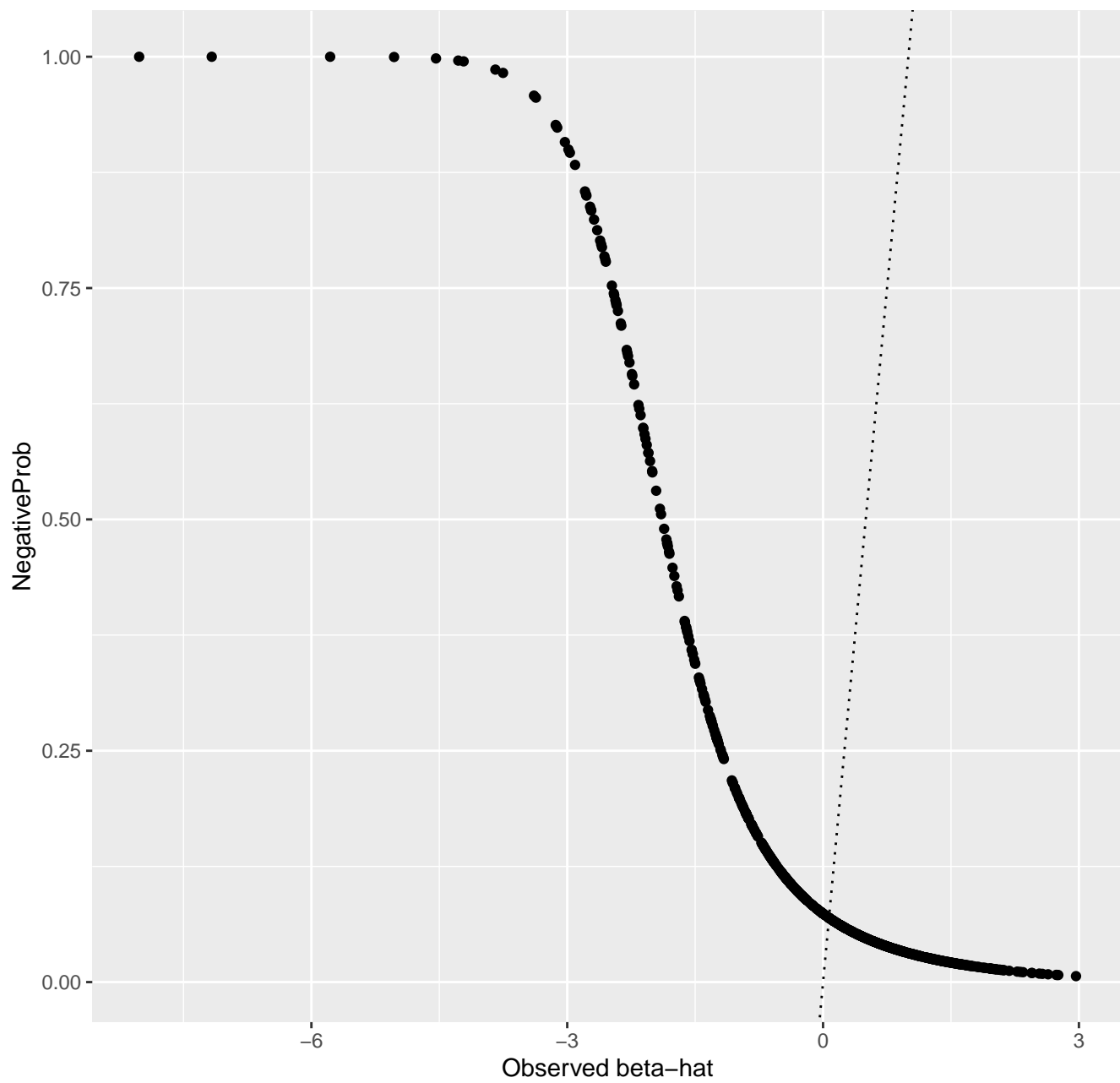
WDR44_(54521) negative



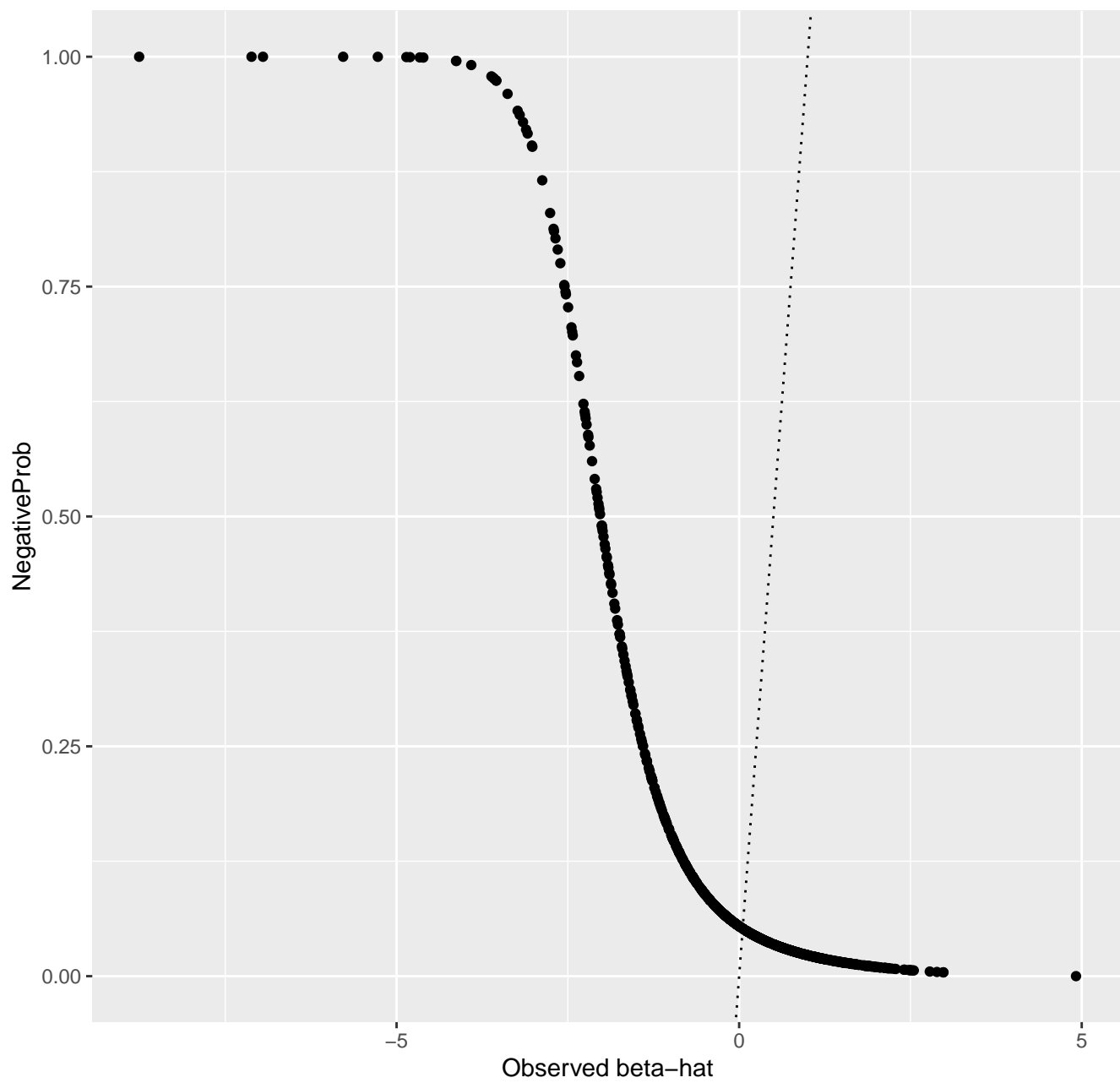
SRD5A3_(79644) negative



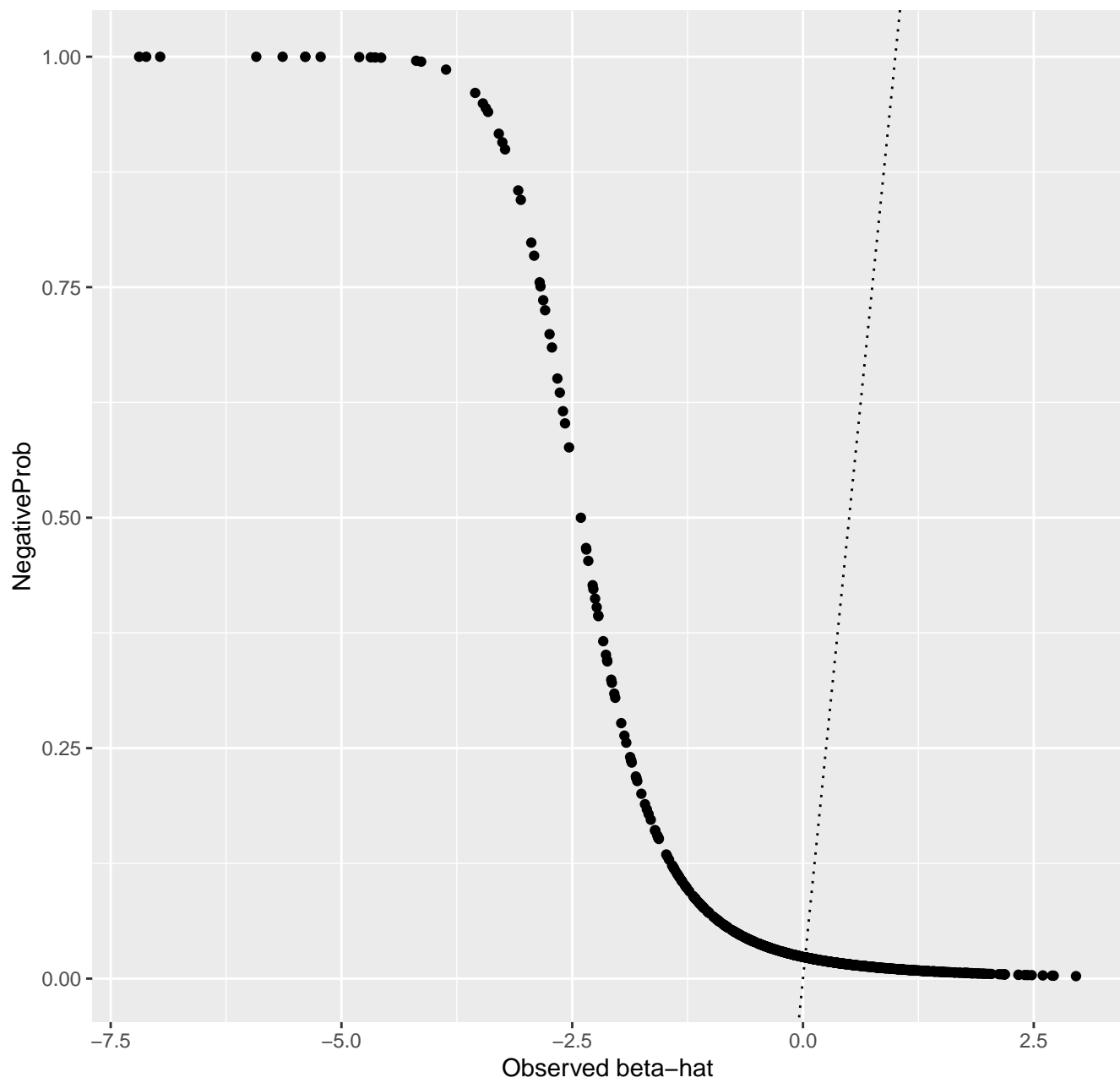
IKBKG_(8517) negative



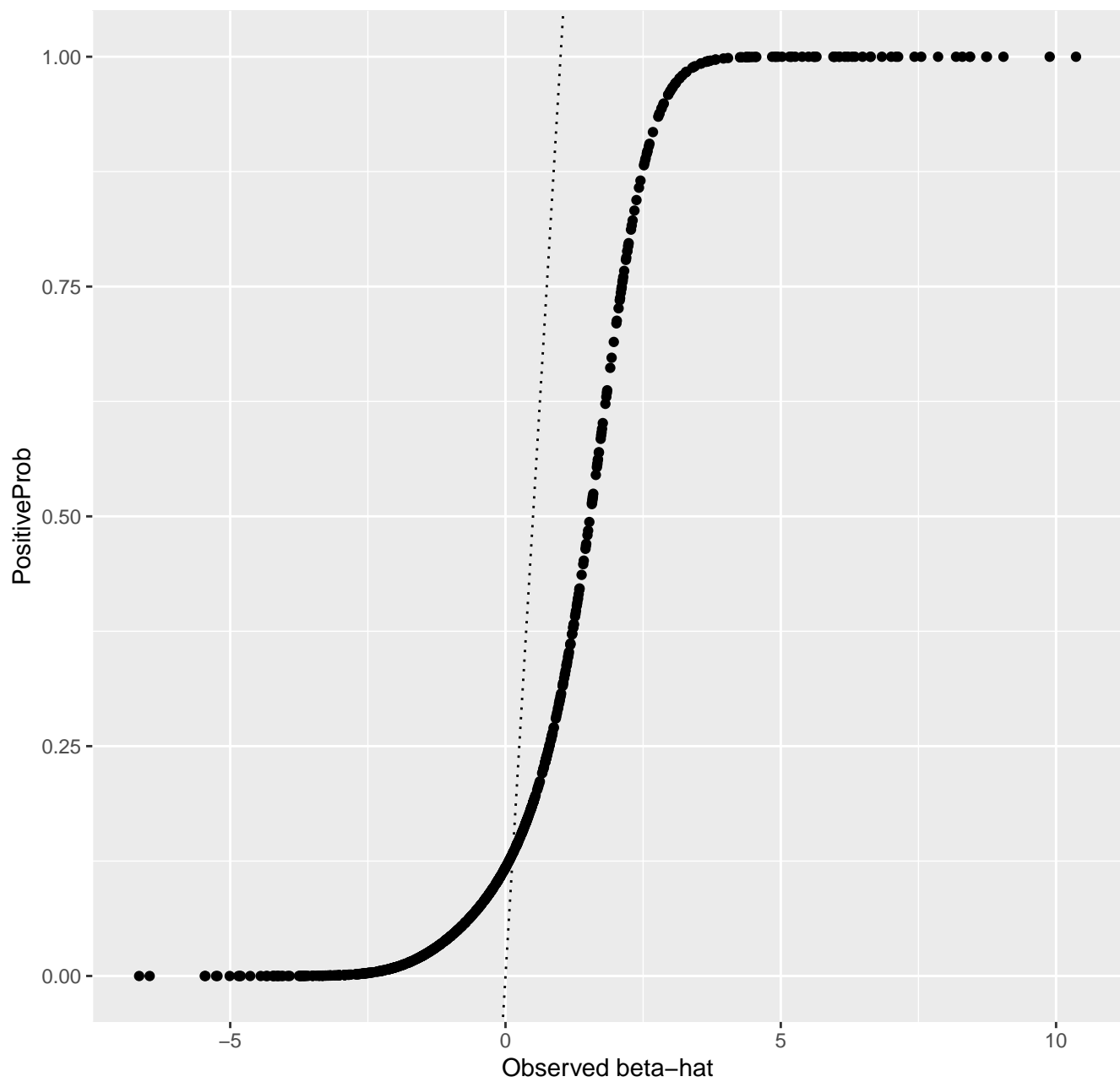
TRIO_(7204) negative



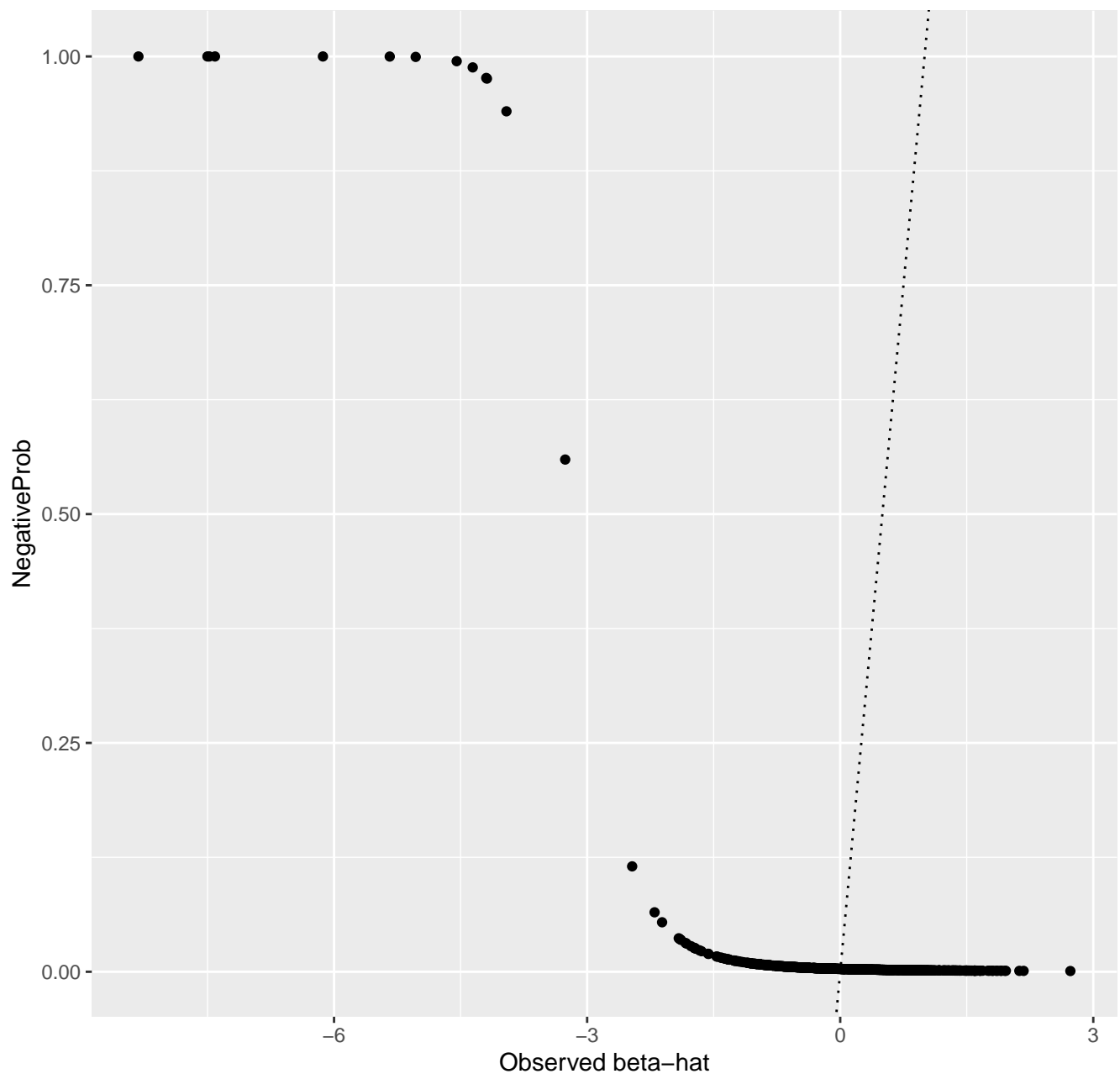
CWF19L1_(55280) negative



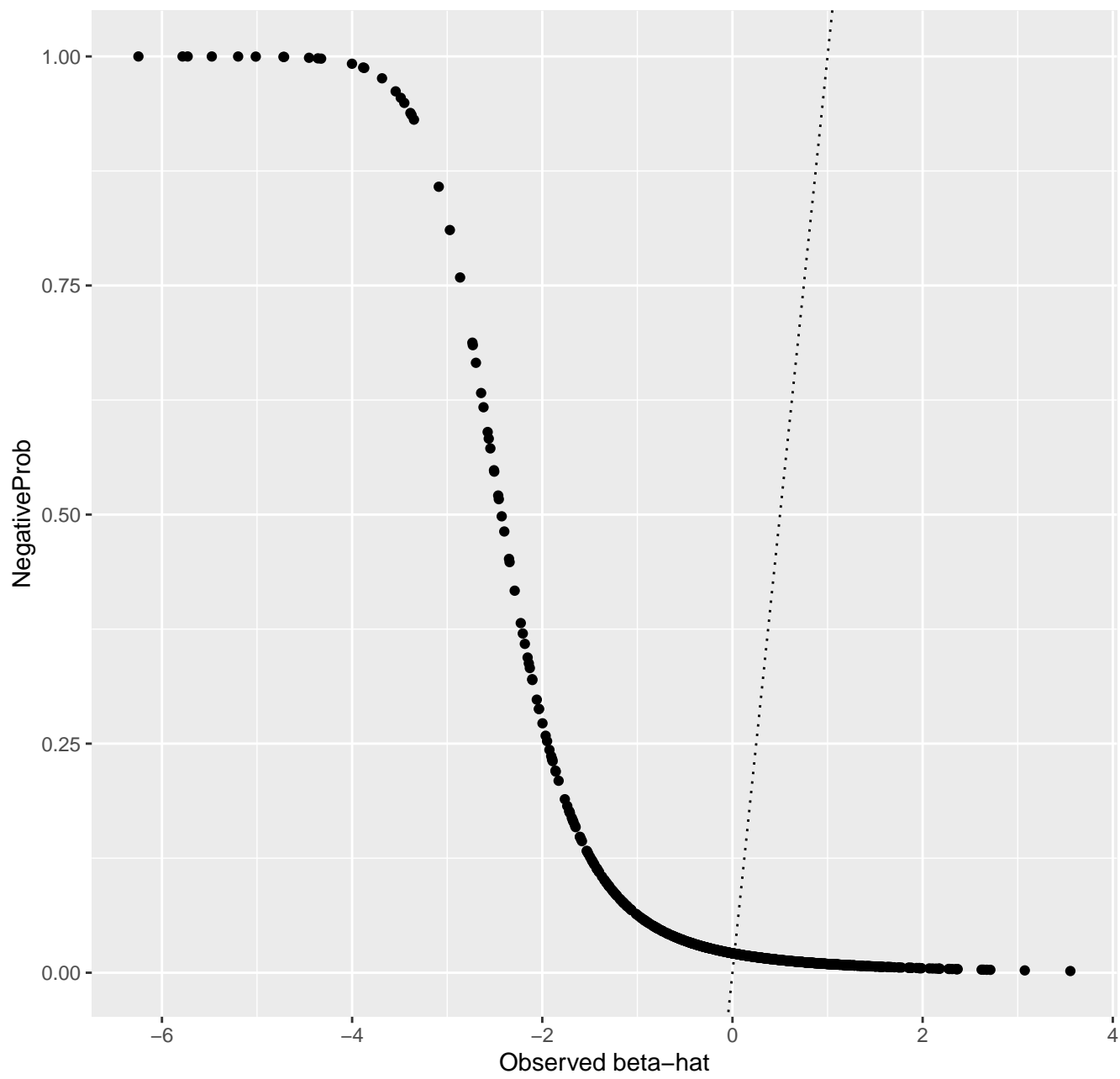
DTYMK_(1841) positive



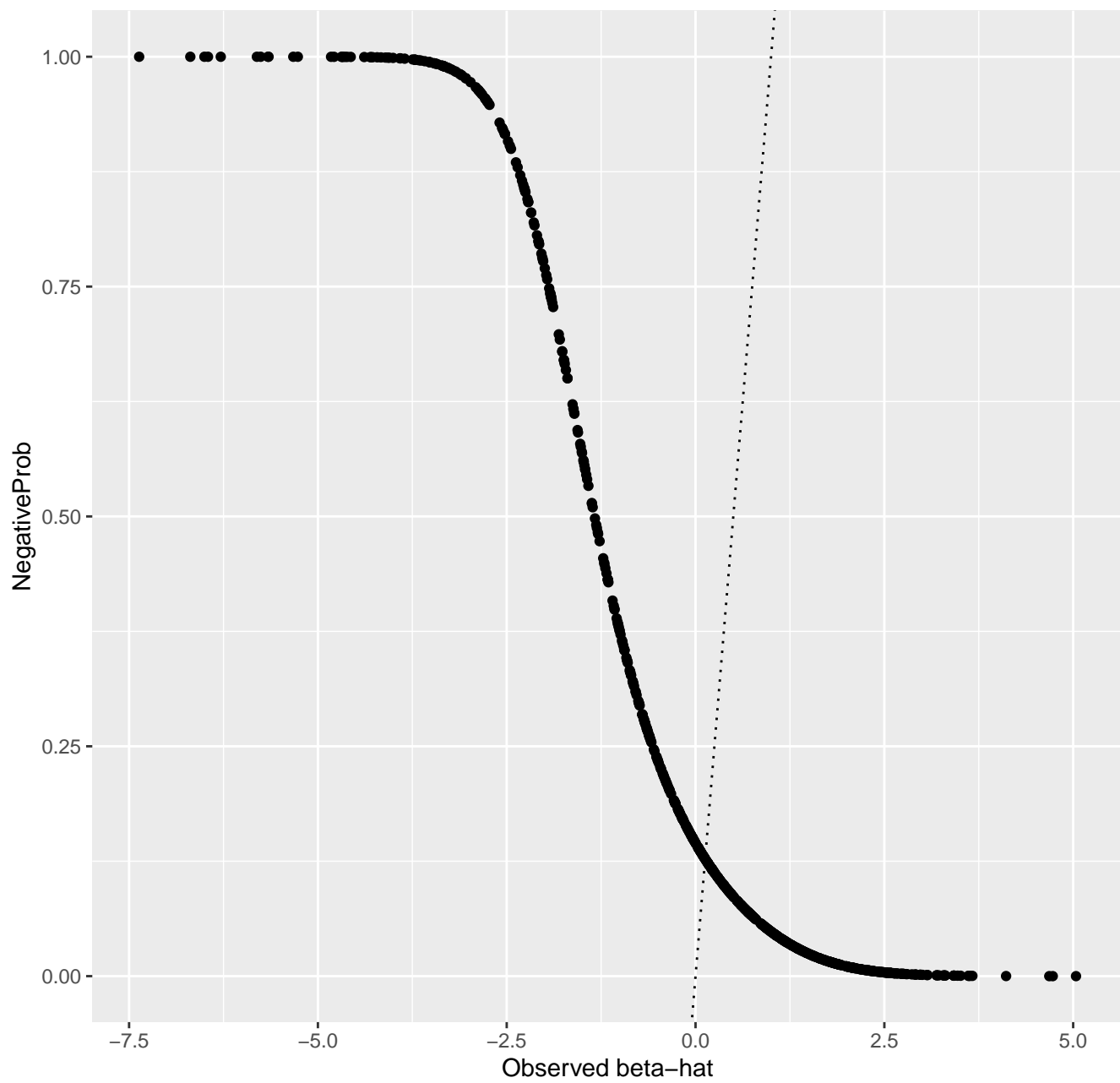
IRF8_(3394) negative



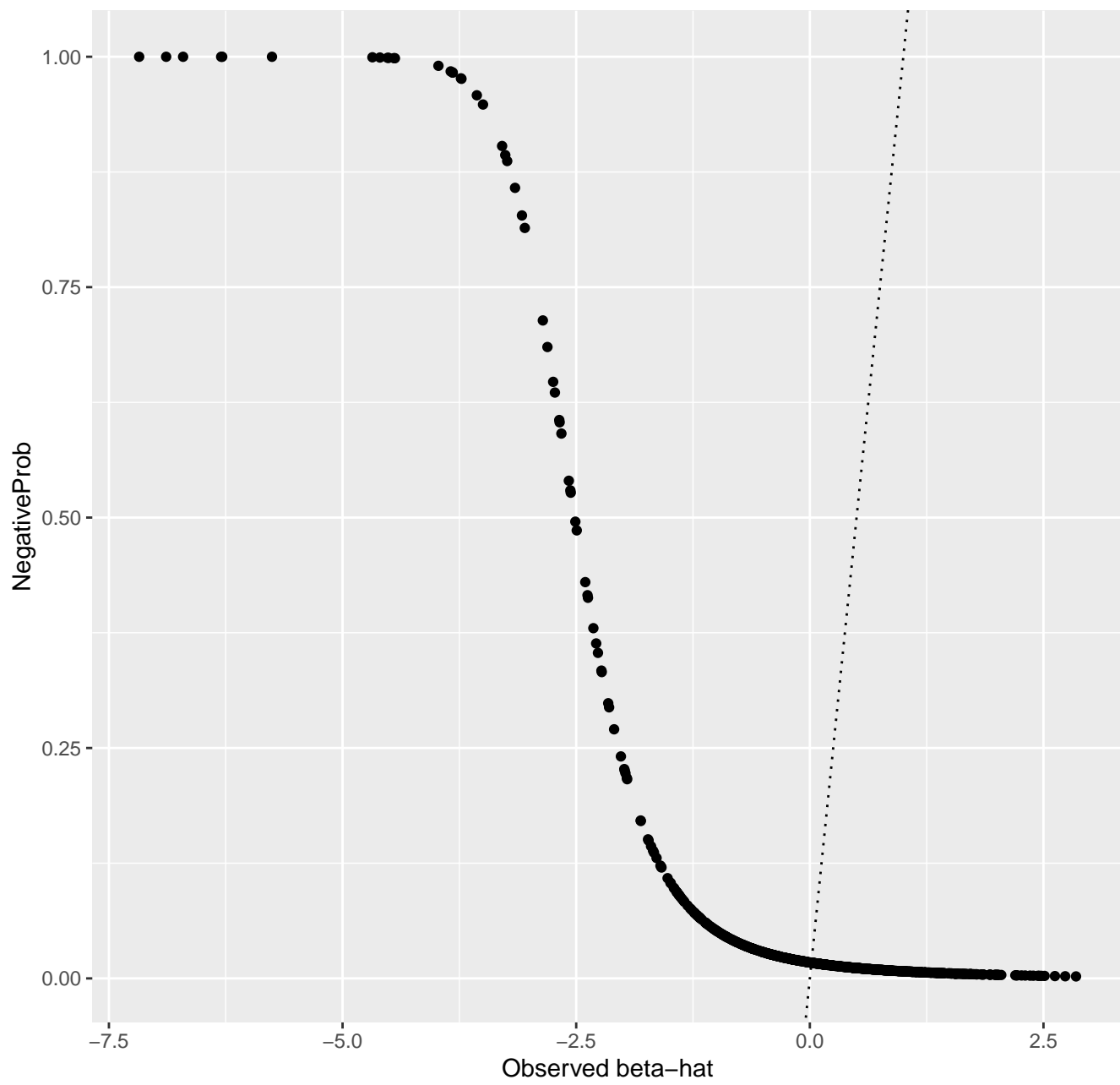
SAR1A_(56681) negative



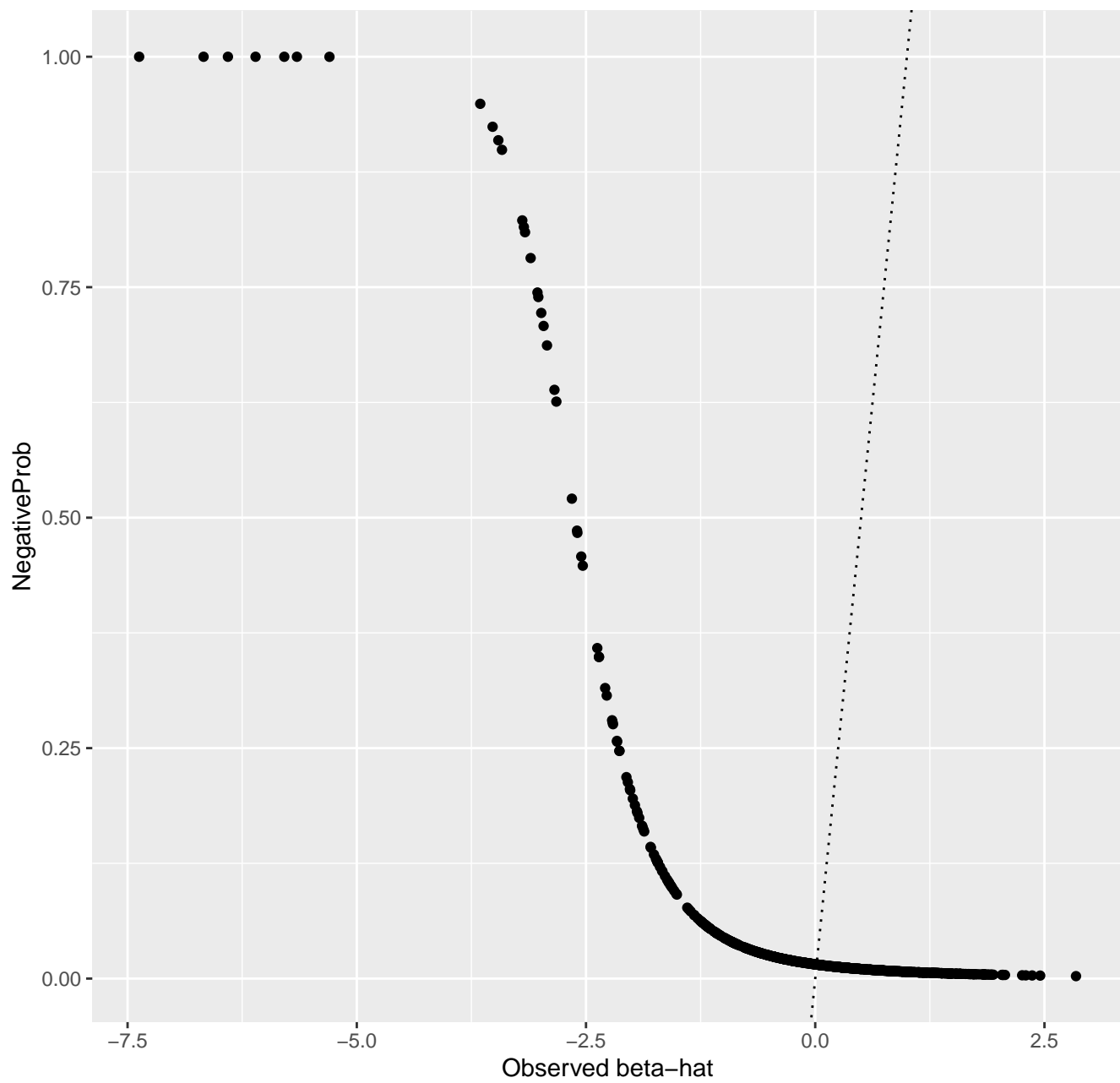
TAZ_(6901) negative



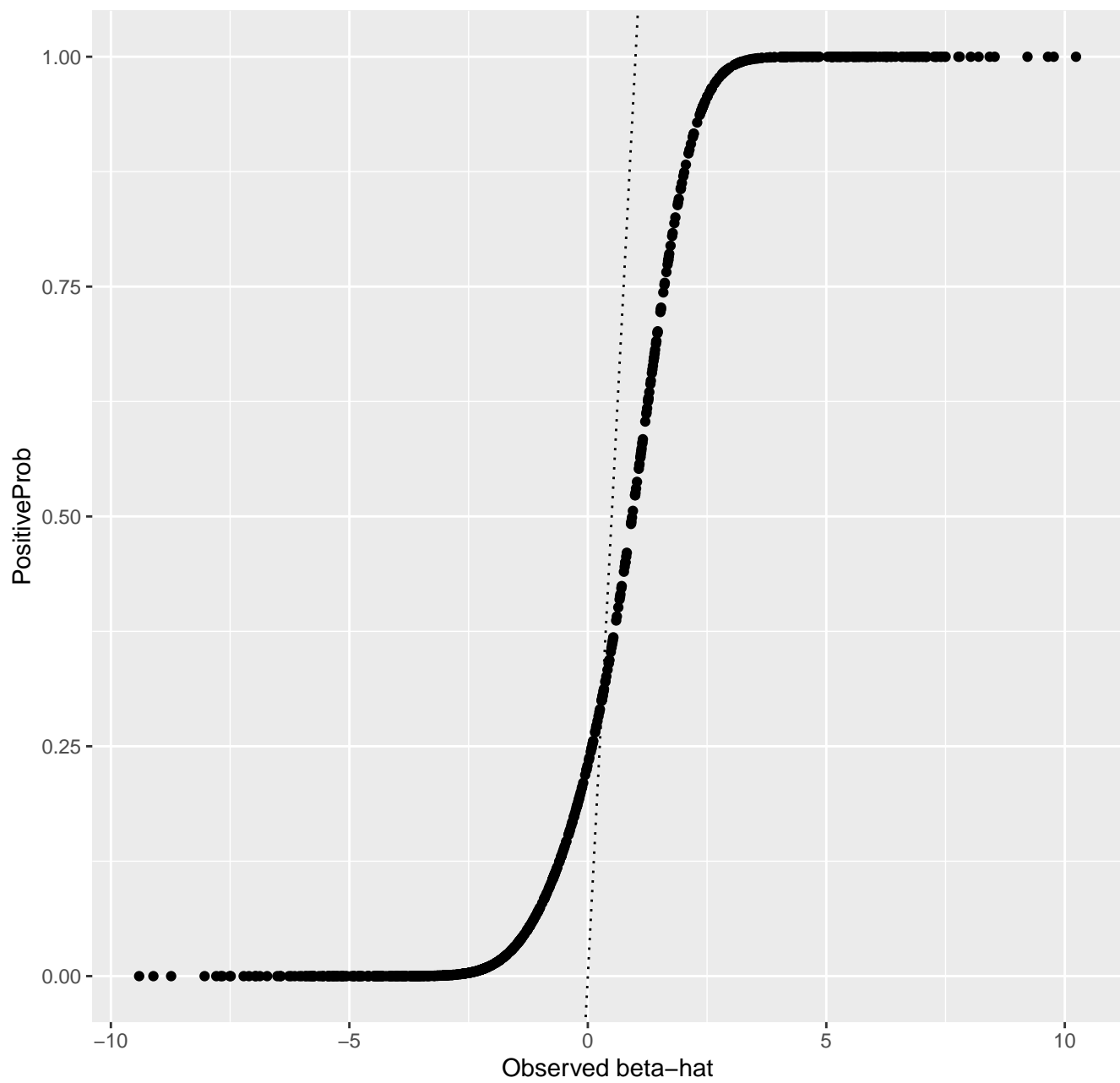
SREBF2_(6721) negative



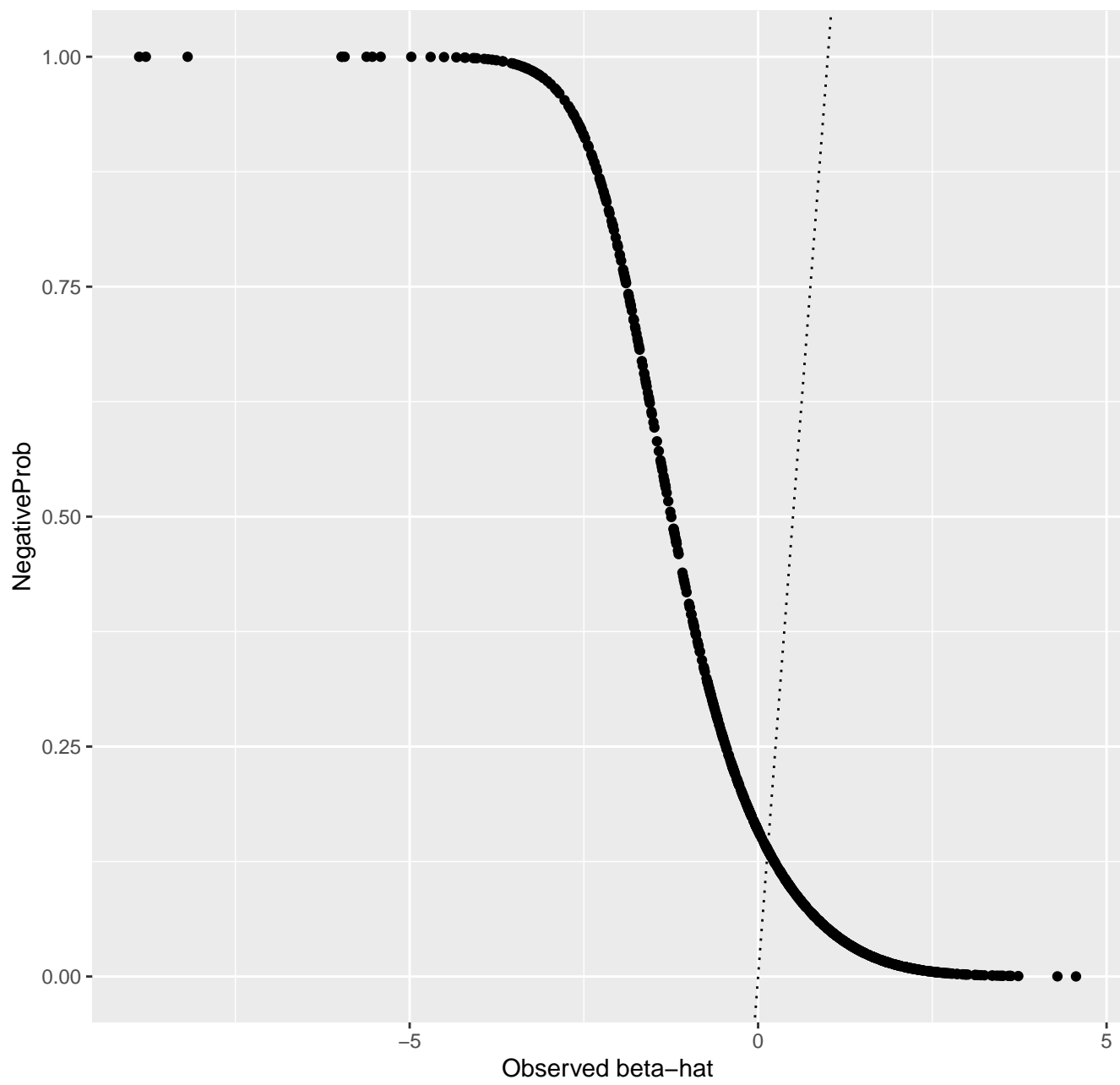
RAB1A_(5861) negative



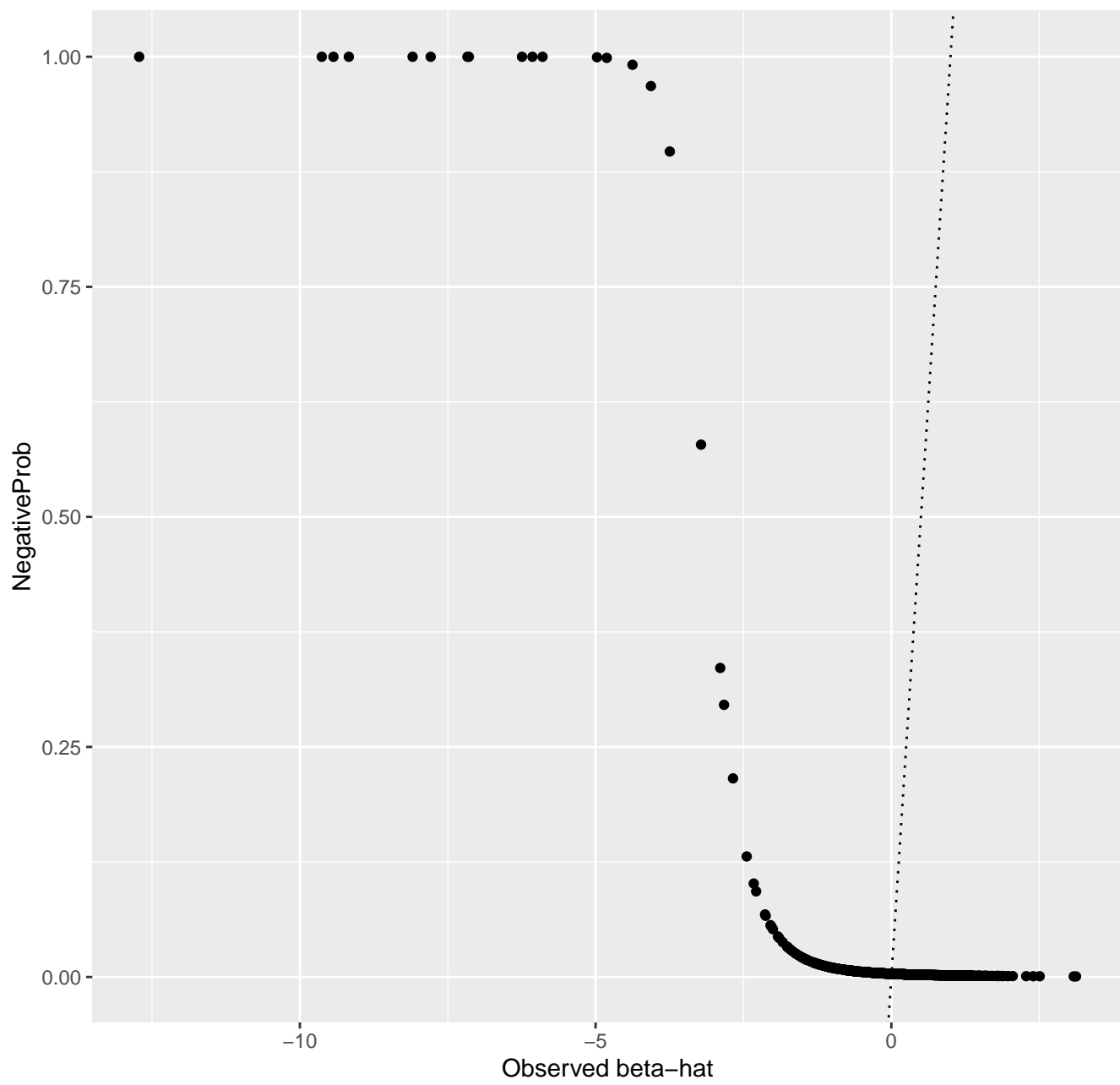
SCAP_(22937) positive



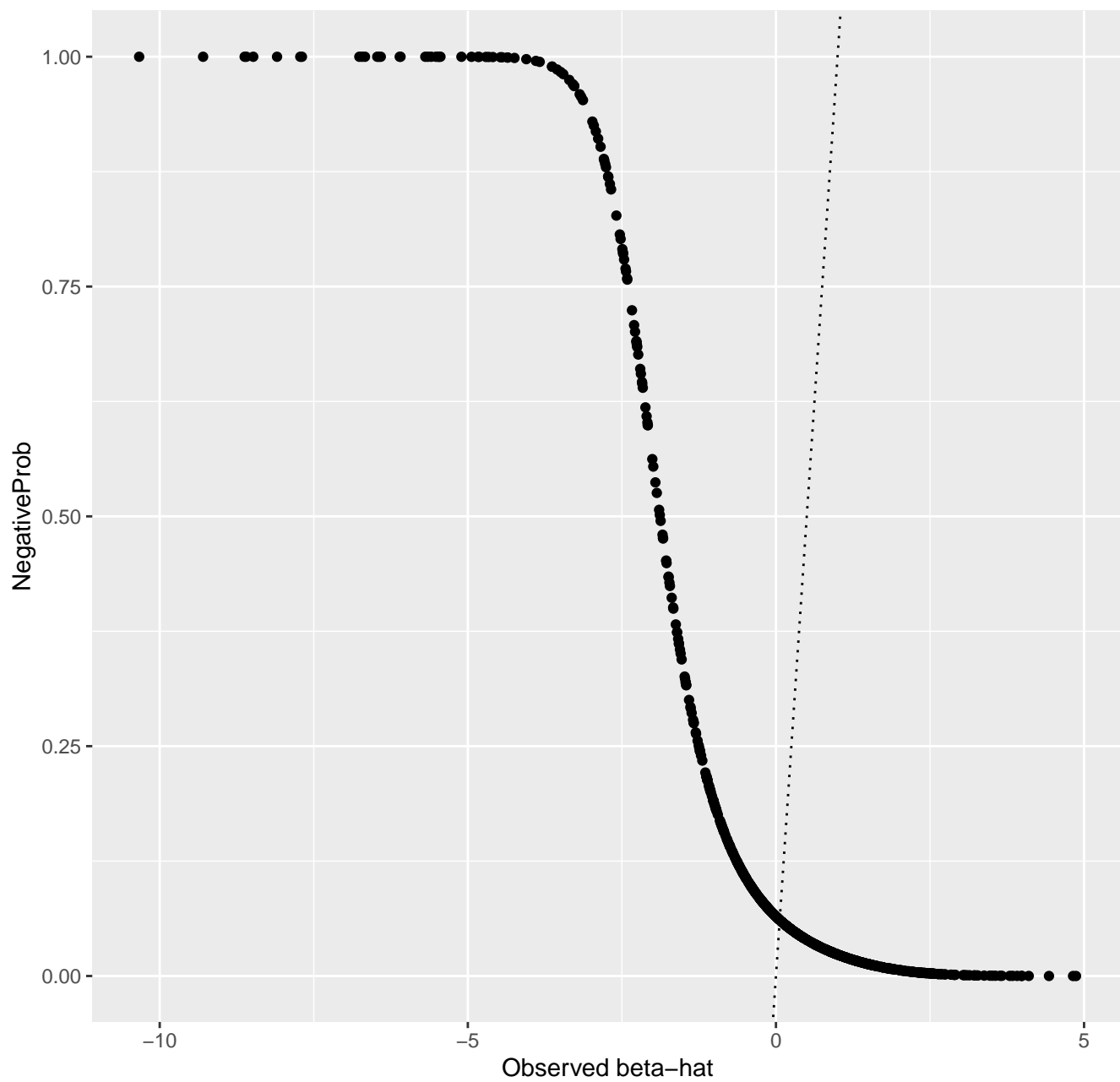
DUX4_(100288687) negative



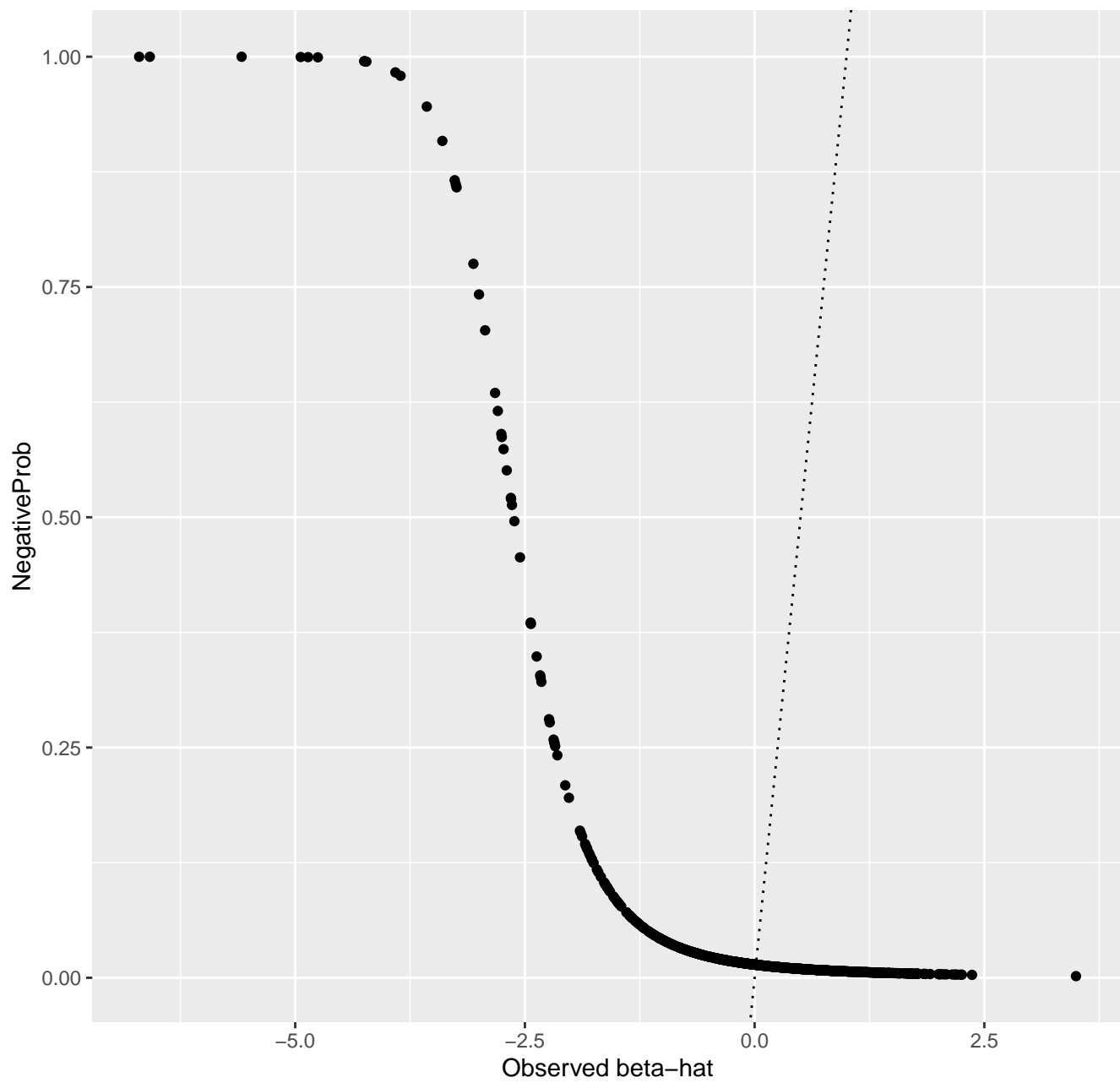
TNFSF10_(8743) negative



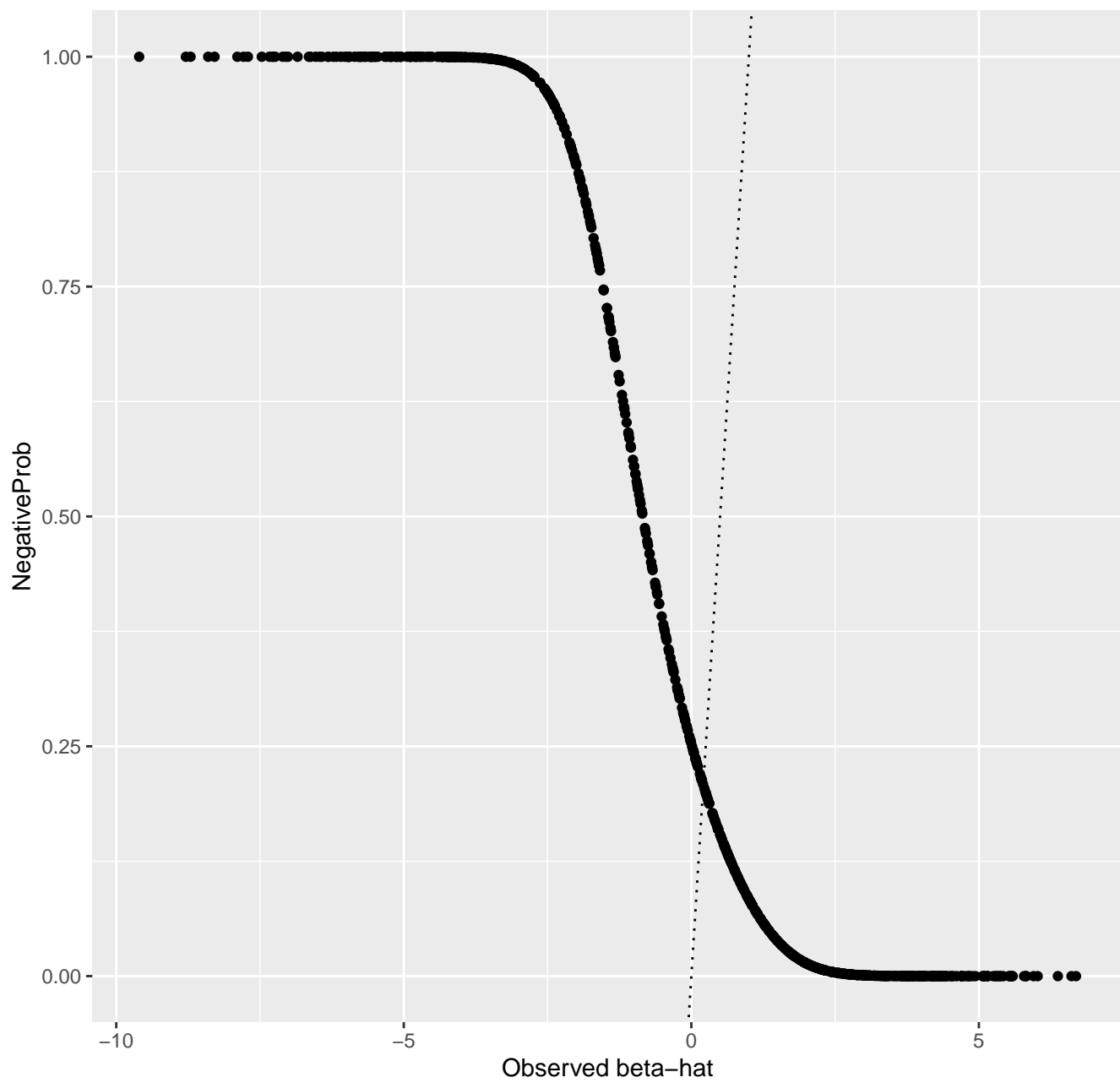
MAPK1_(5594) negative



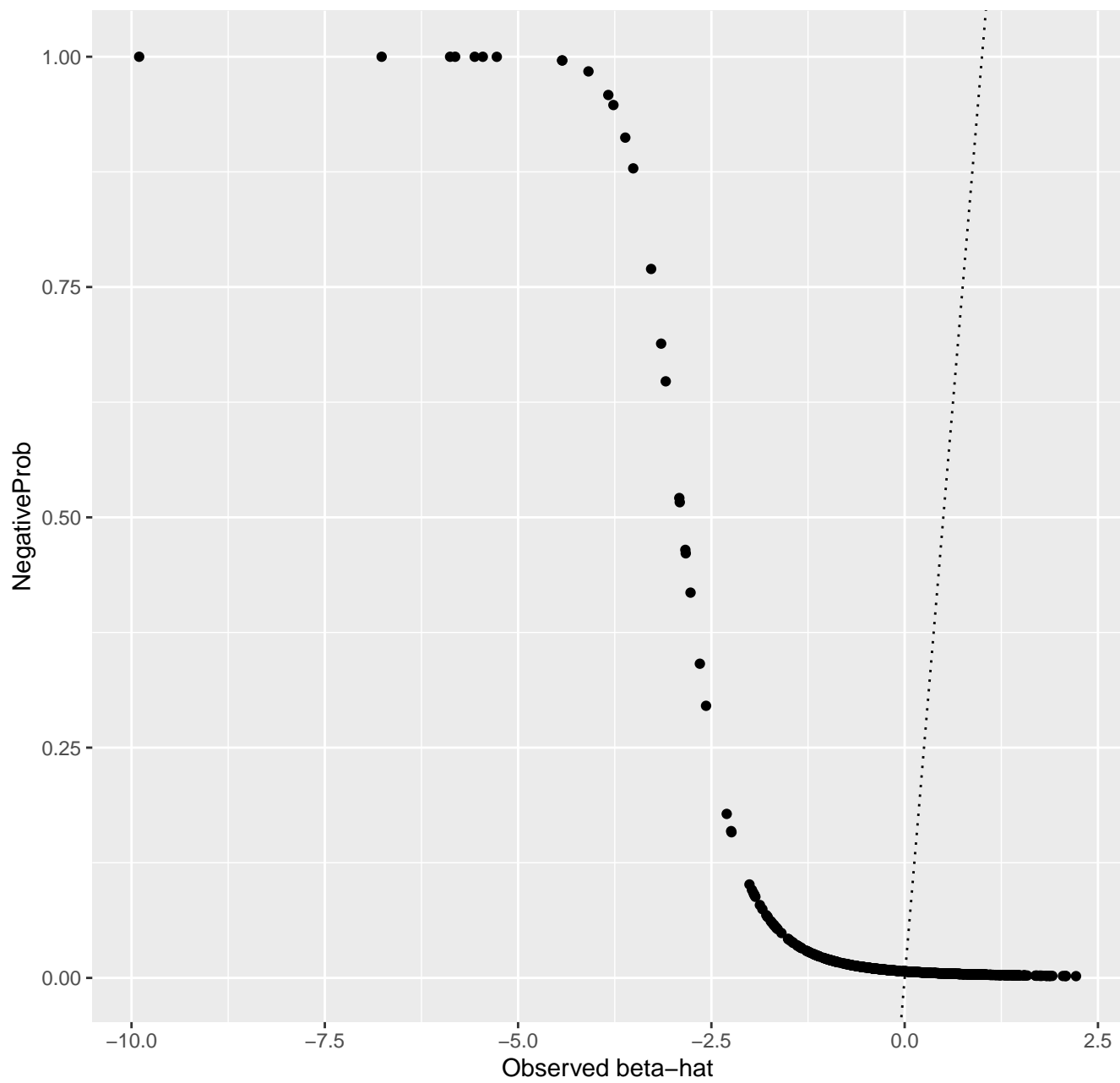
SLC1A5_(6510) negative



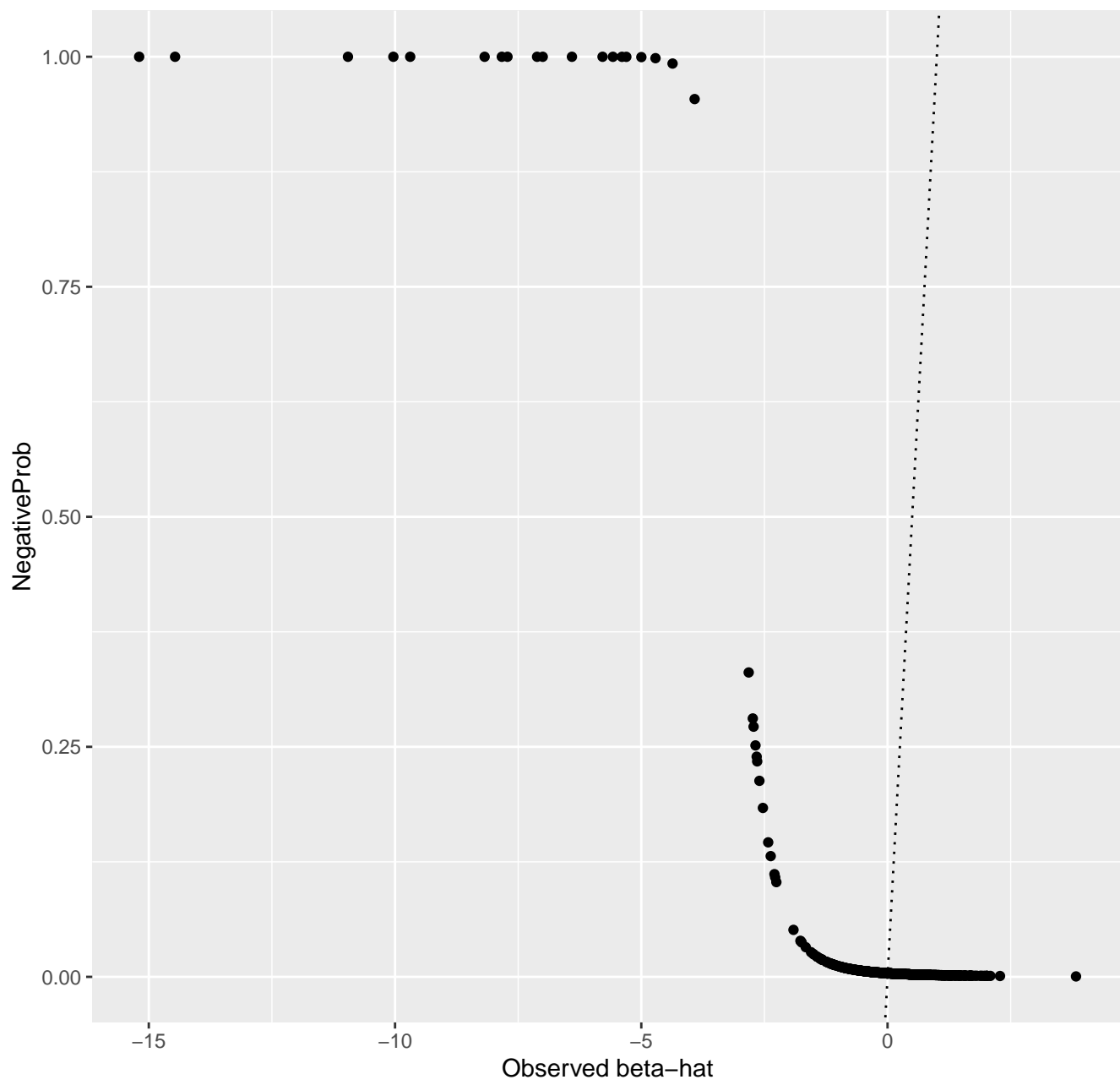
CFLAR_(8837) negative



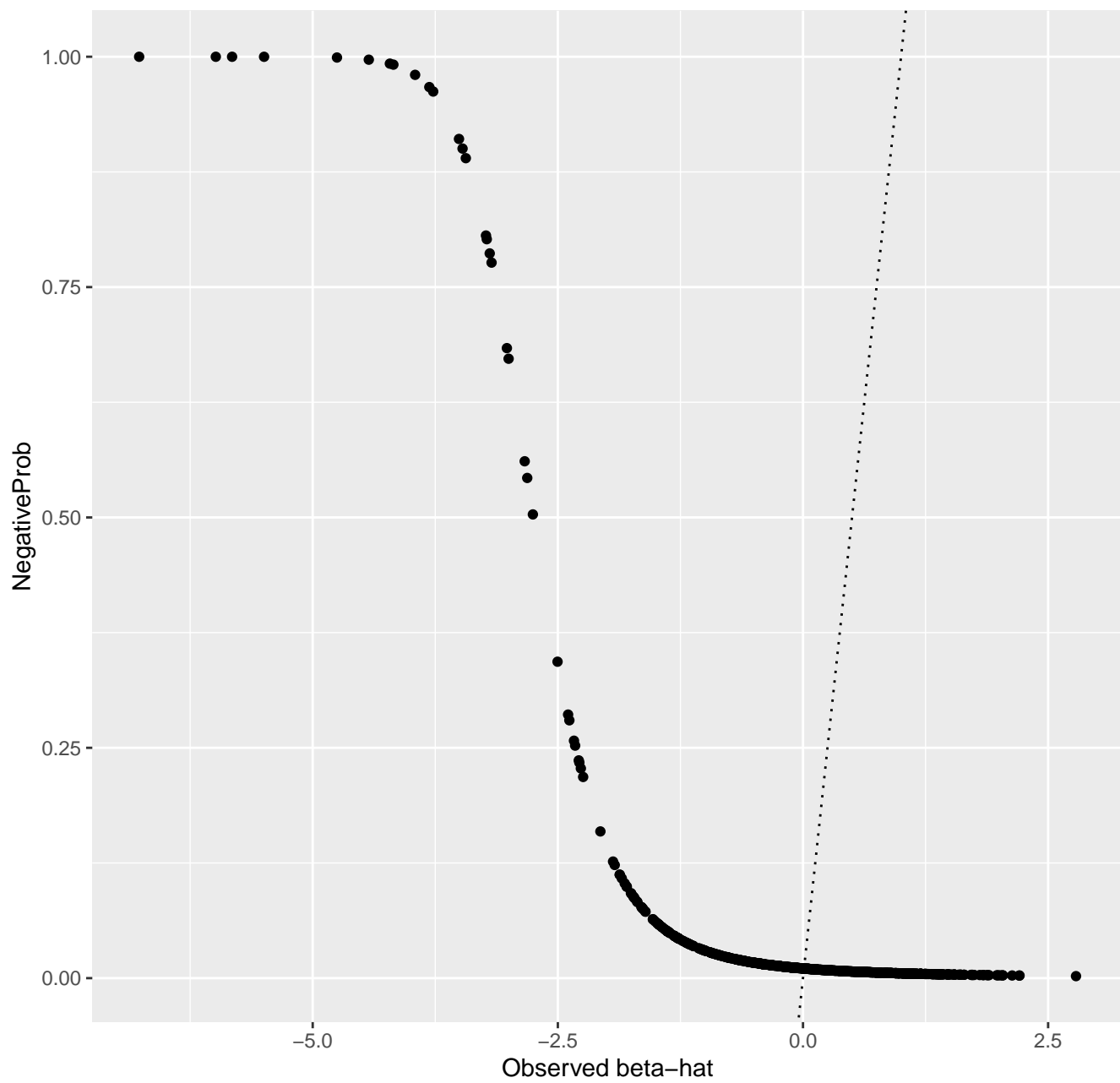
DMXL1_(1657) negative



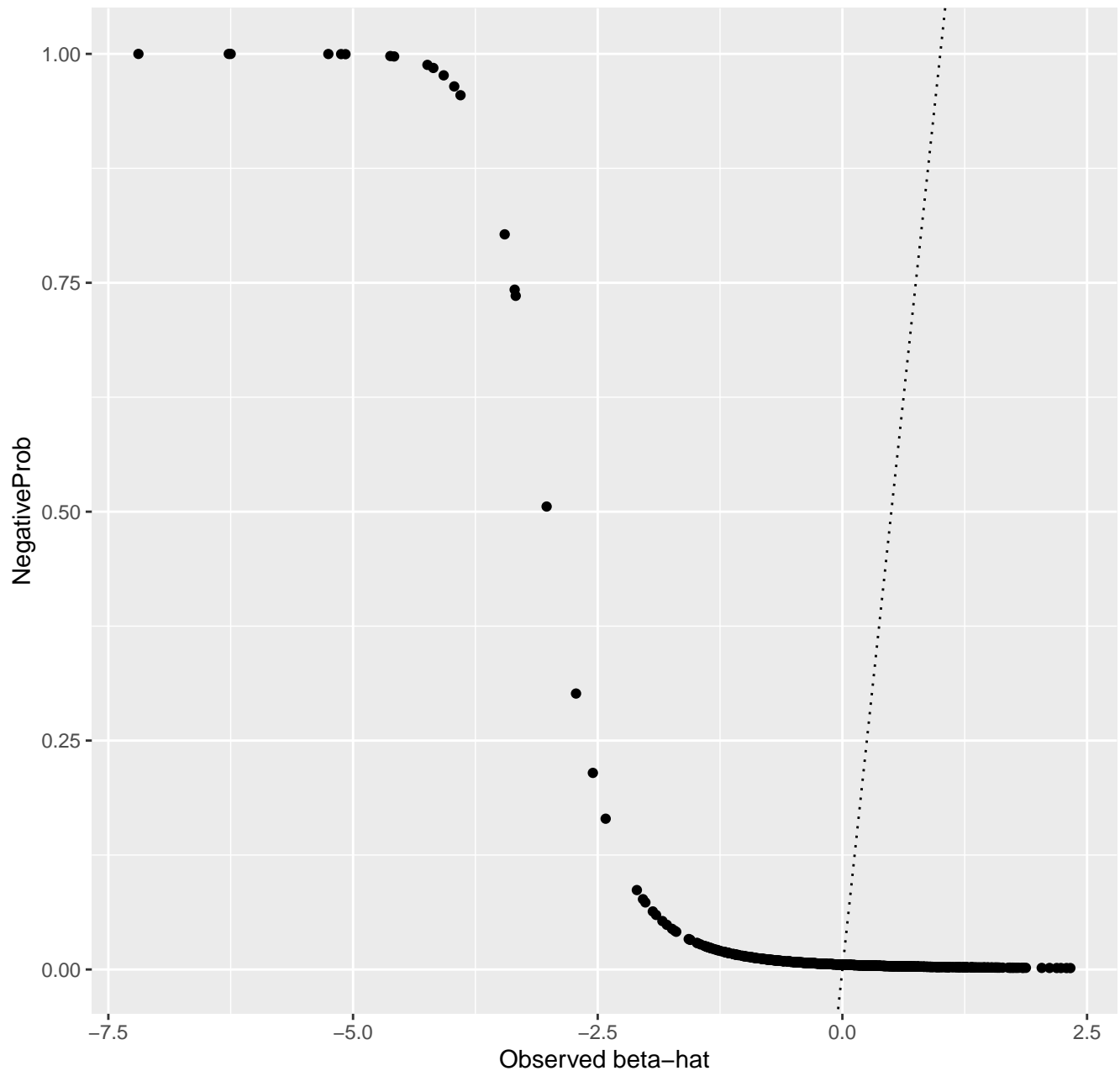
MYCN_(4613) negative



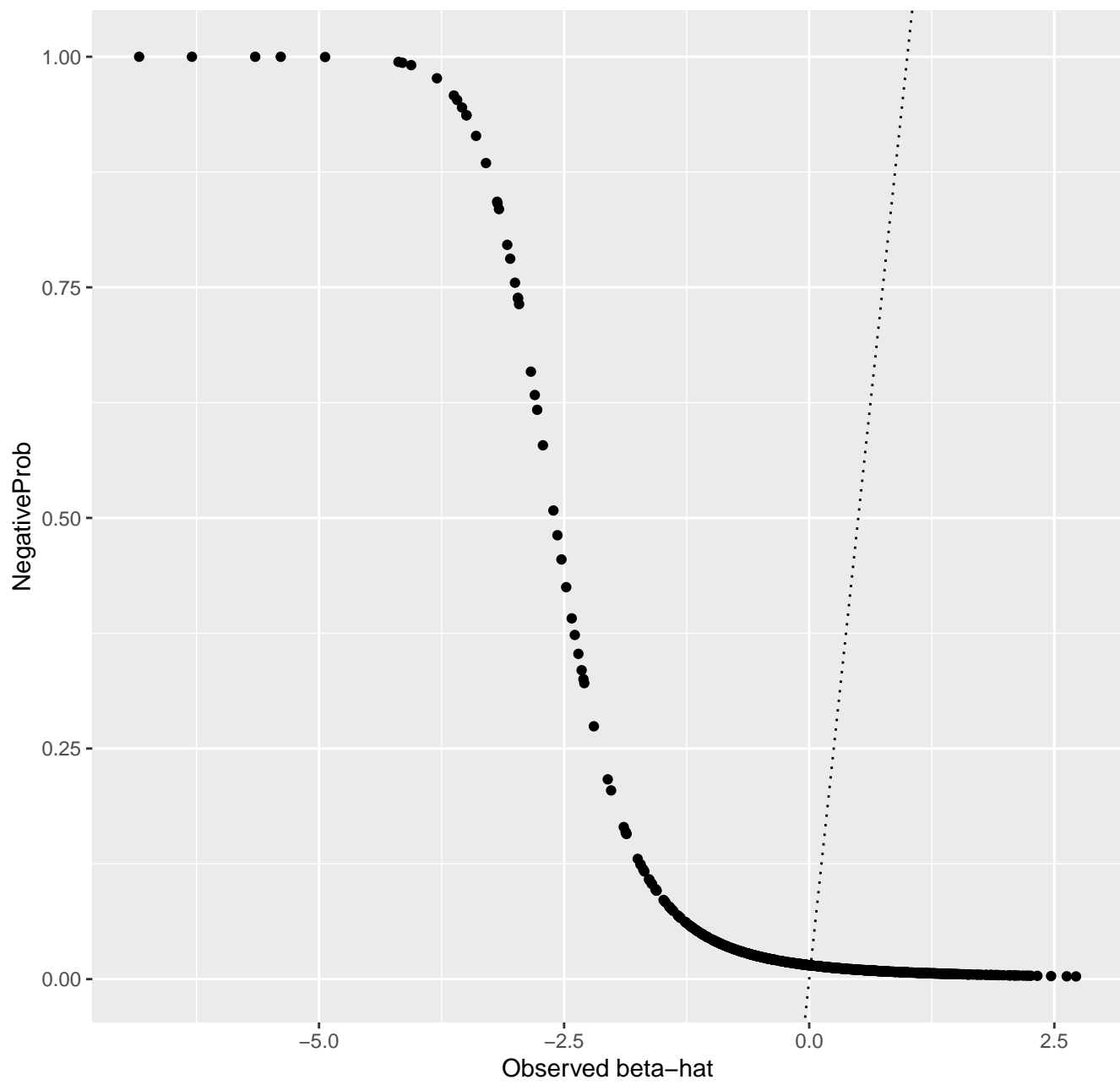
ATOX1_(475) negative



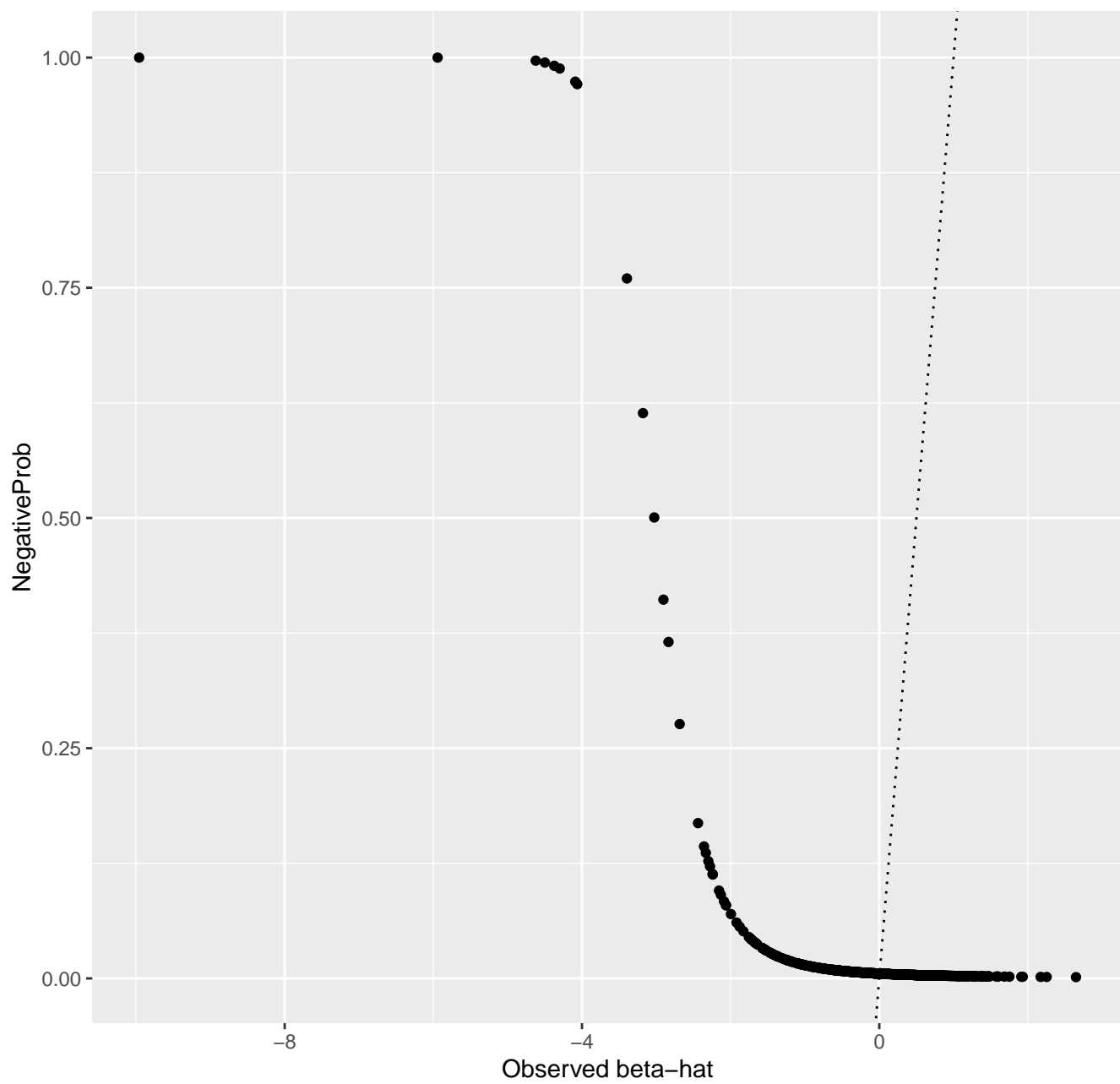
PAX5_(5079) negative



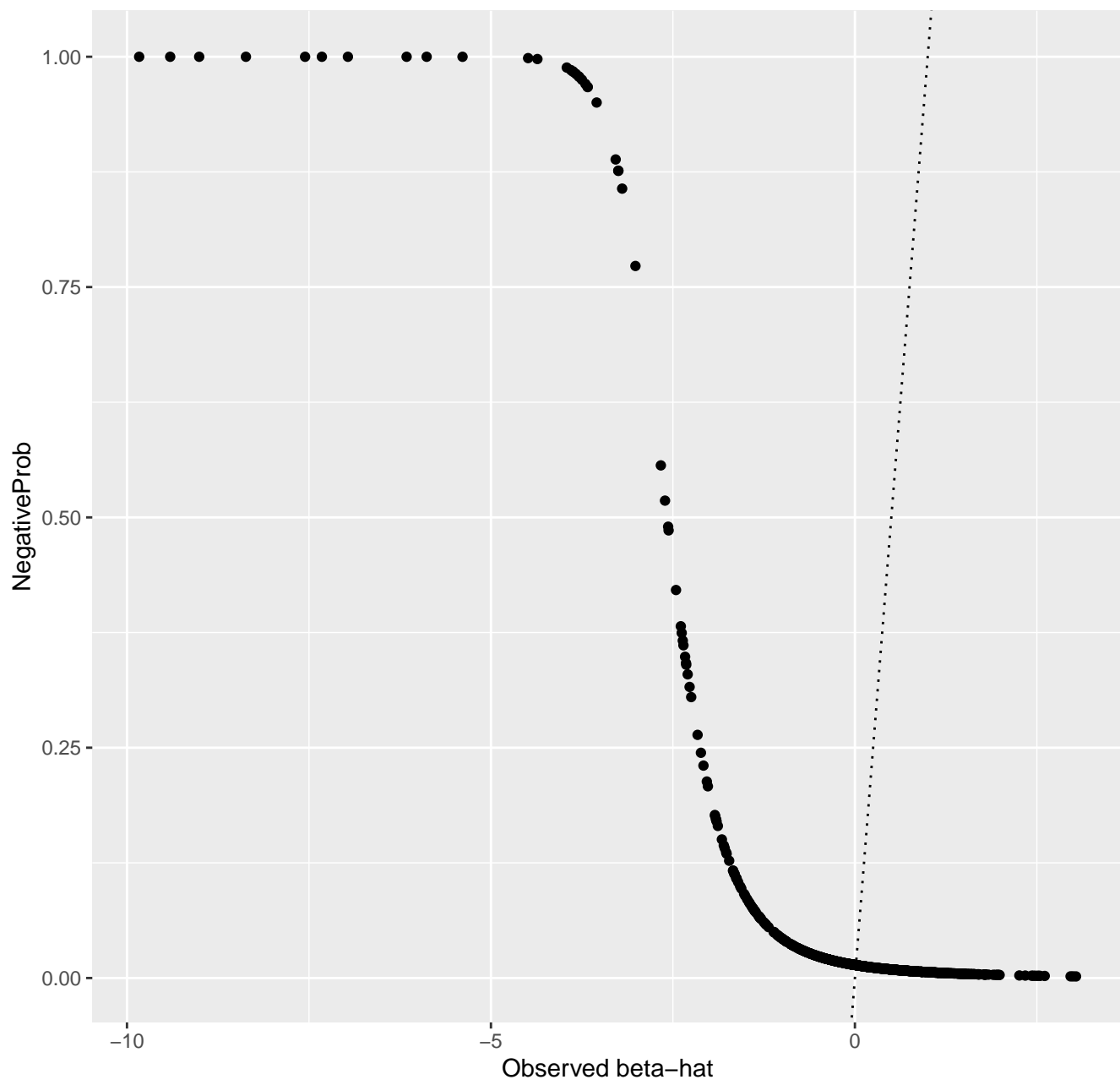
FAM136A_(84908) negative



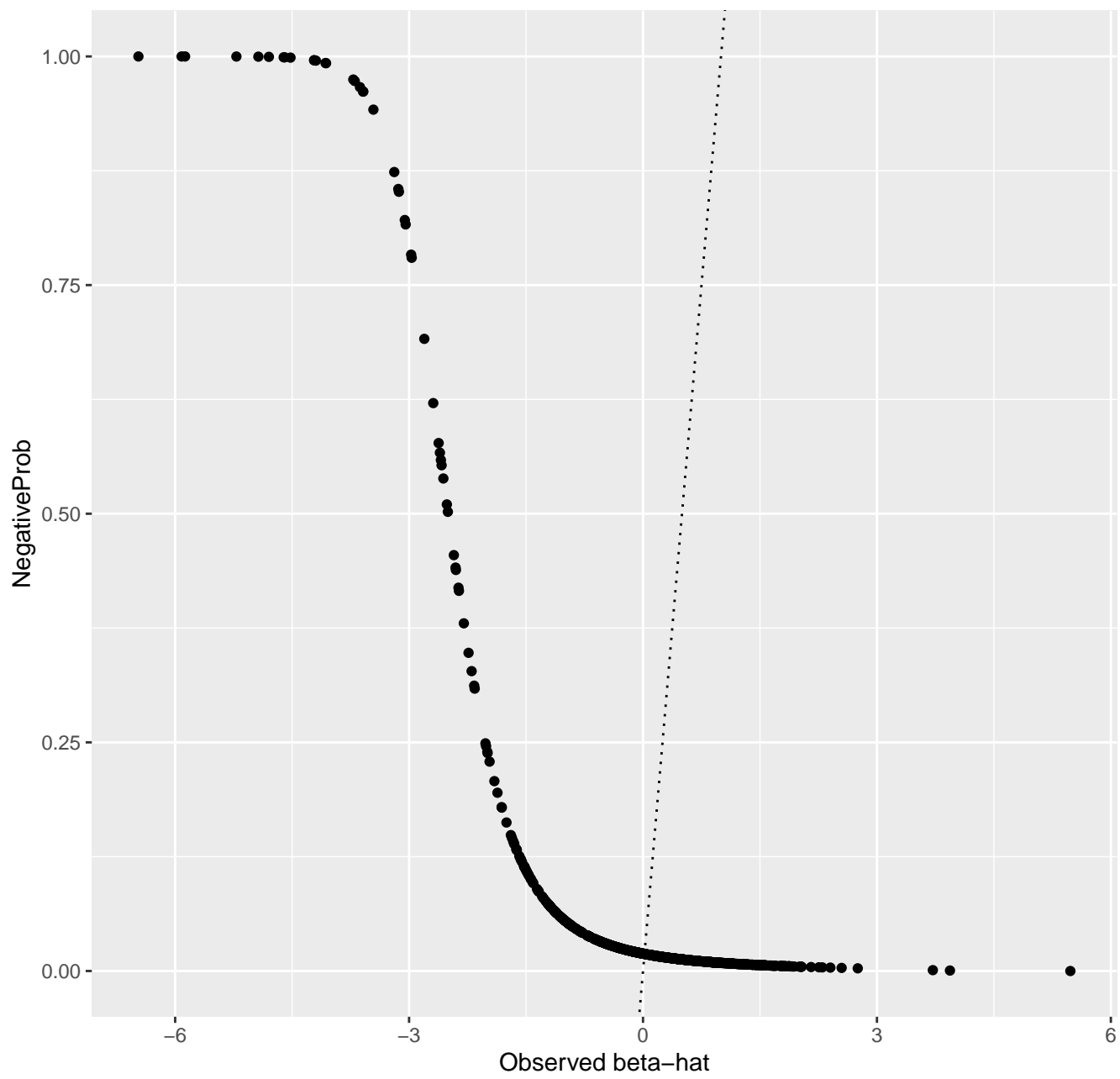
WDCP_(80304) negative



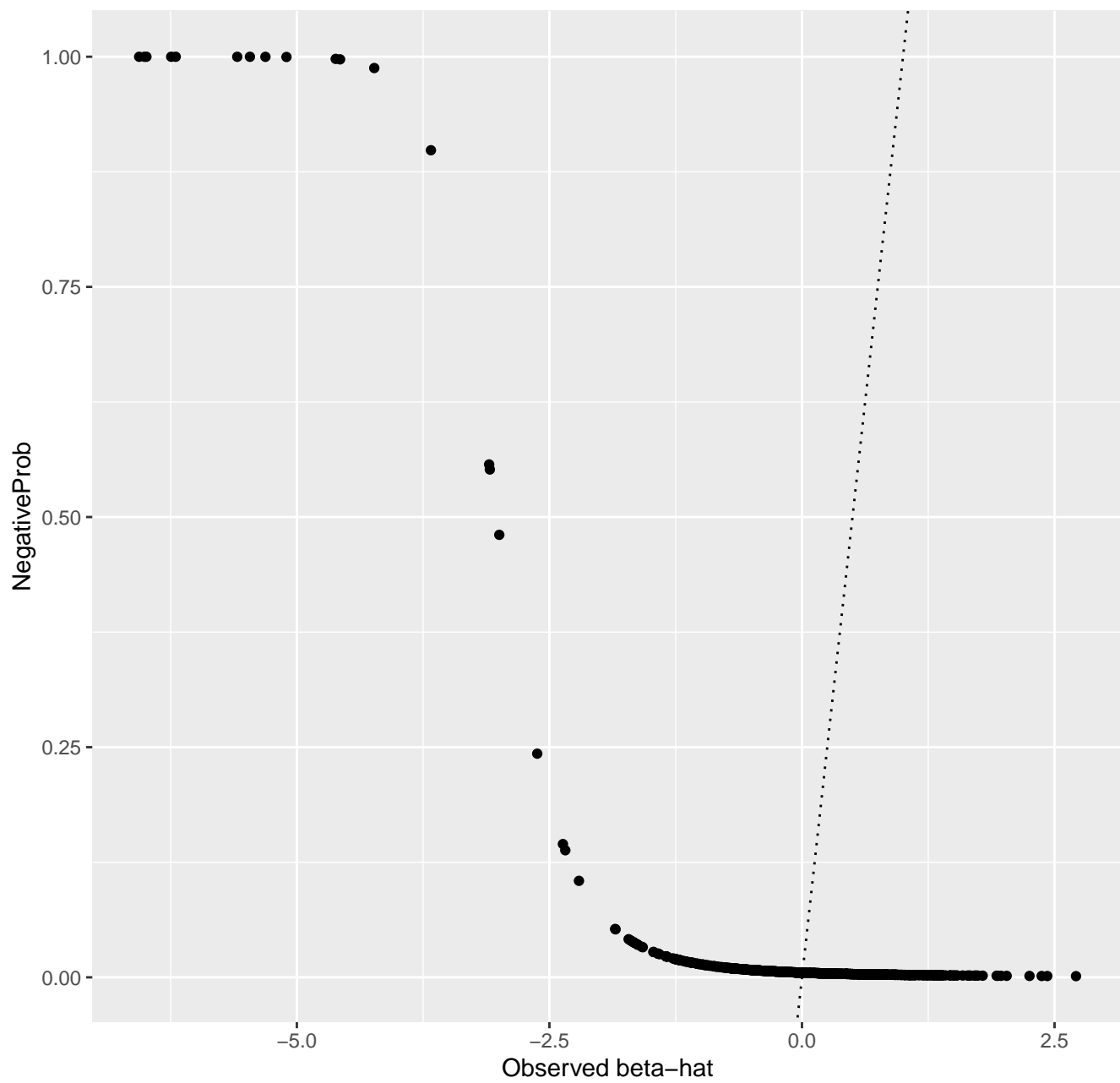
BAMBI_(25805) negative



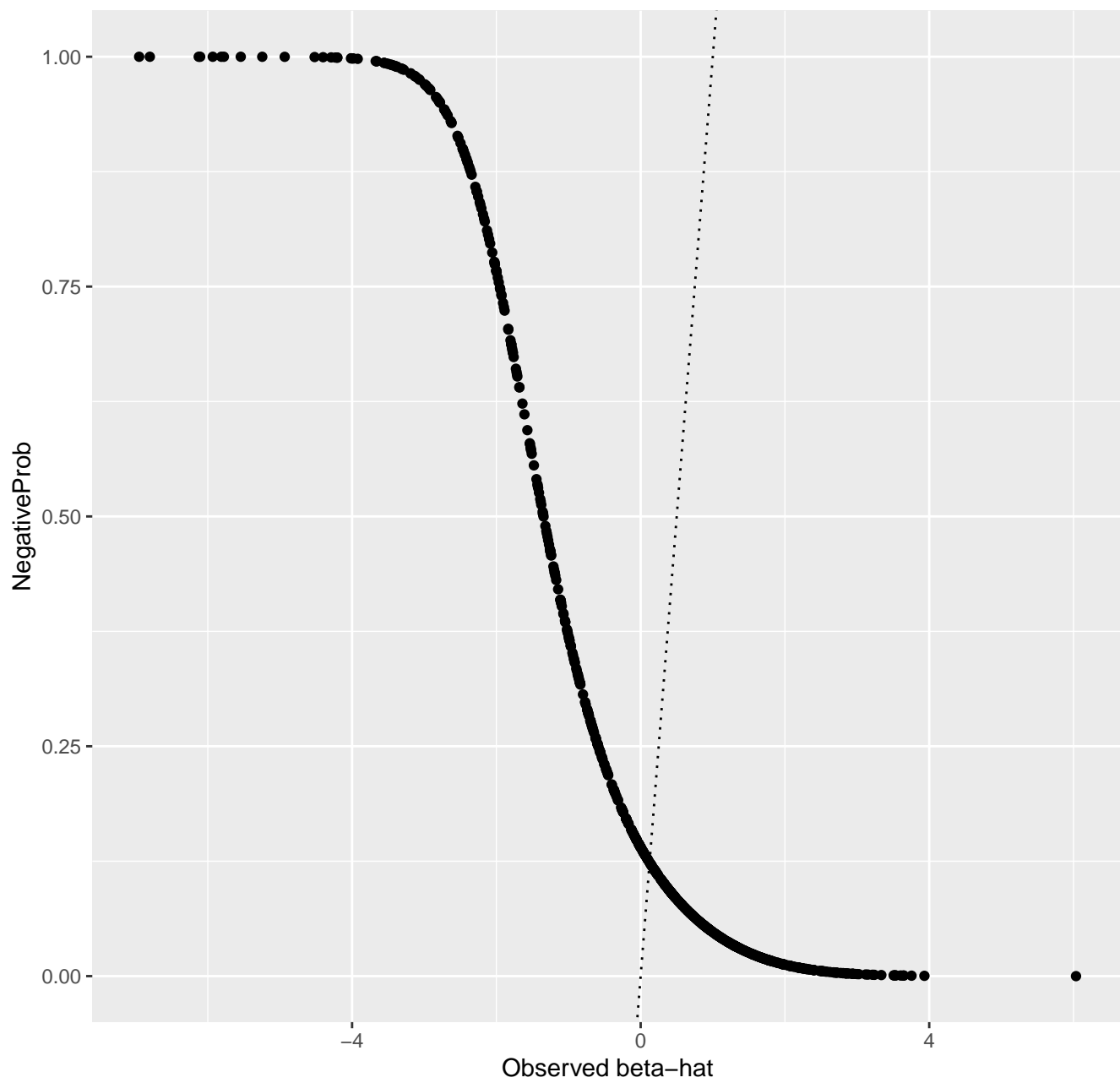
TNFAIP3_(7128) negative



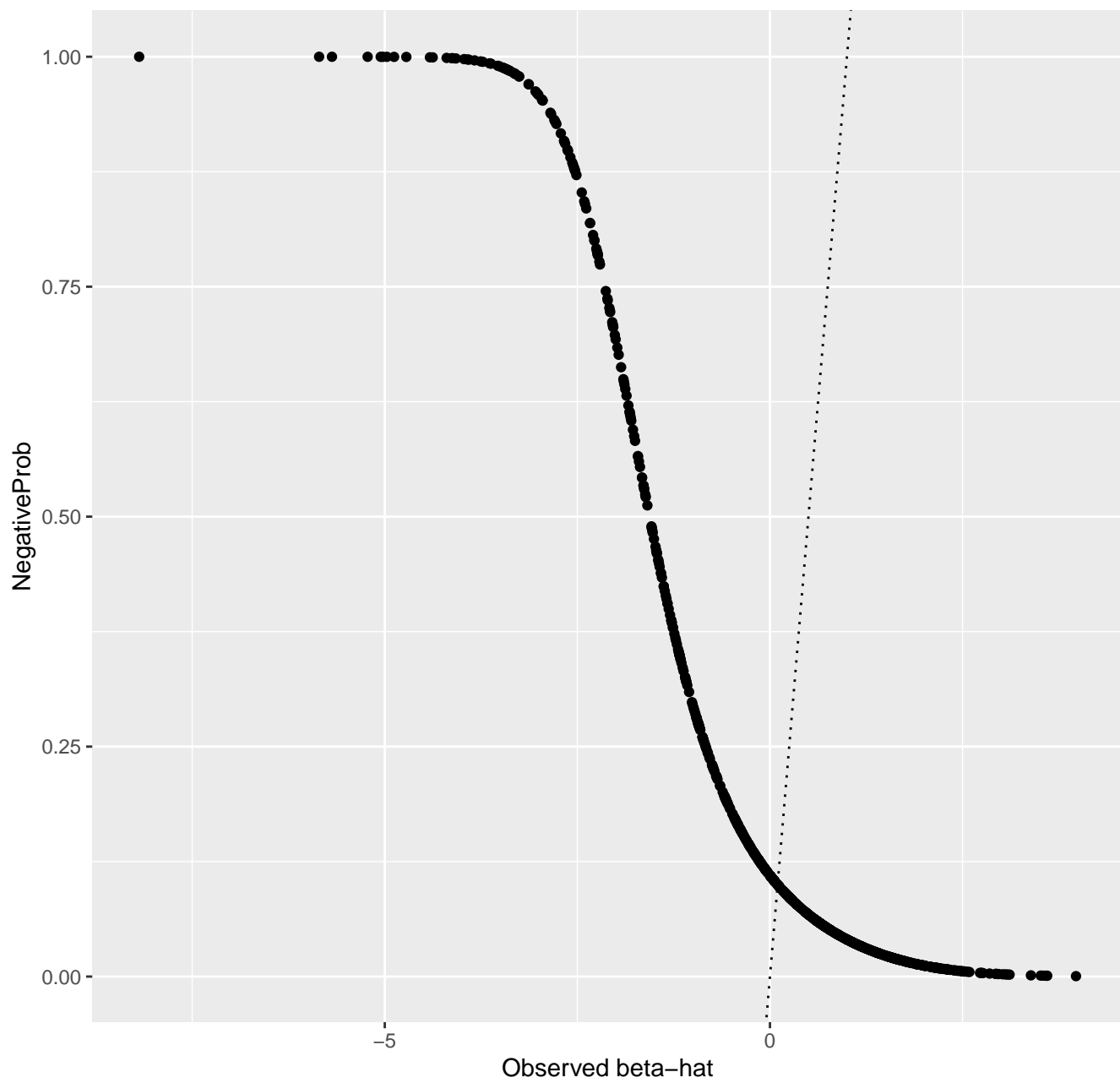
GF11_(2672) negative



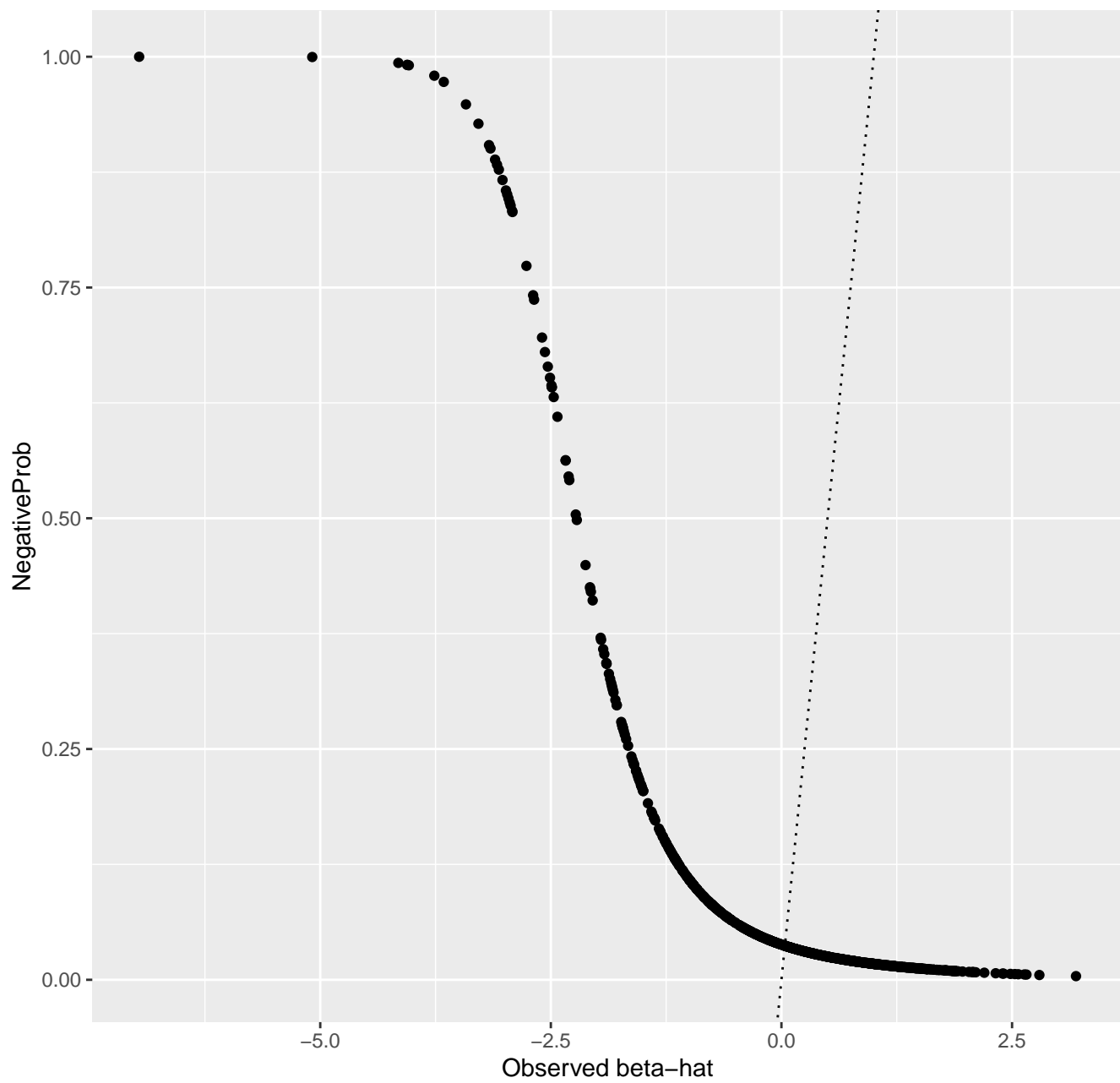
SPTSSA_(171546) negative



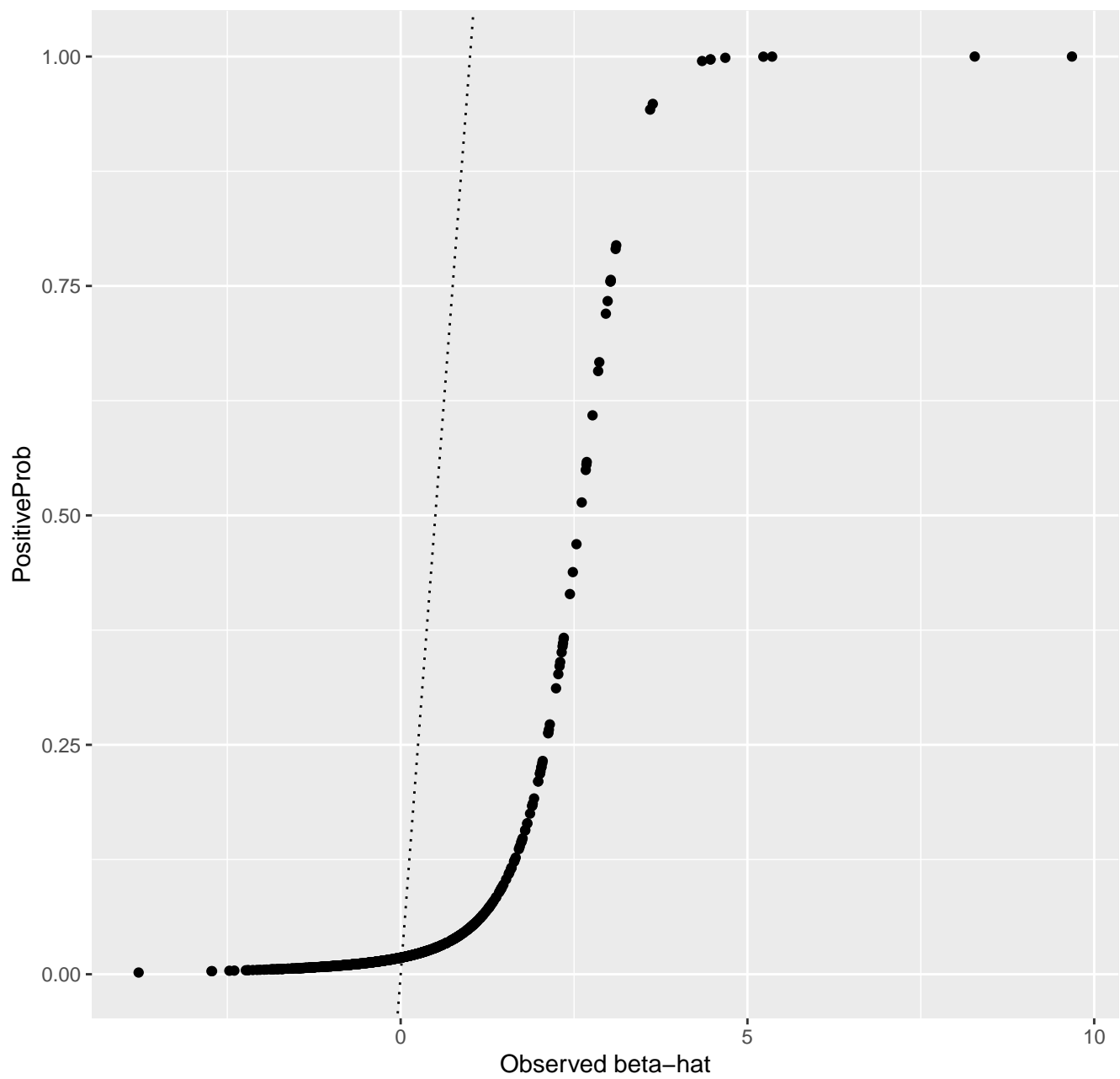
GRSF1_(2926) negative



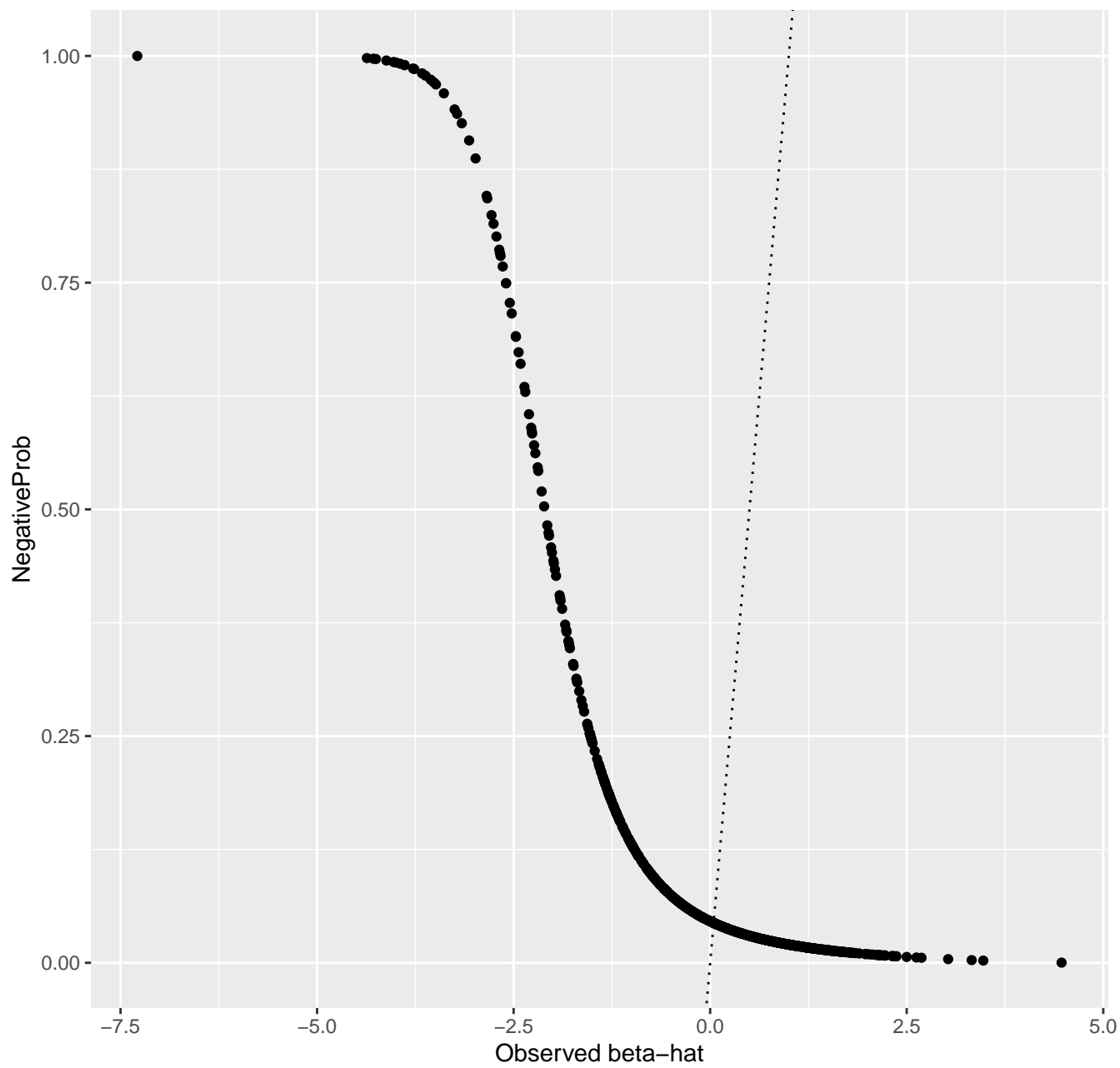
CLPX_(10845) negative



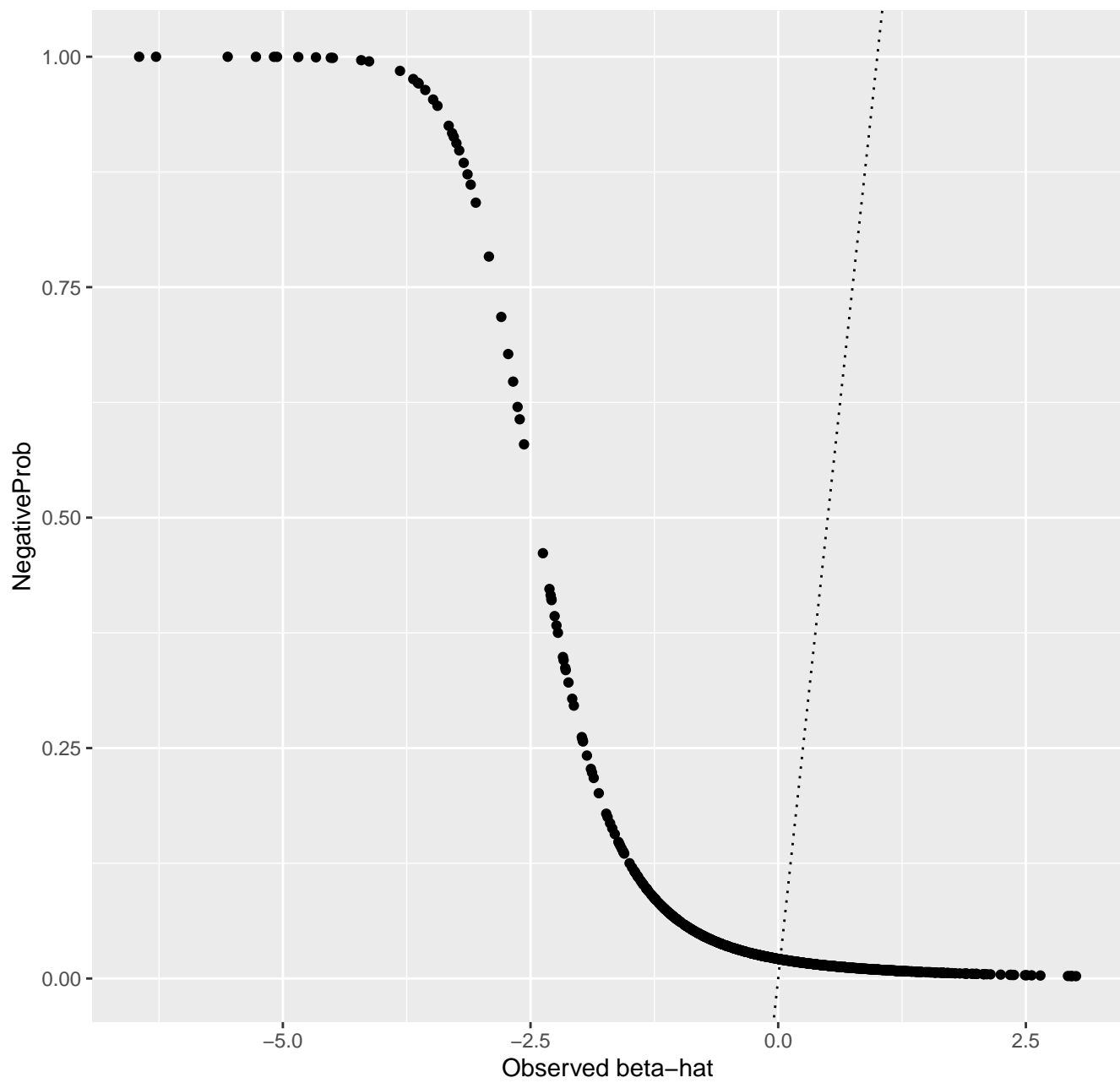
PLEKHA1_(59338) positive



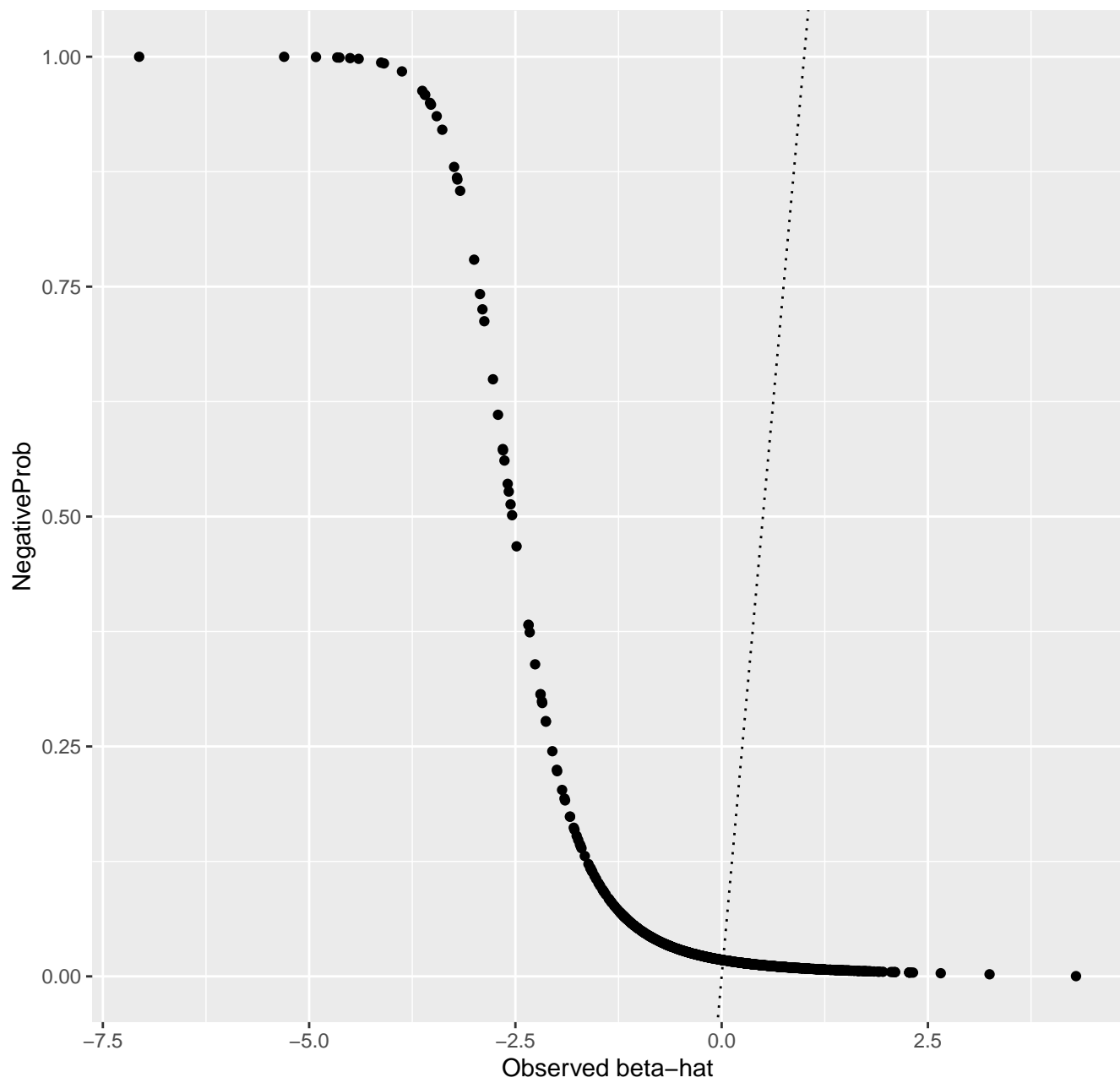
FNDC3B_(64778) negative



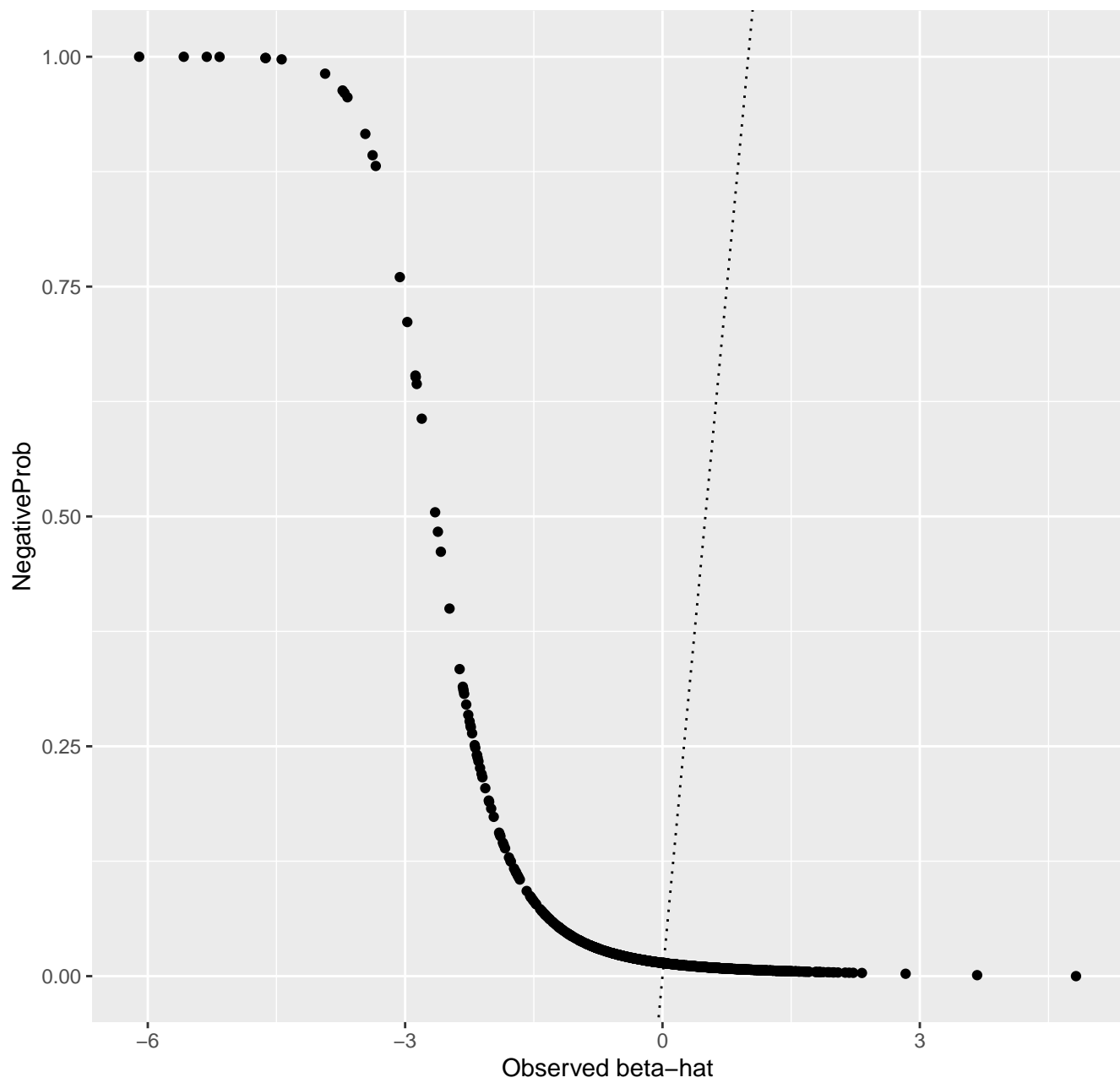
PEX12_(5193) negative



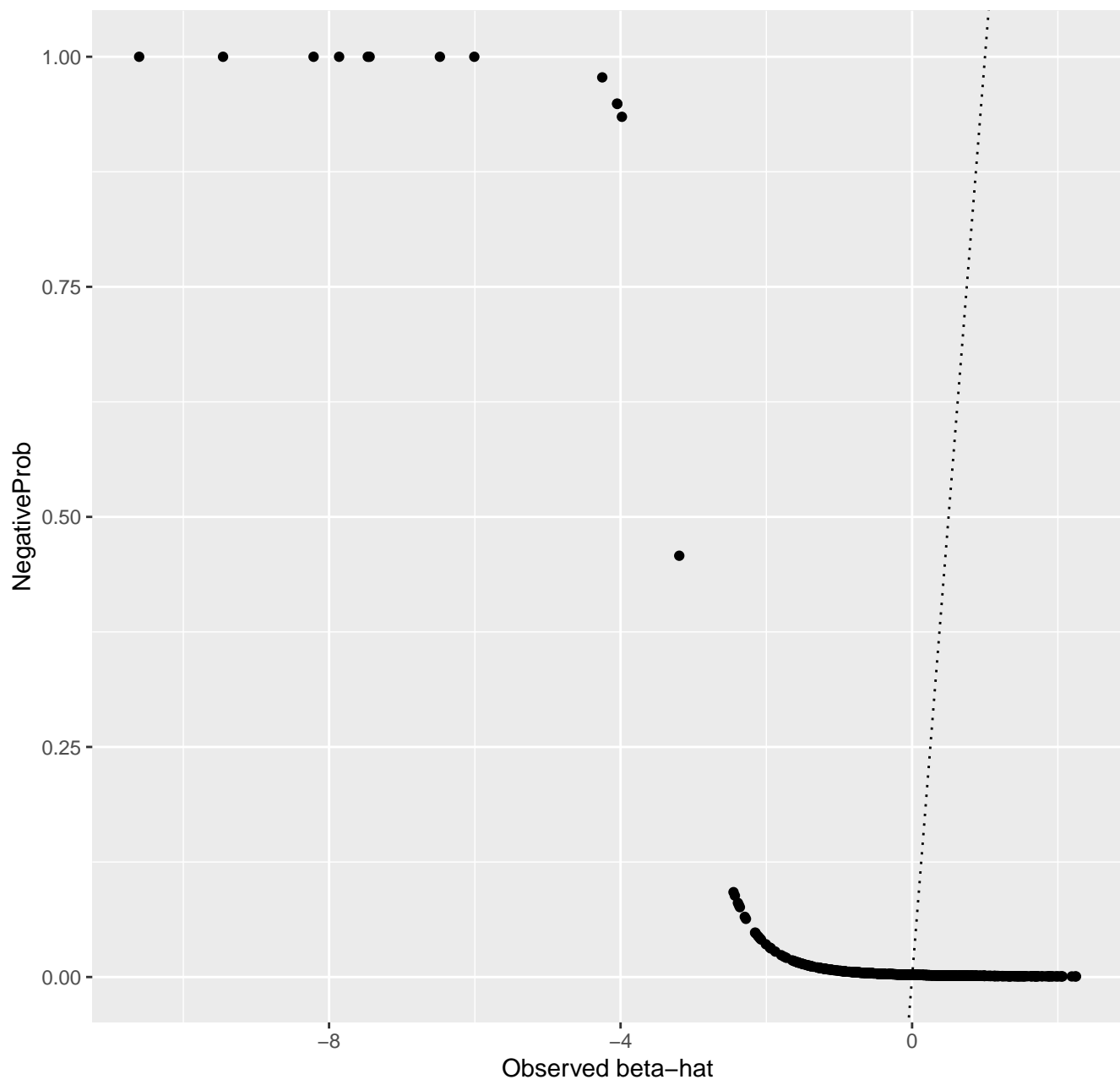
ATRN_(8455) negative



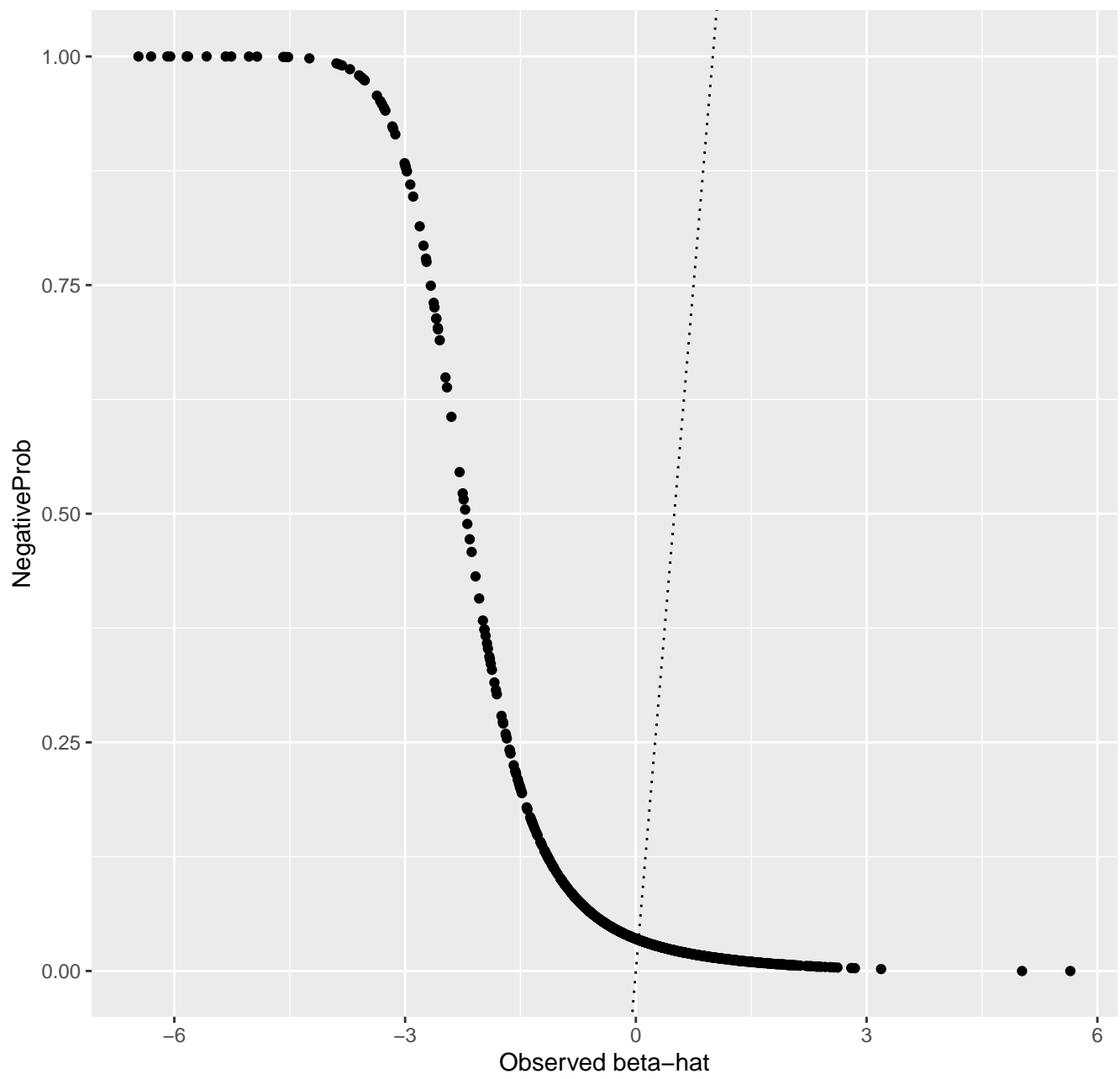
DNAJC3_(5611) negative



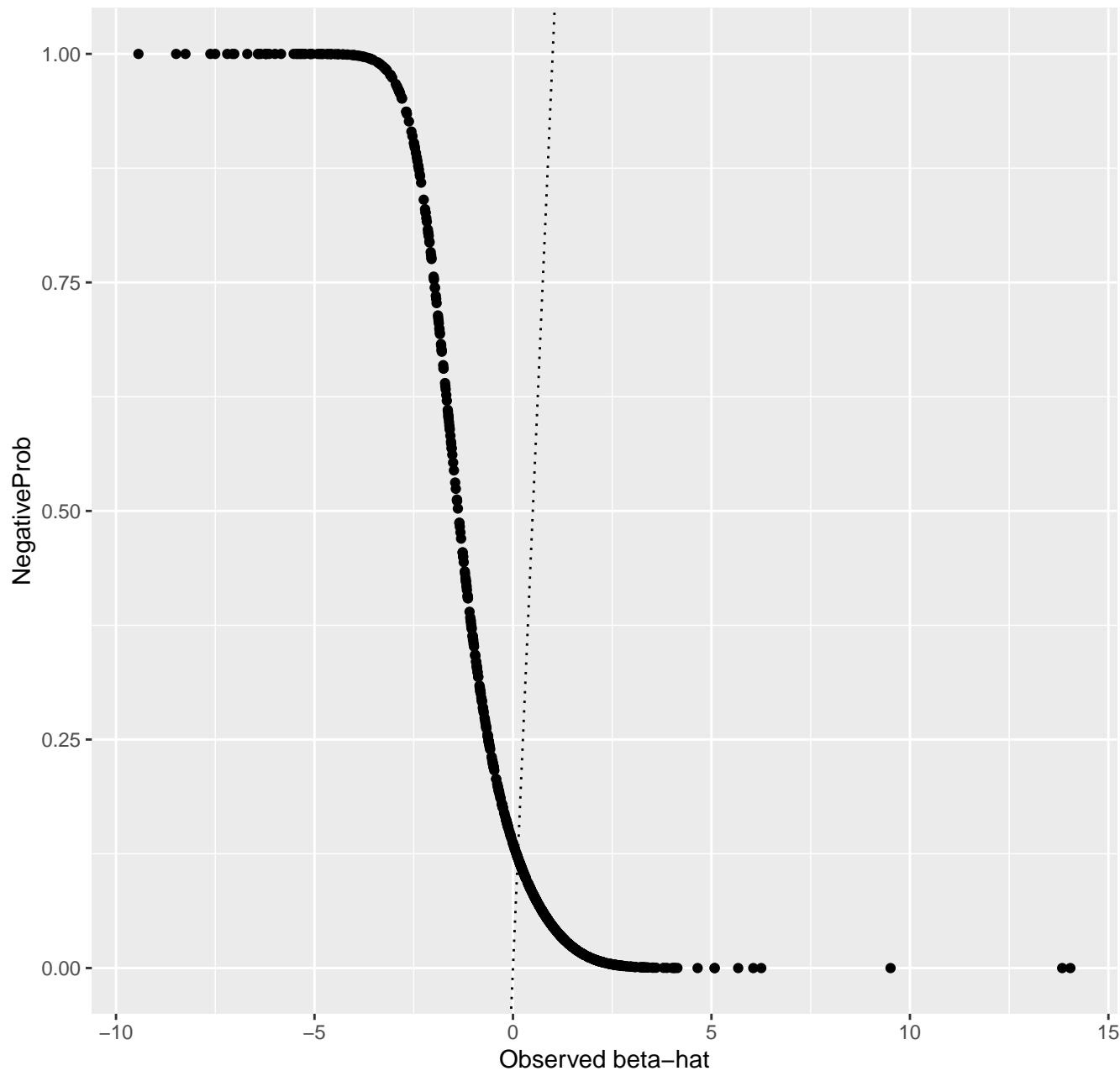
MYRF_(745) negative



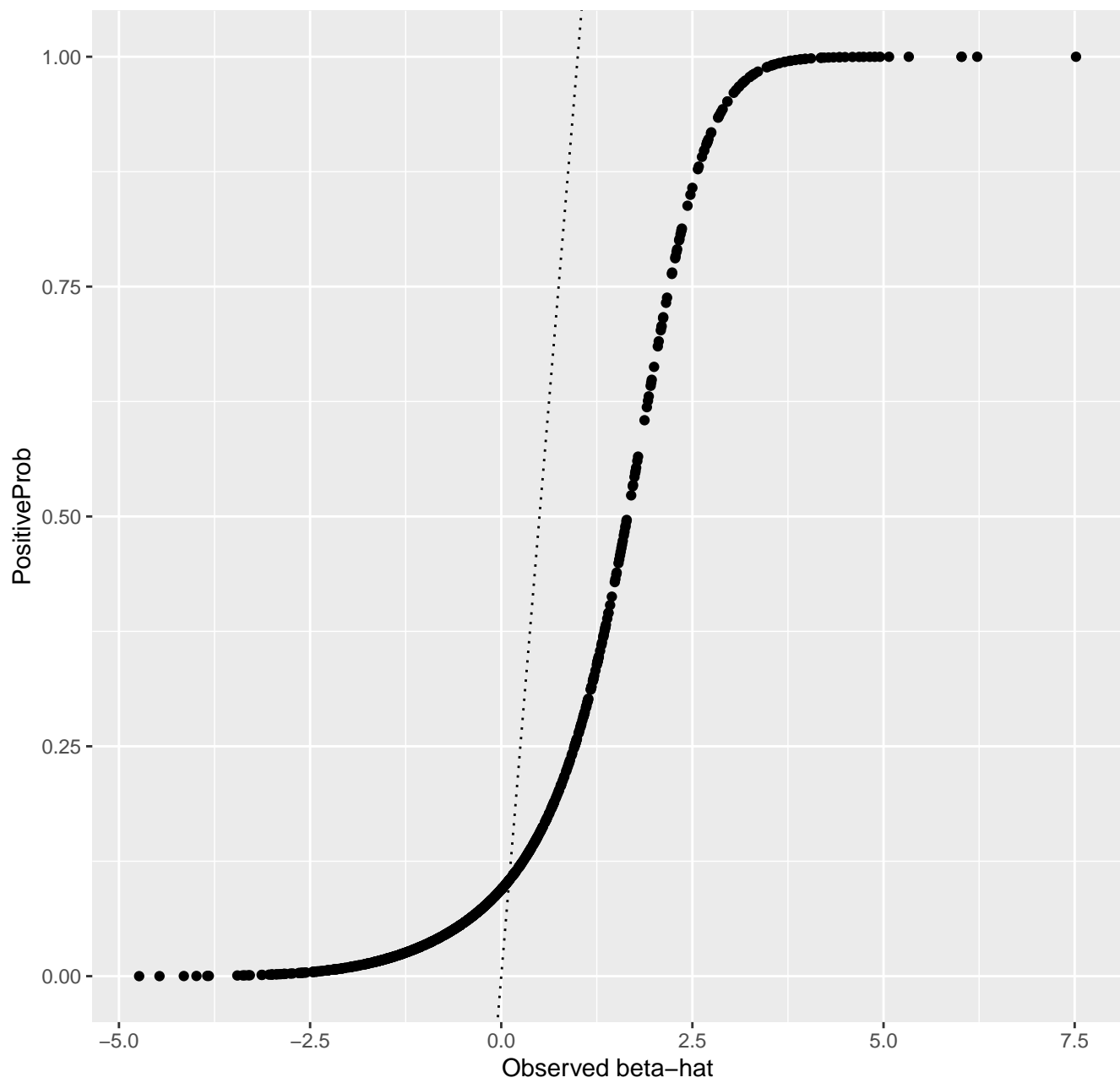
PEX26_(55670) negative



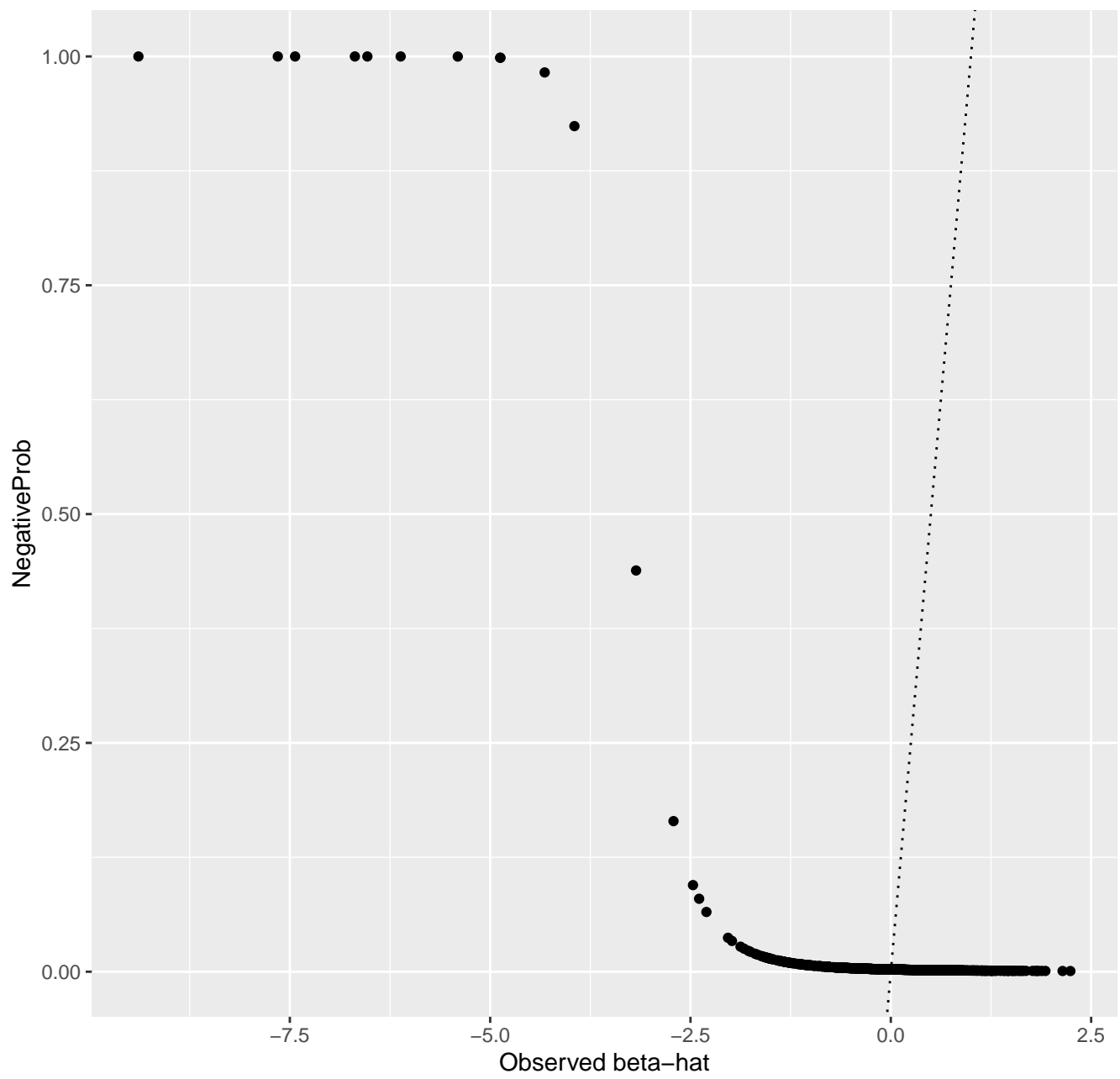
FASN_(2194) negative



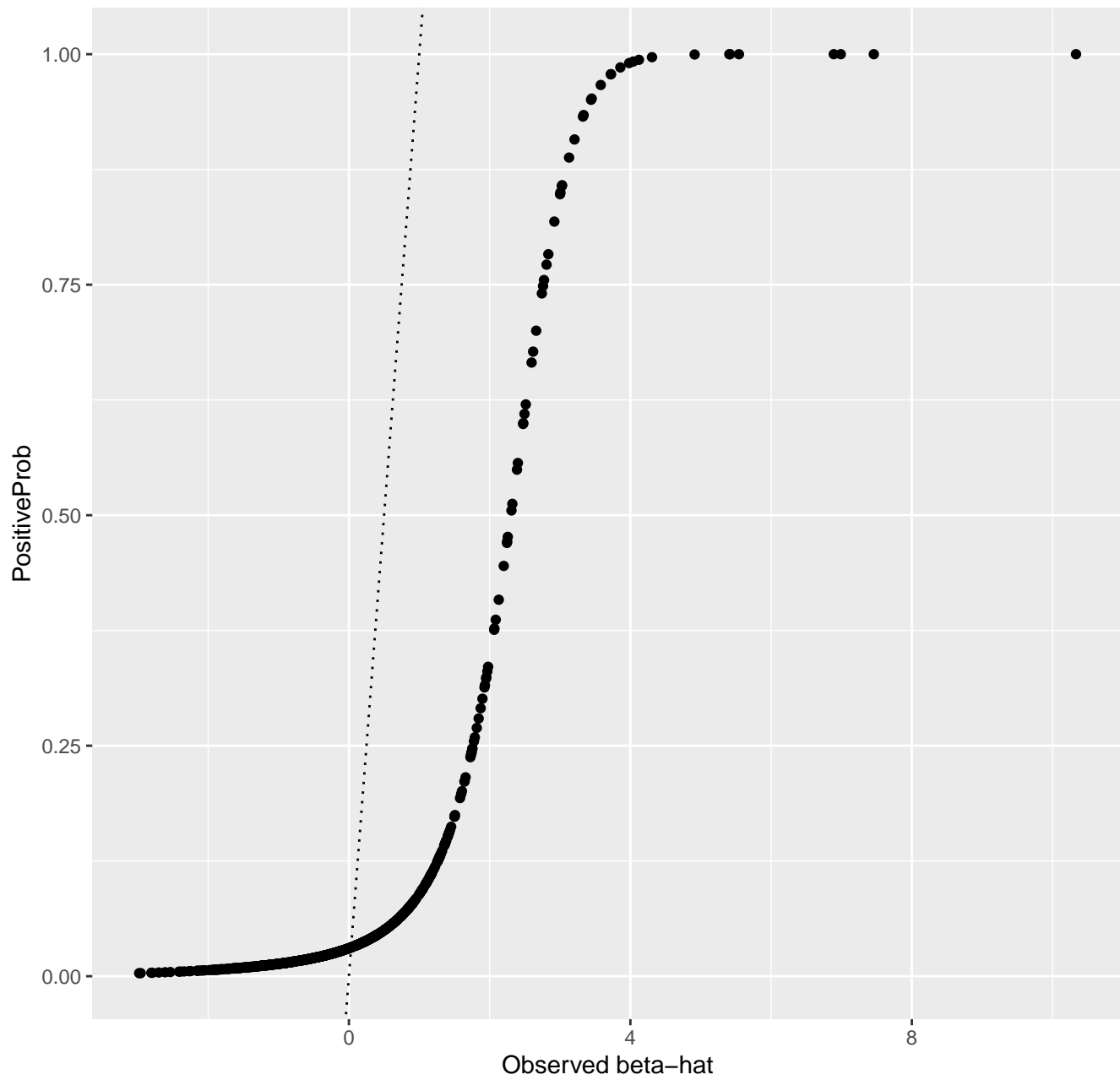
COPE_(11316) positive



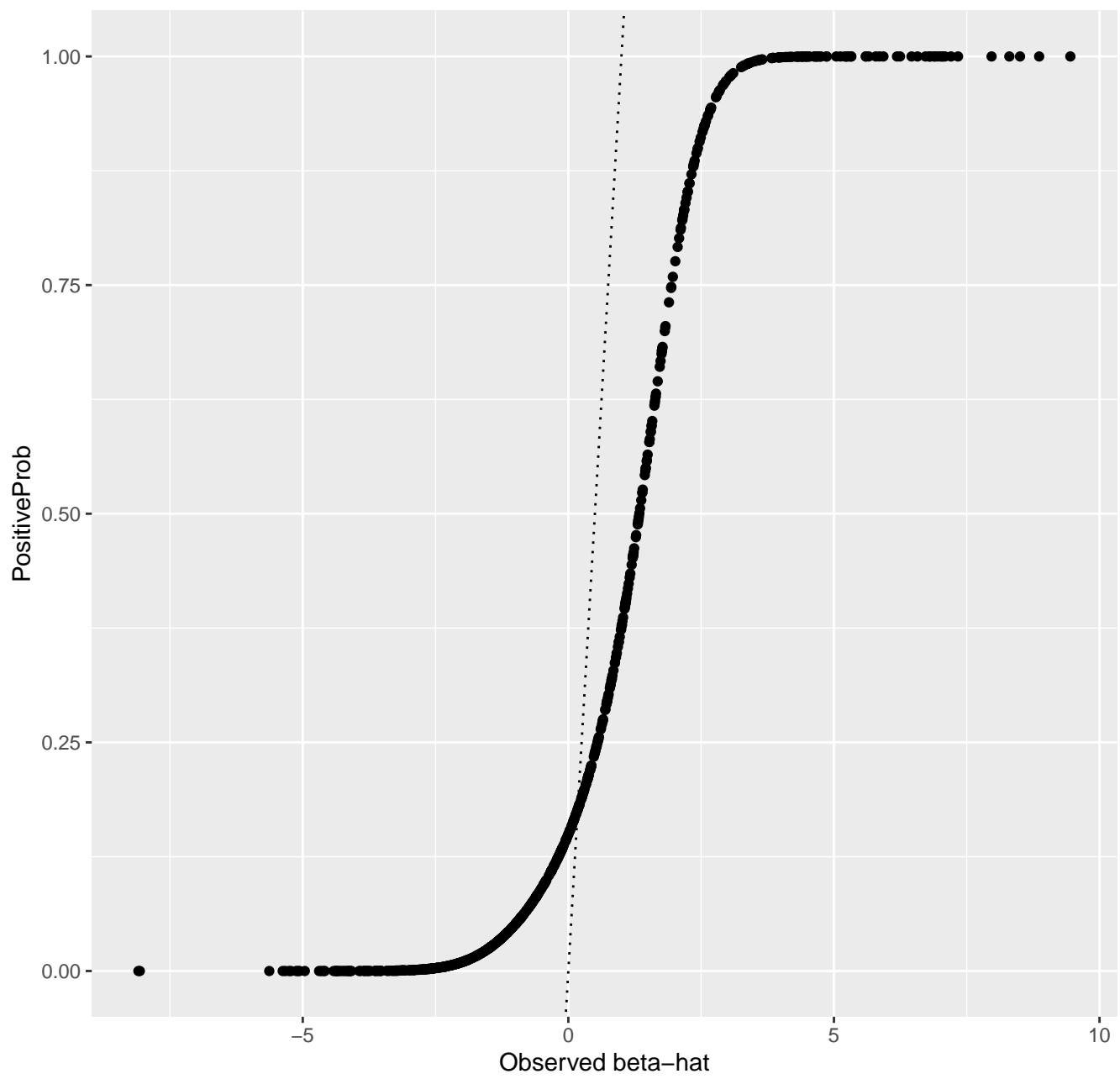
TFAP2B_(7021) negative



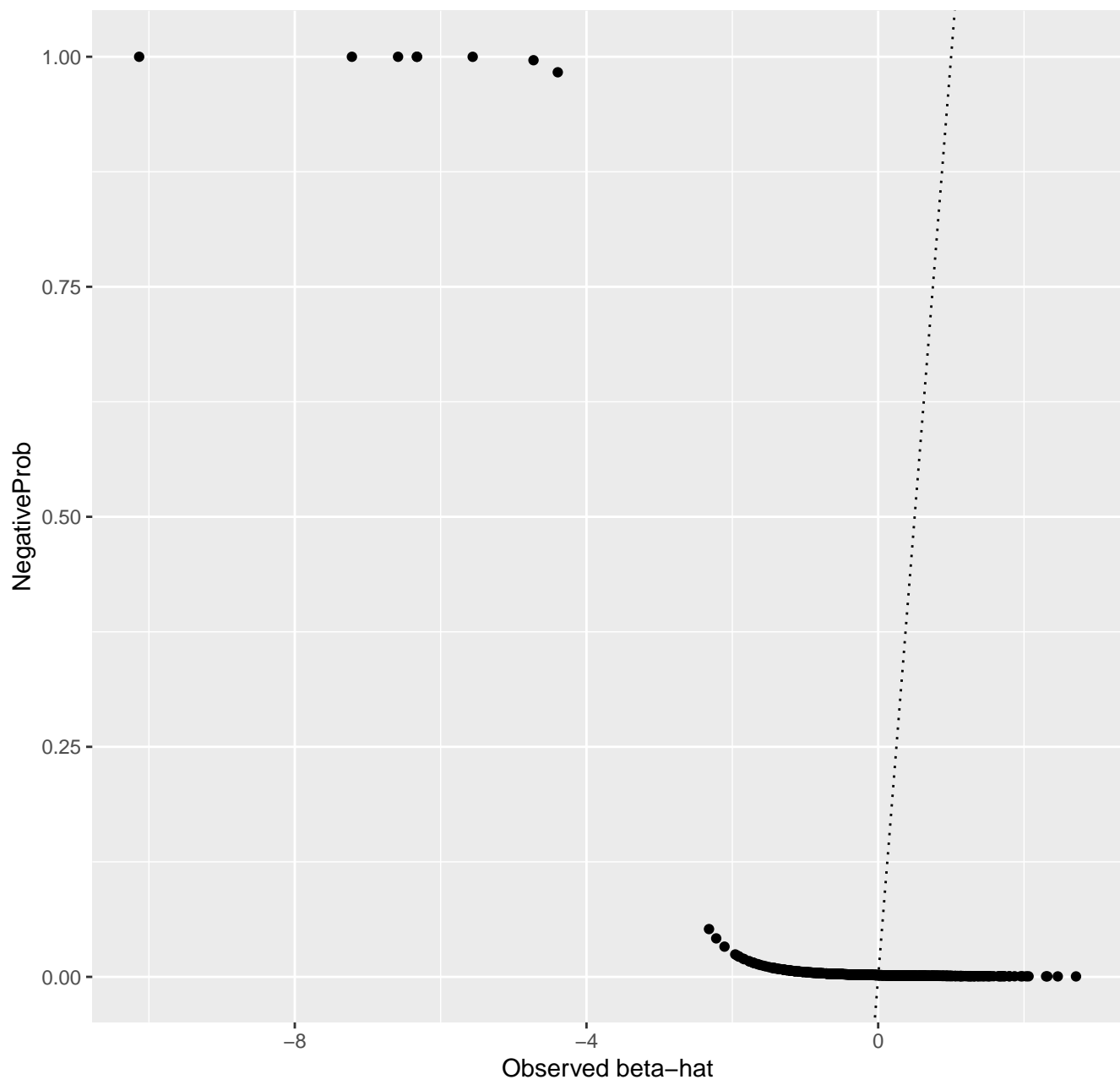
EPHA2_(1969) positive



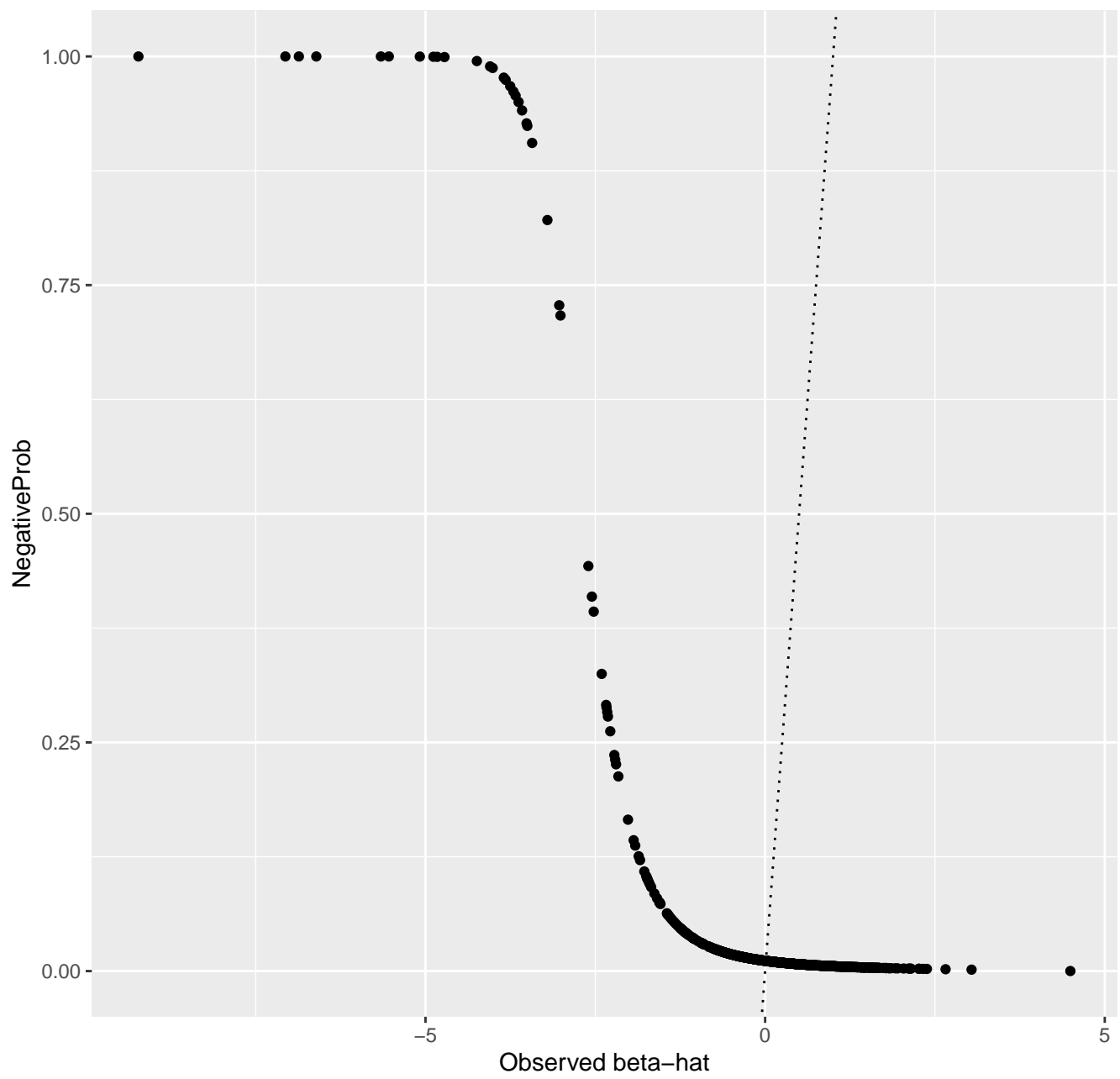
THAP1_(55145) positive



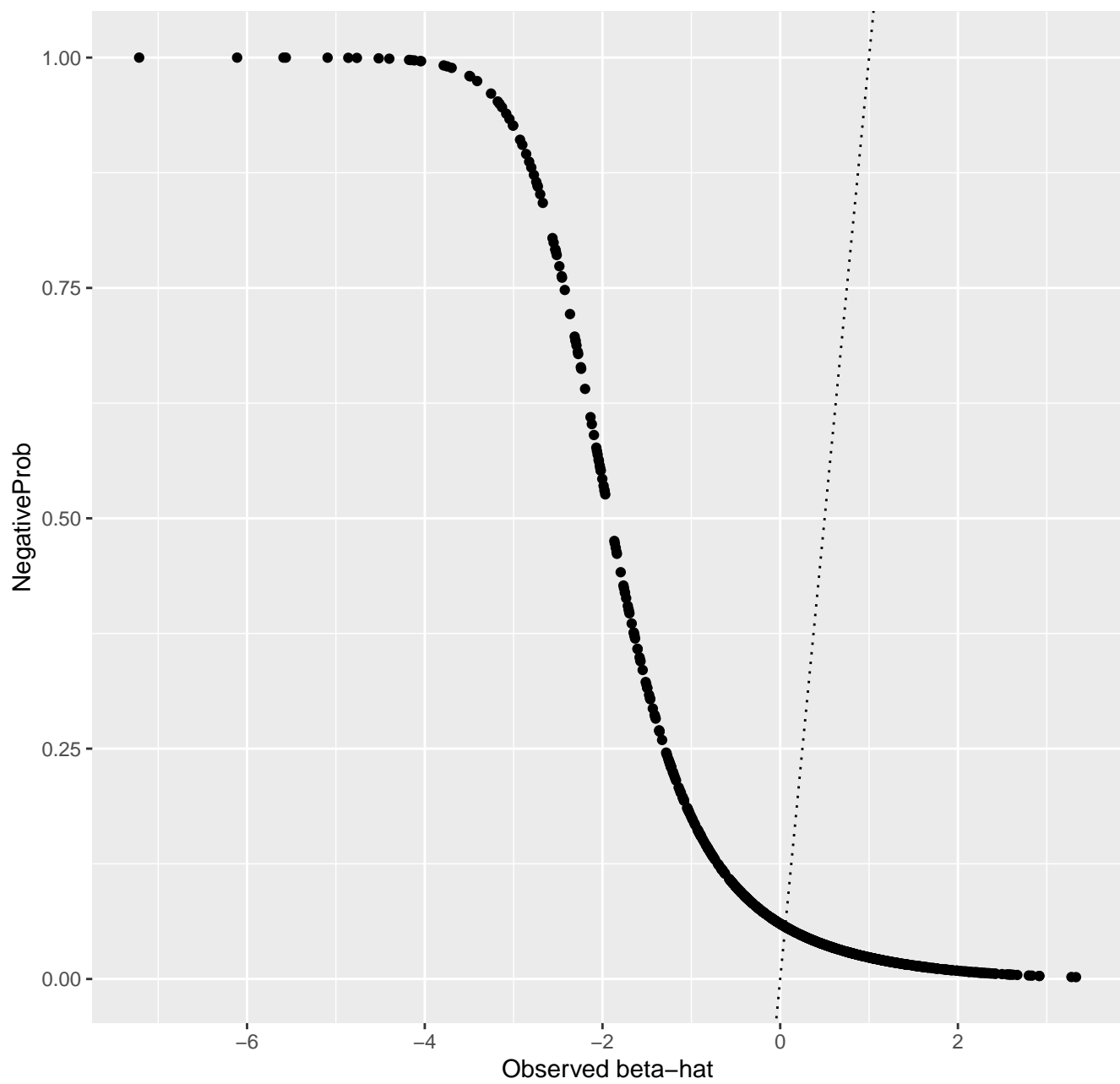
ESR1_(2099) negative



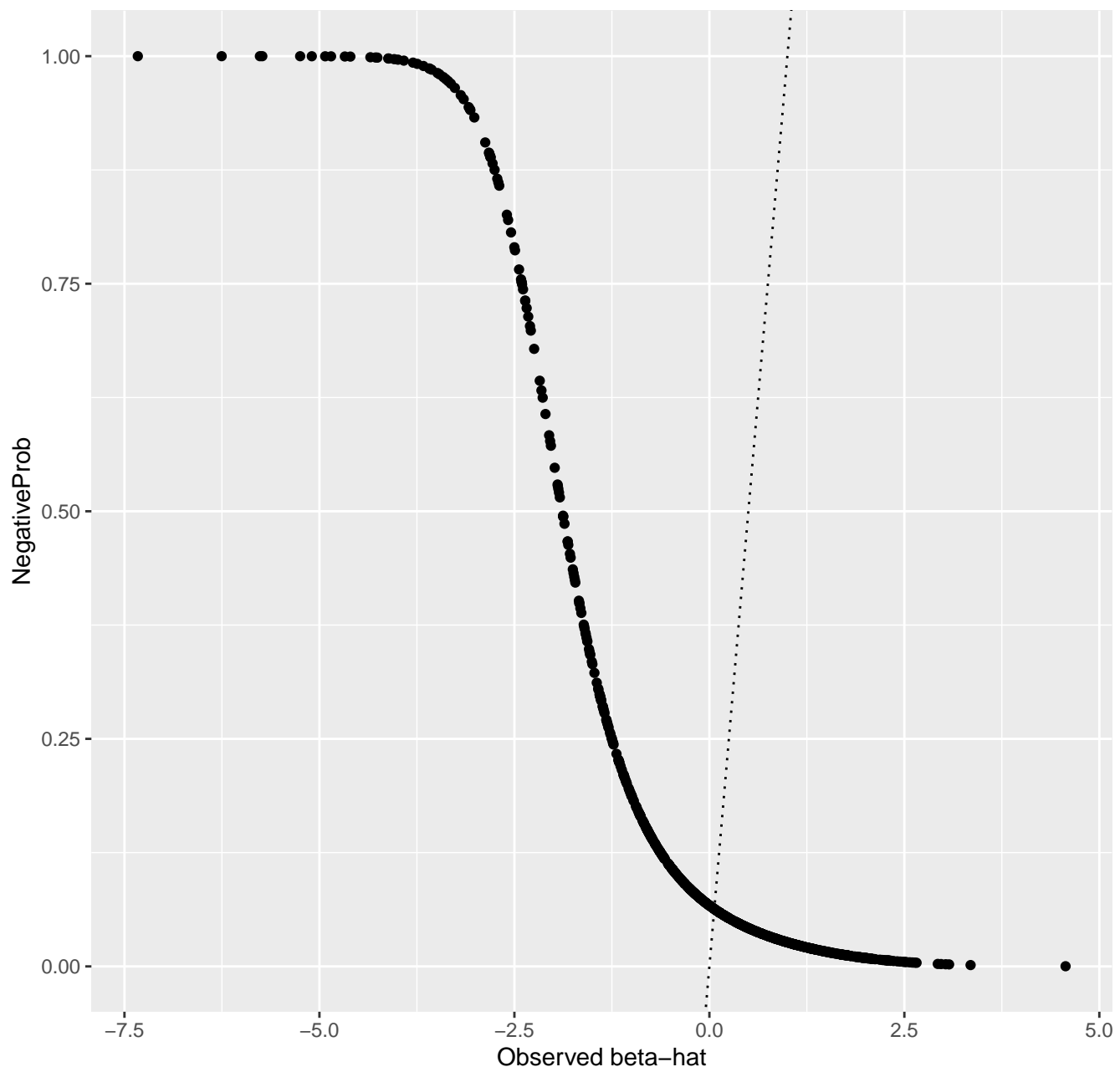
IKZF1_(10320) negative



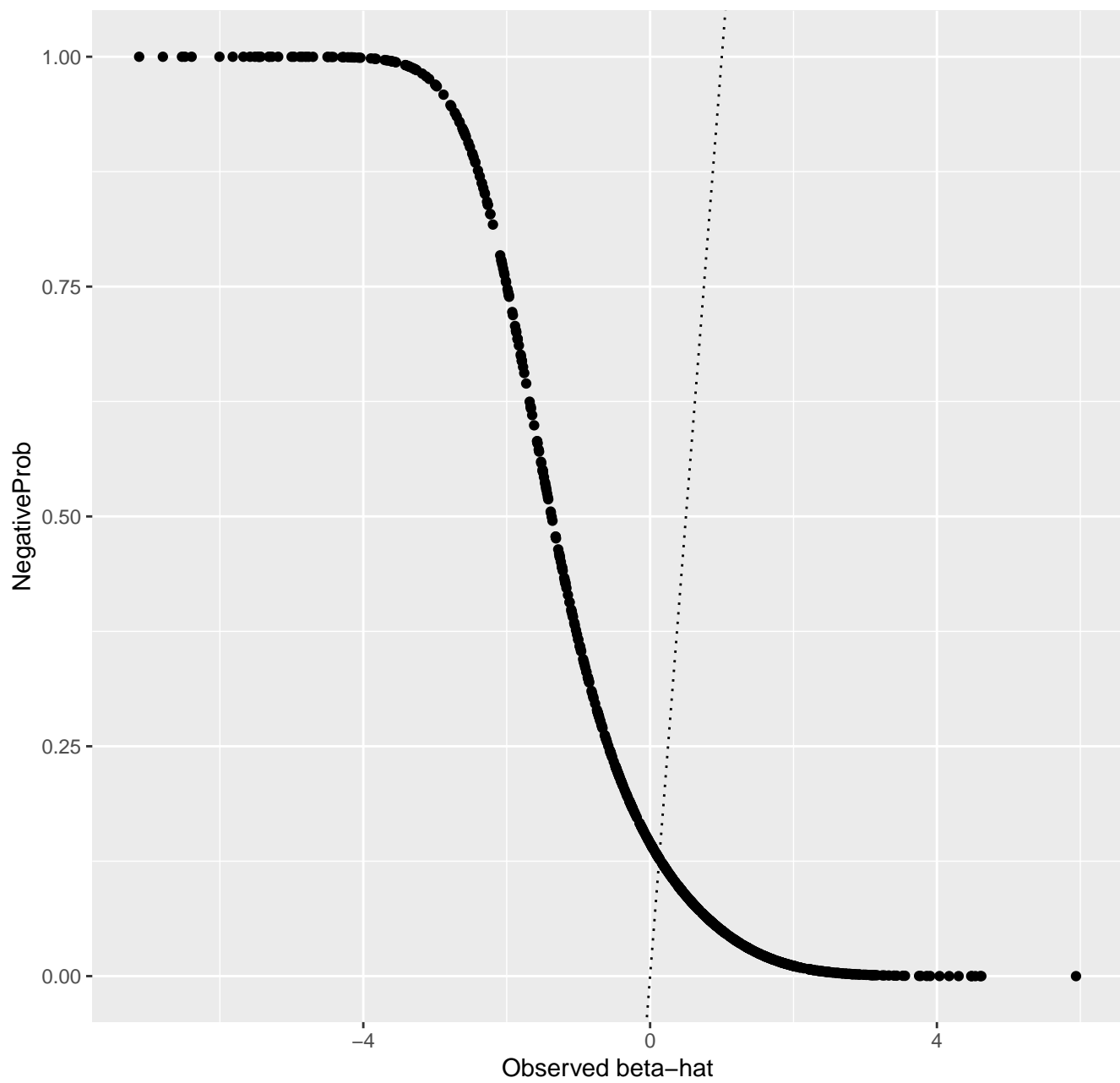
NGLY1_(55768) negative



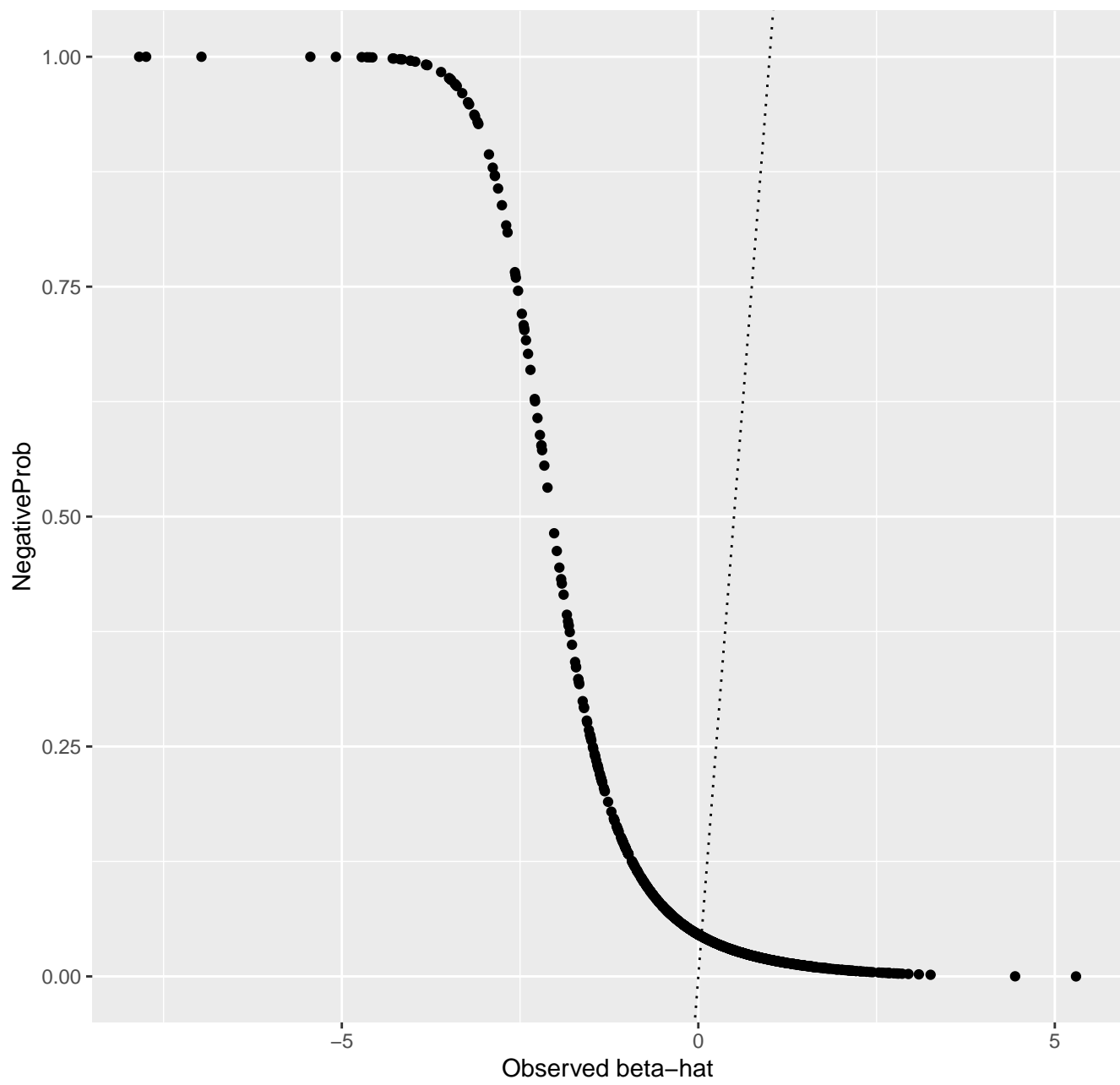
TMEM167A_(153339) negative



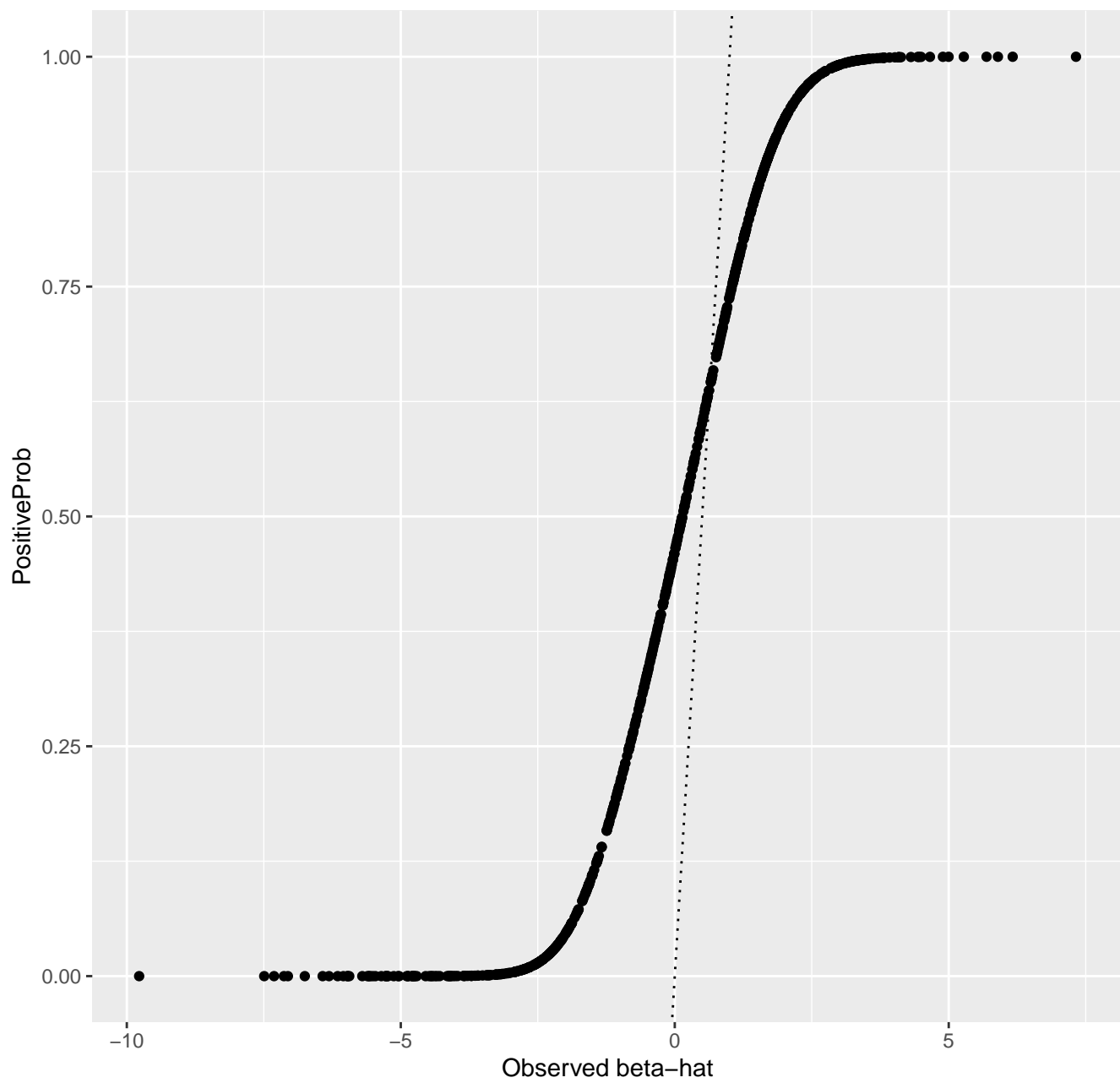
NDUFA8_(4702) negative



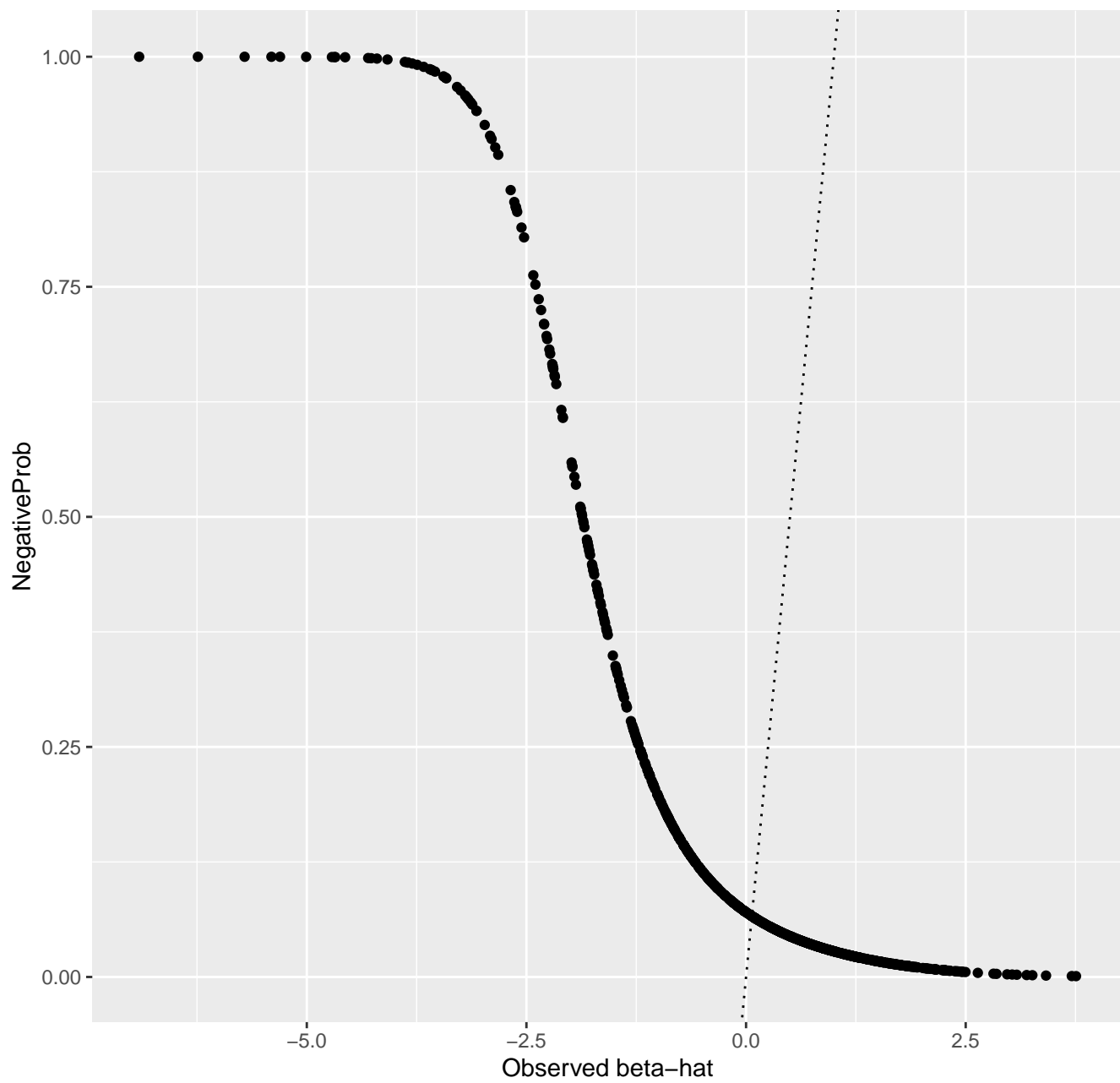
GCLC_(2729) negative



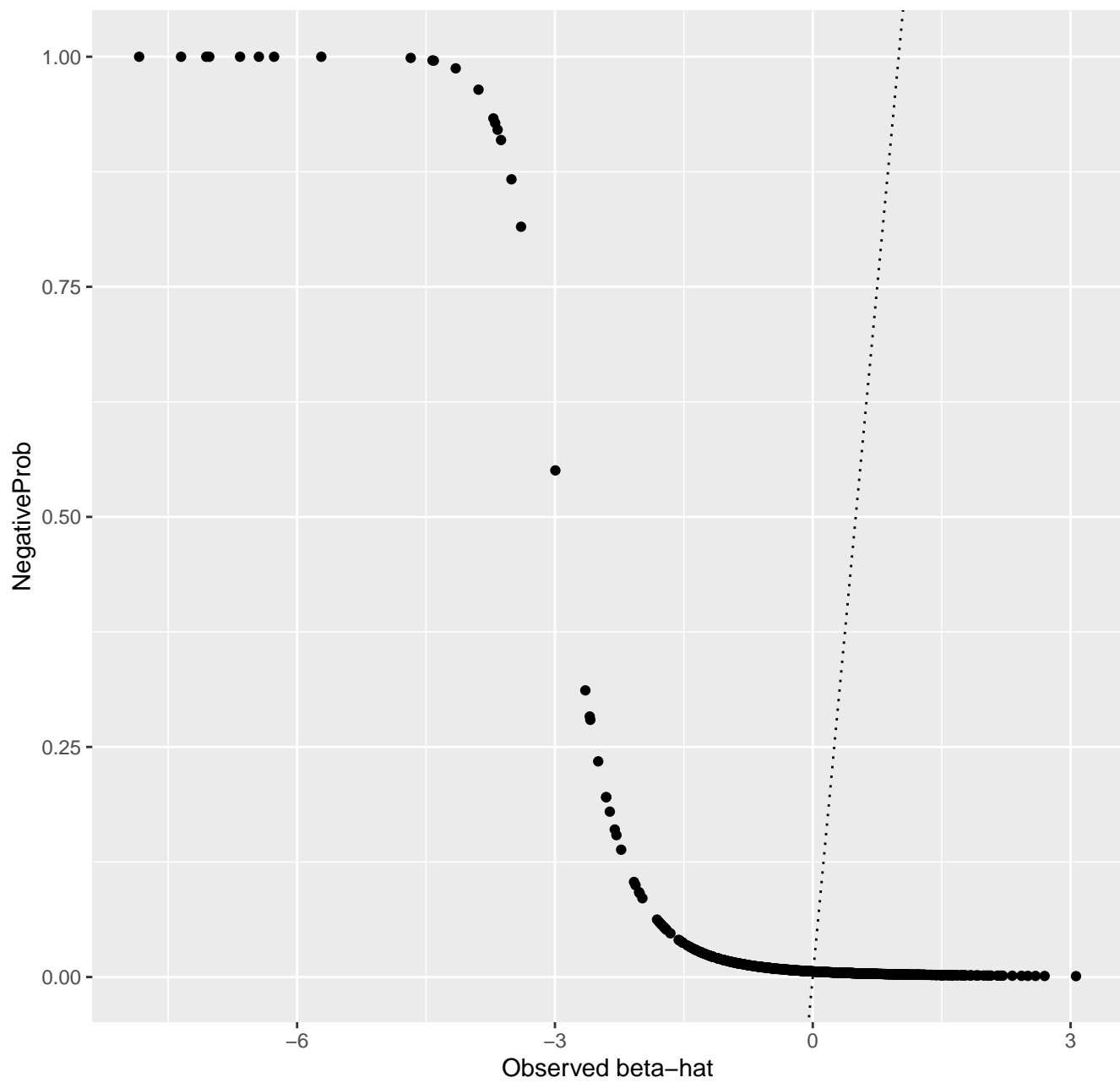
PGM3_(5238) positive



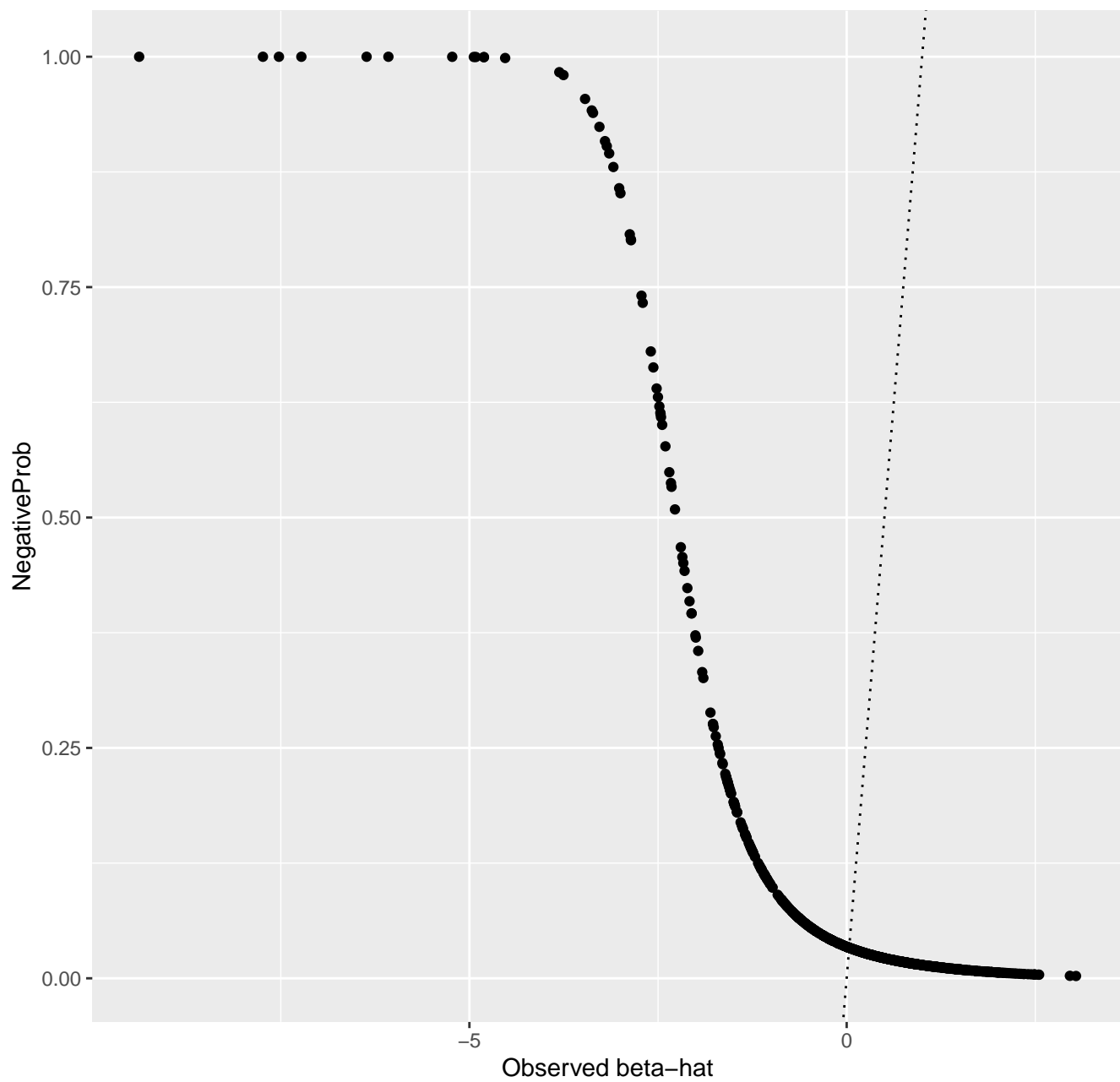
TENT4B_(64282) negative



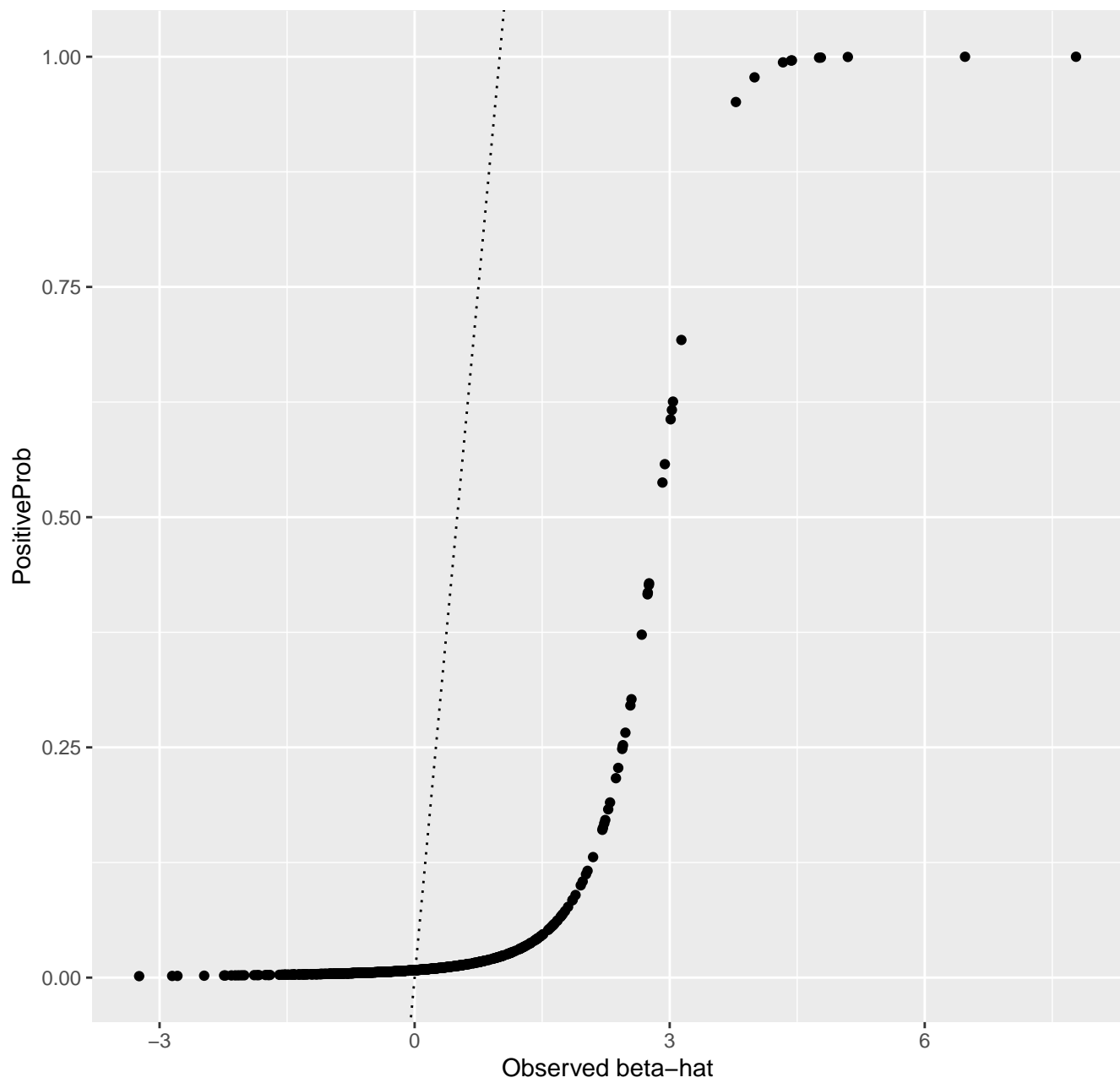
HNF1A_(6927) negative



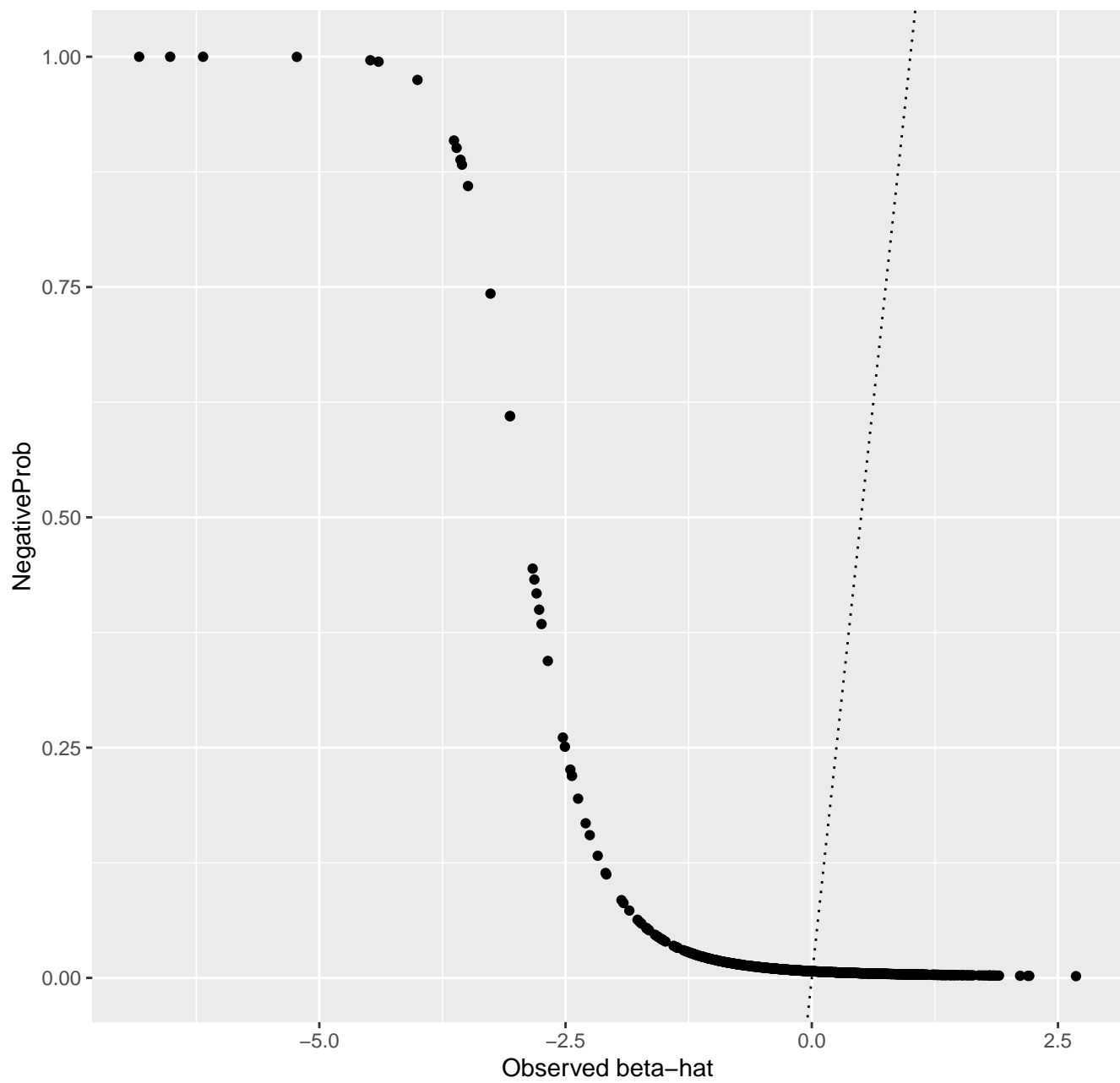
CUL4B_(8450) negative



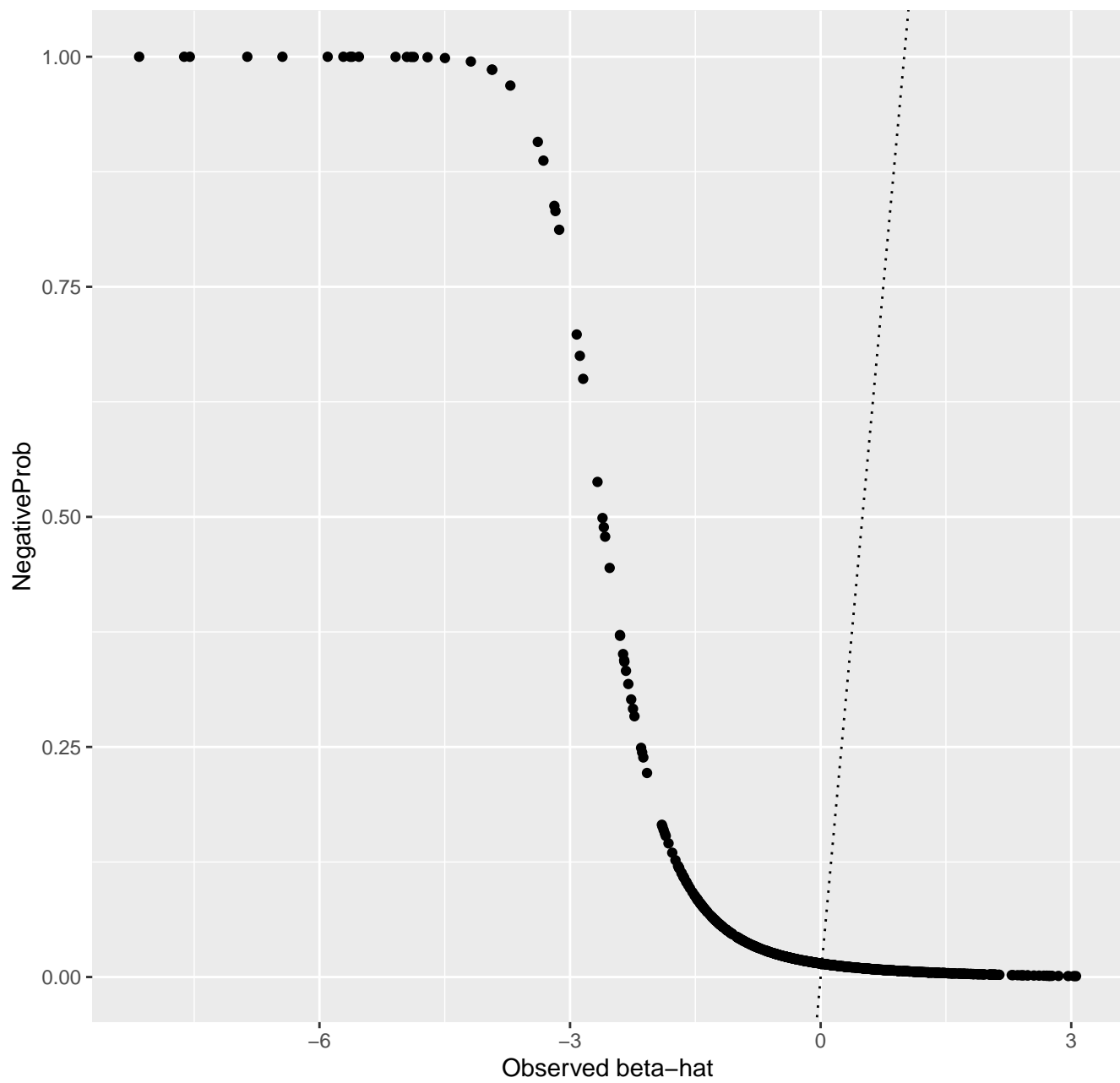
PLXNA1_(5361) positive



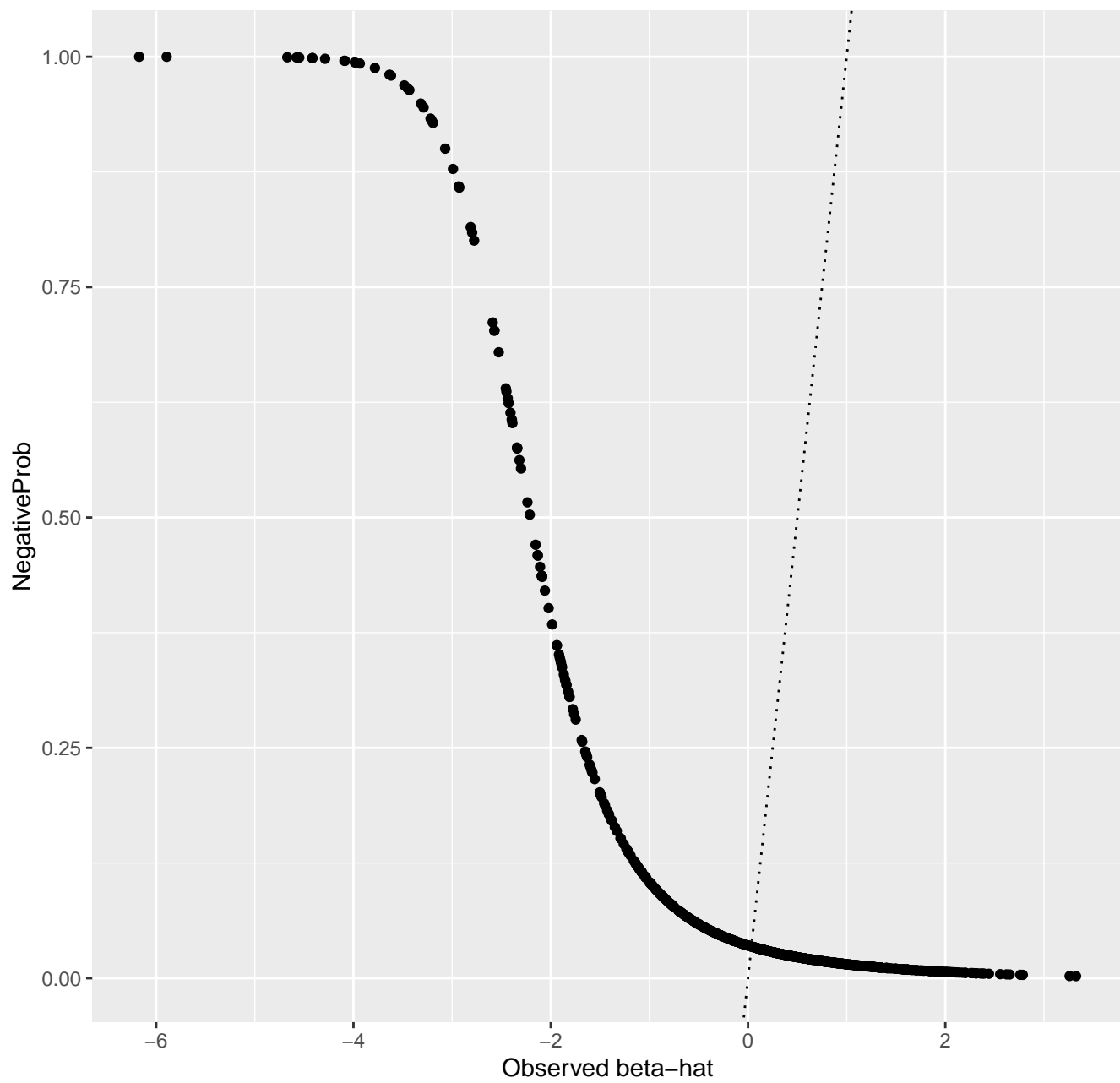
RAB3GAP1_(22930) negative



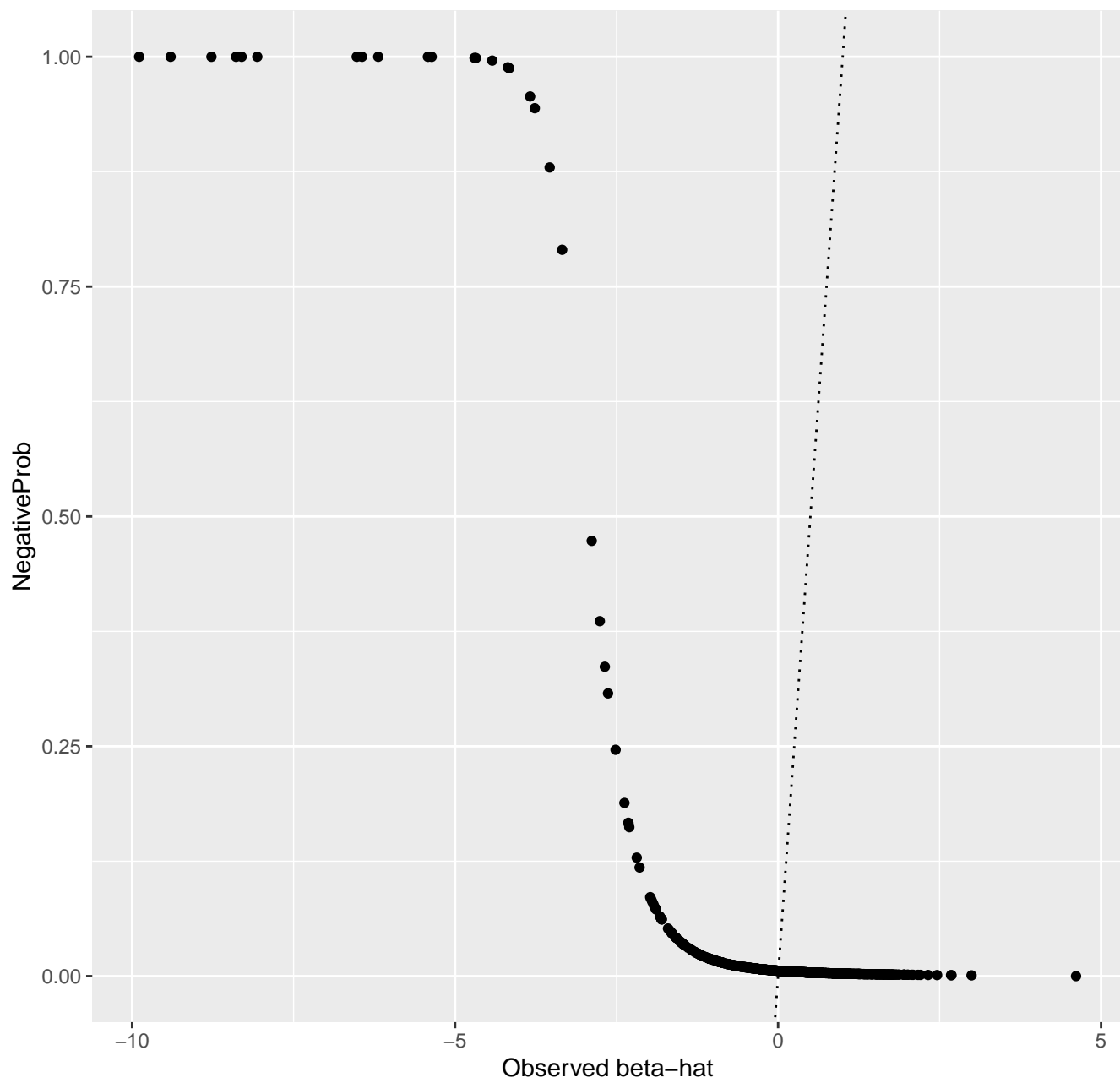
STAG1_(10274) negative



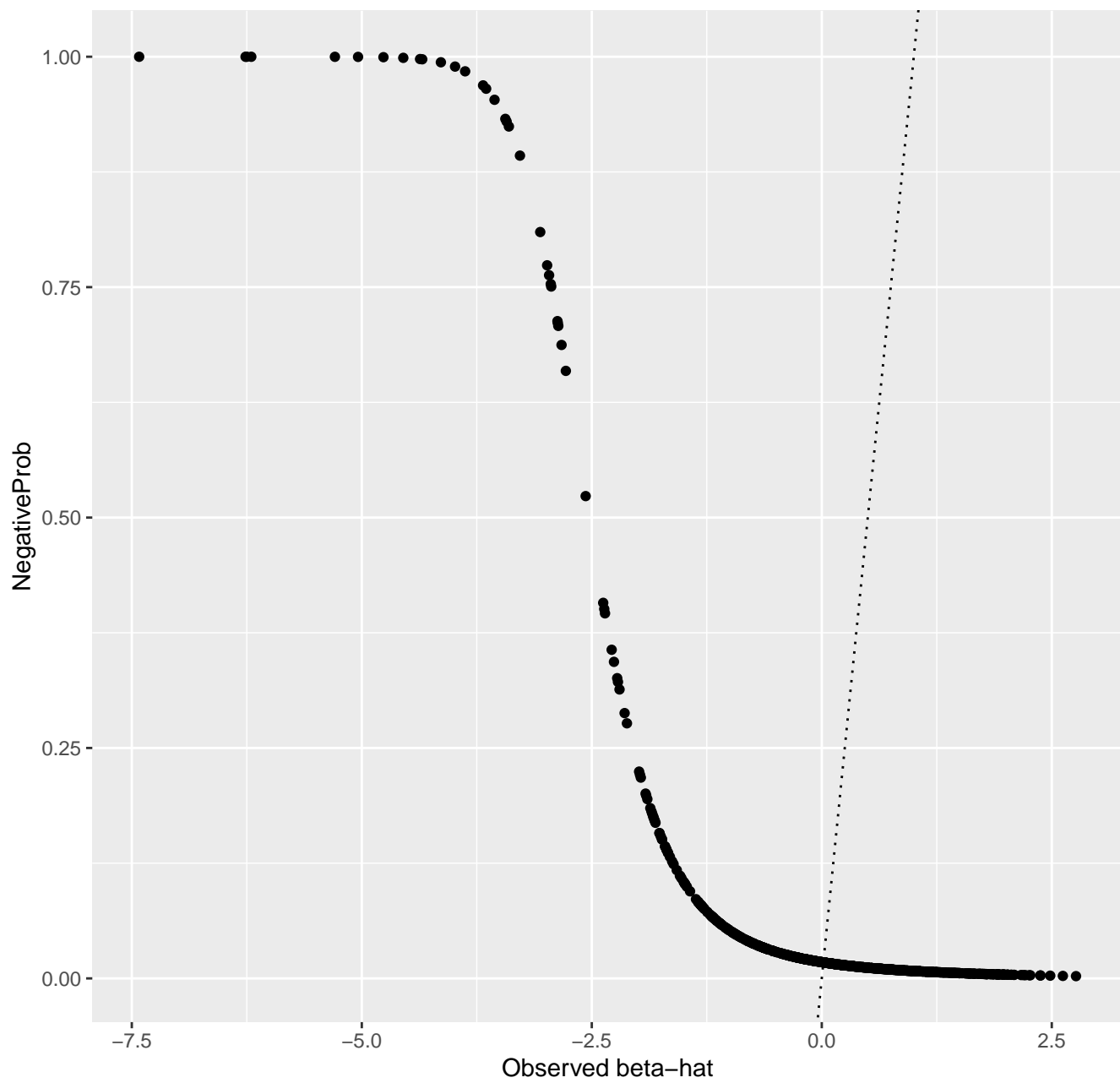
INPP5K_(51763) negative



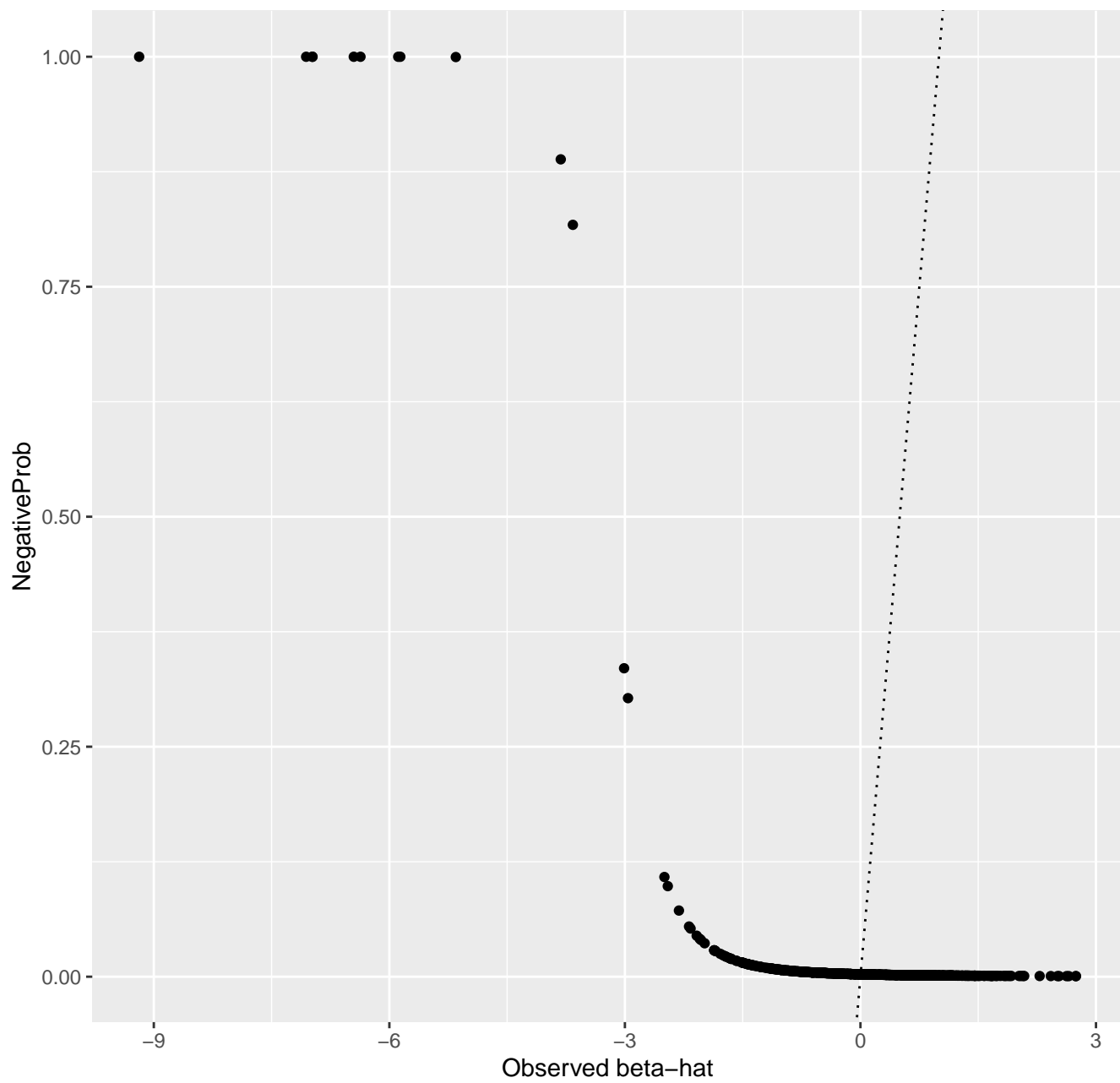
SPI1_(6688) negative



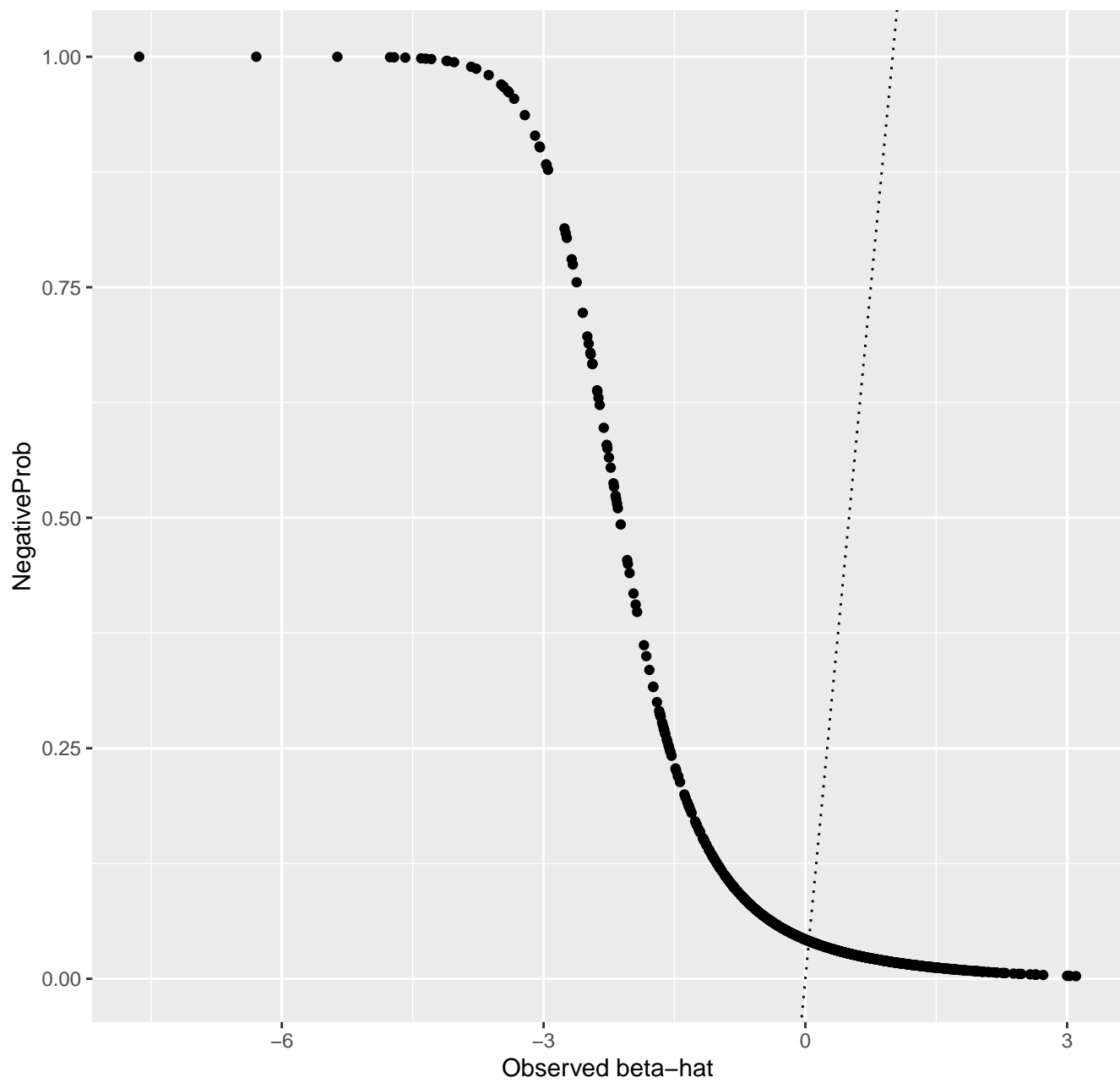
EXO1_(9156) negative



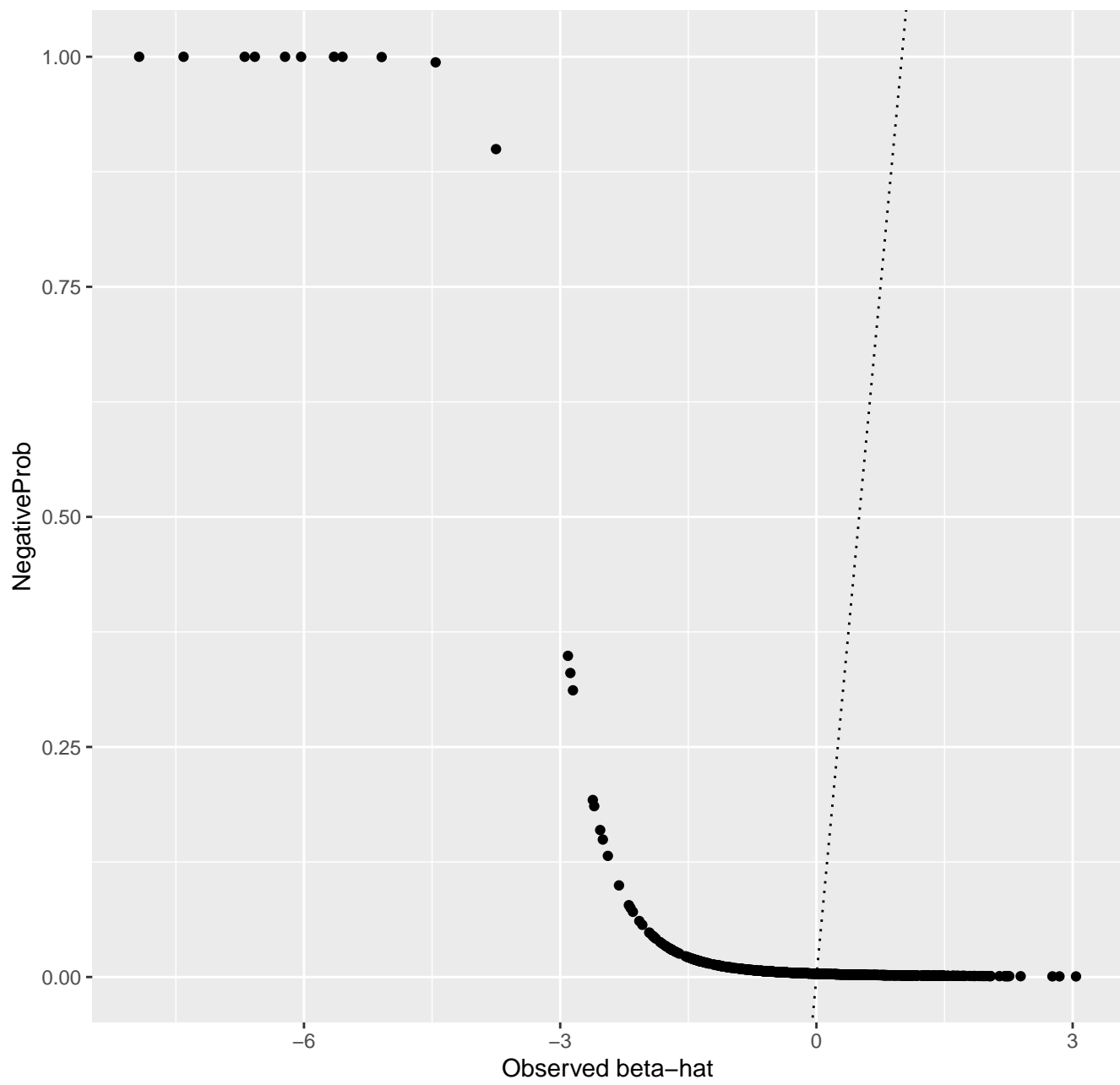
UBE3A_(7337) negative



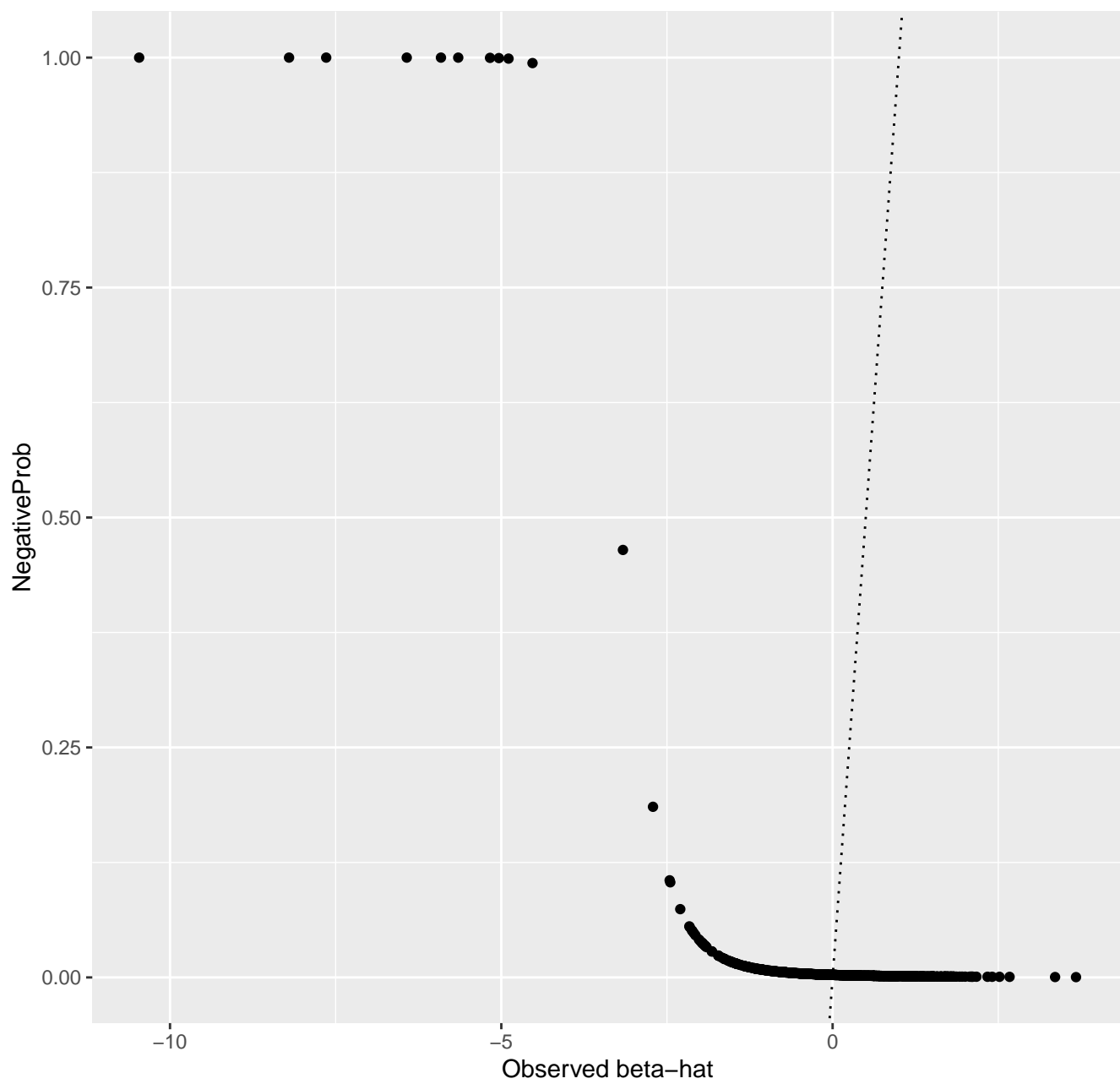
RIMBP3_(85376) negative



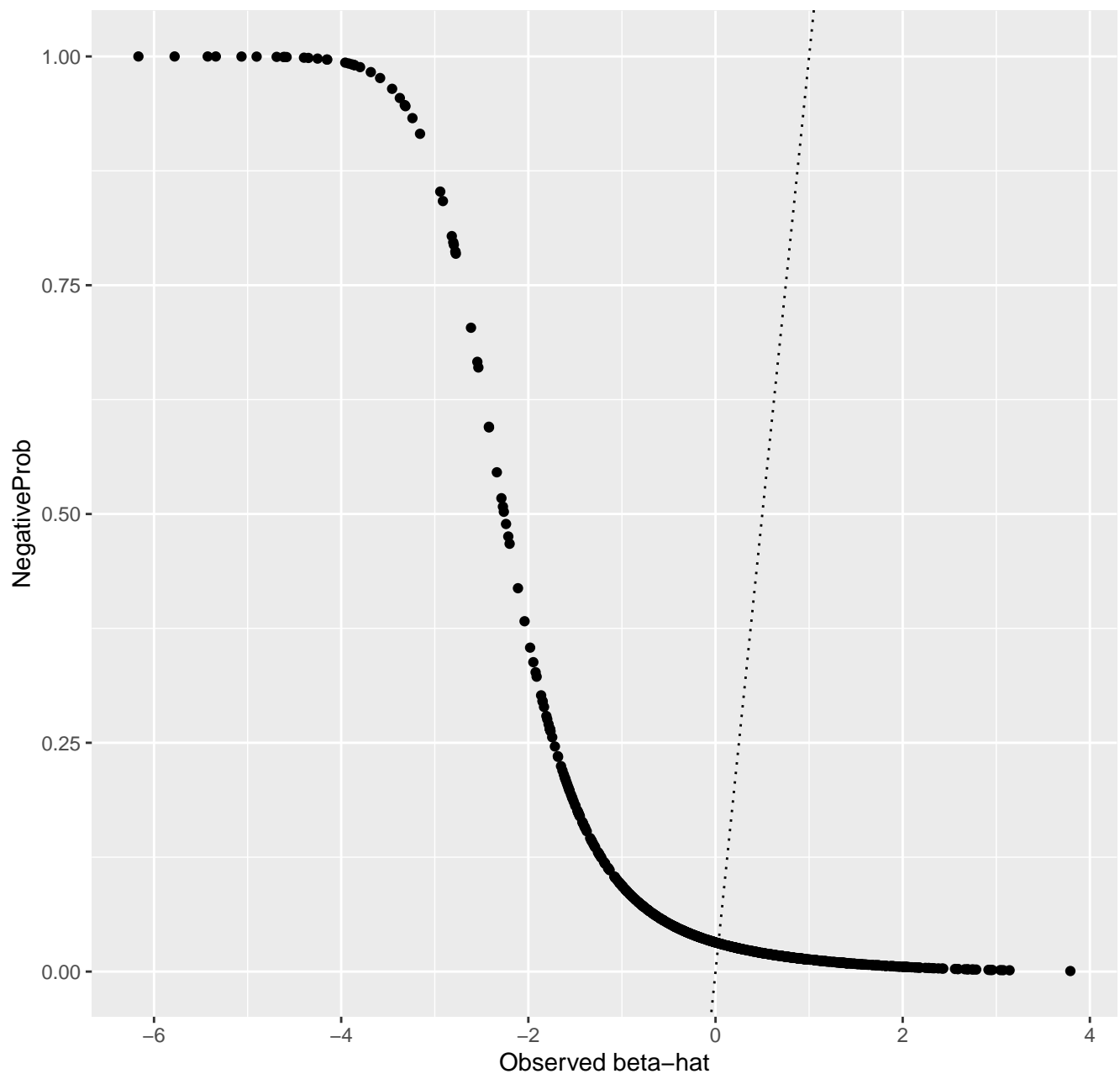
ZER1_(10444) negative



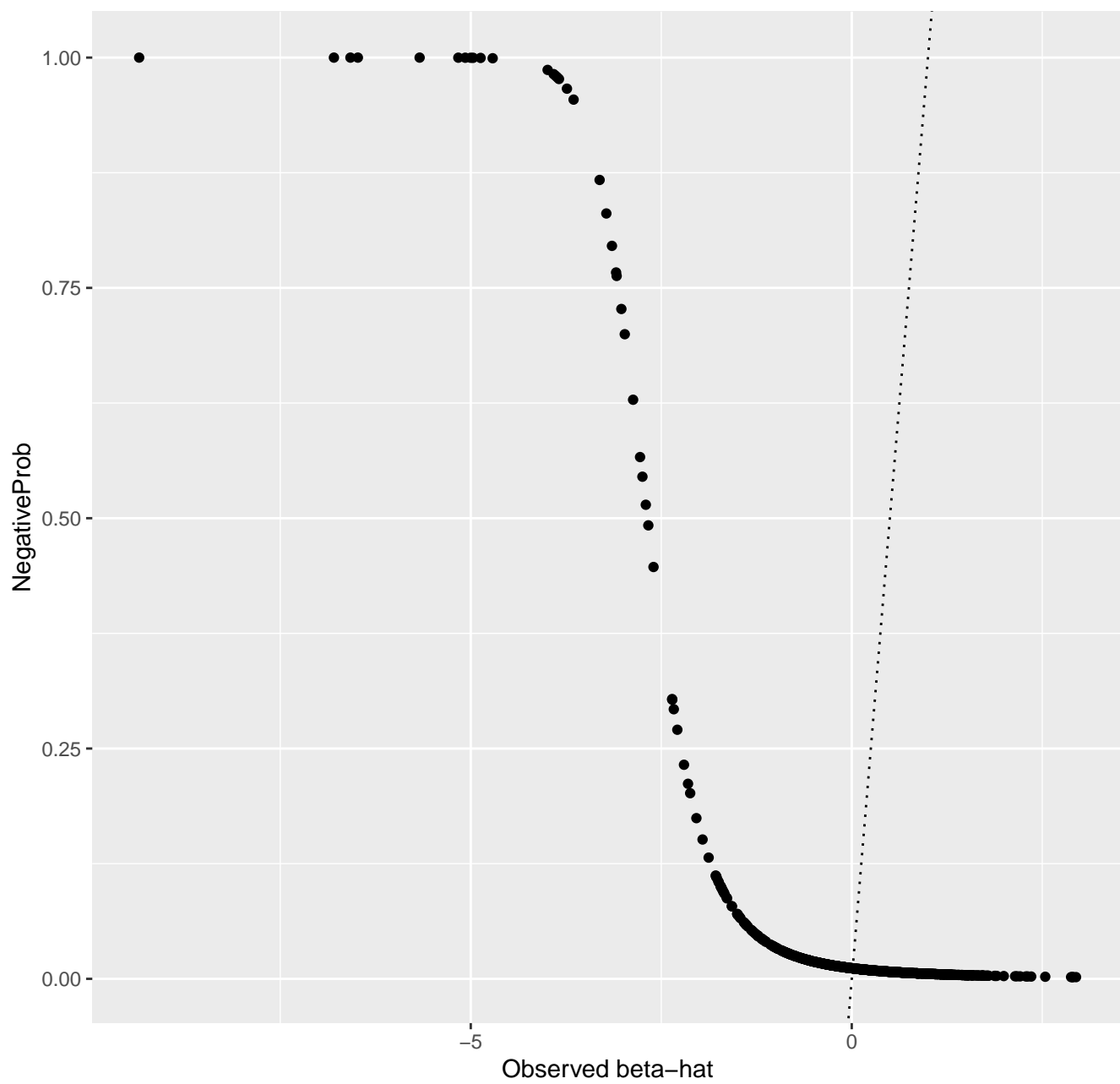
HRAS_(3265) negative



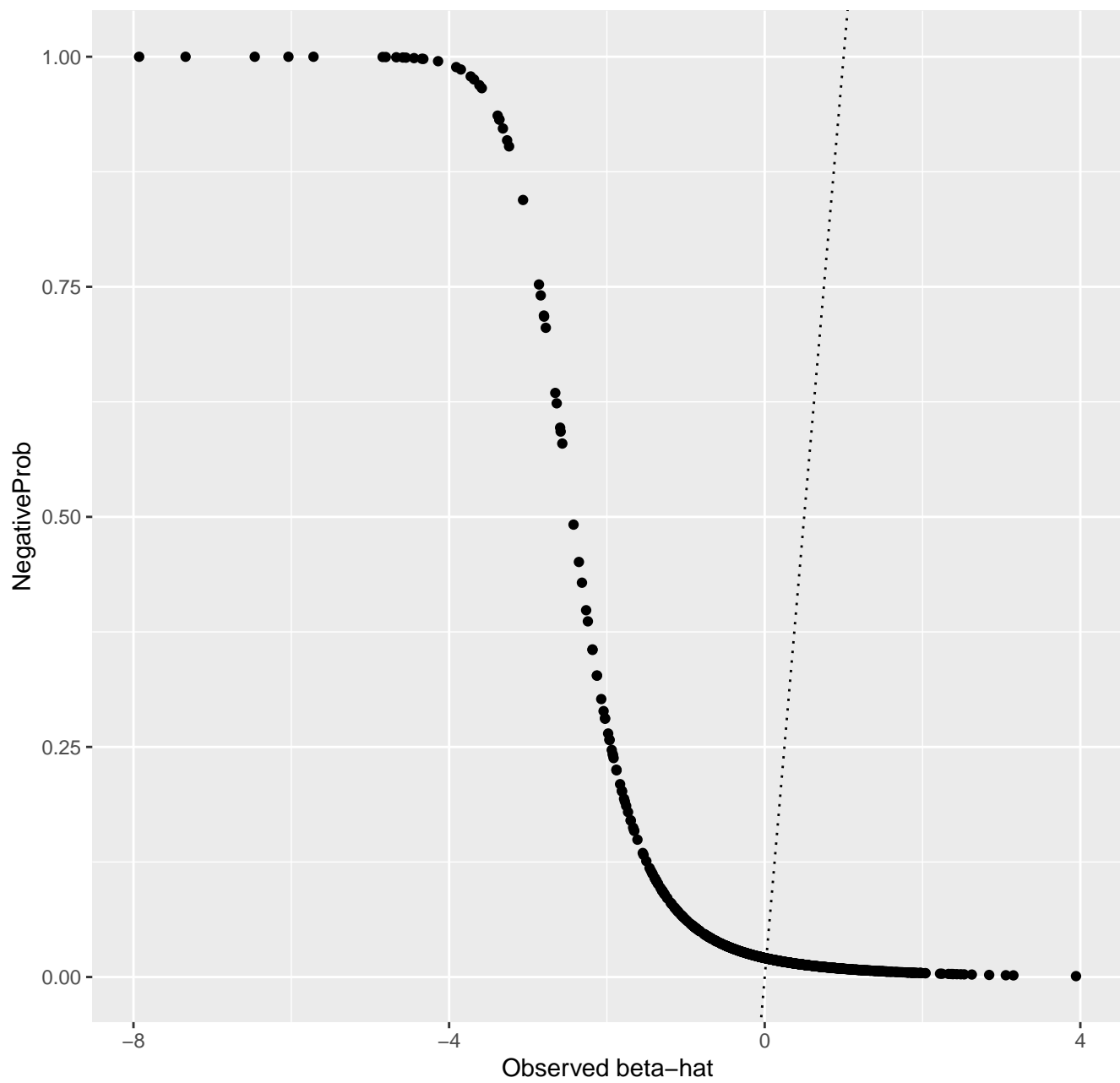
ALG12_(79087) negative



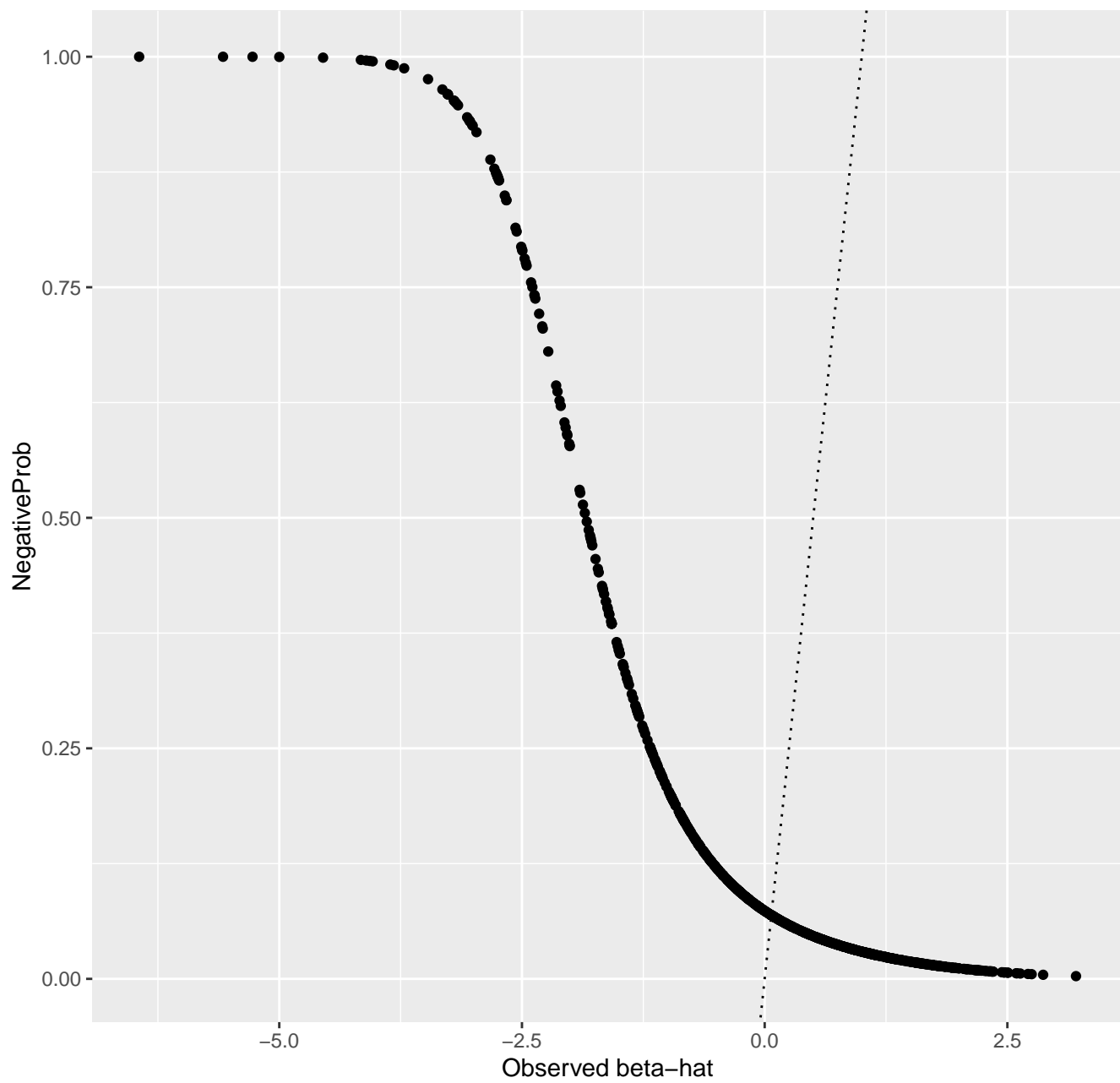
PEA15_(8682) negative



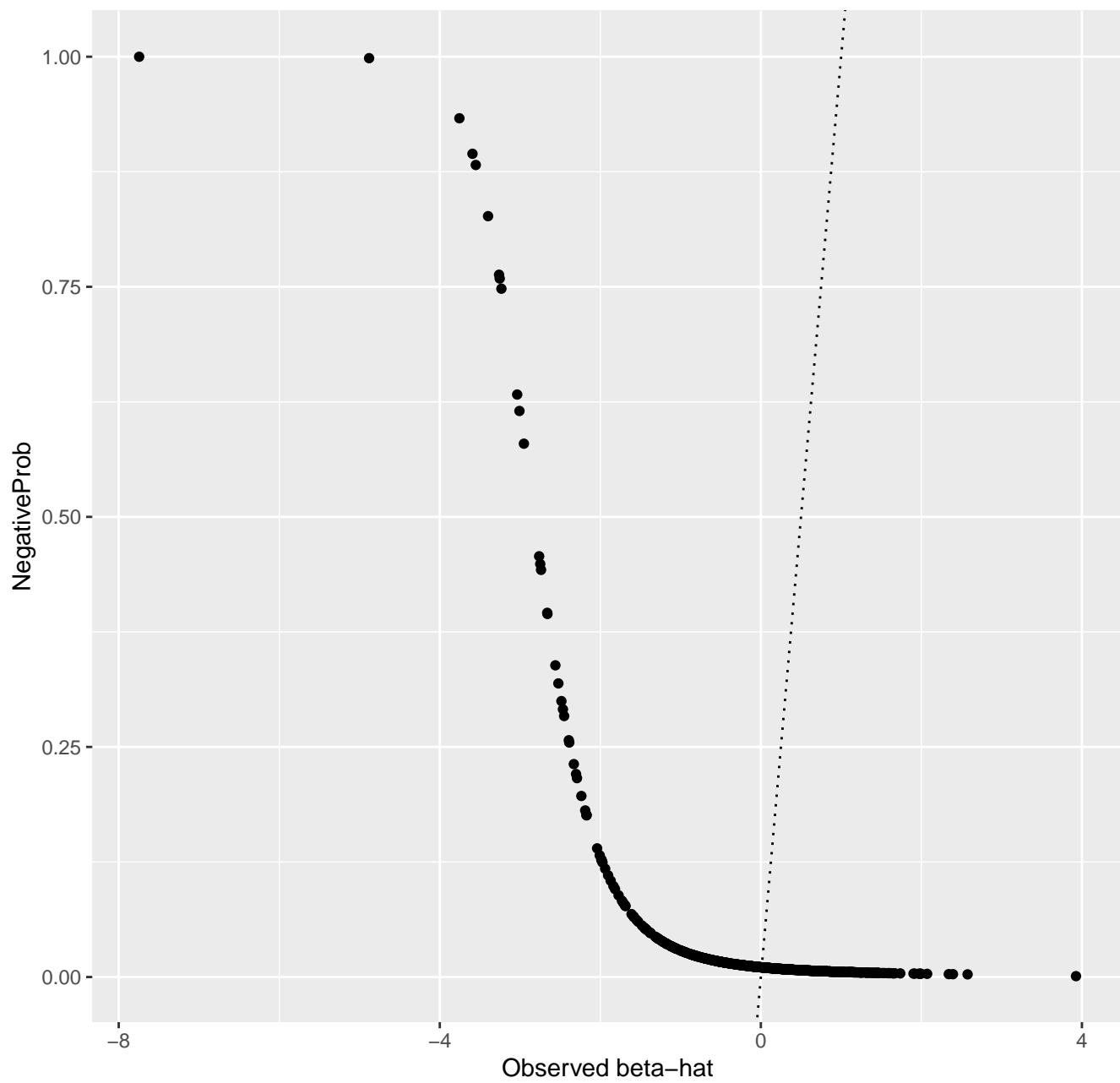
FLVCR1_(28982) negative



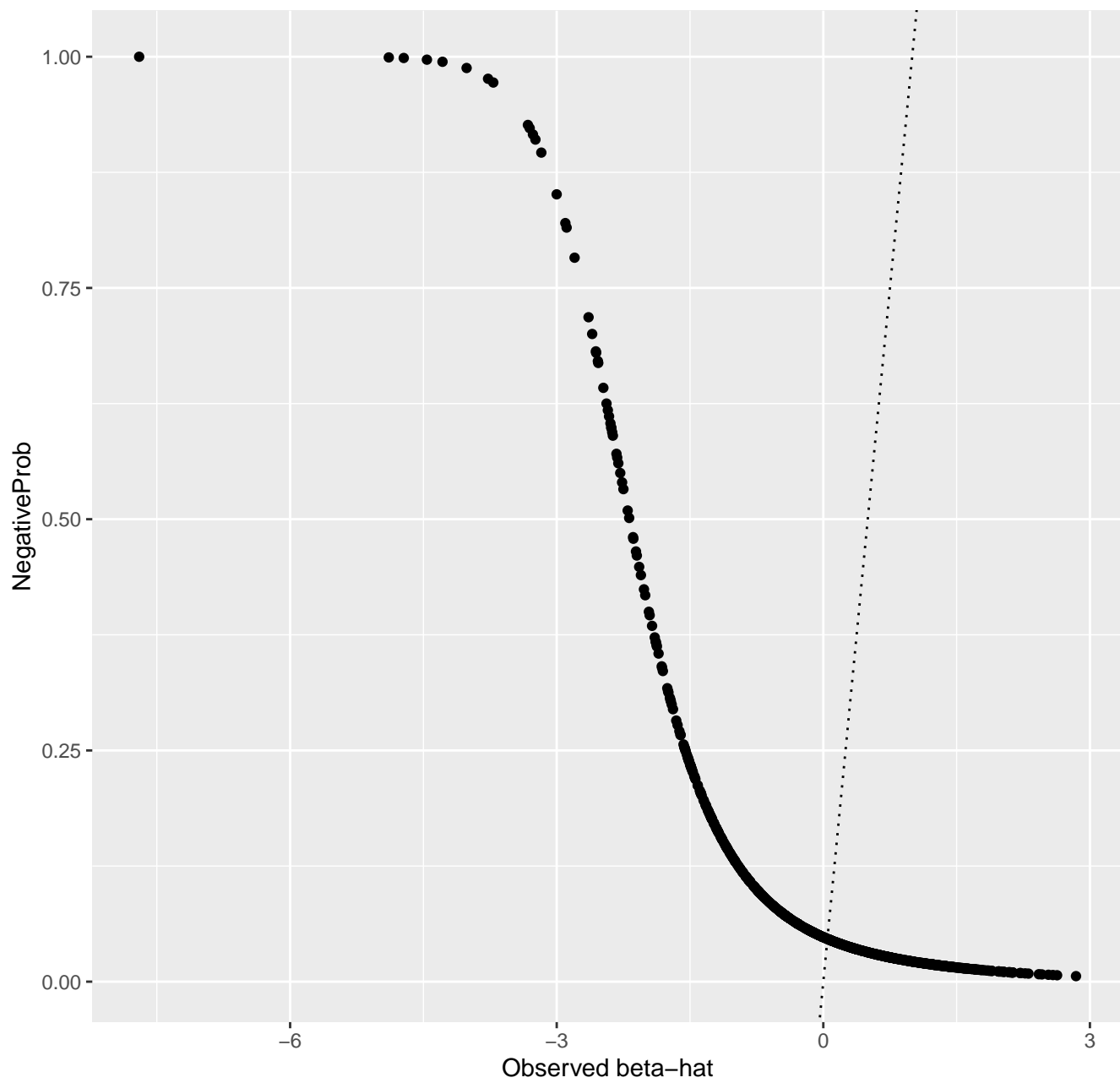
NEK1_(4750) negative



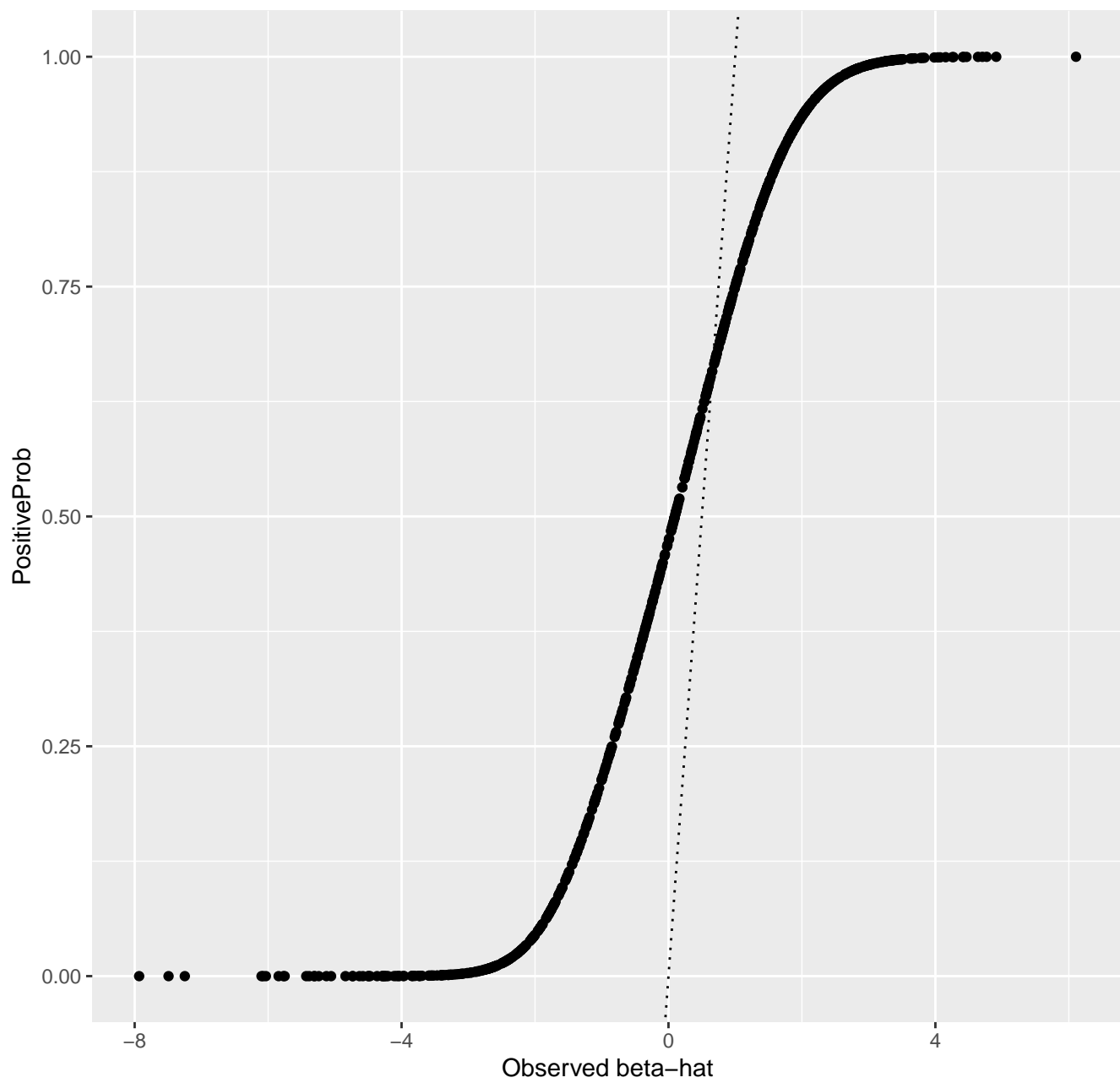
TAB2_(23118) negative



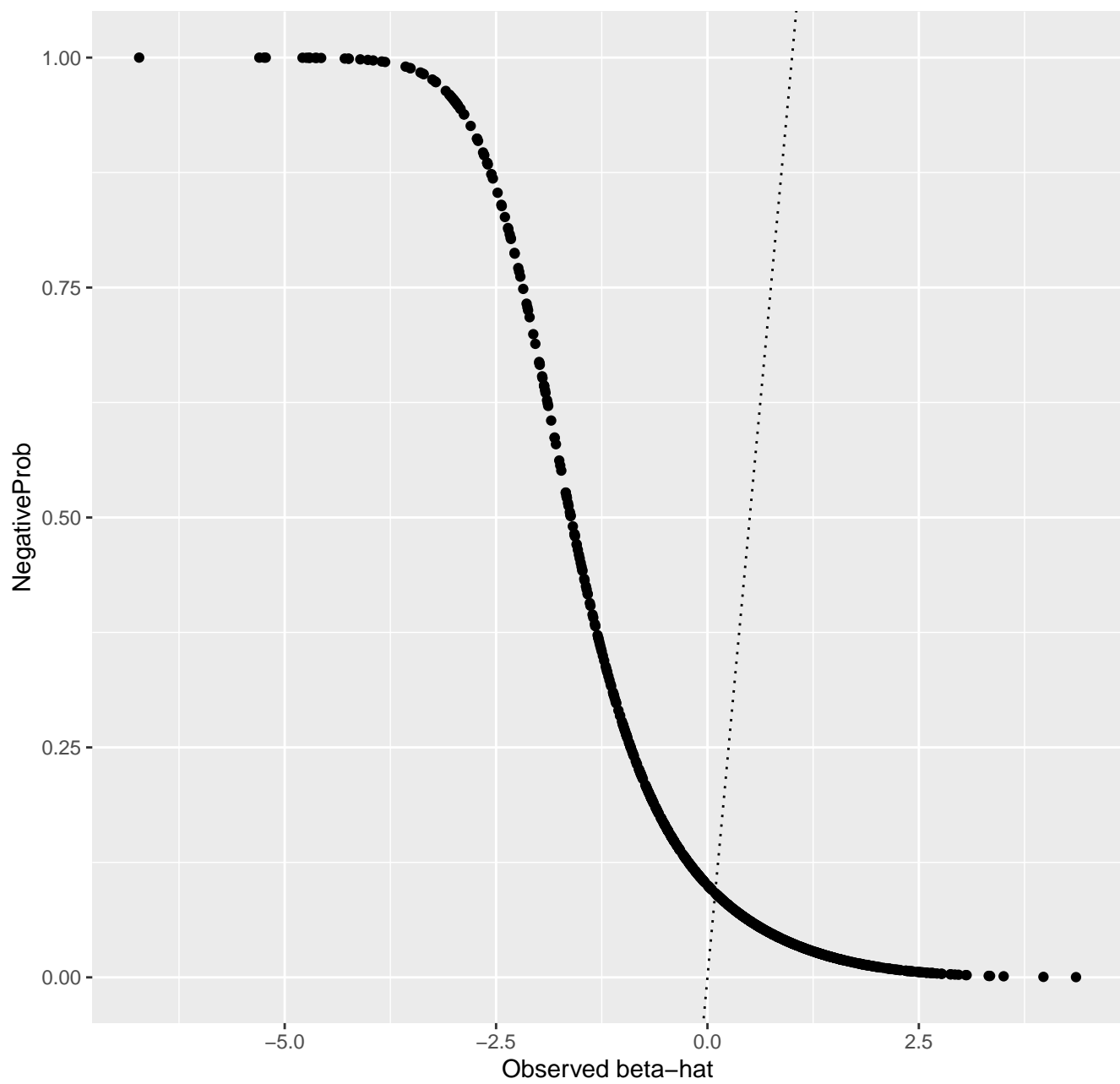
WASL_(8976) negative



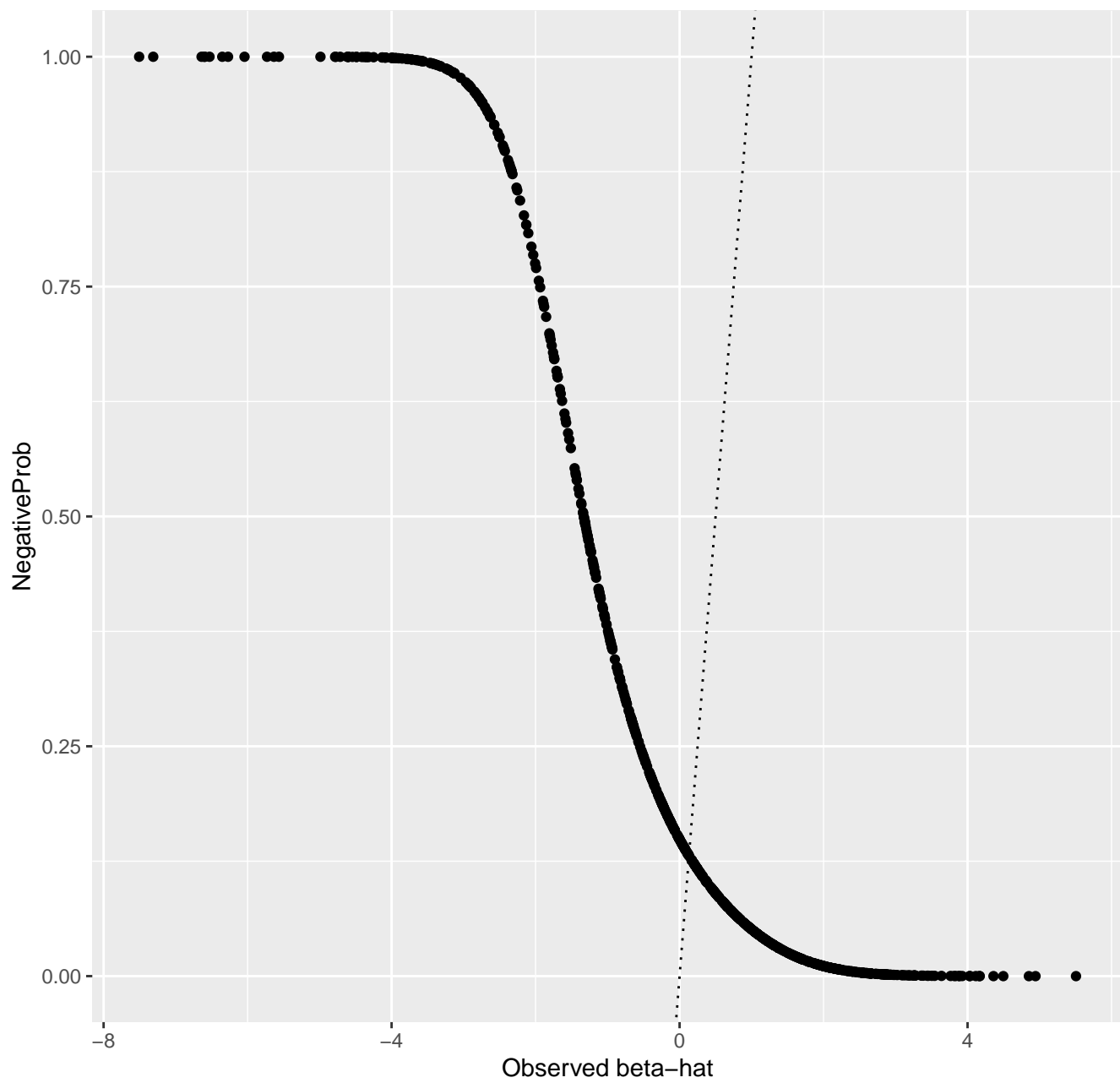
GFPT1_(2673) positive



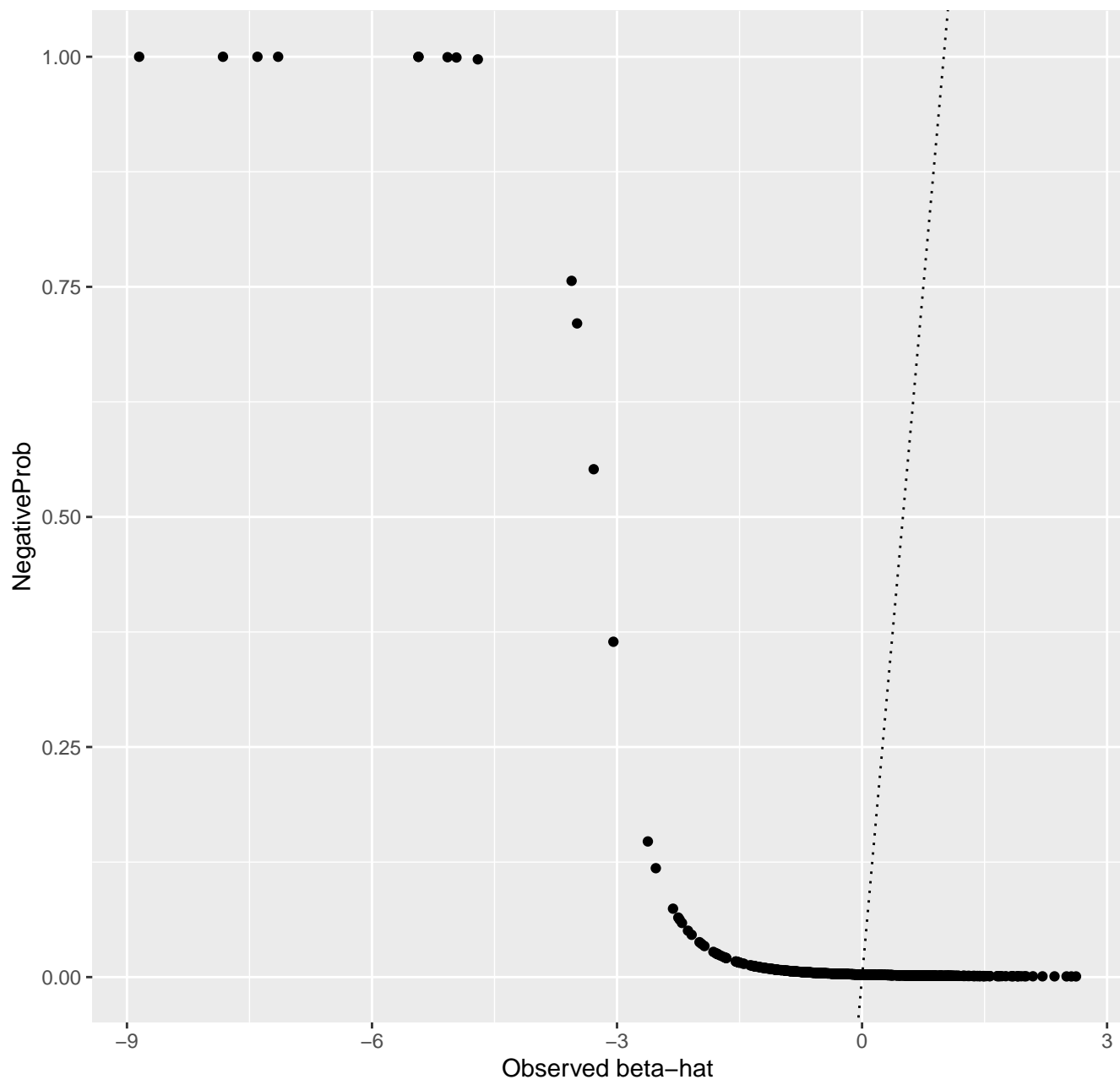
NDUFAF1_(51103) negative



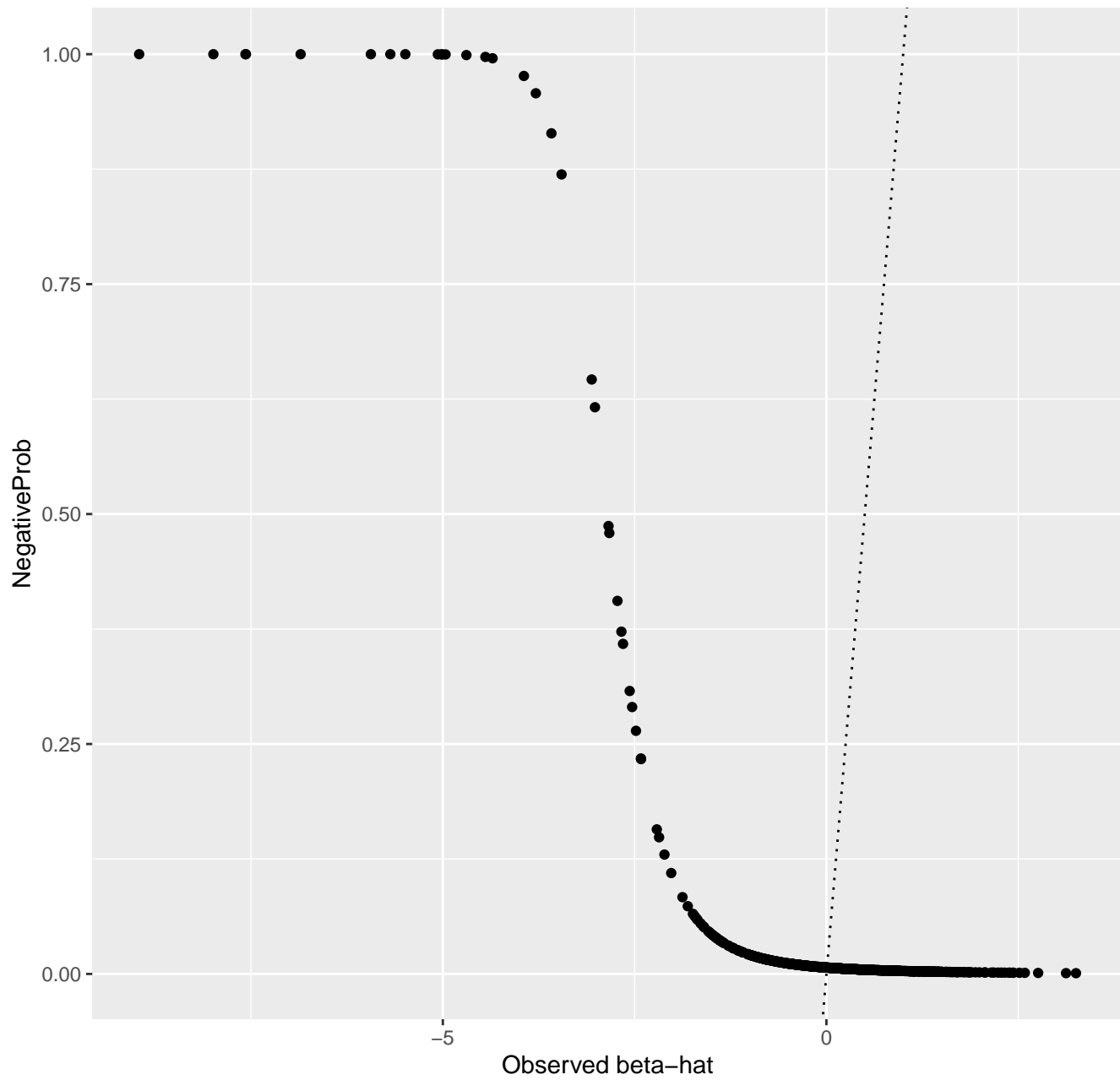
MINOS1_(440574) negative



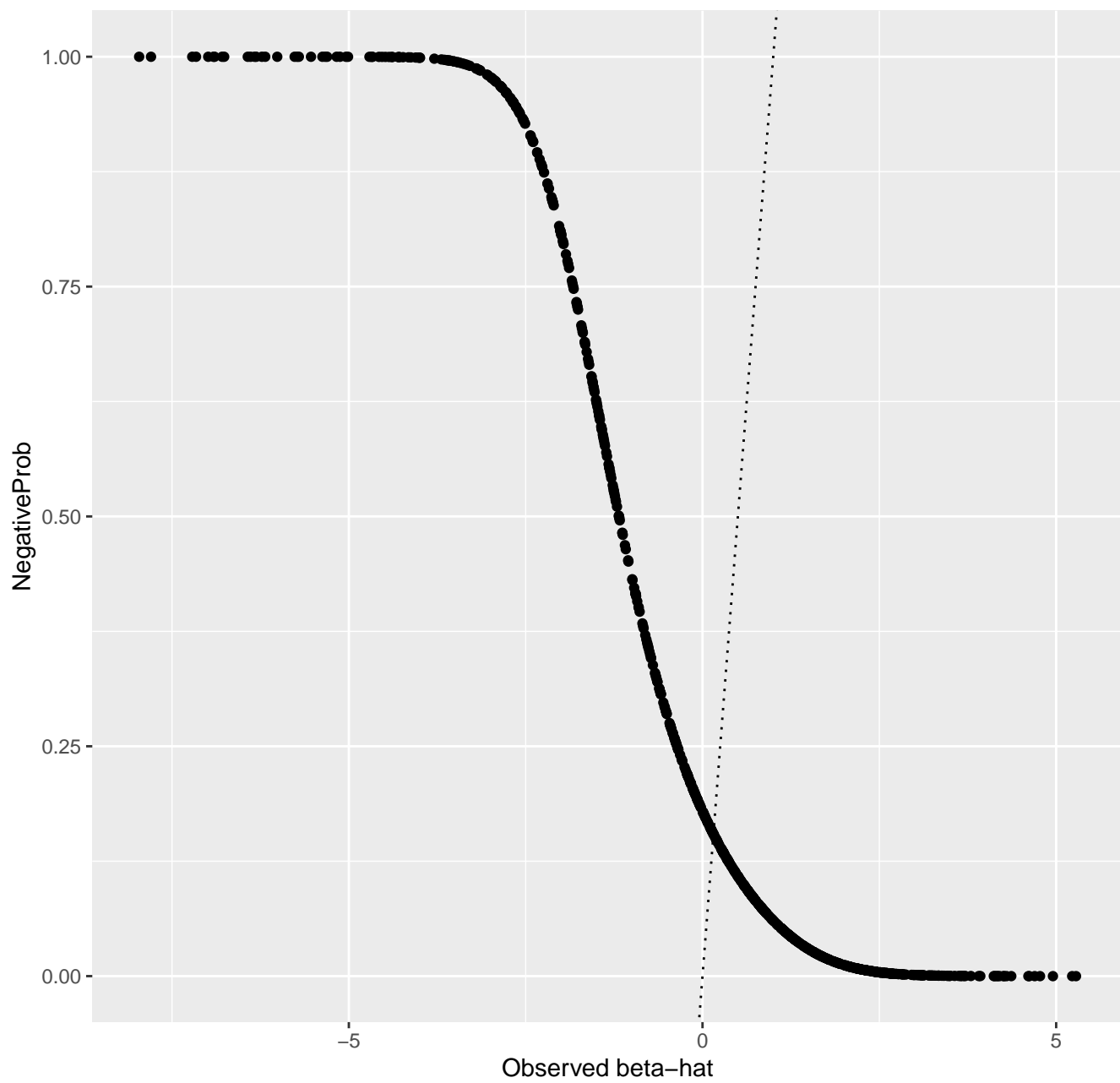
ITPA_(3704) negative



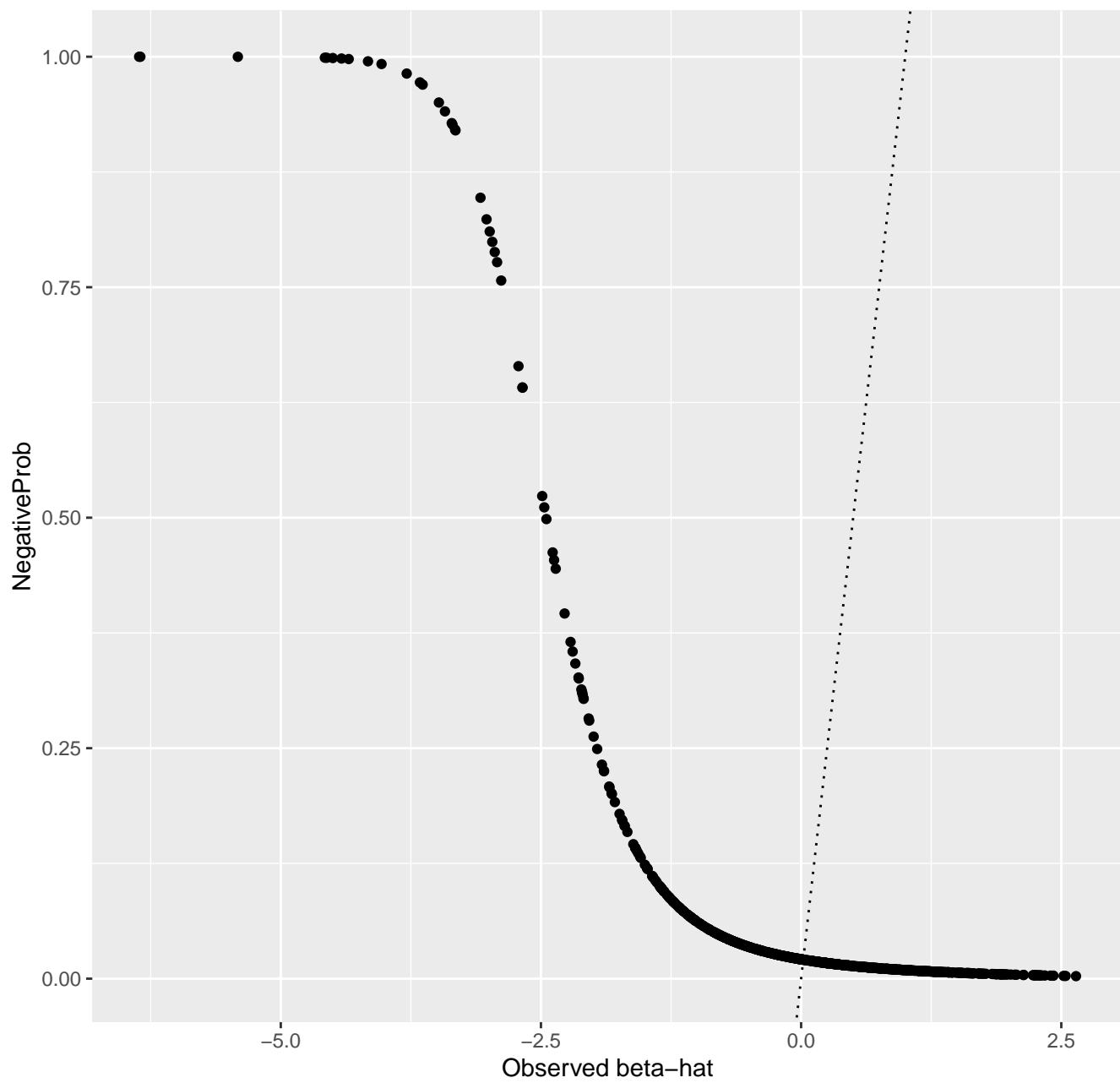
TCF3_(6929) negative



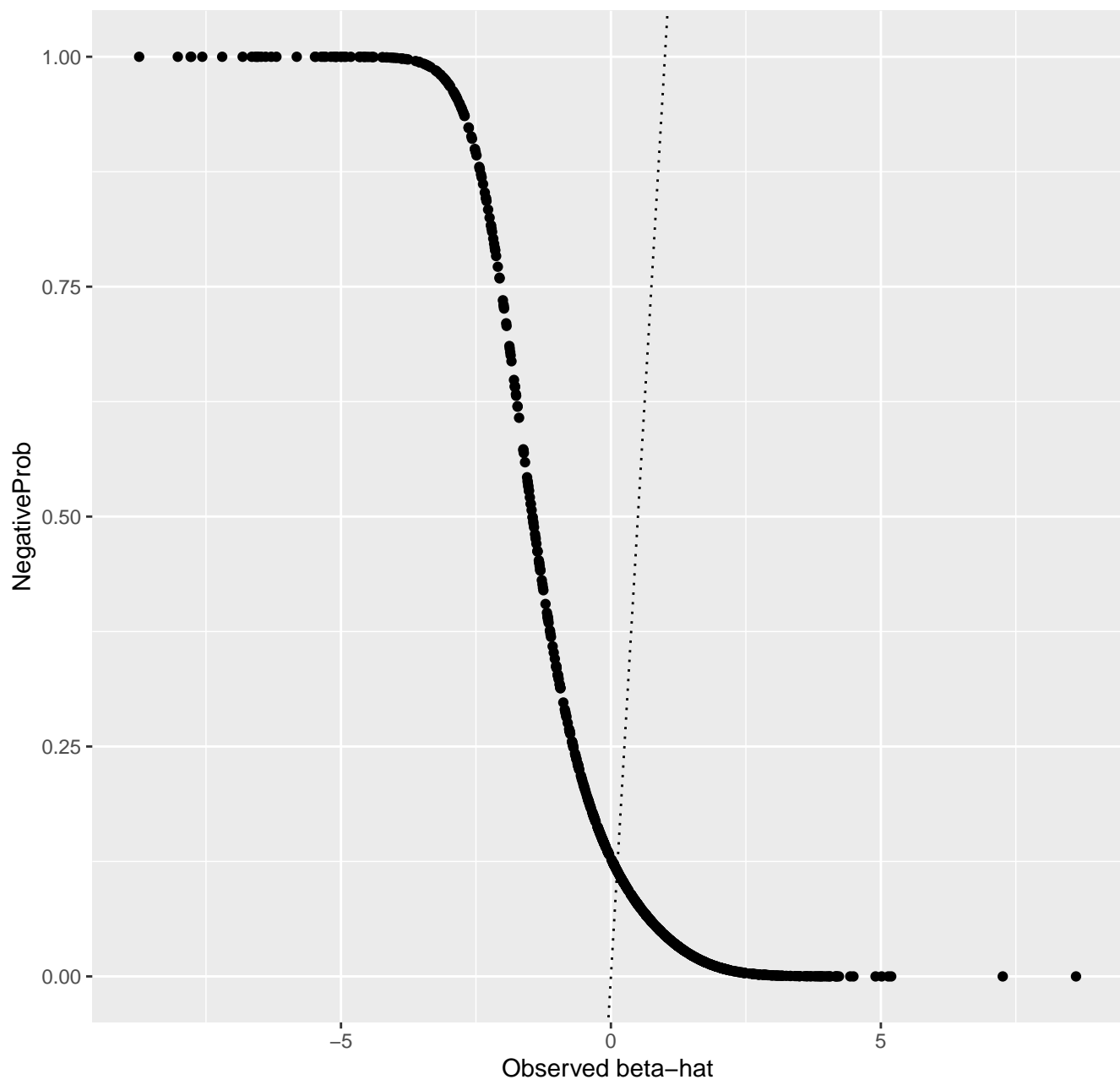
AHCYL1_(10768) negative



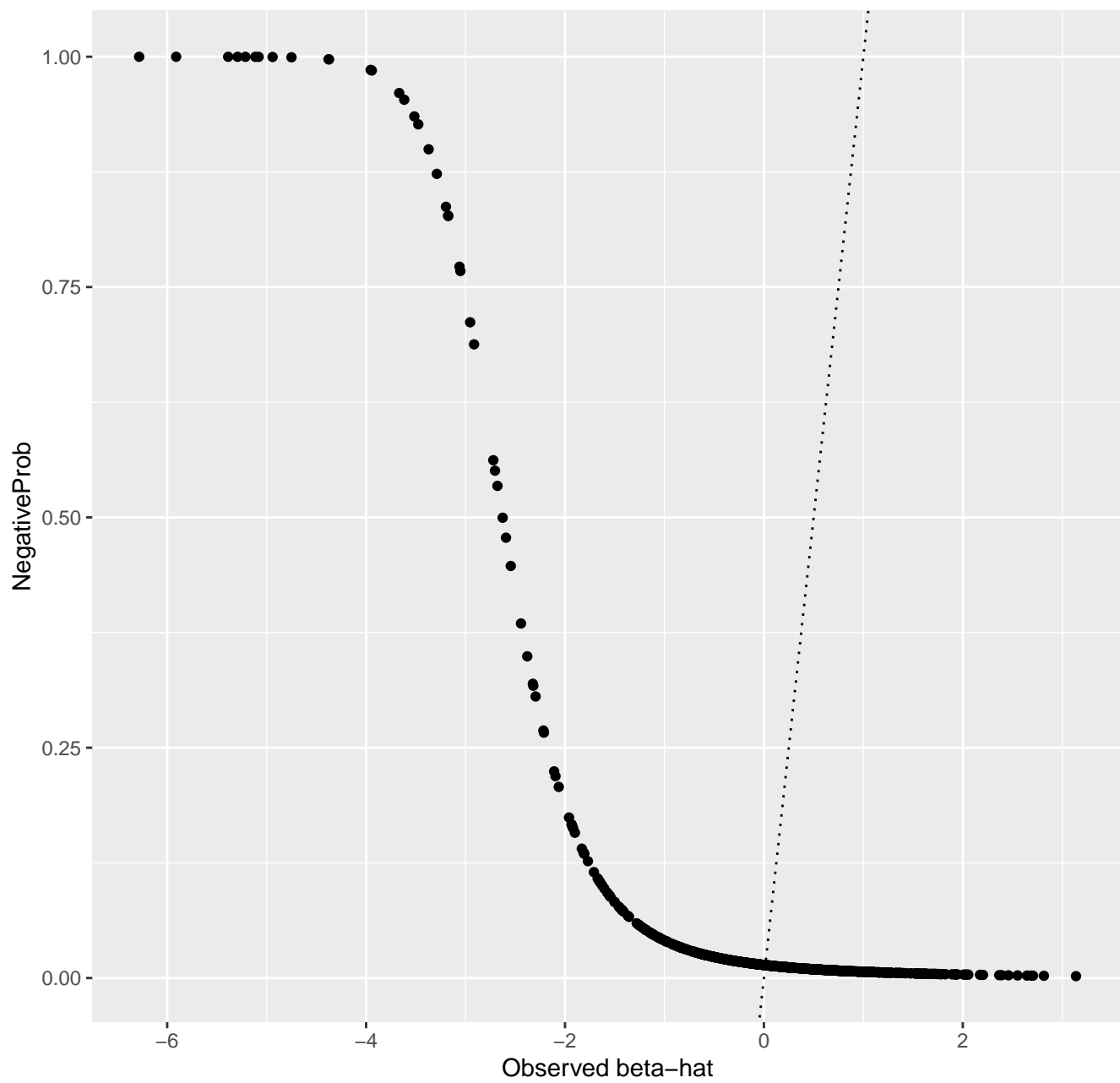
IKBKB_(3551) negative



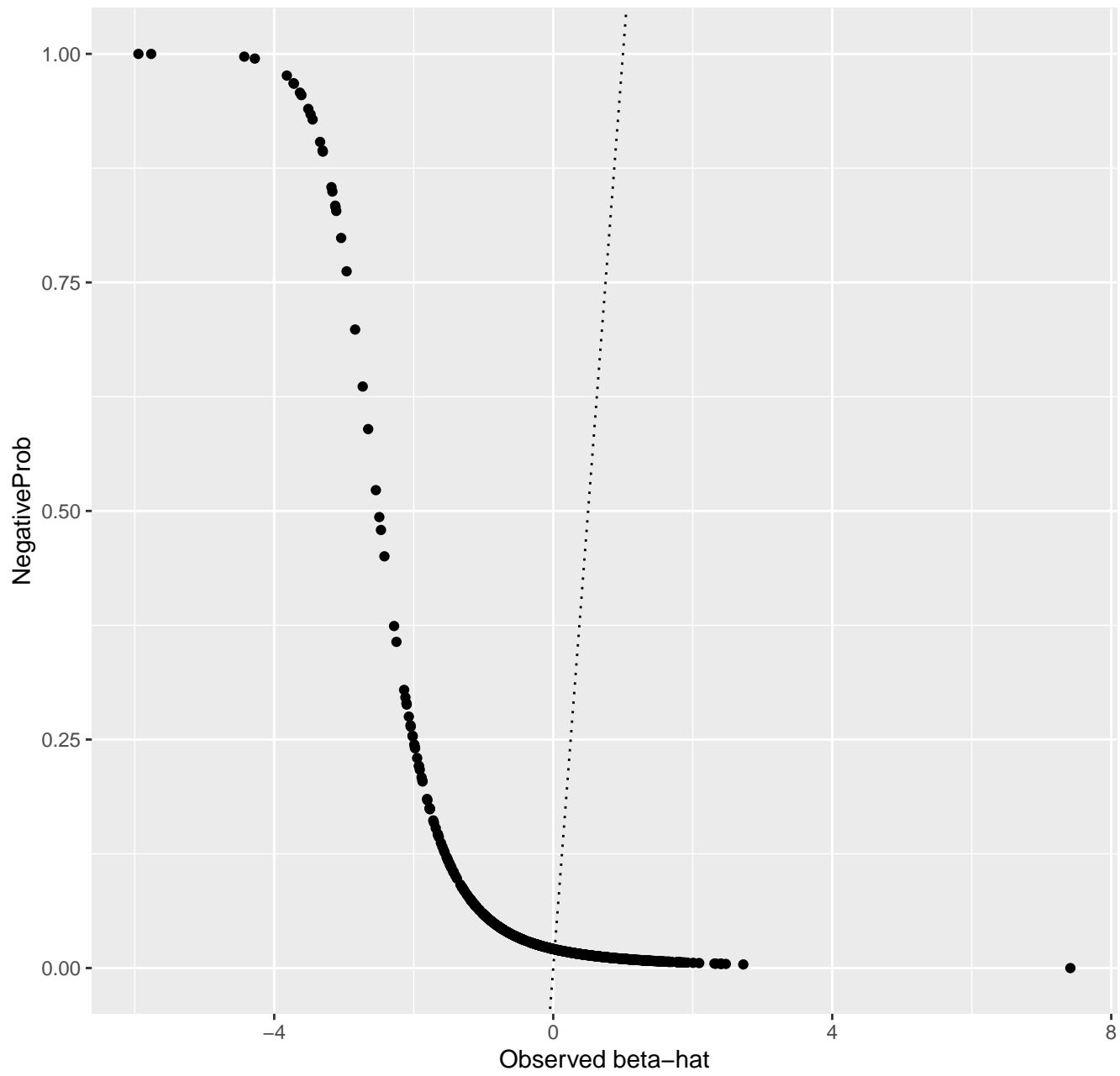
ITGB1_(3688) negative



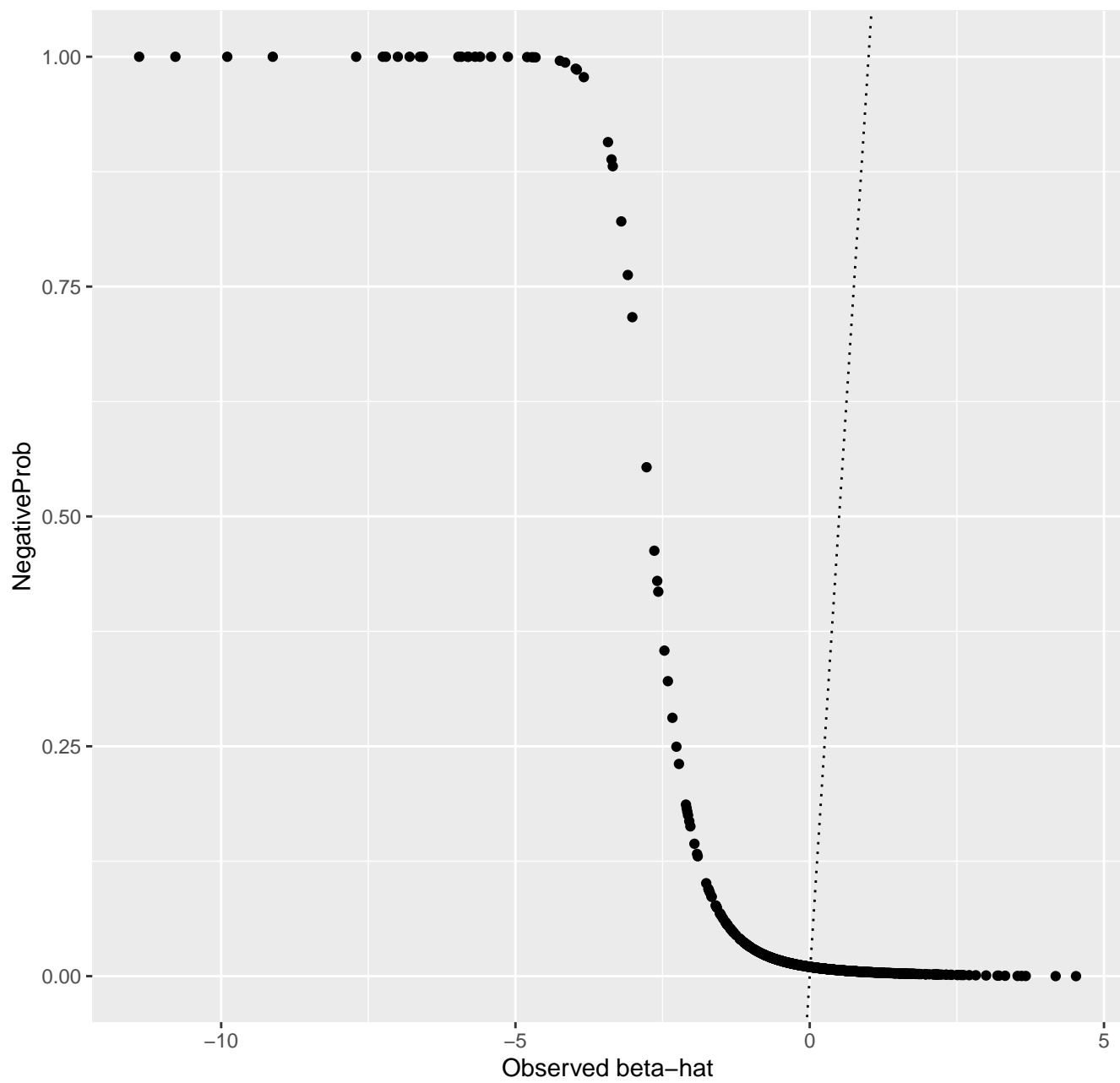
DDC_(1644) negative



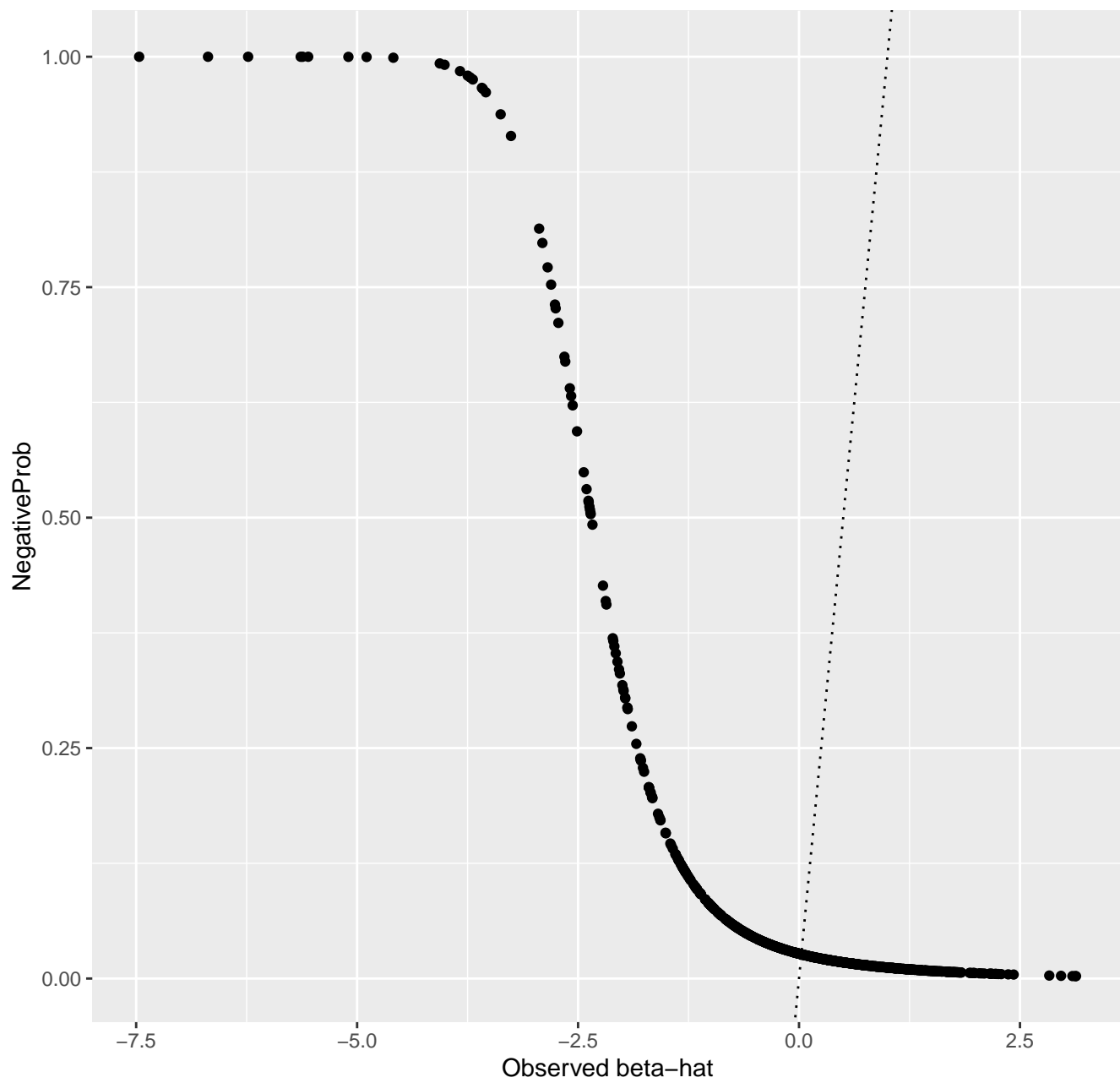
PPTC7_(160760) negative



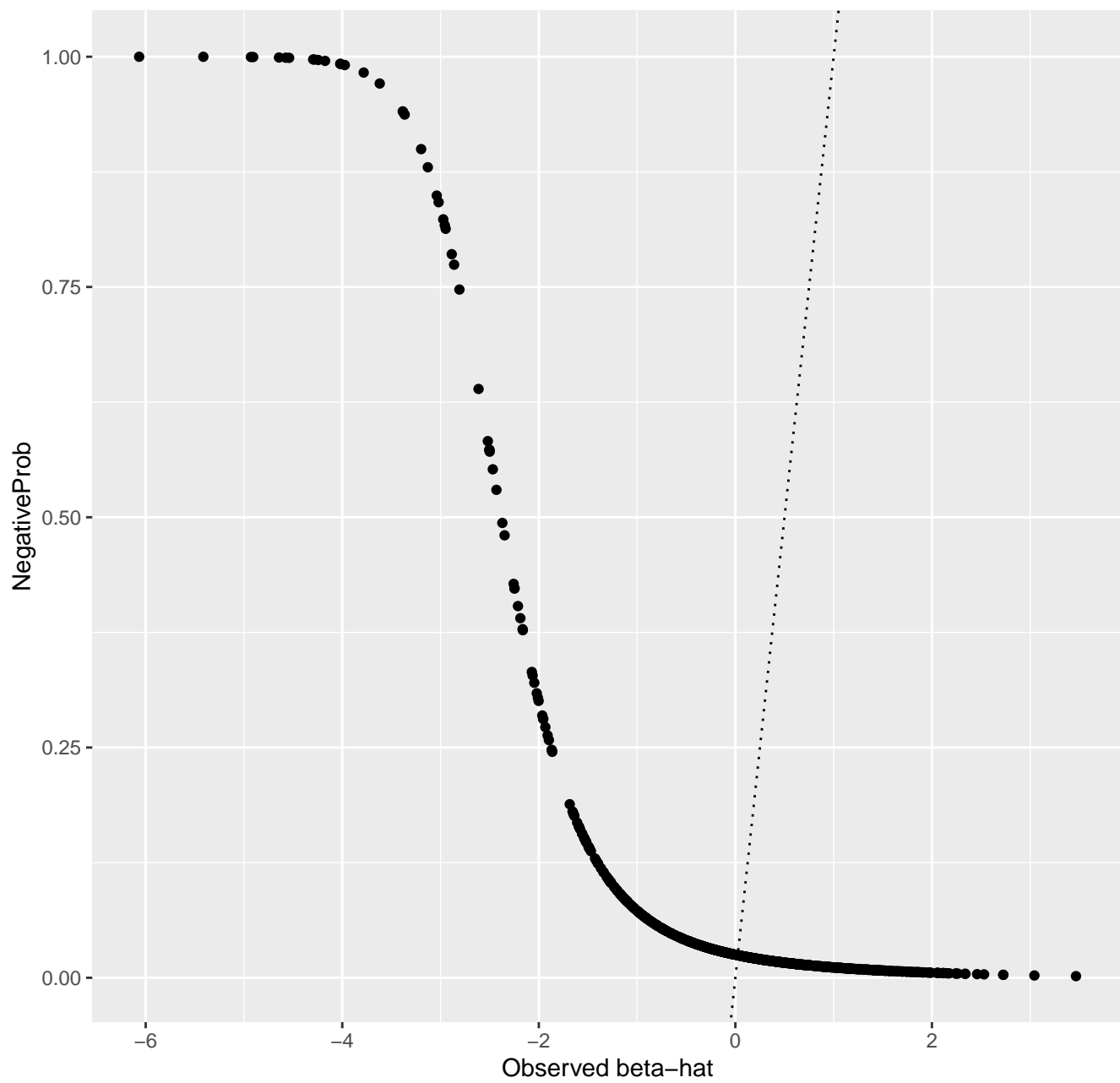
FOXA1_(3169) negative



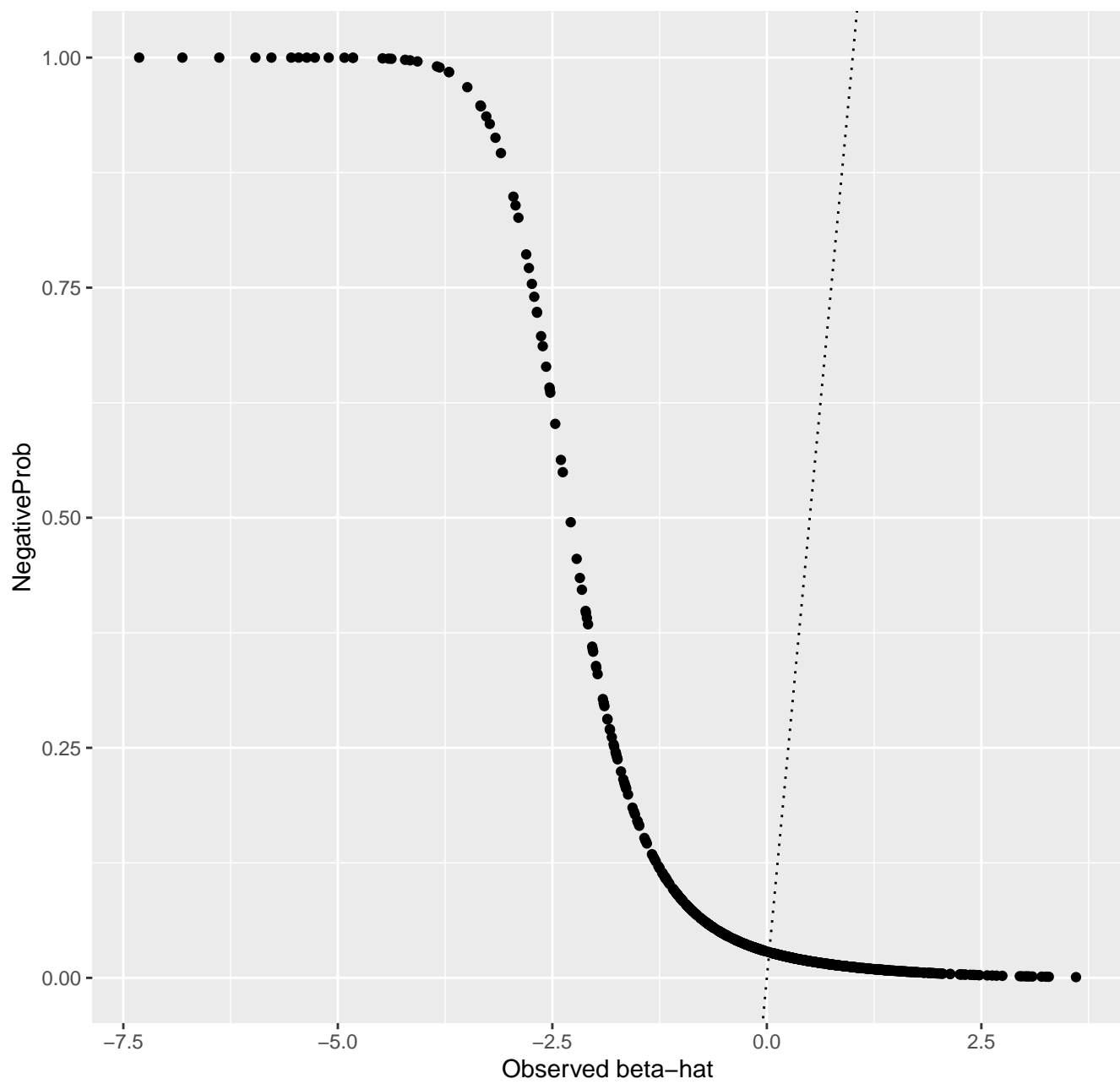
VDAC2_(7417) negative



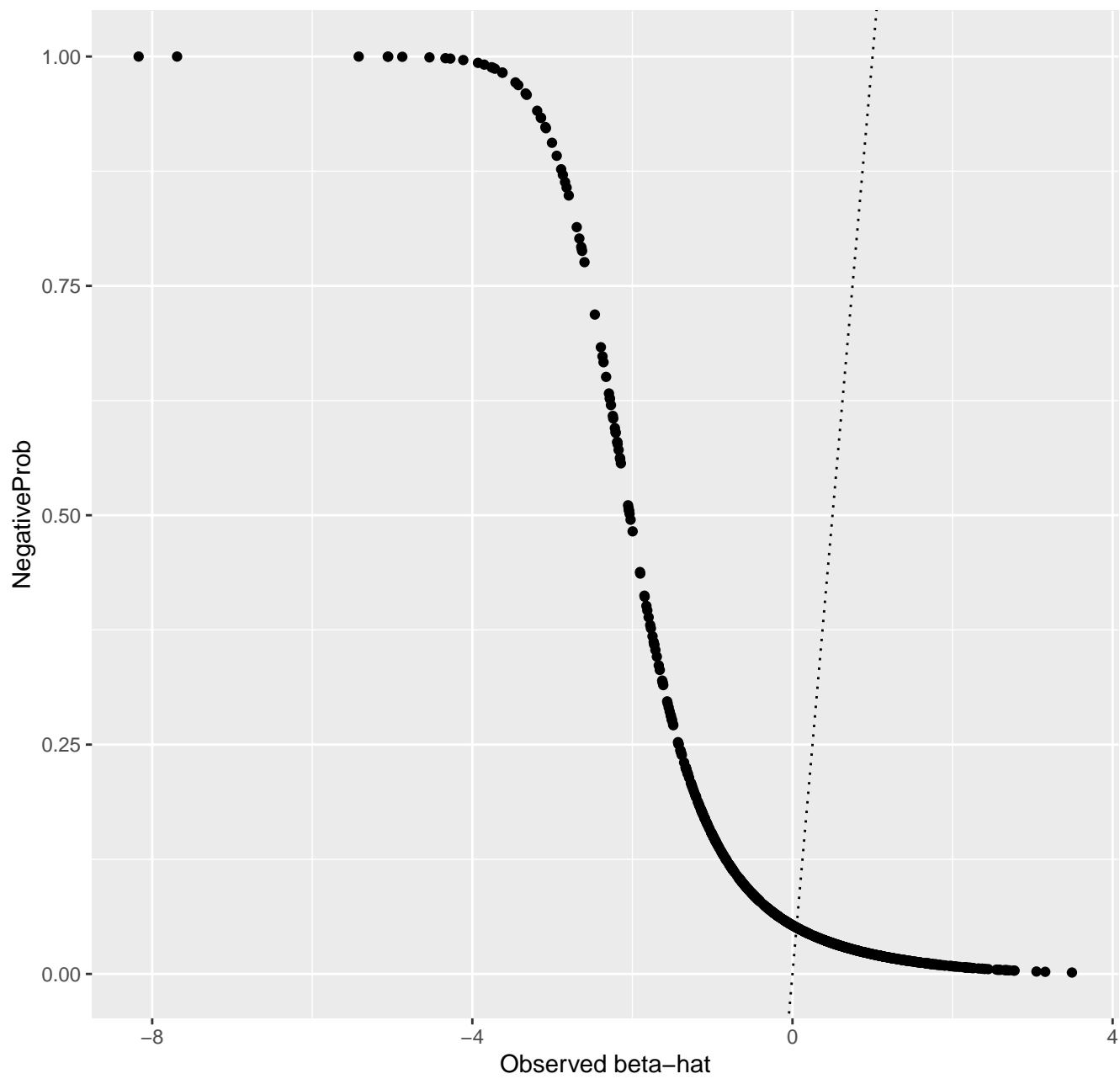
SEC31A_(22872) negative



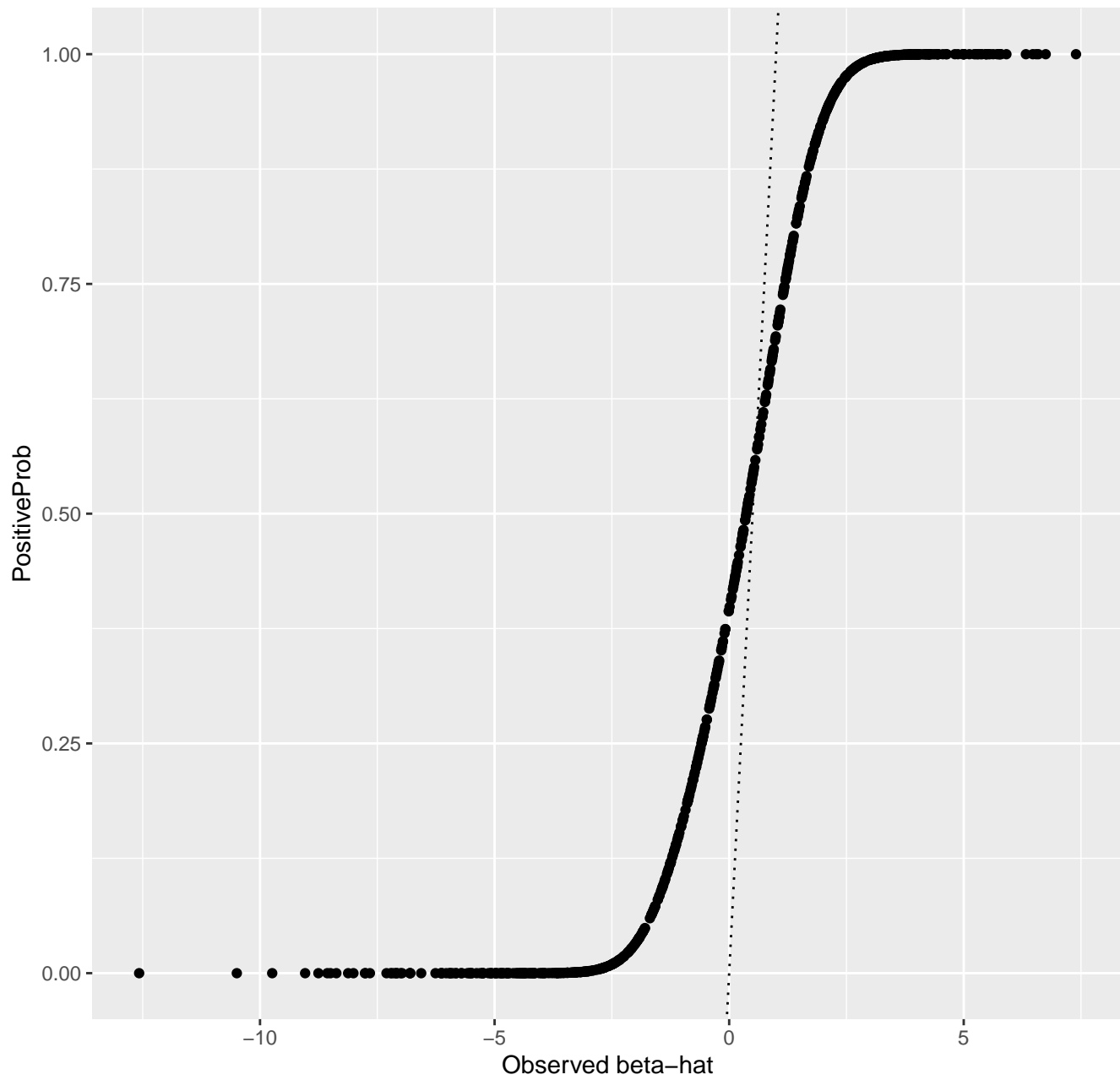
ARID1B_(57492) negative



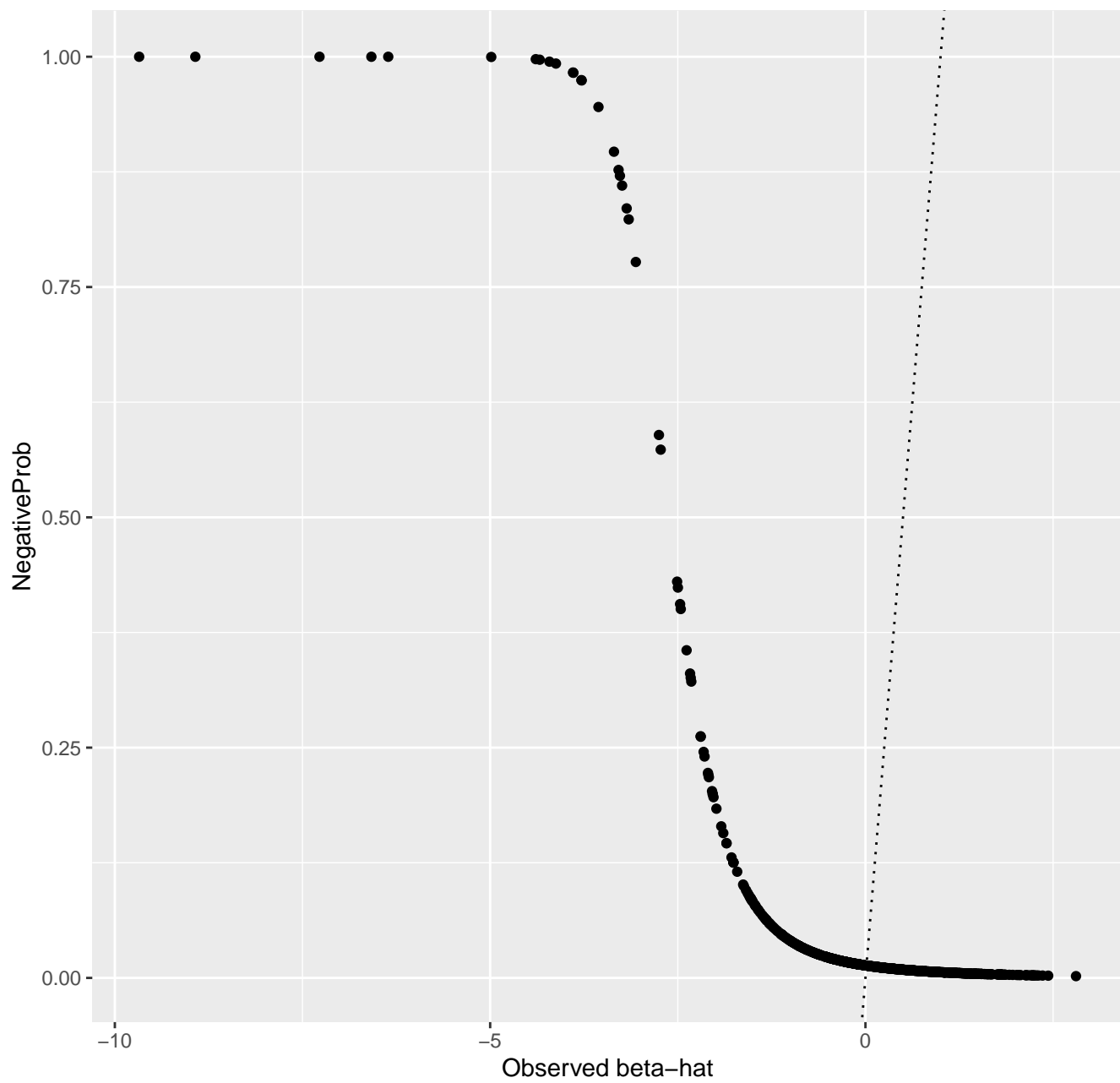
FANCG_(2189) negative



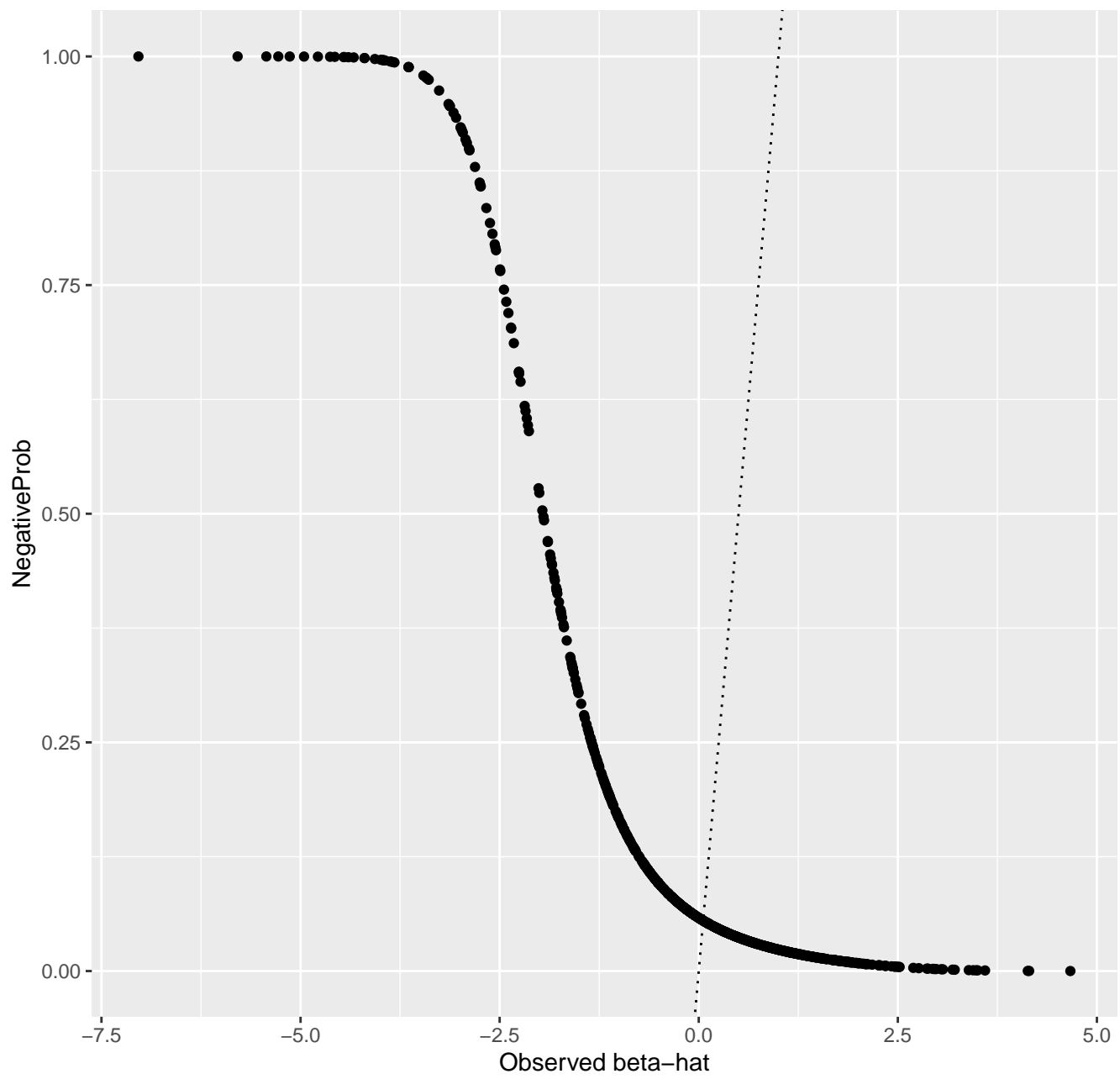
BCL2L1_(598) positive



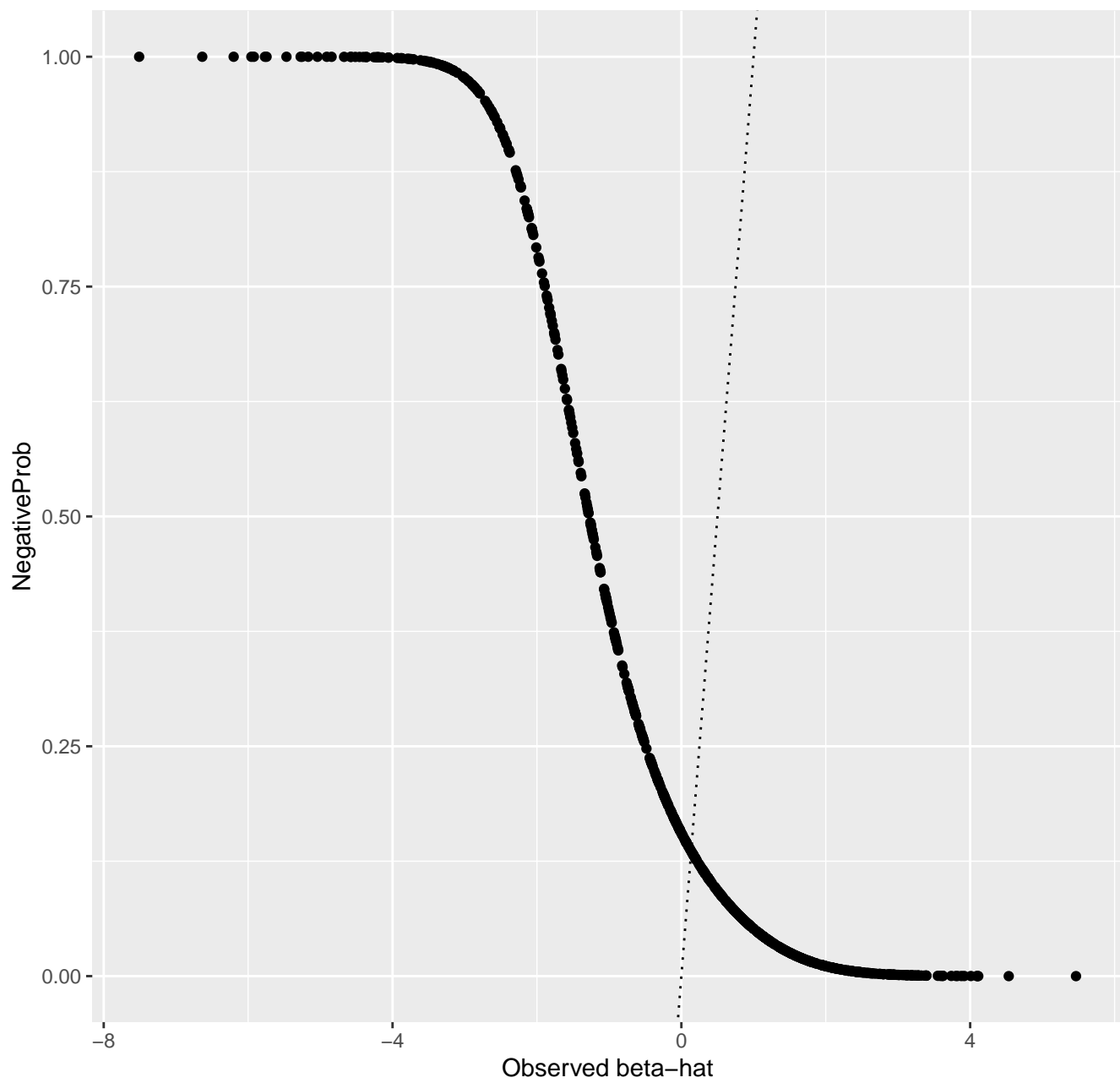
CSTF2_(1478) negative



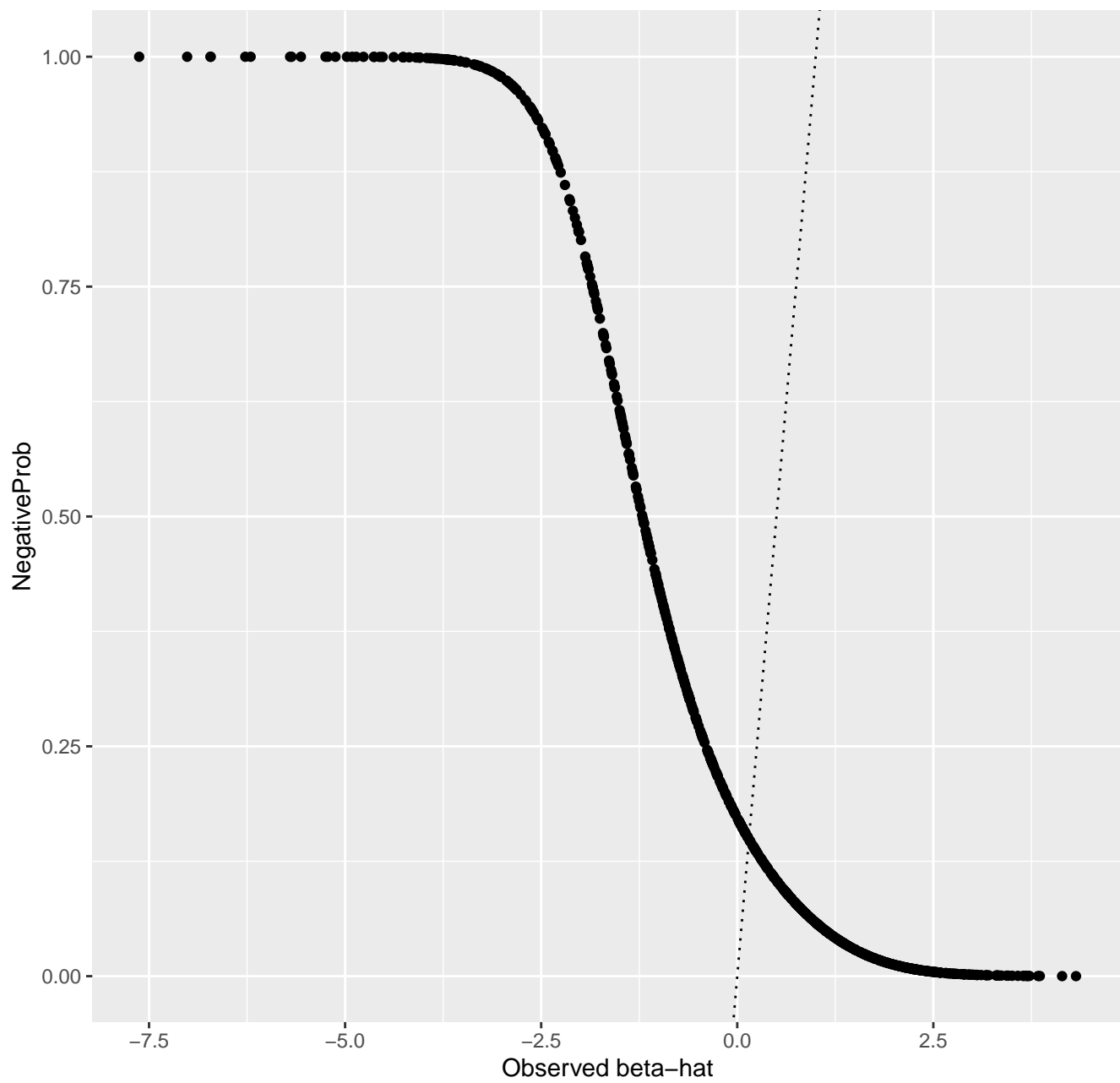
UGCG_(7357) negative



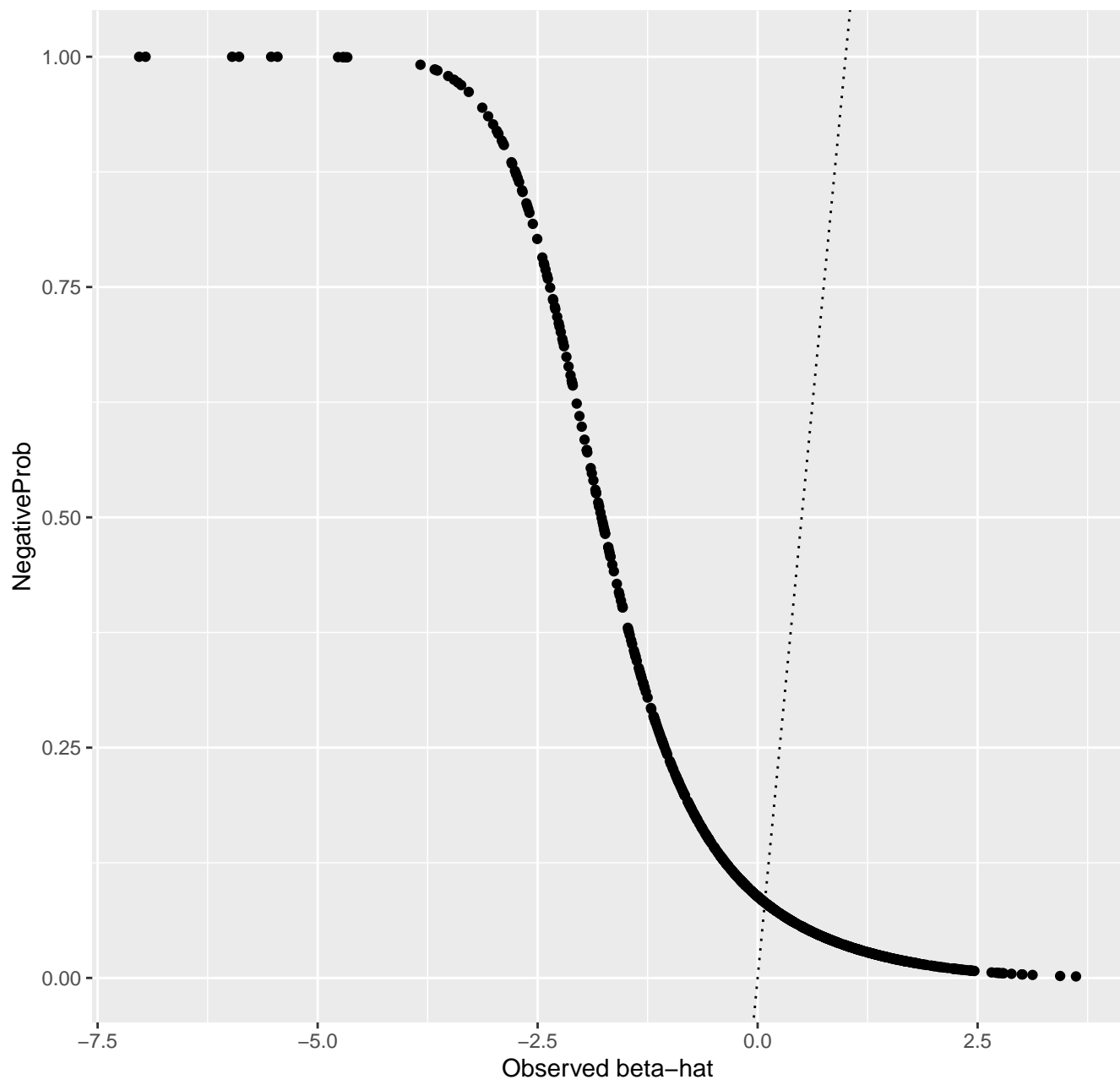
RFK_(55312) negative



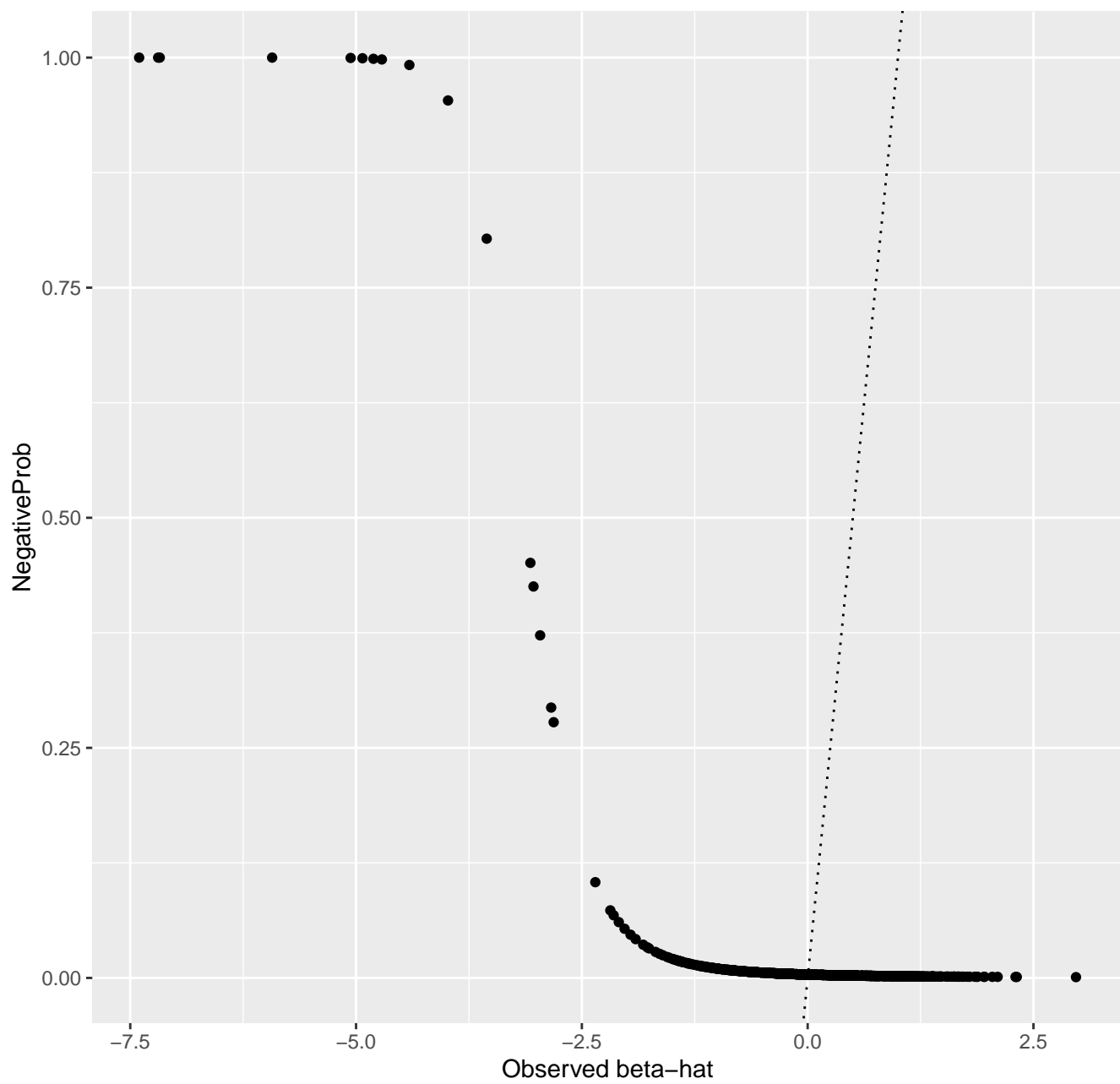
POP7_(10248) negative



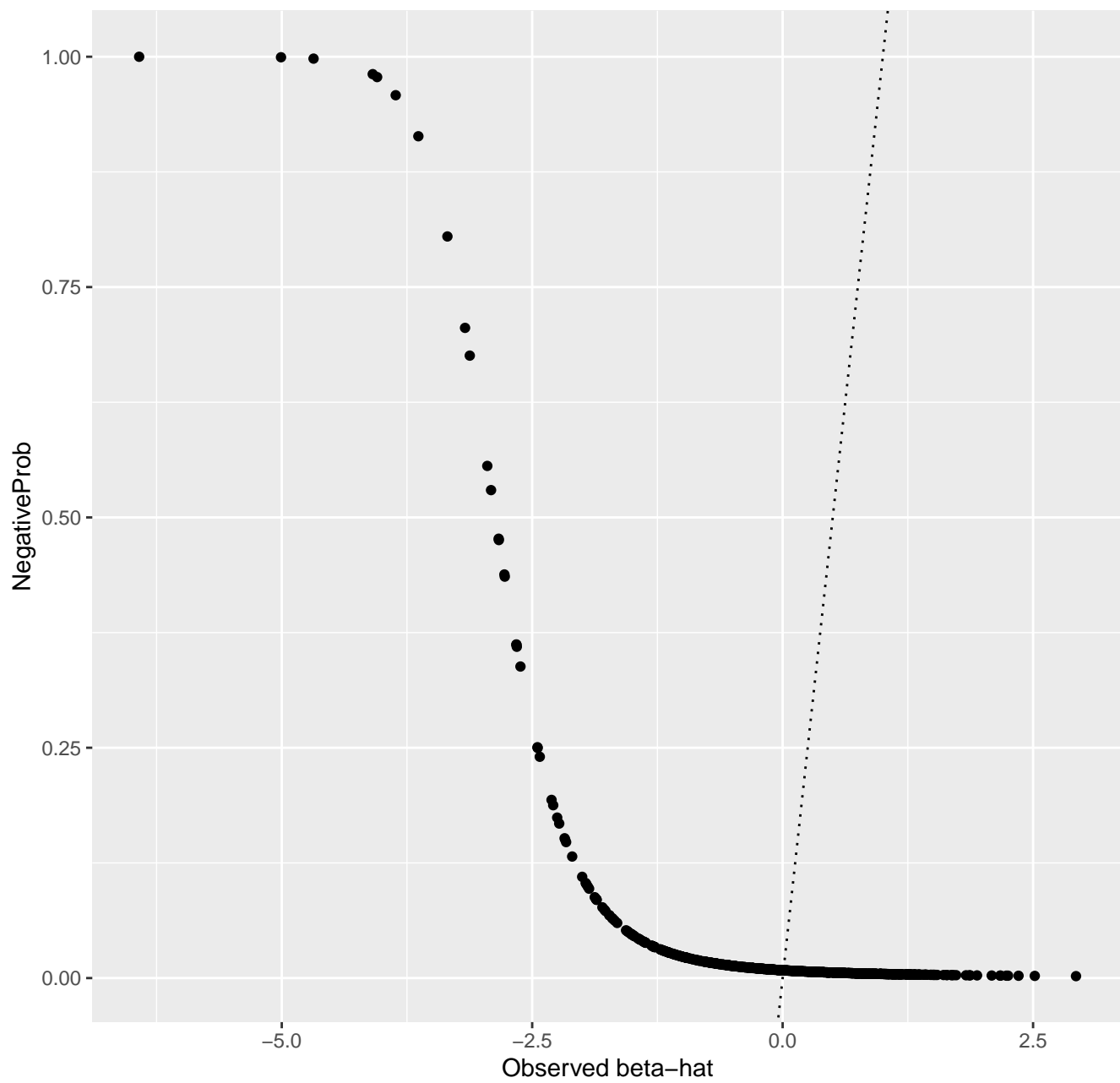
RNH1_(6050) negative



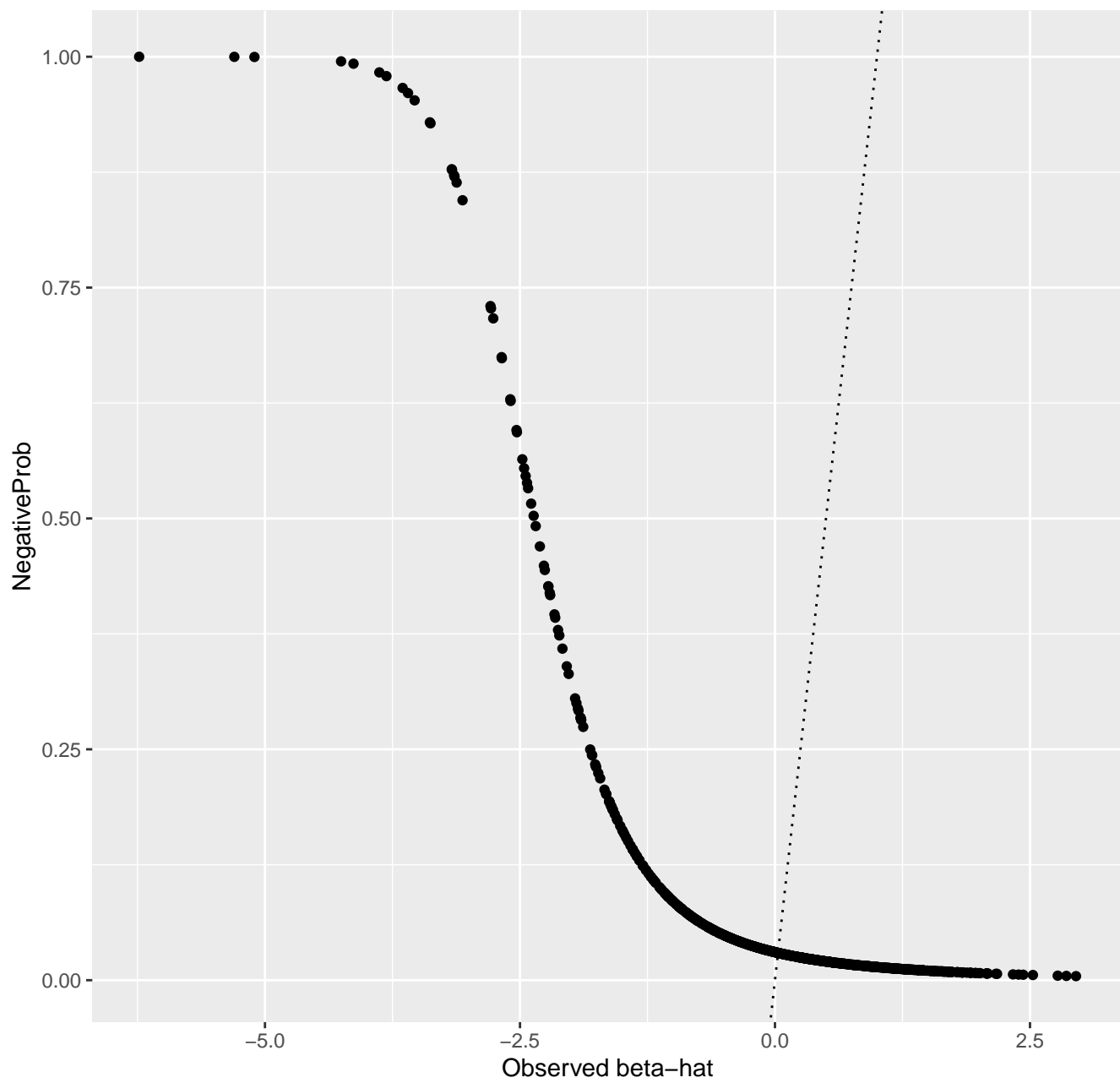
FGFR2_(2263) negative



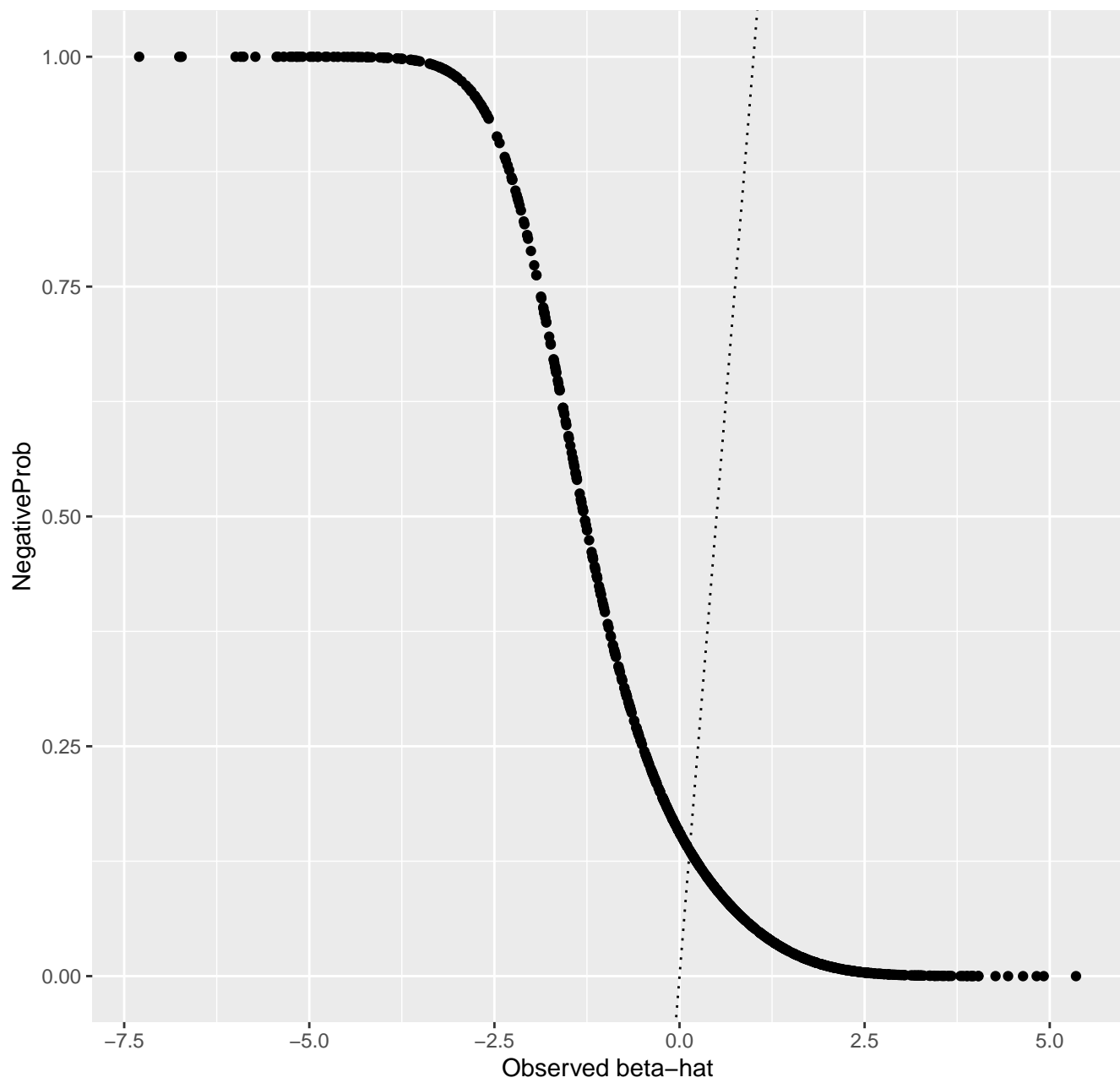
WDR89_(112840) negative



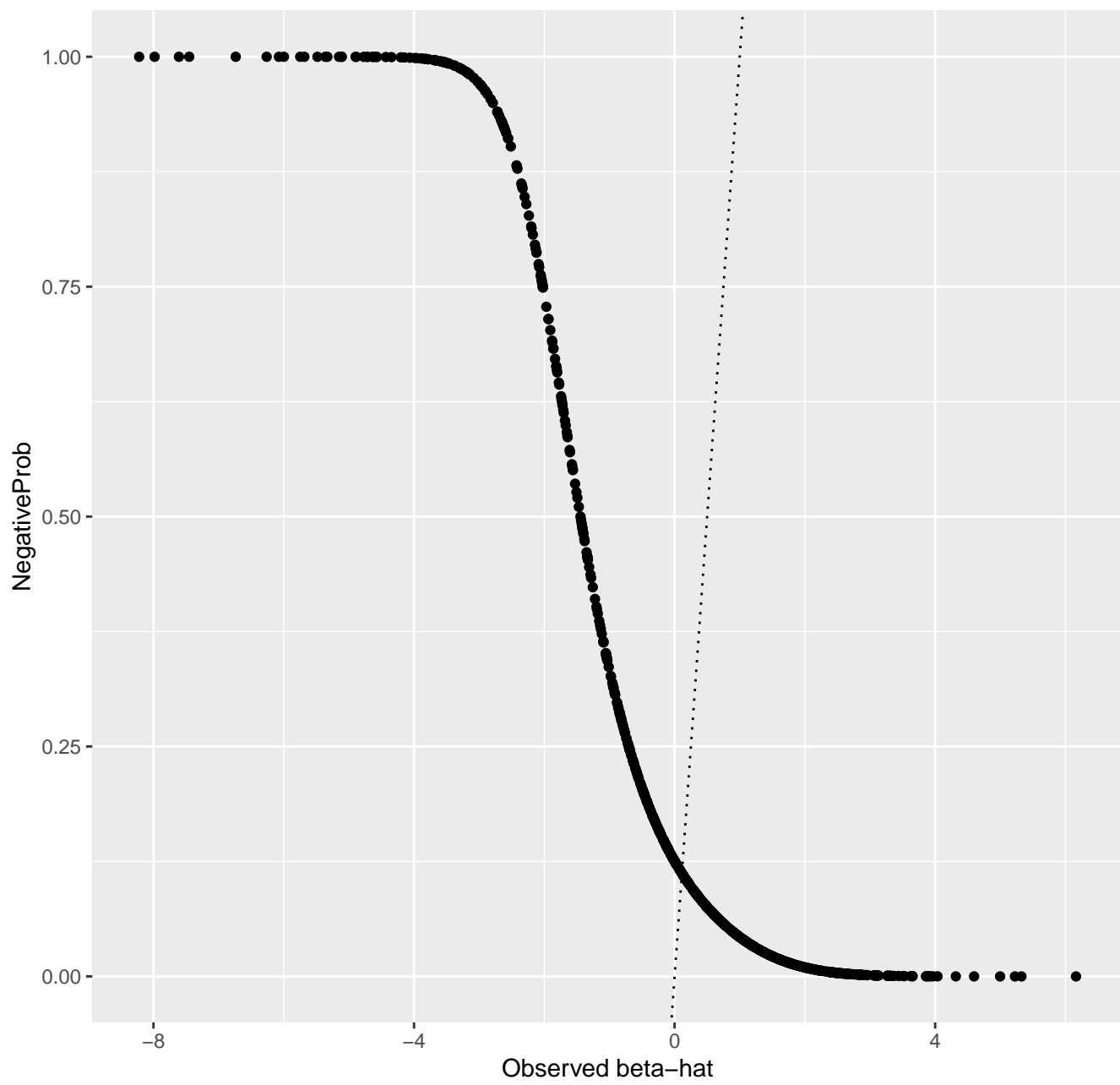
MTFMT_(123263) negative



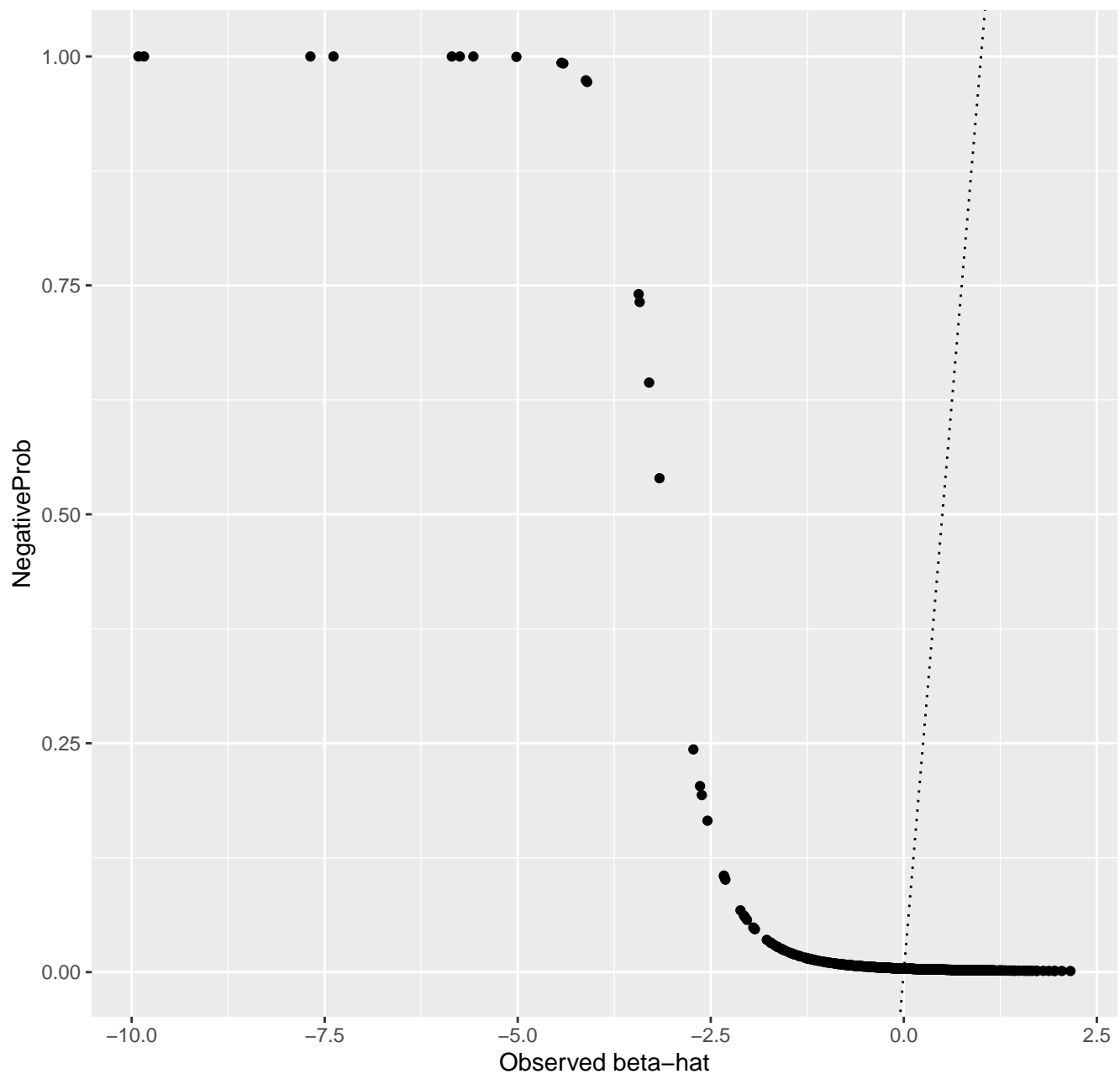
TKT_(7086) negative



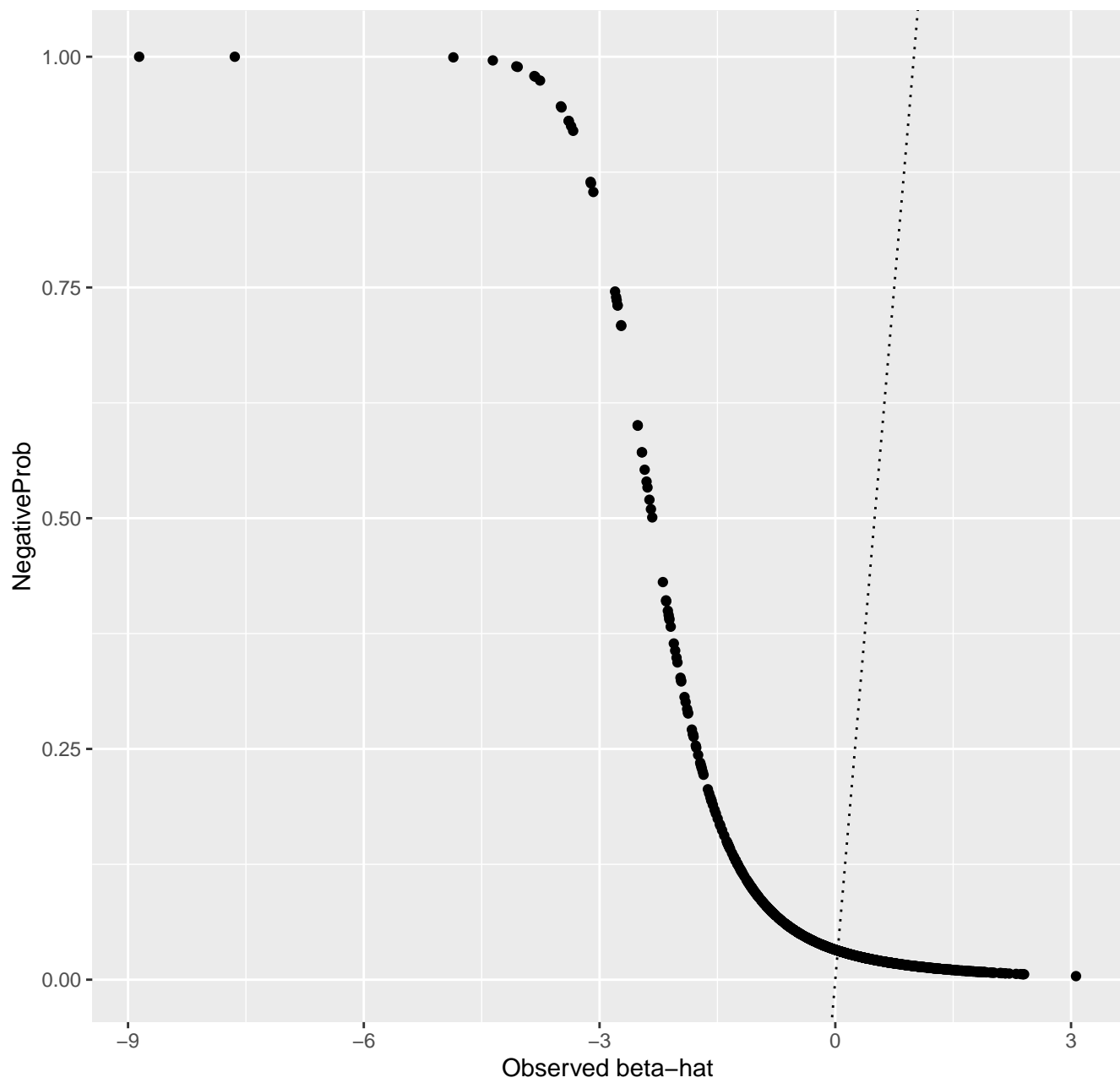
SLC4A7_(9497) negative



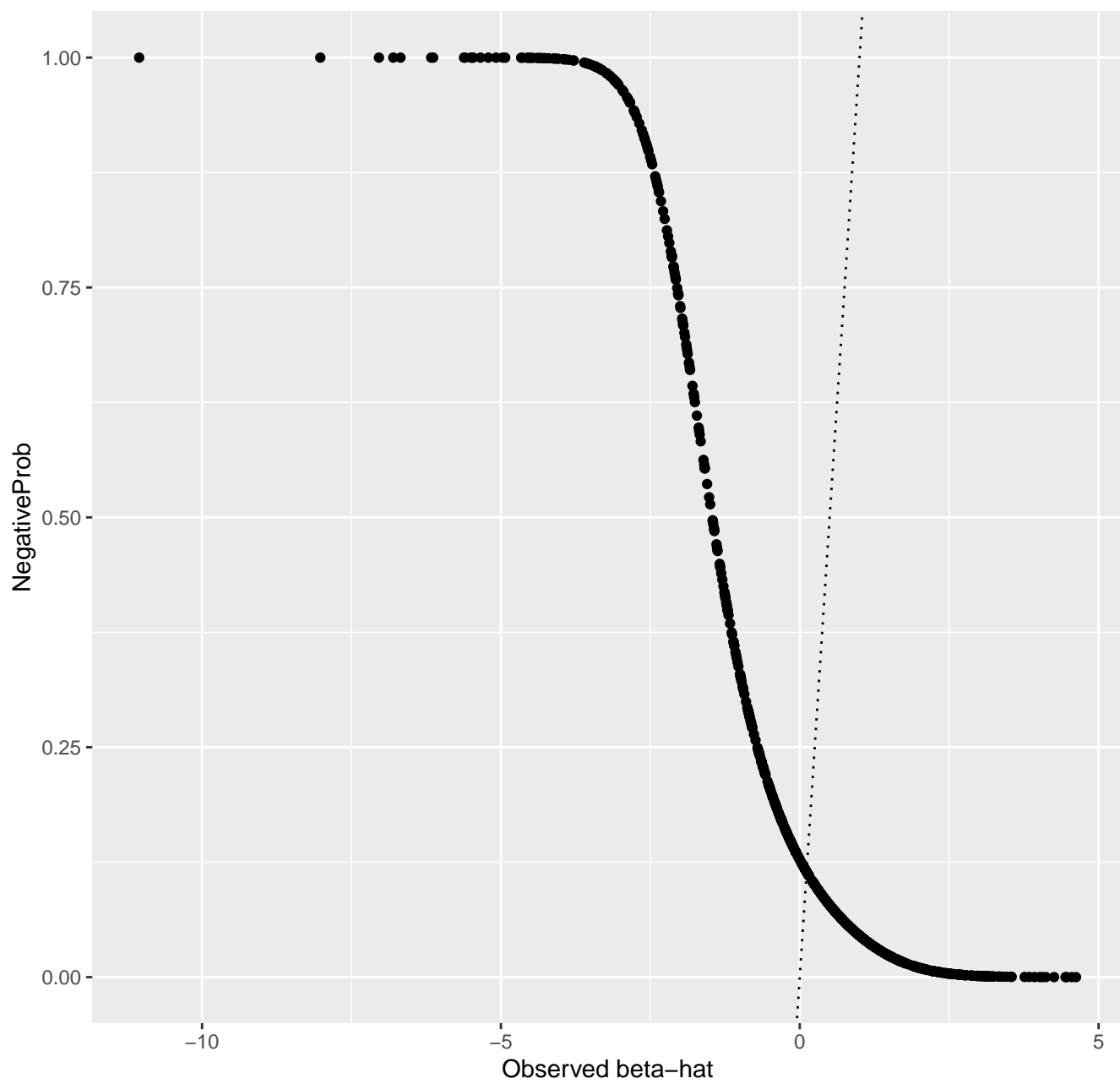
CD46_(4179) negative



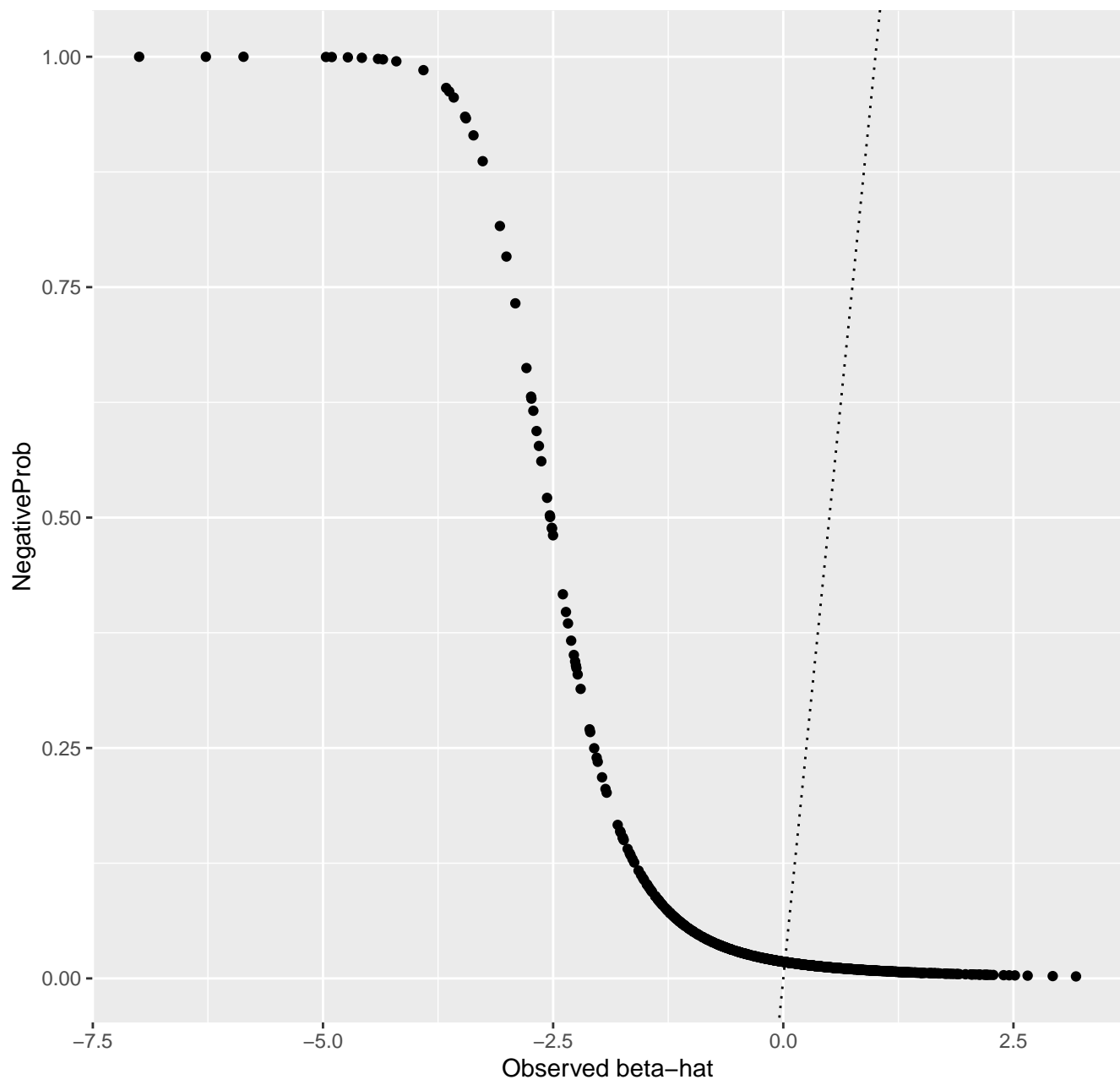
ARHGAP1_(392) negative



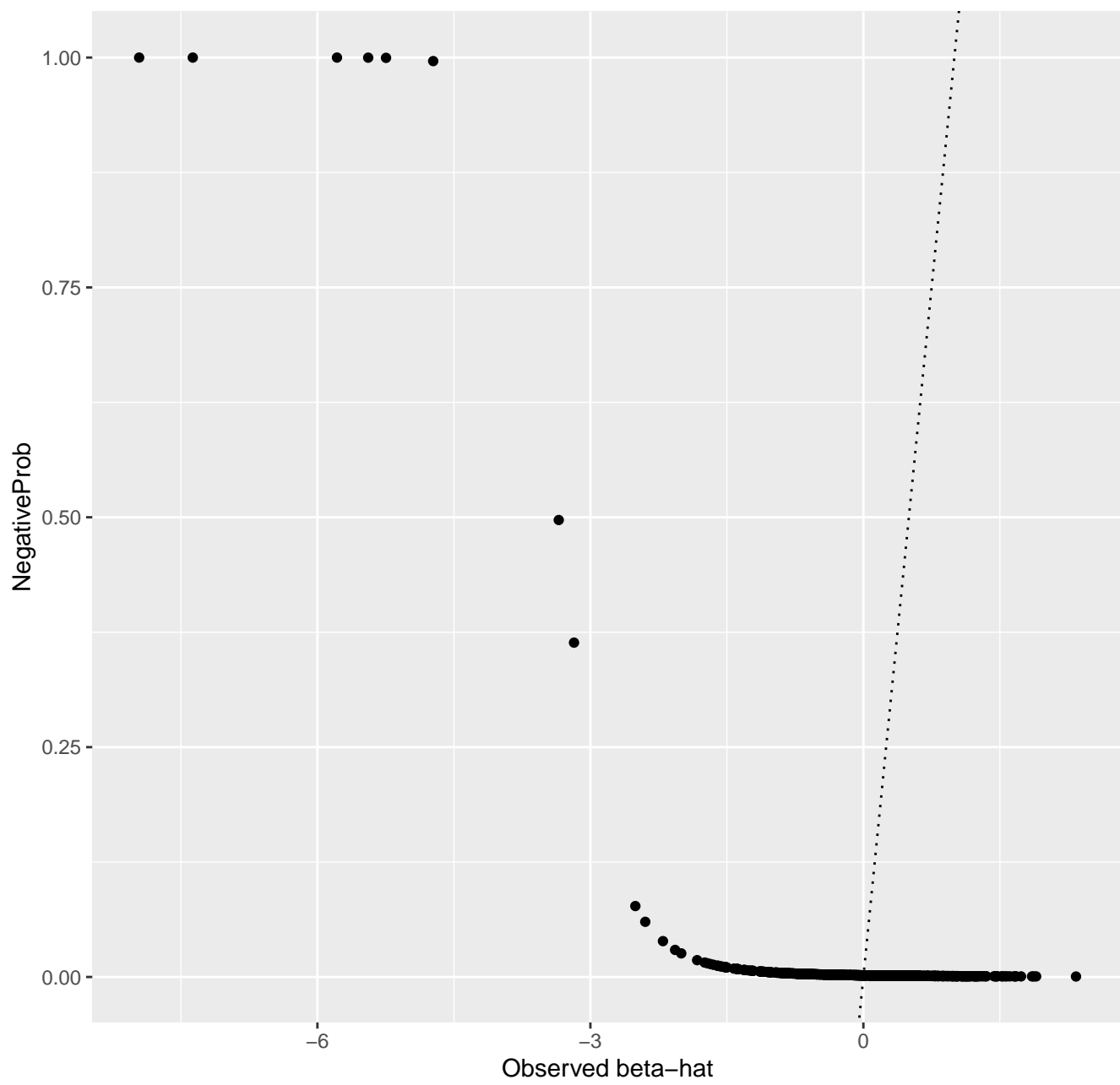
CMIP_(80790) negative



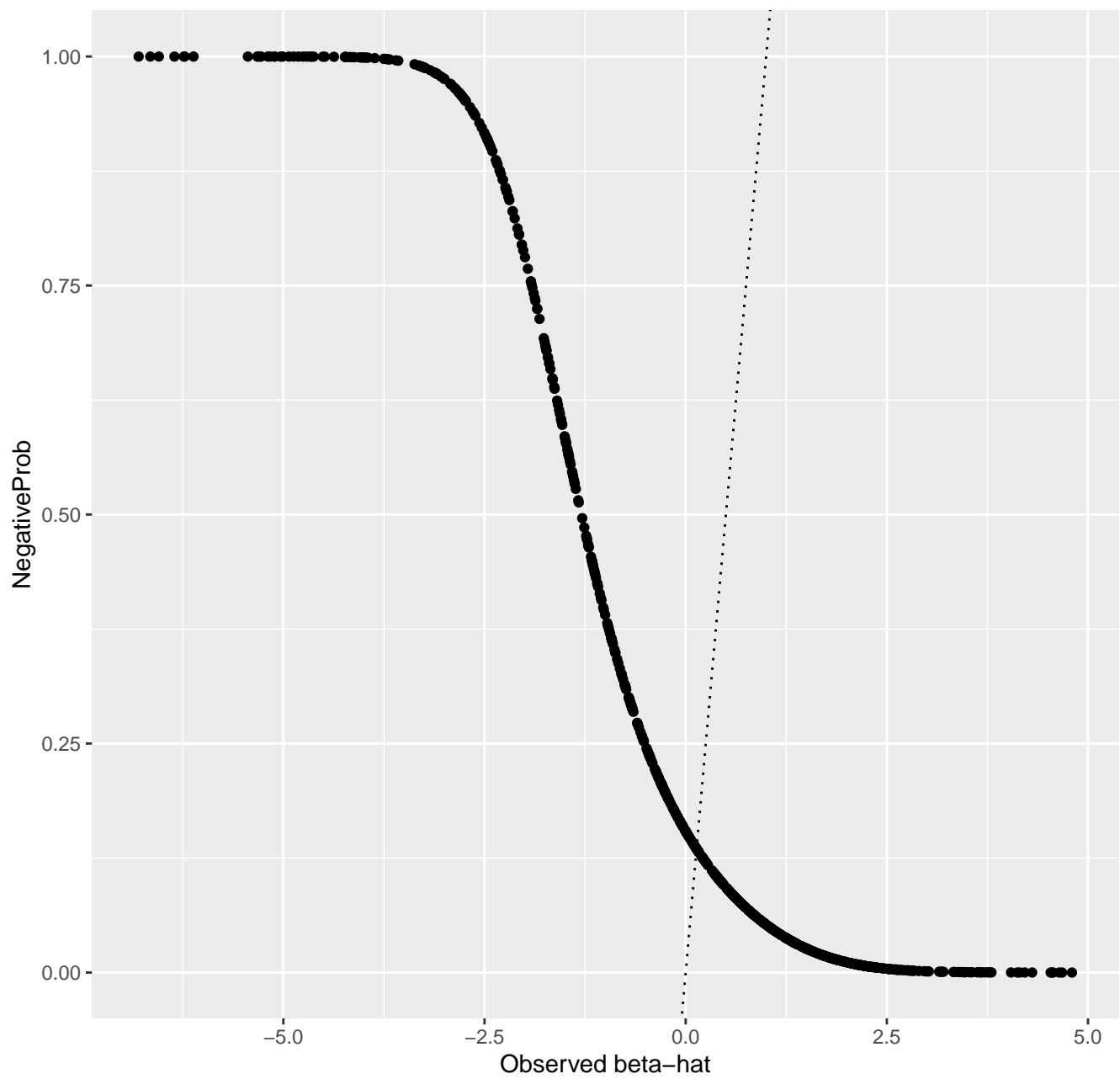
PEX13_(5194) negative



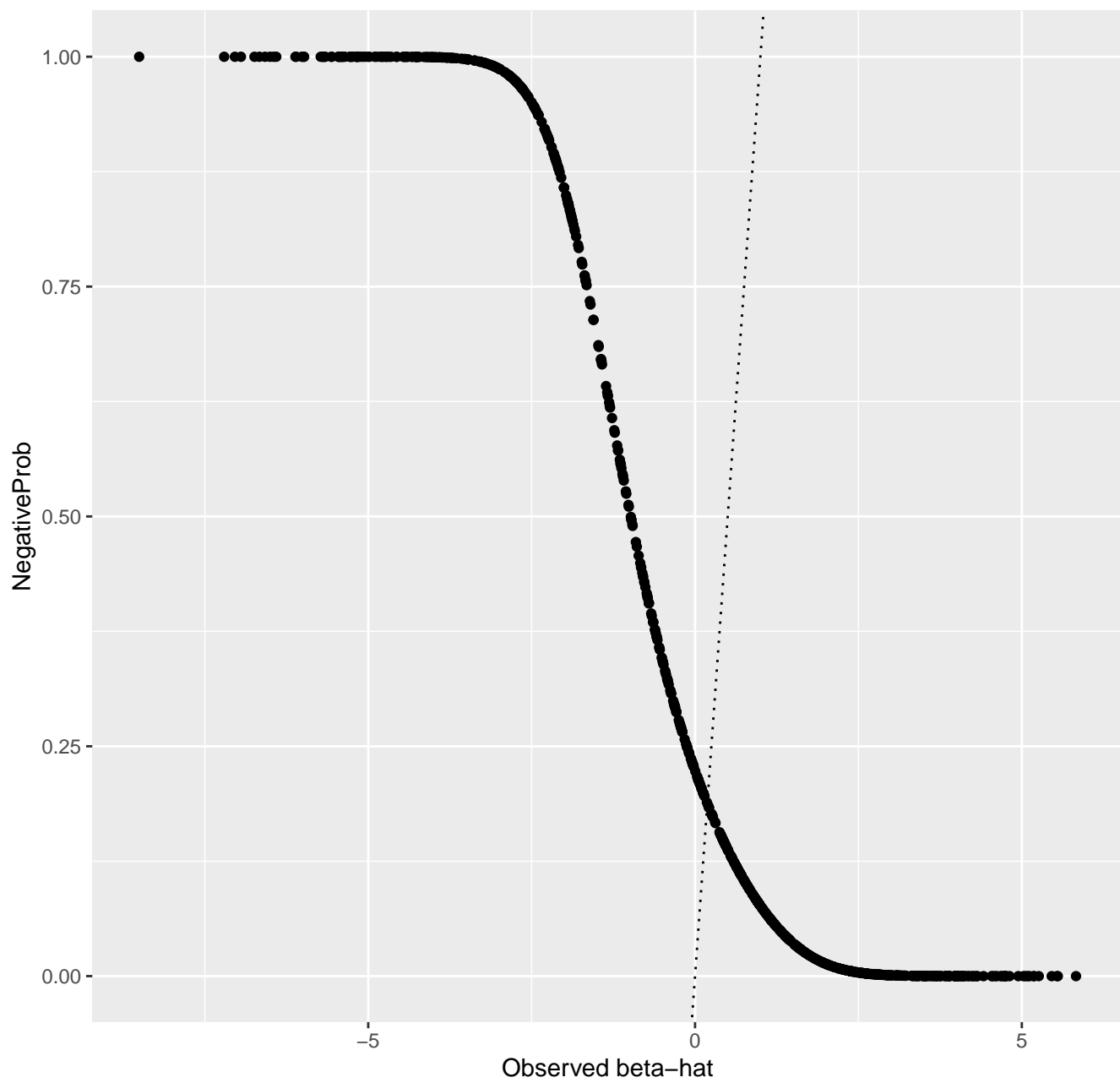
CYB561A3_(220002) negative



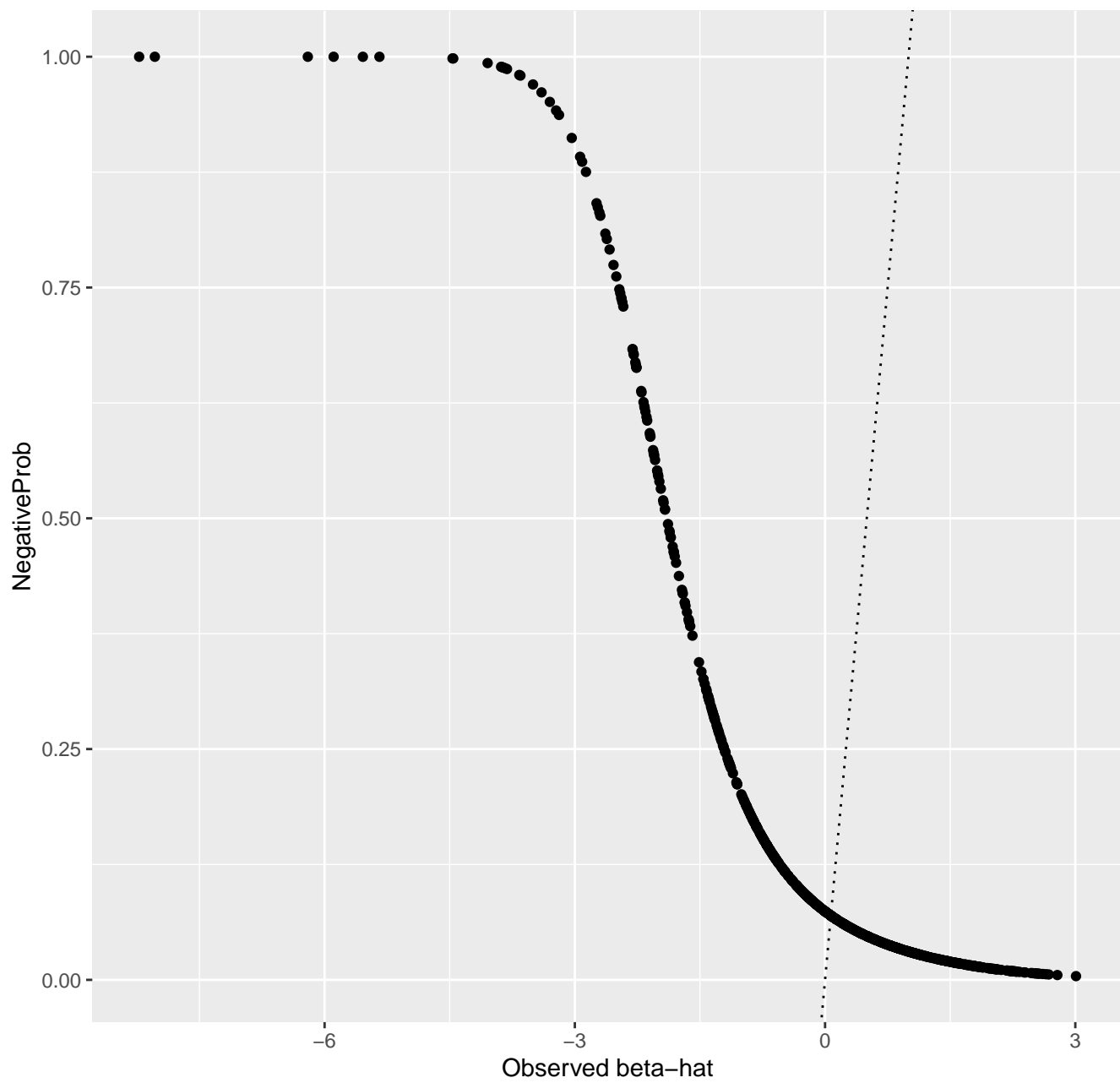
GET4_(51608) negative



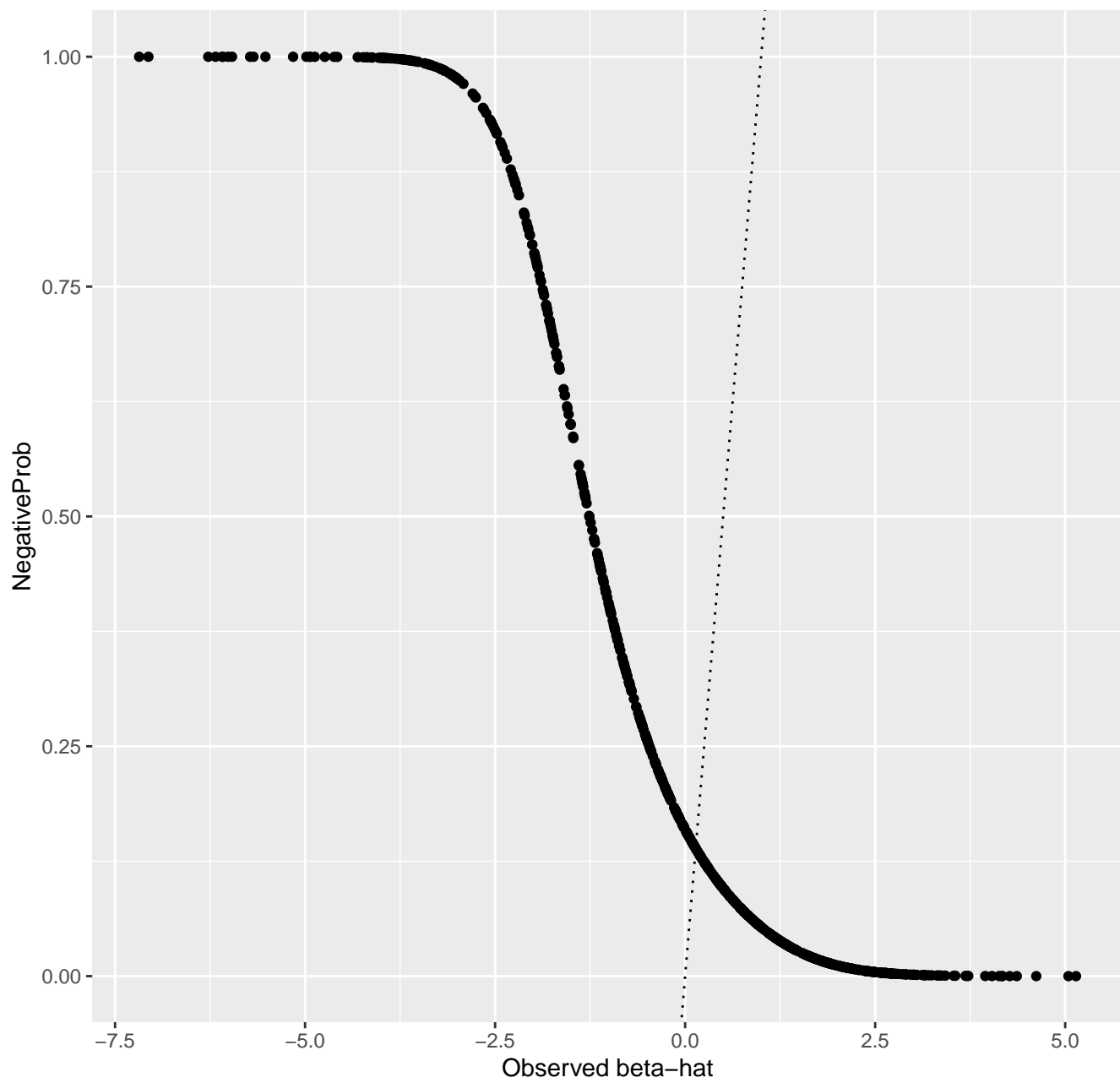
WDR73_(84942) negative



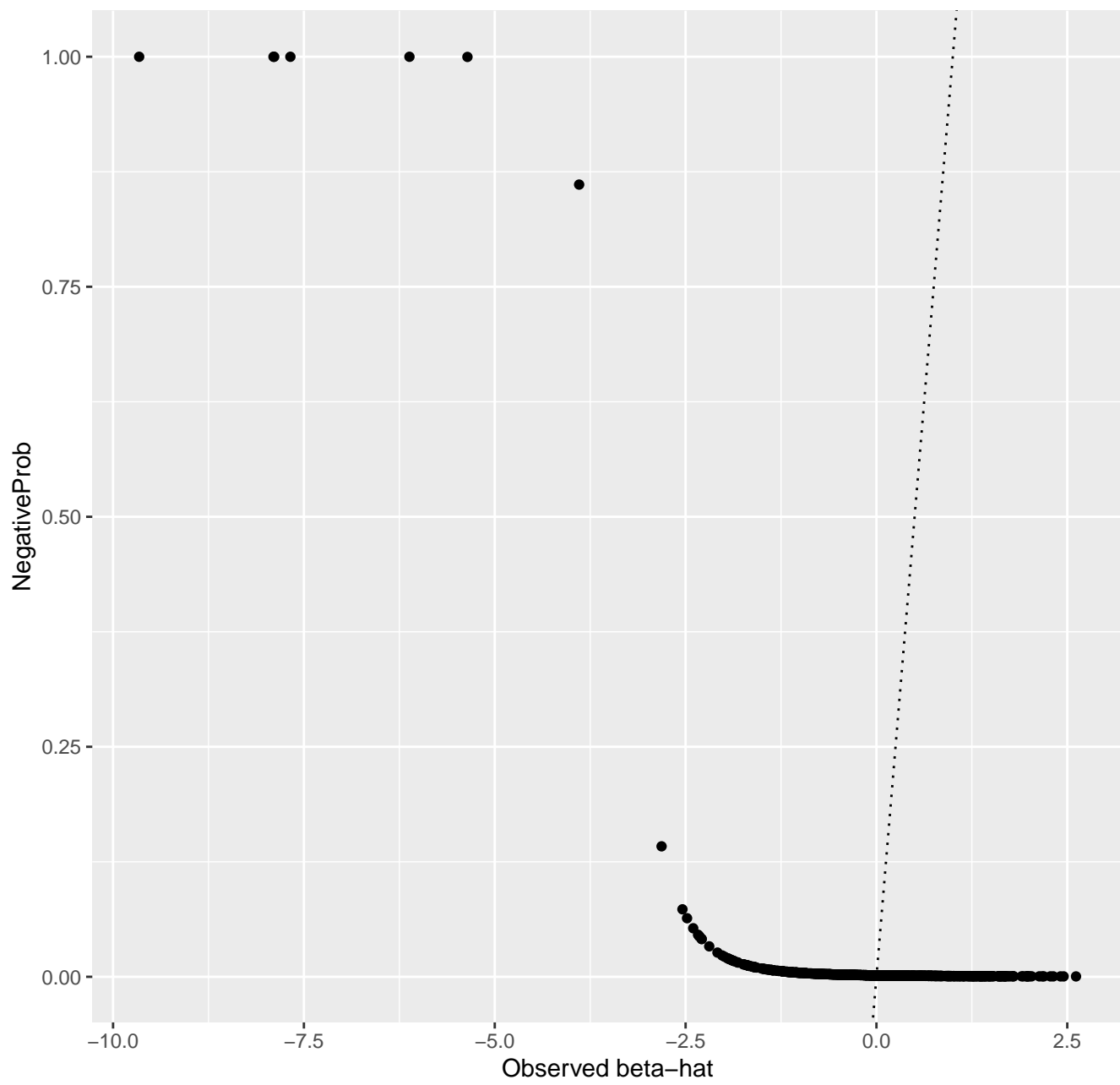
CIP2A_(57650) negative



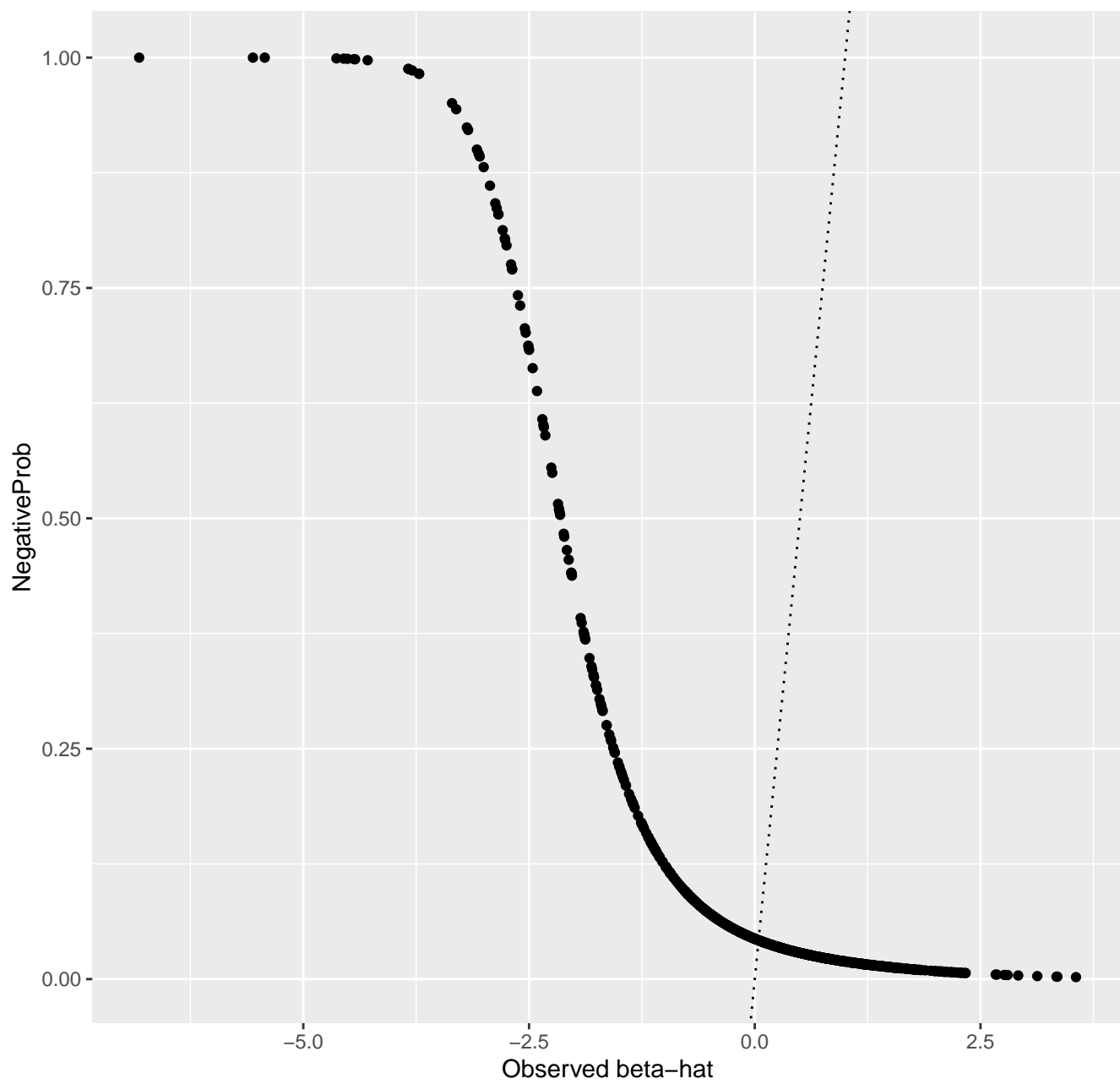
RBM18_(92400) negative



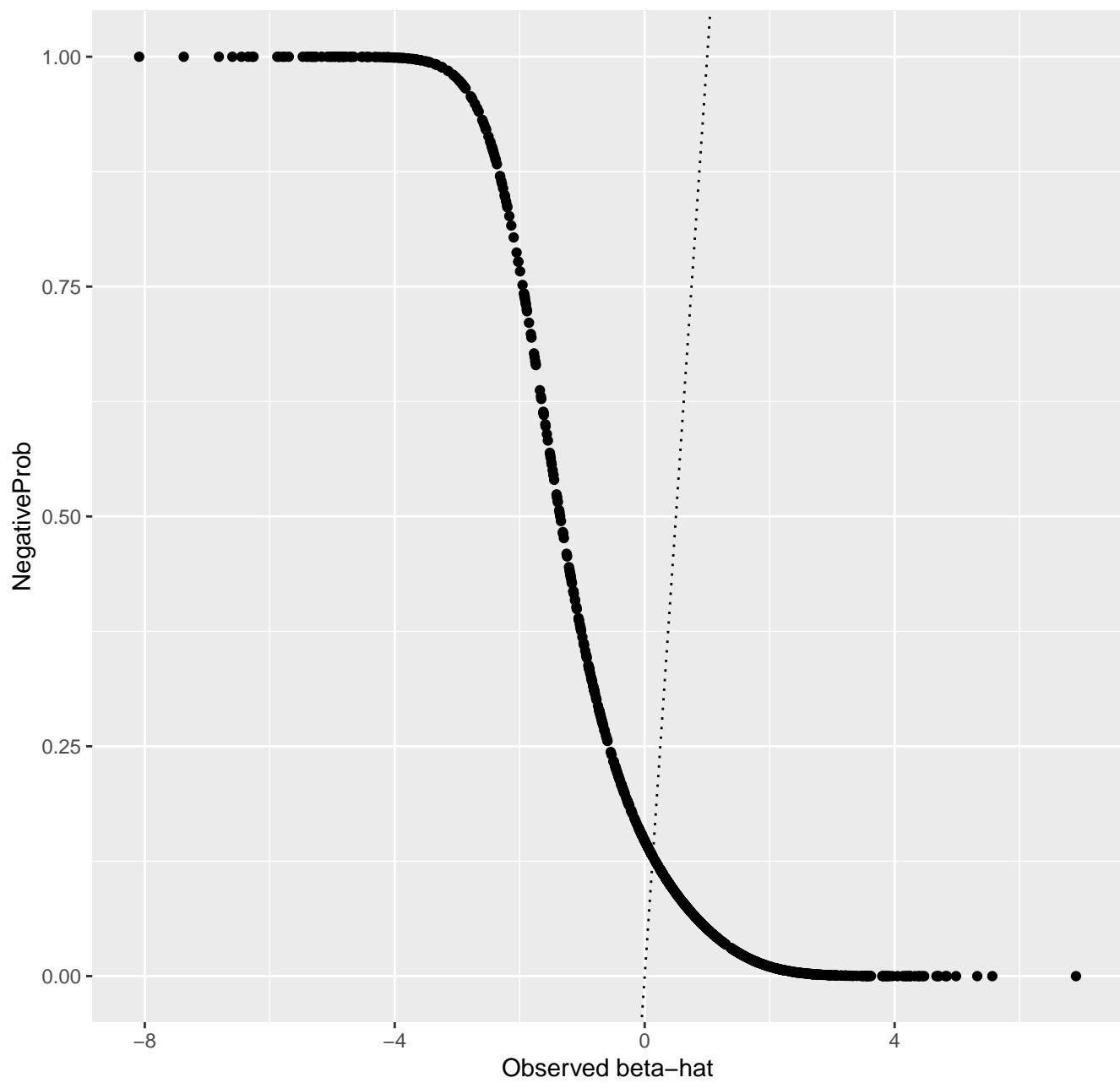
BCR_(613) negative



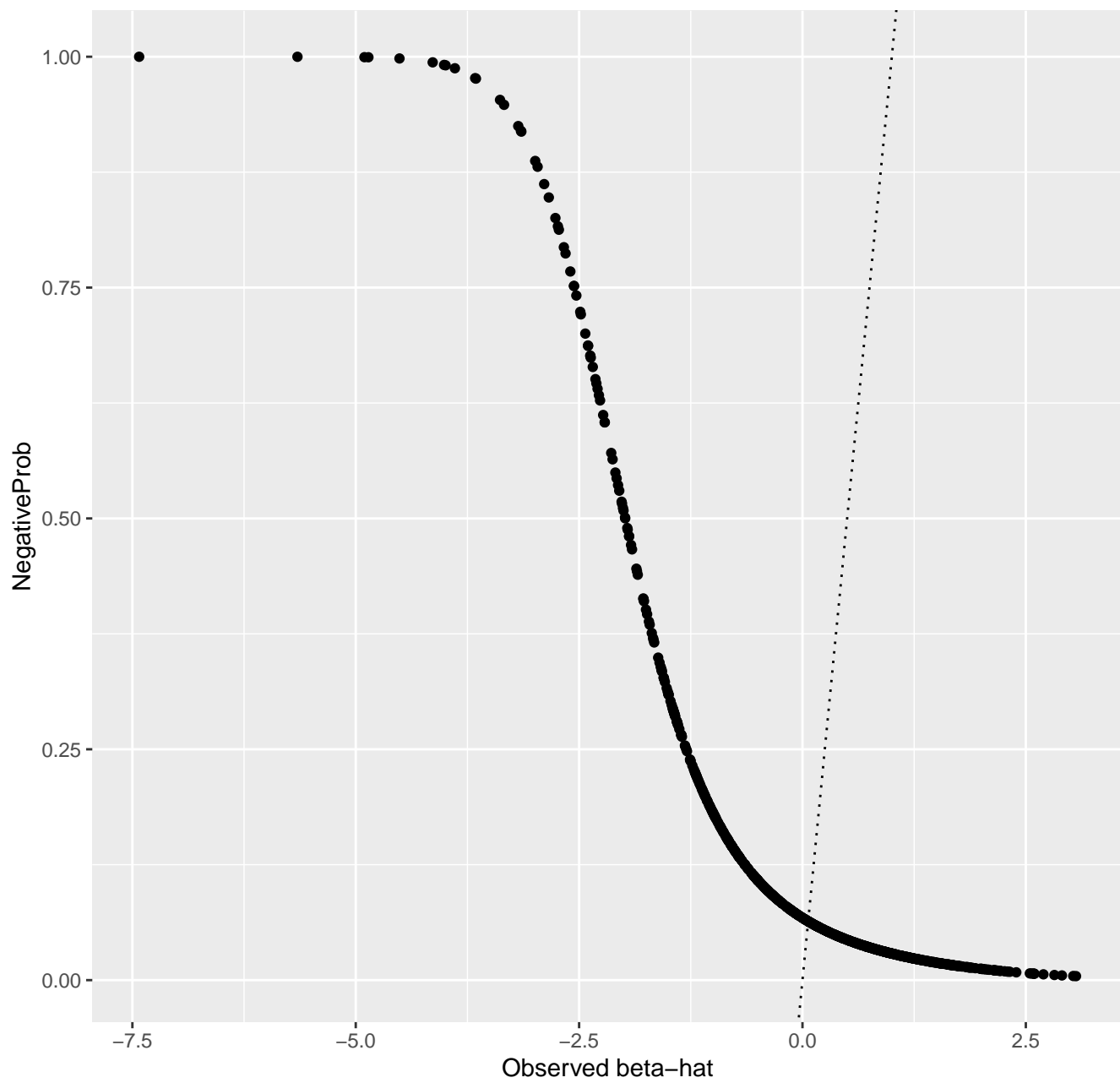
C19orf70_(125988) negative



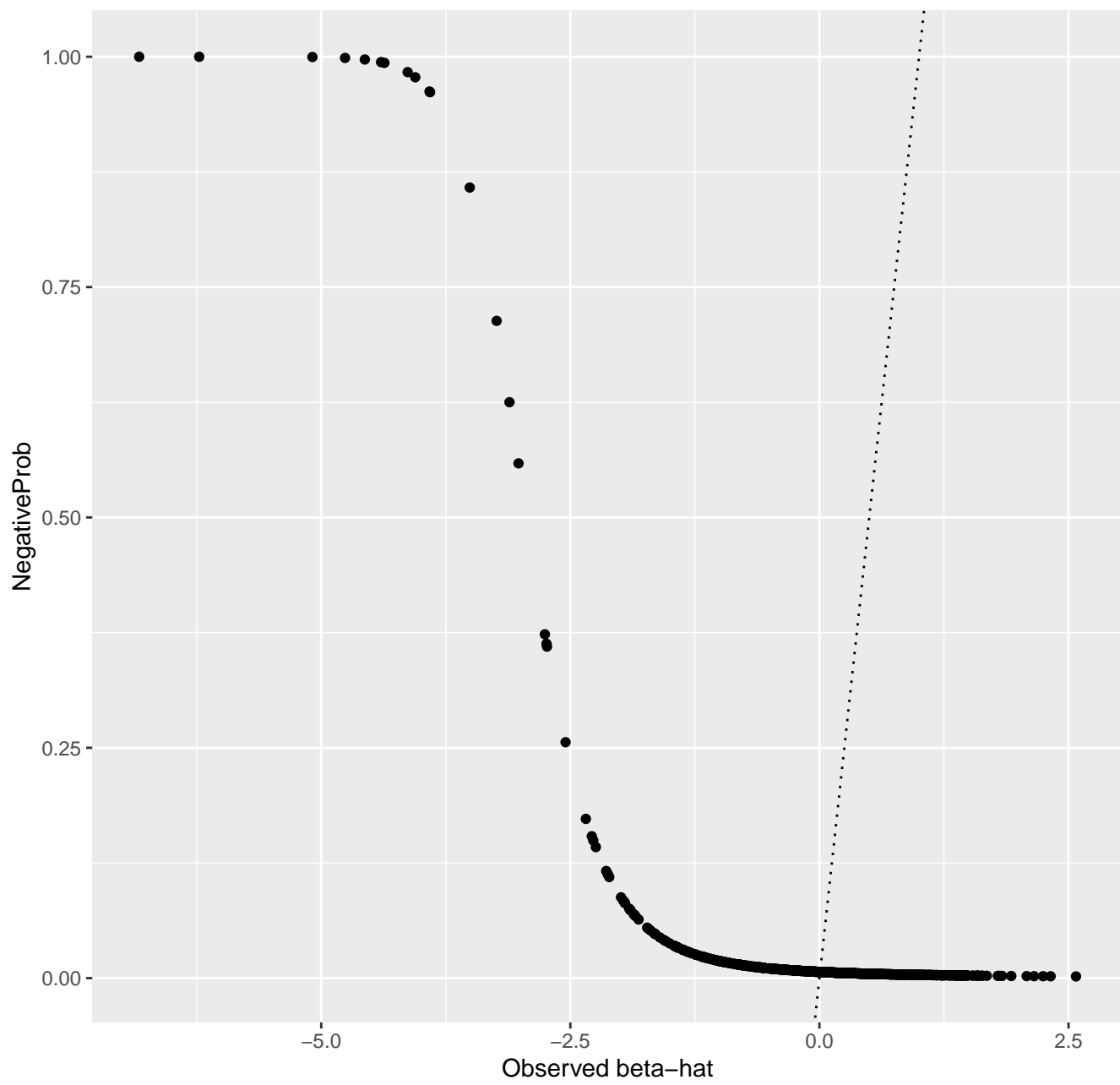
RAB10_(10890) negative



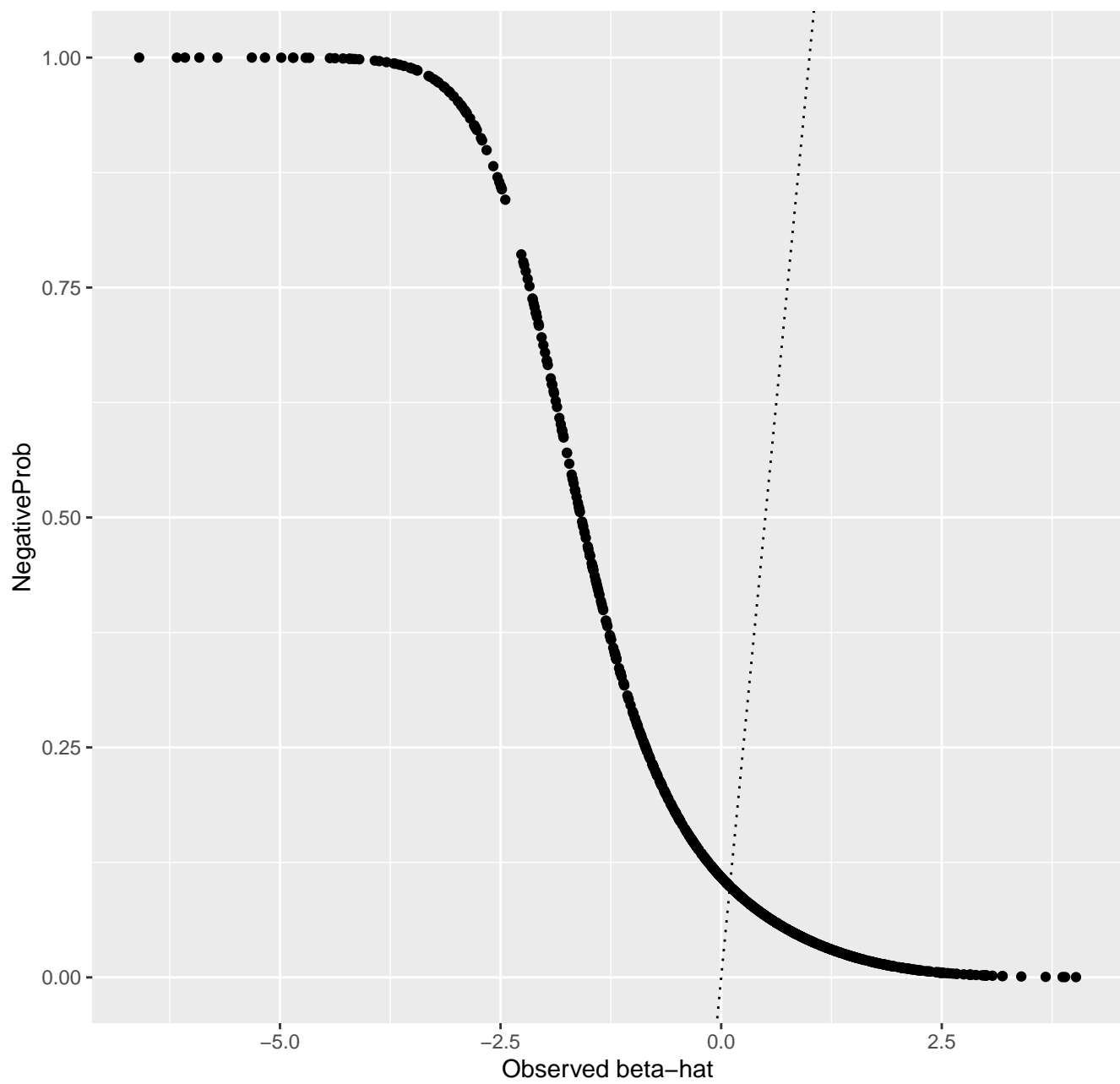
COL4A3BP_(10087) negative



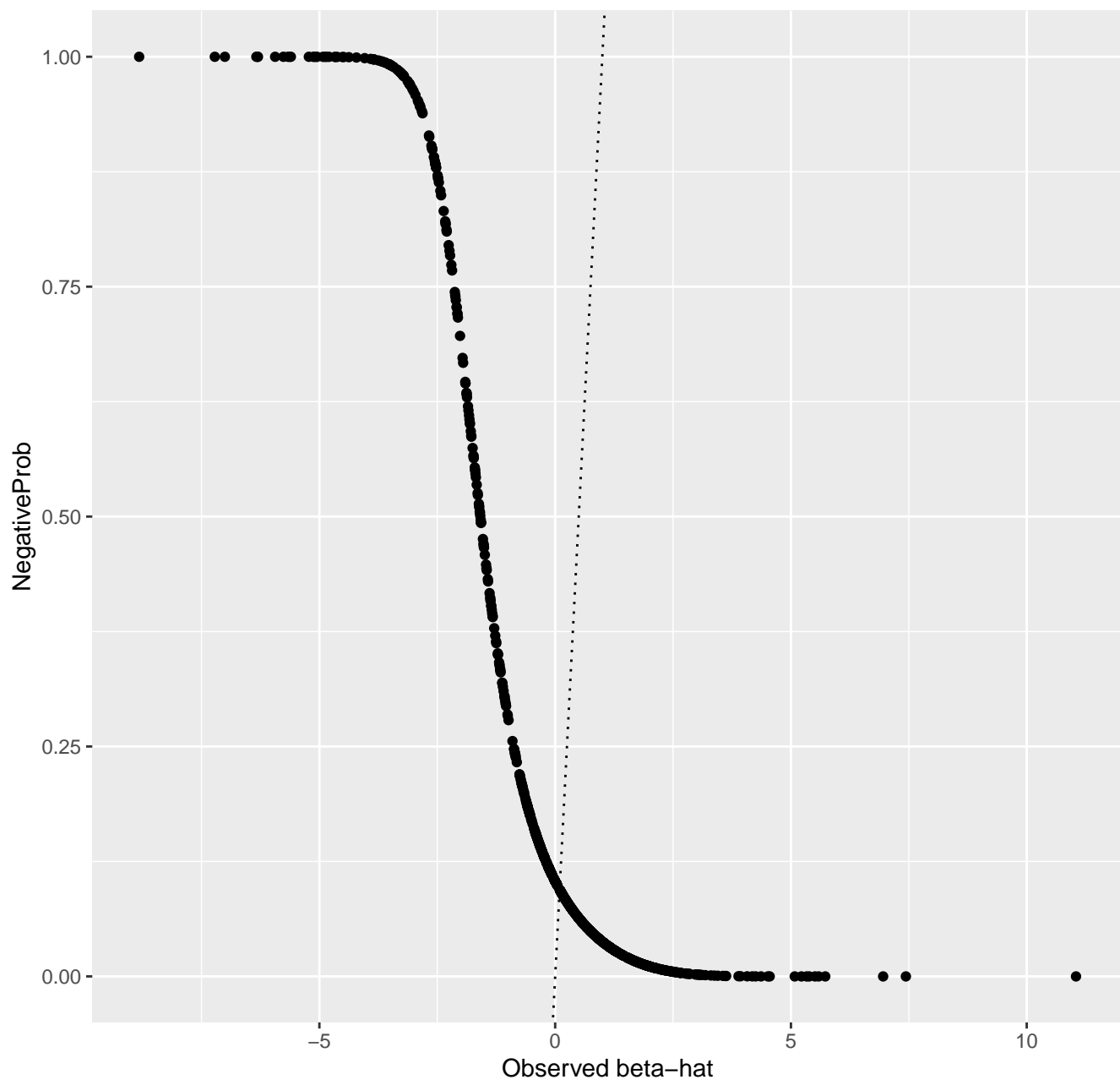
DHRS7B_(25979) negative



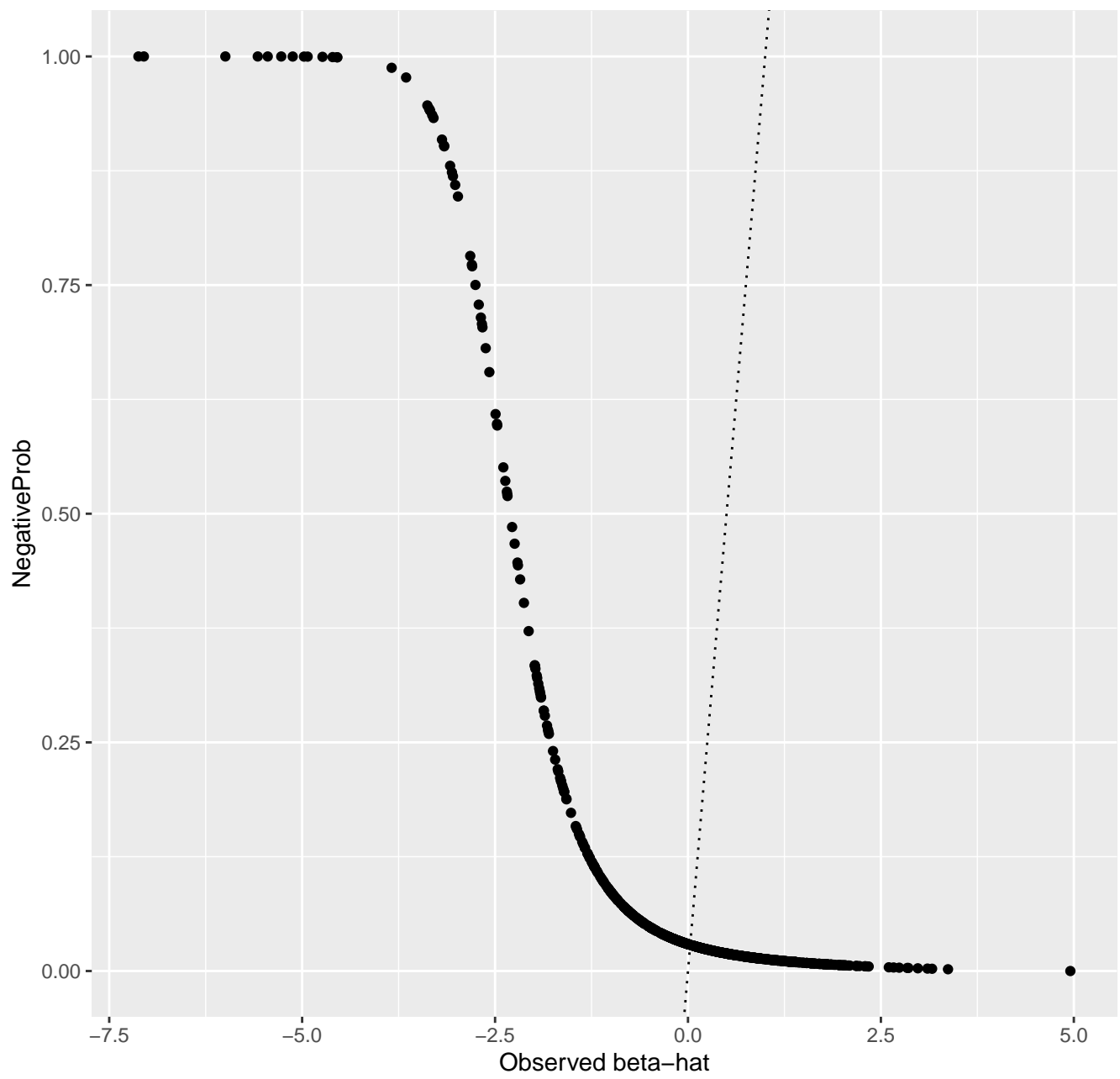
DNAJC21_(134218) negative



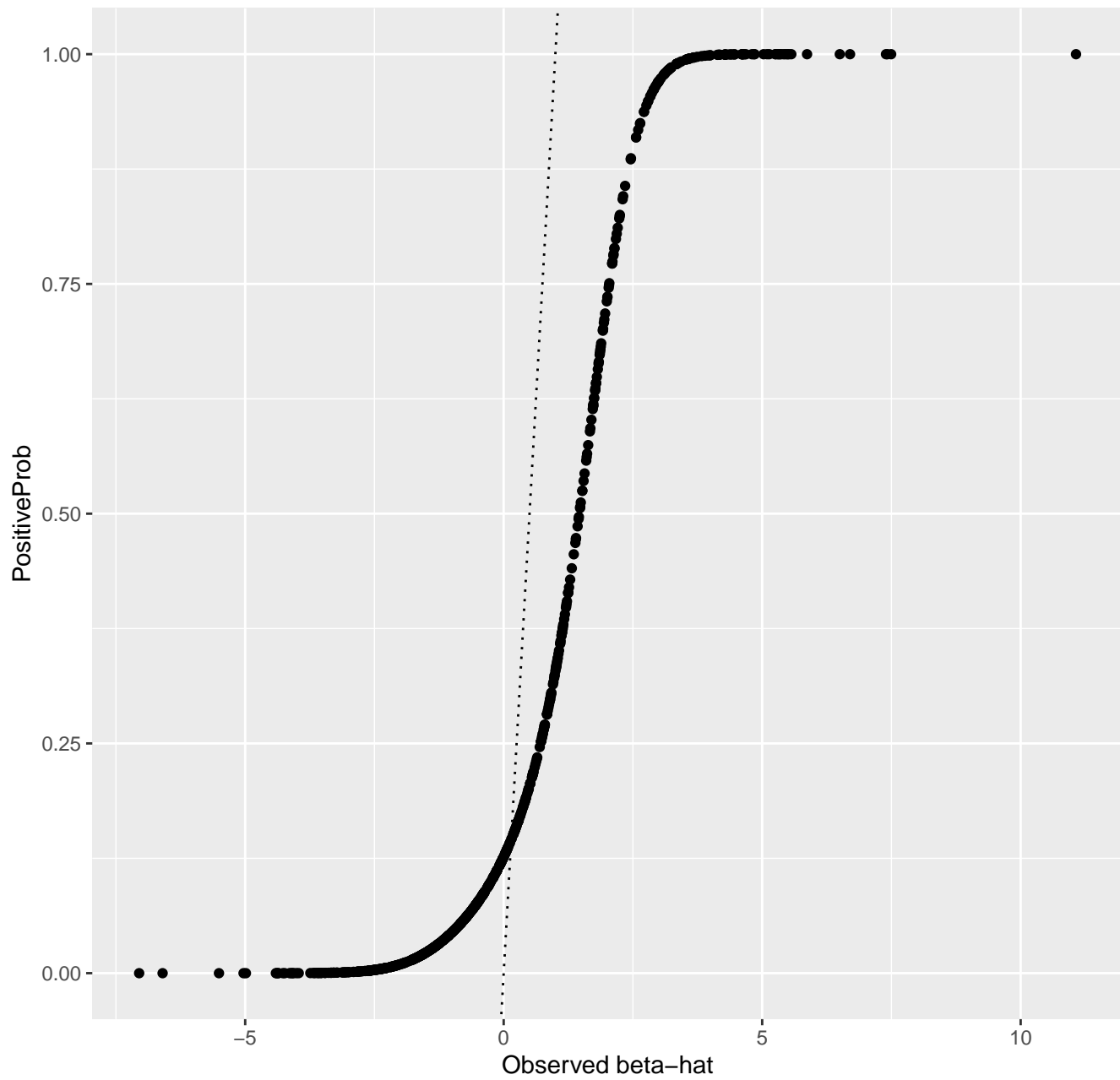
CTNNA1_(1495) negative



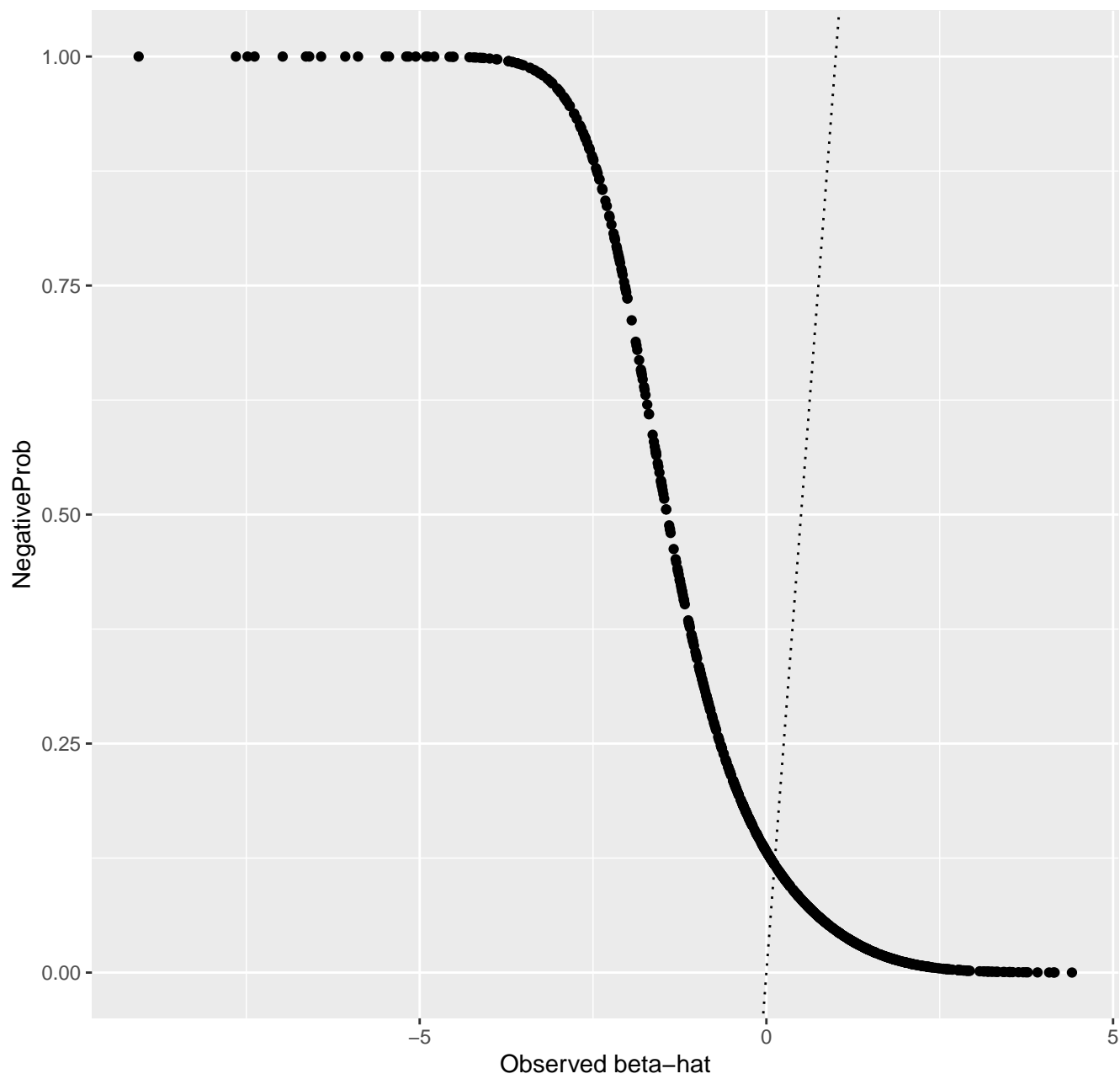
RXRA_(6256) negative



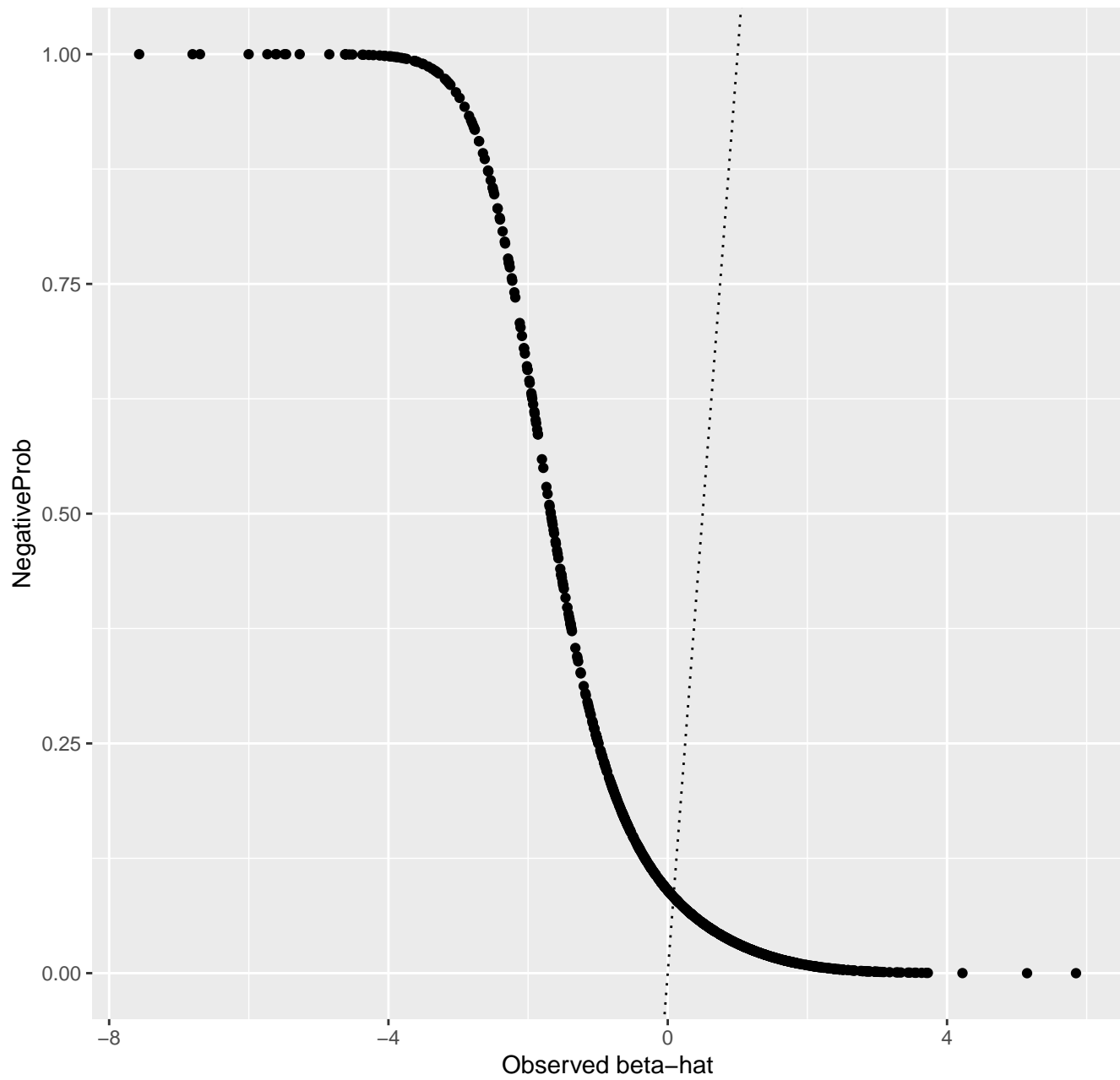
UBE2N_(7334) positive



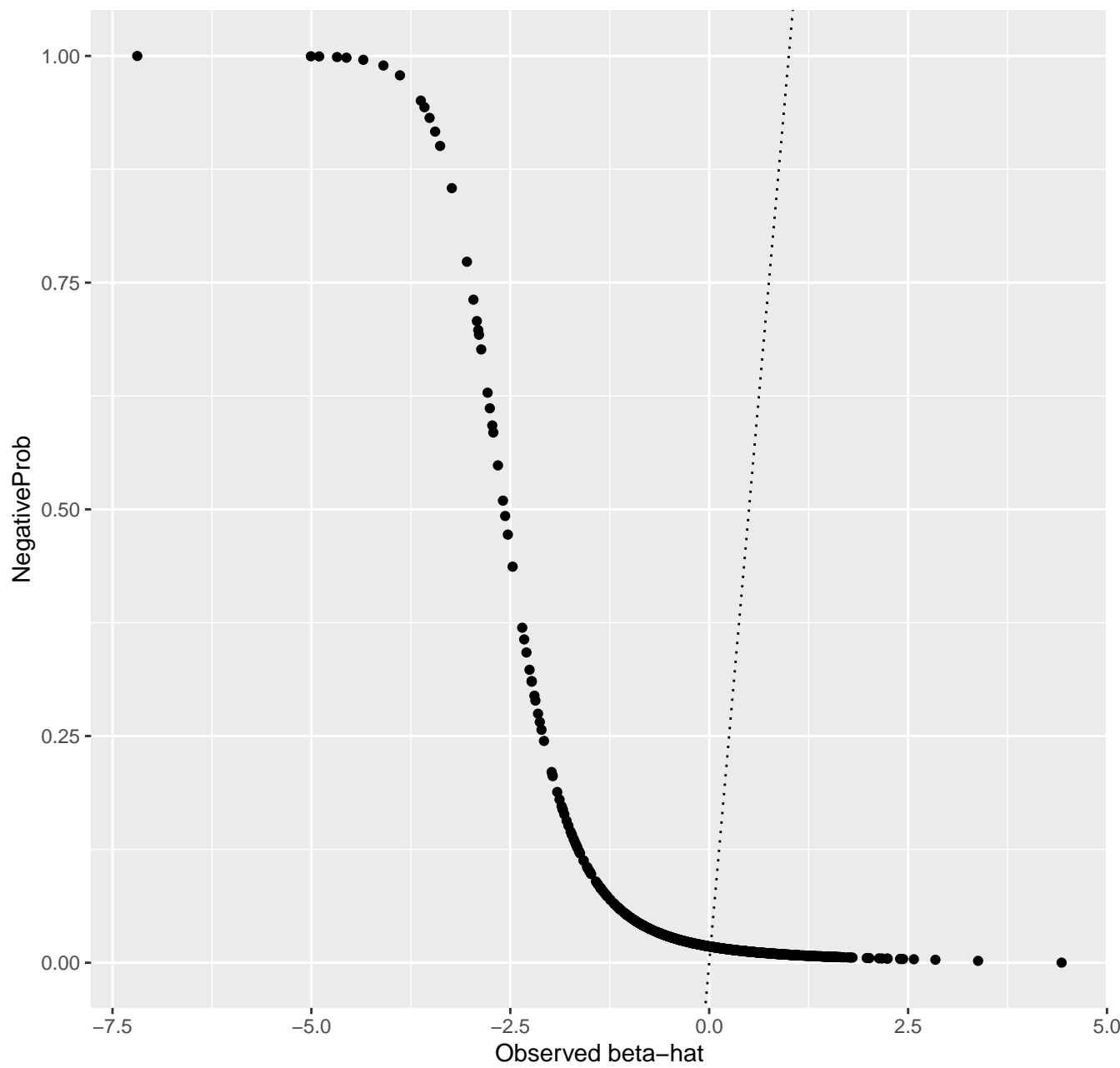
ARF1_(375) negative



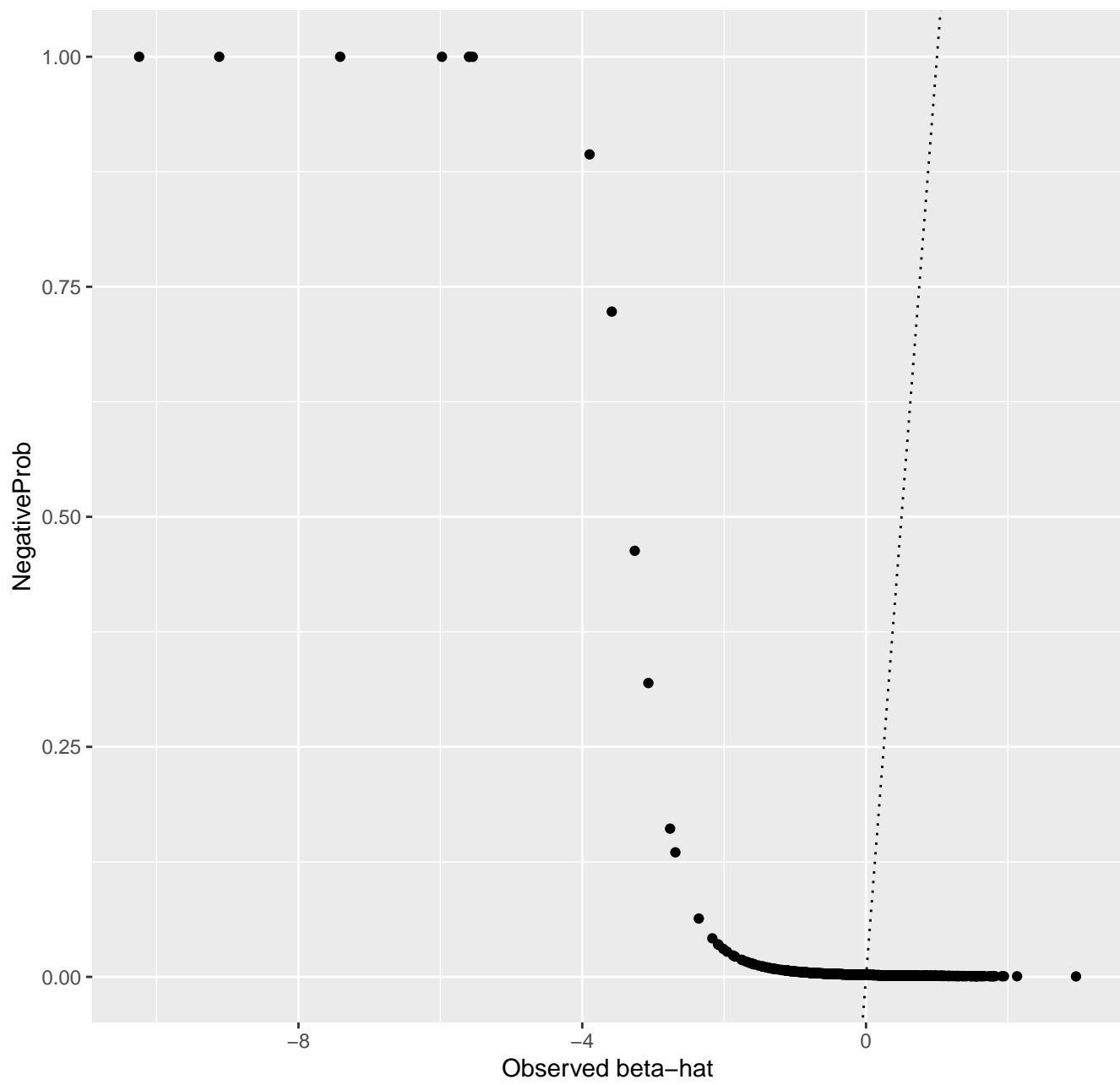
UBE2G2_(7327) negative



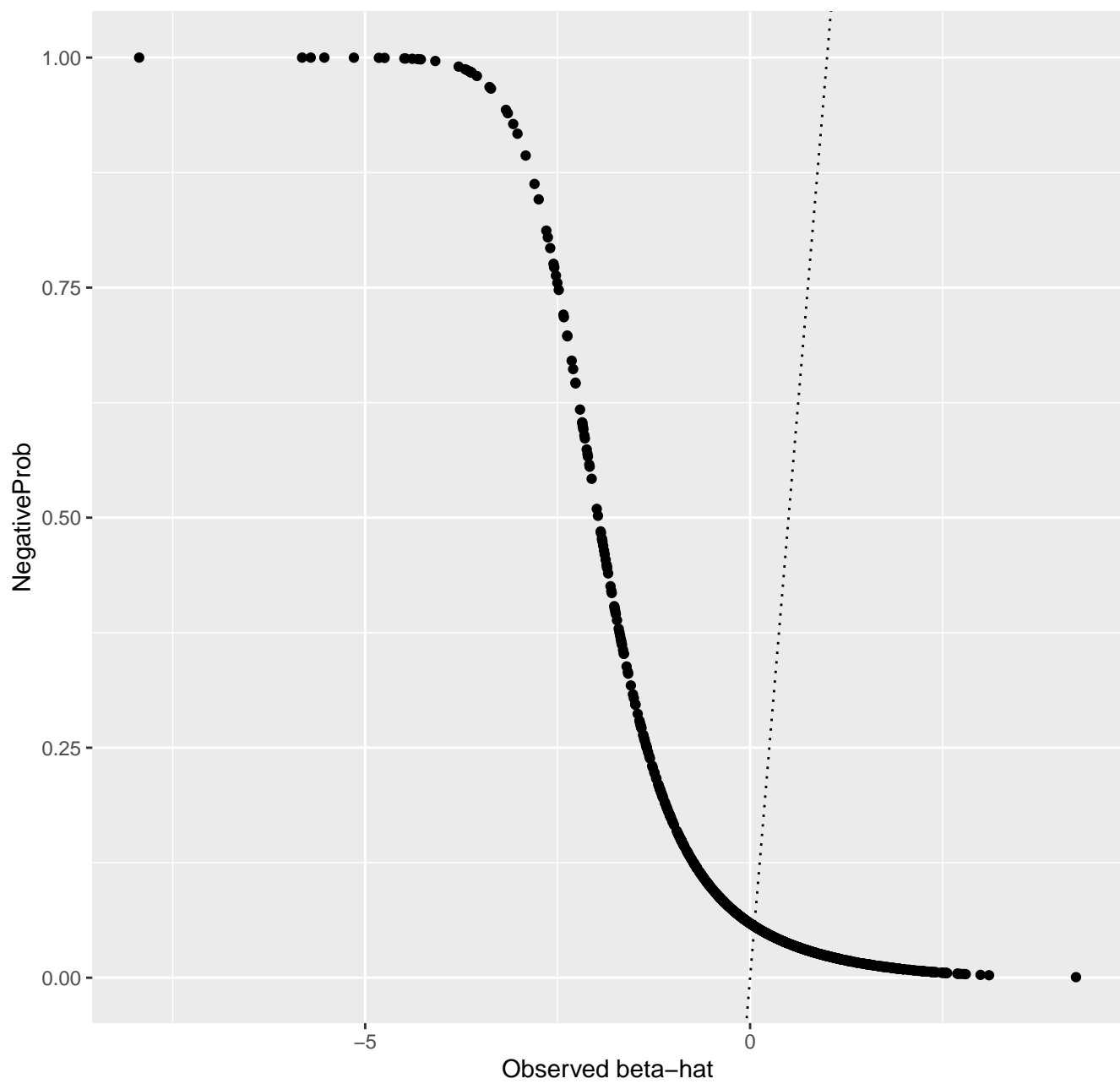
SLC10A7_(84068) negative



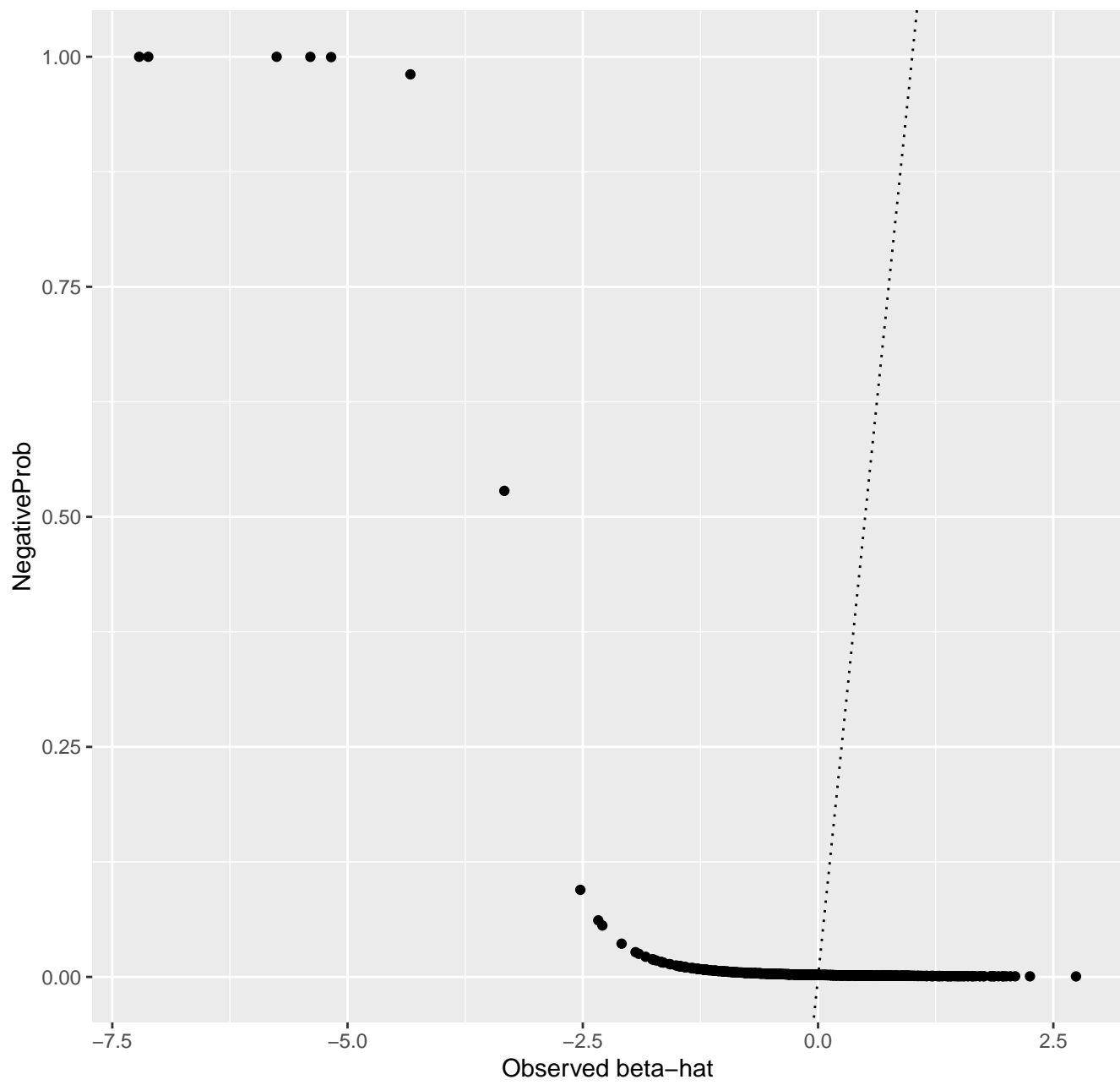
NEUROD1_(4760) negative



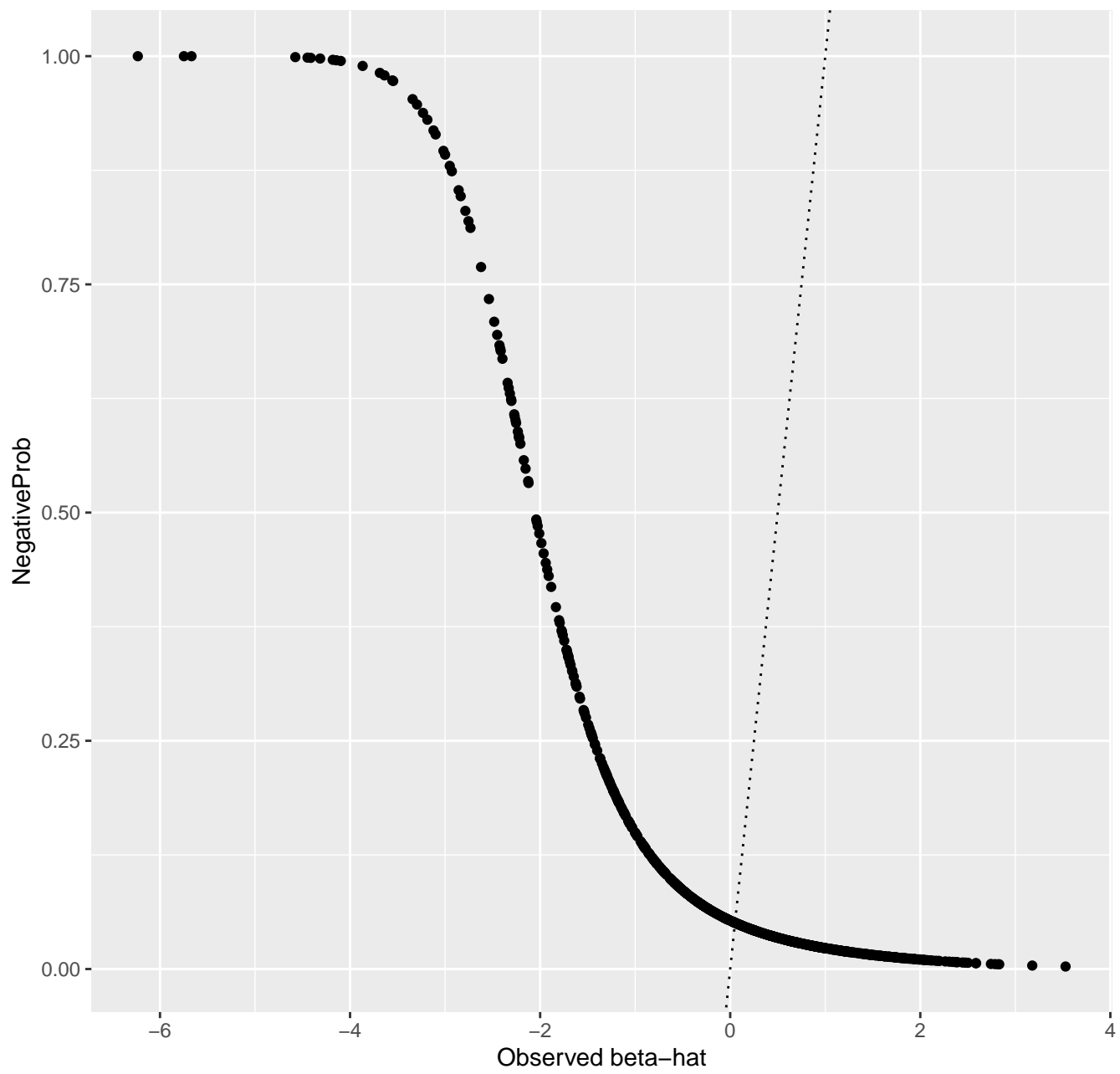
ASPM_(259266) negative



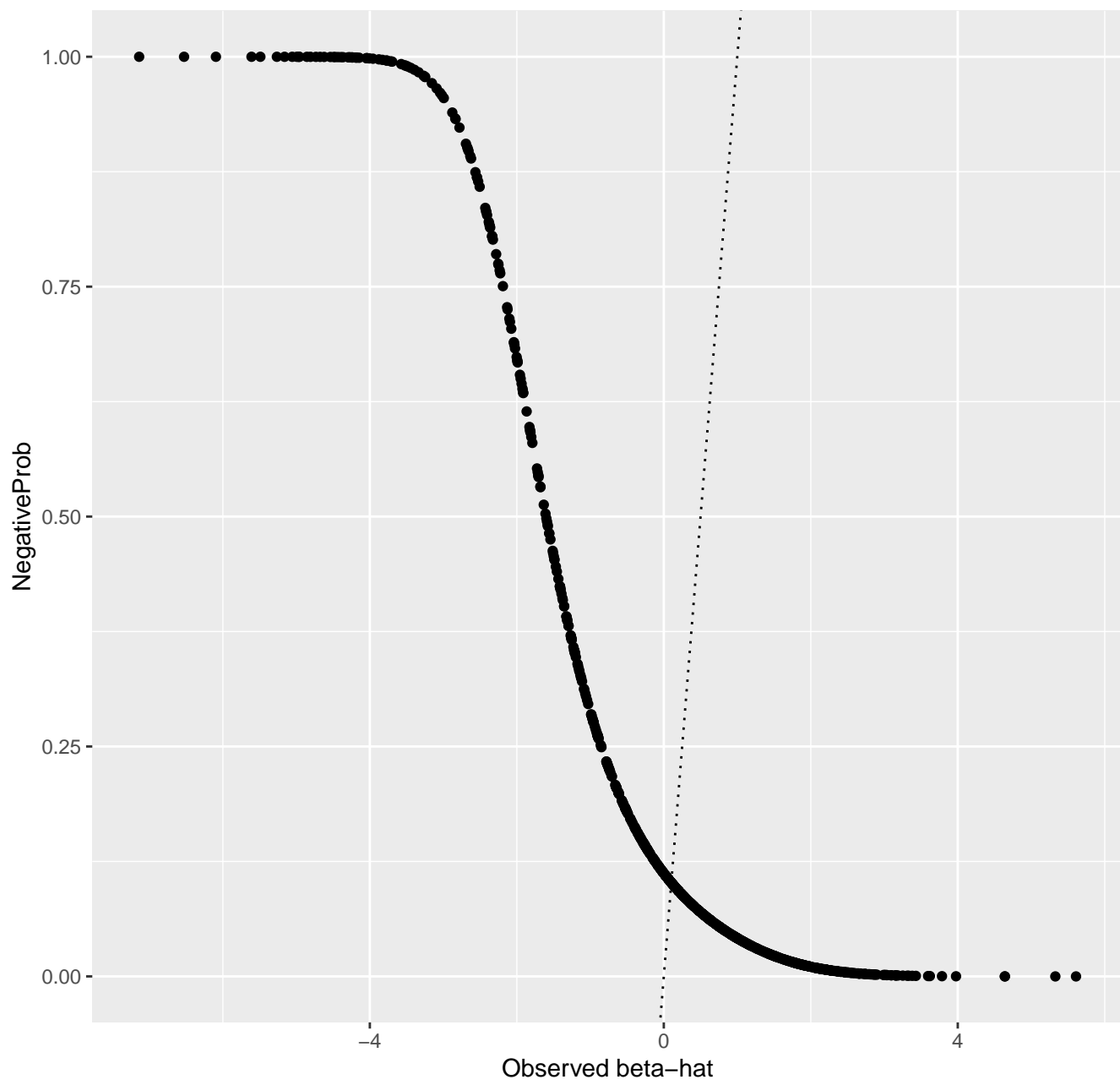
GATA1_(2623) negative



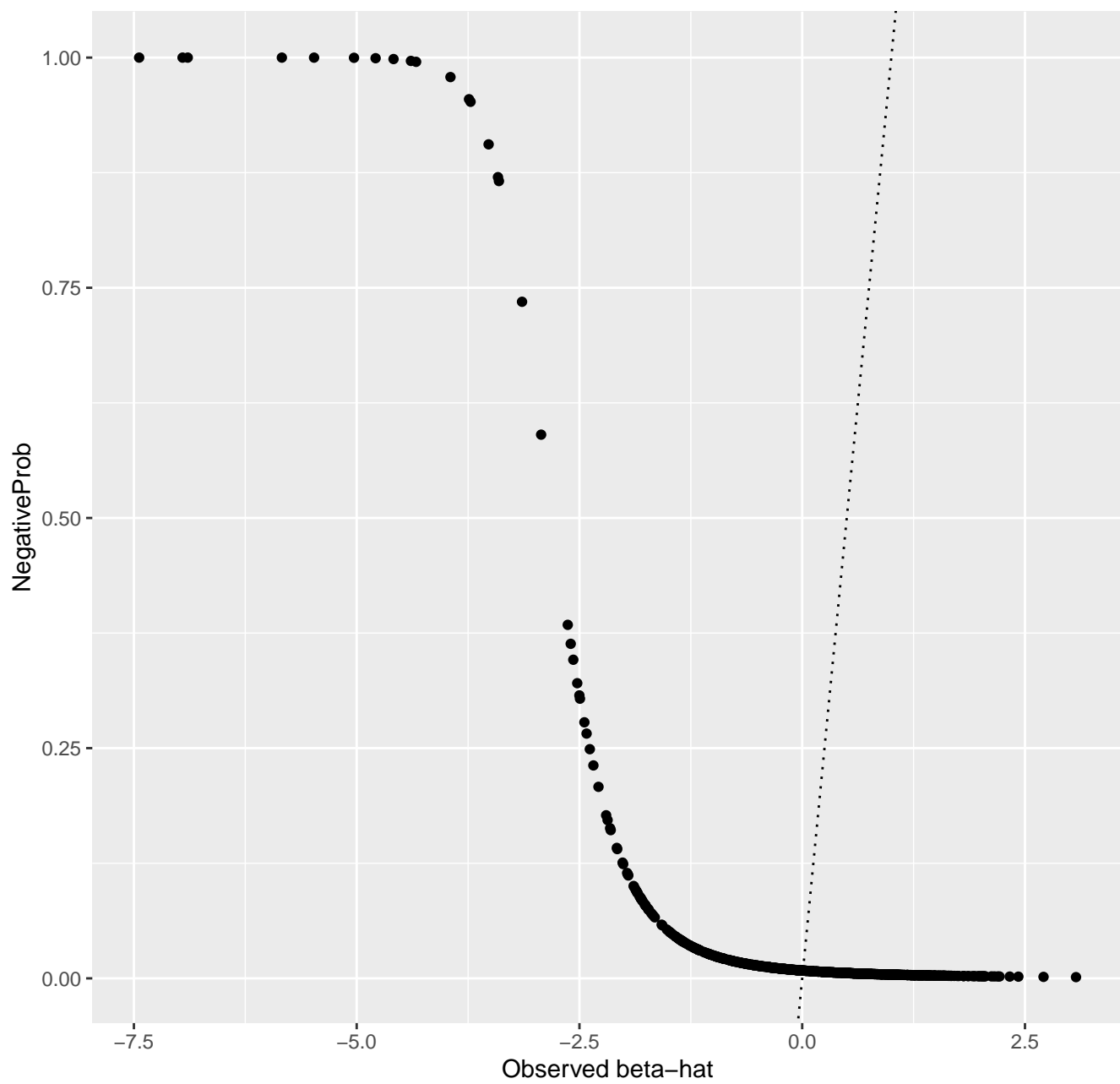
COG6_(57511) negative



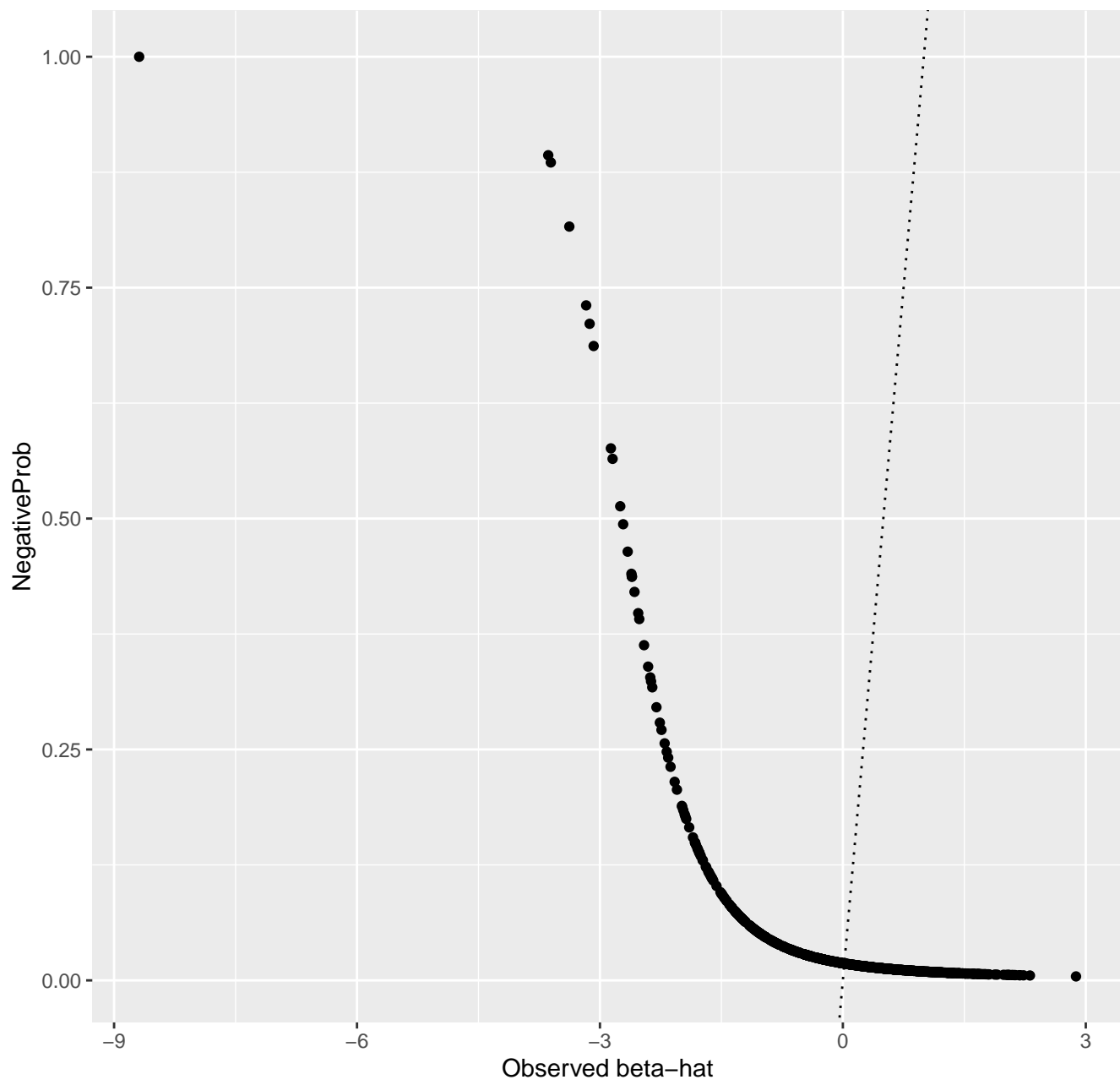
RNF168_(165918) negative



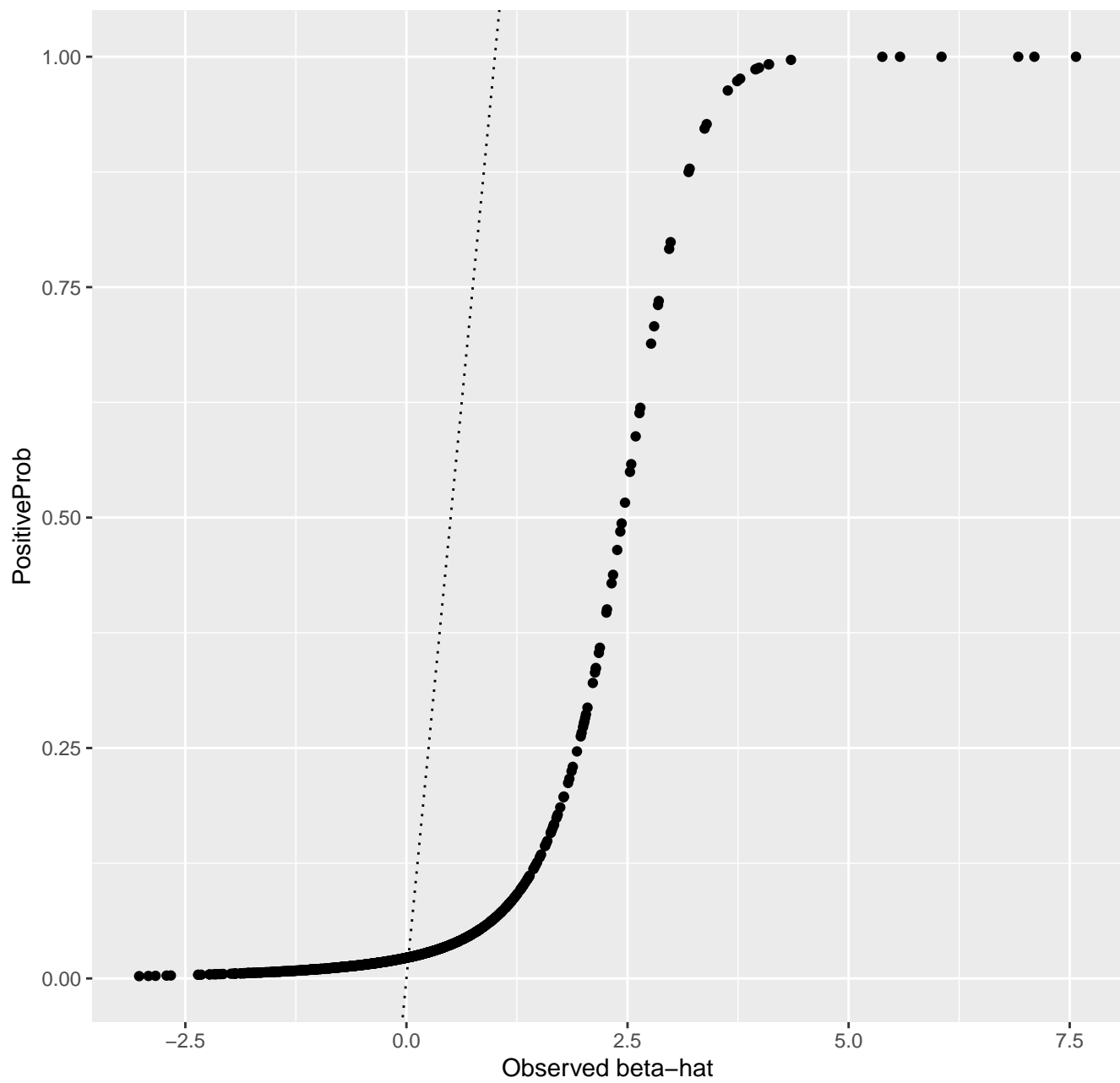
RELB_(5971) negative



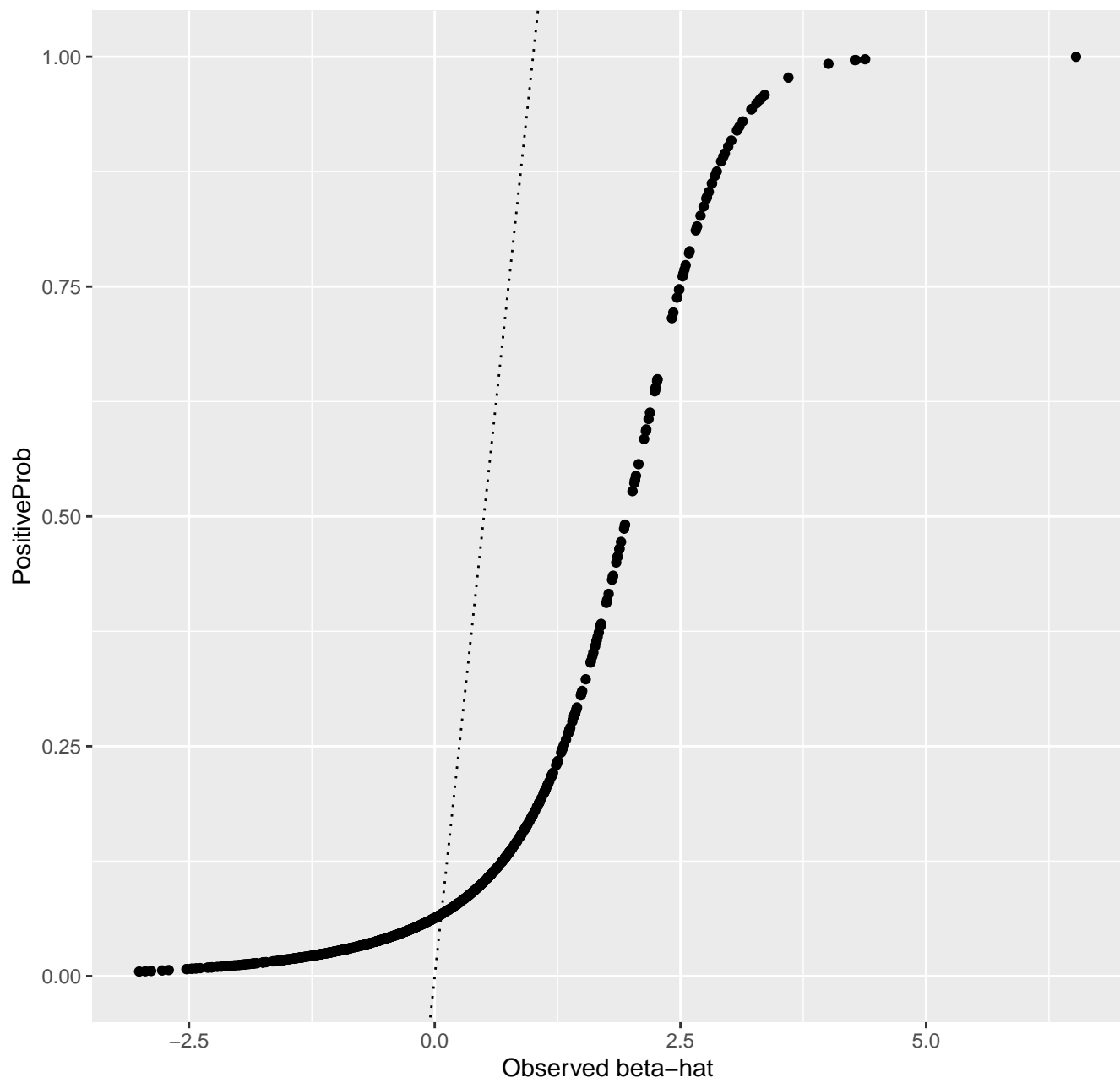
IGHMBP2_(3508) negative



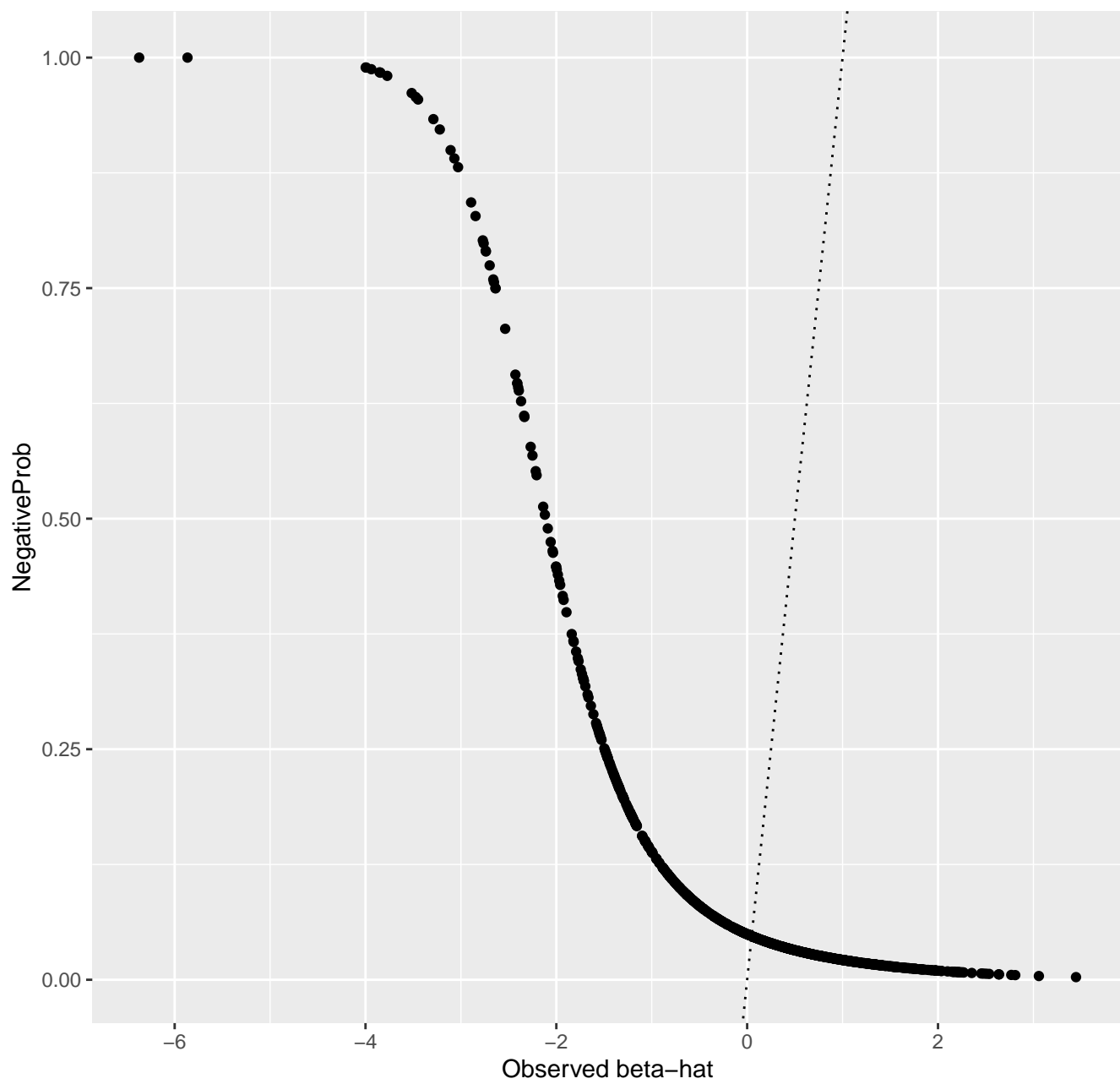
CHEK2_(11200) positive



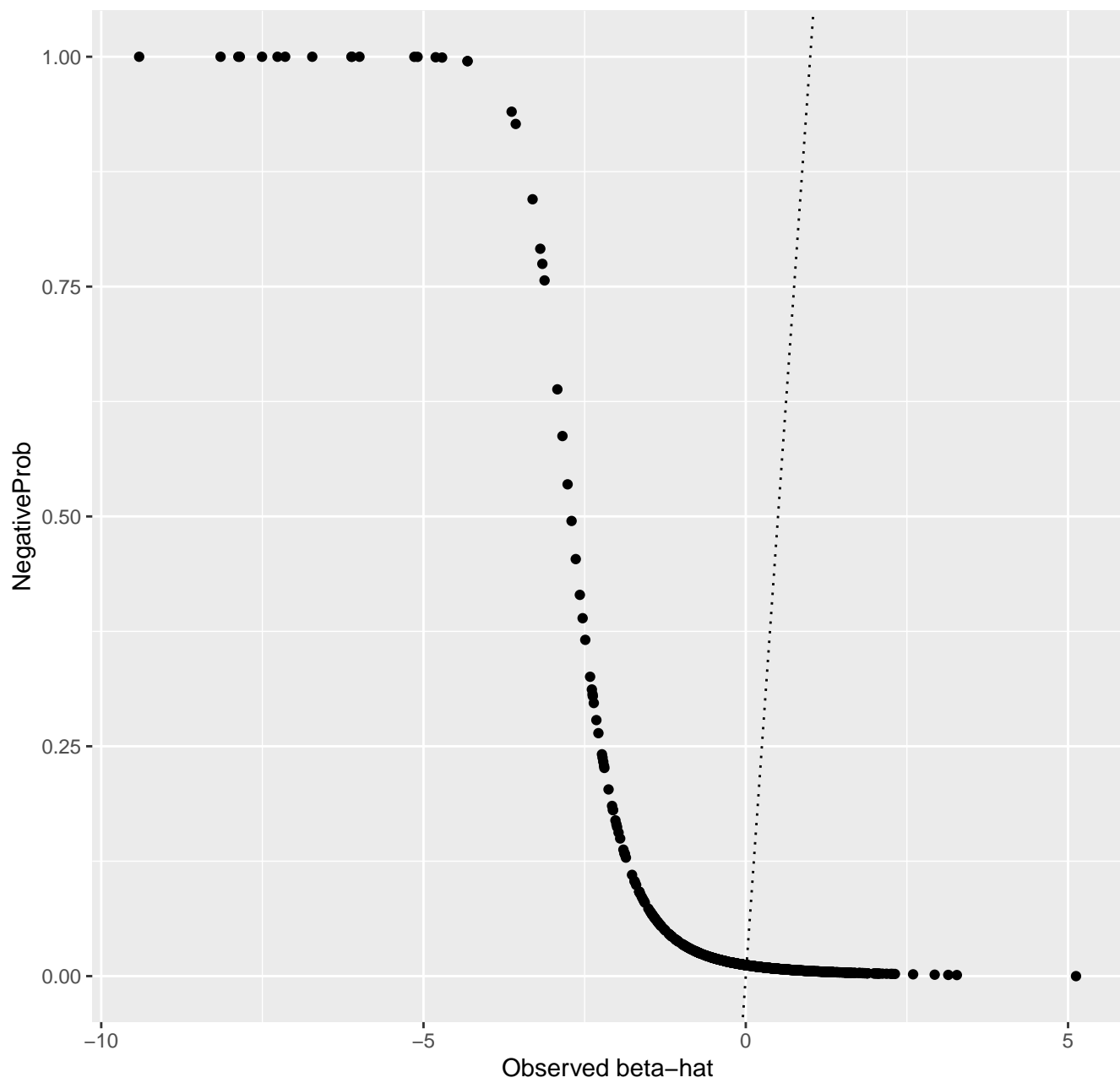
PAWR_(5074) positive



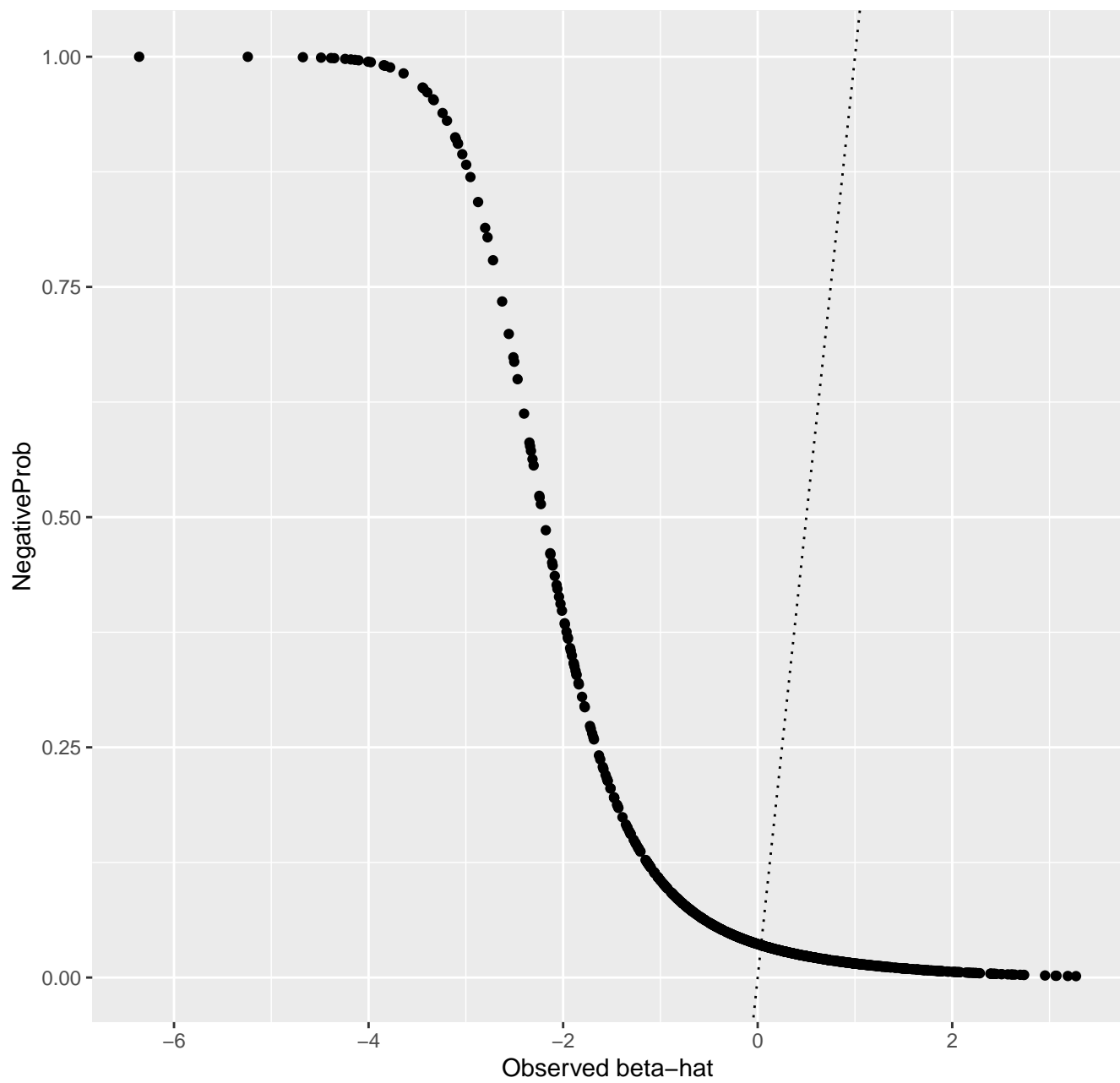
B3GALT6_(126792) negative



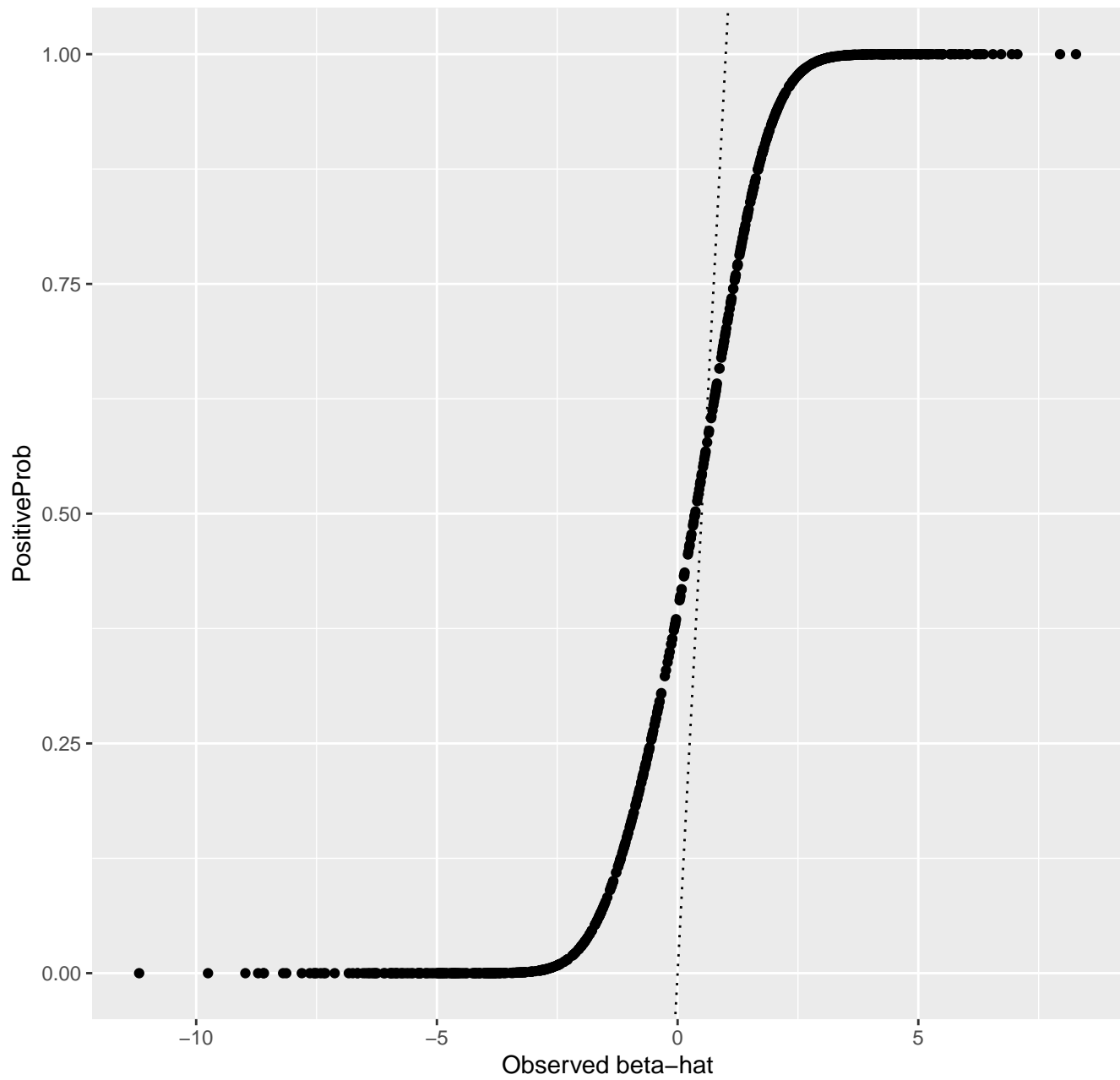
MECOM_(2122) negative



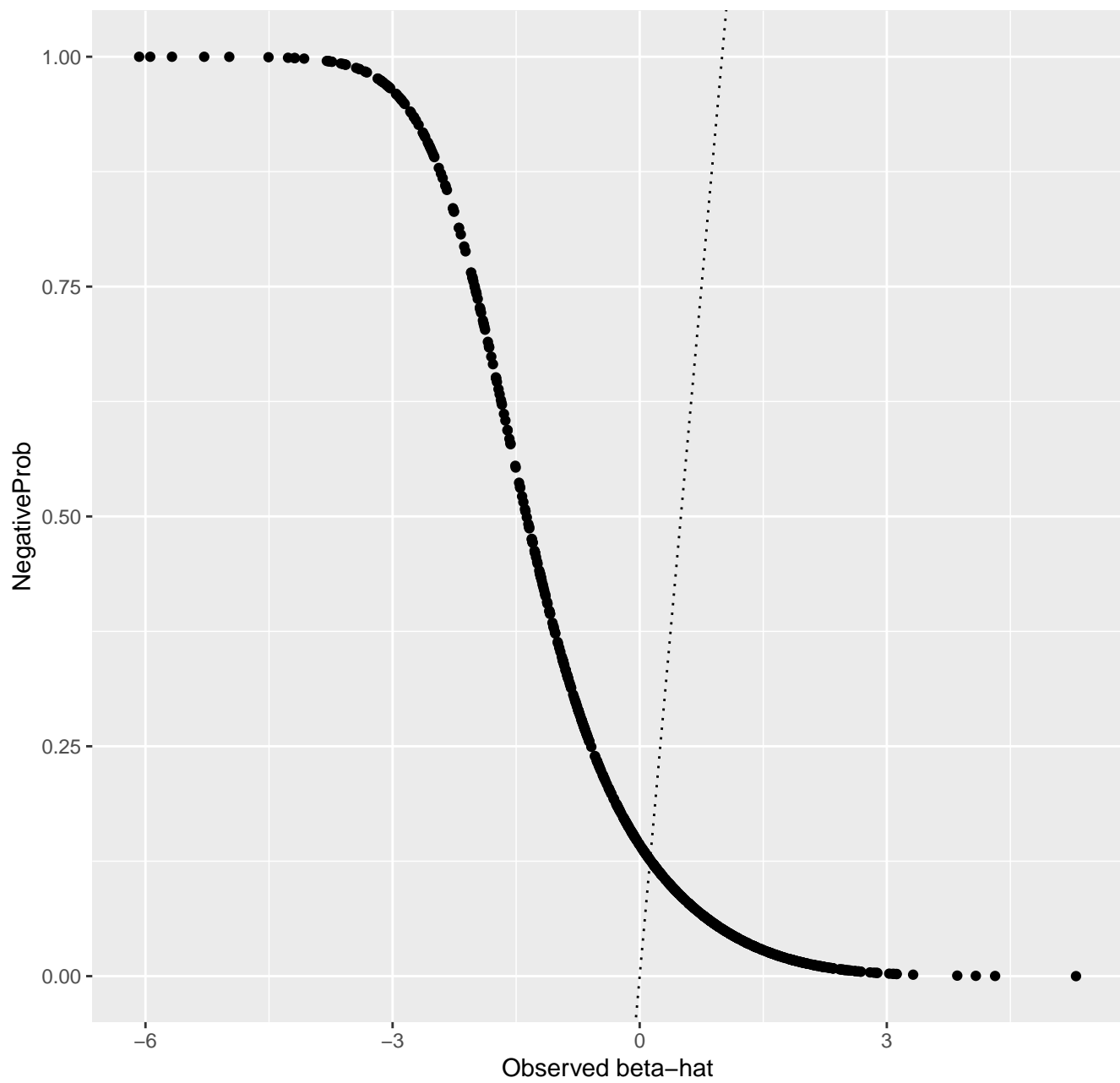
GNPNAT1_(64841) negative



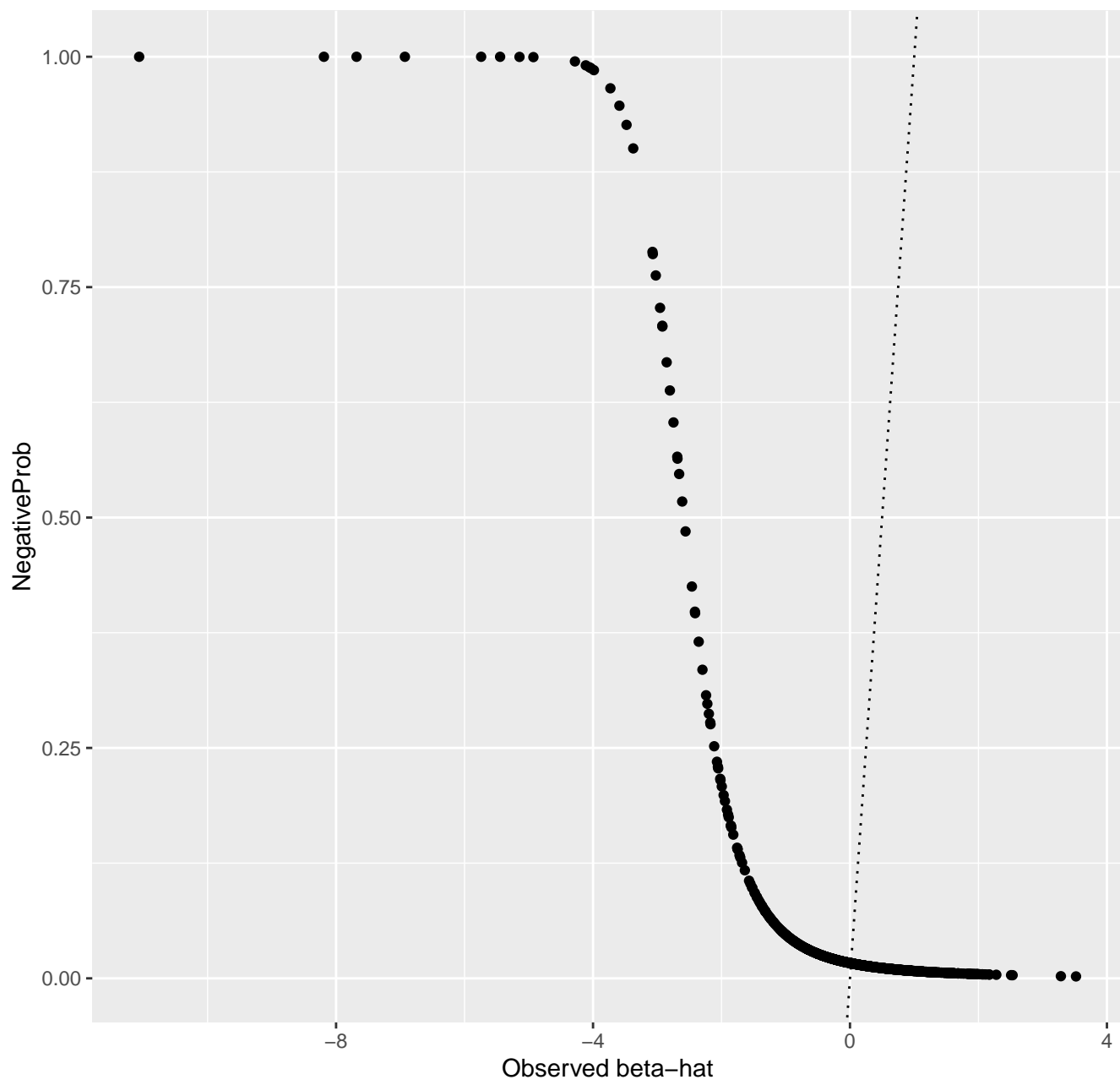
ITGAV_(3685) positive



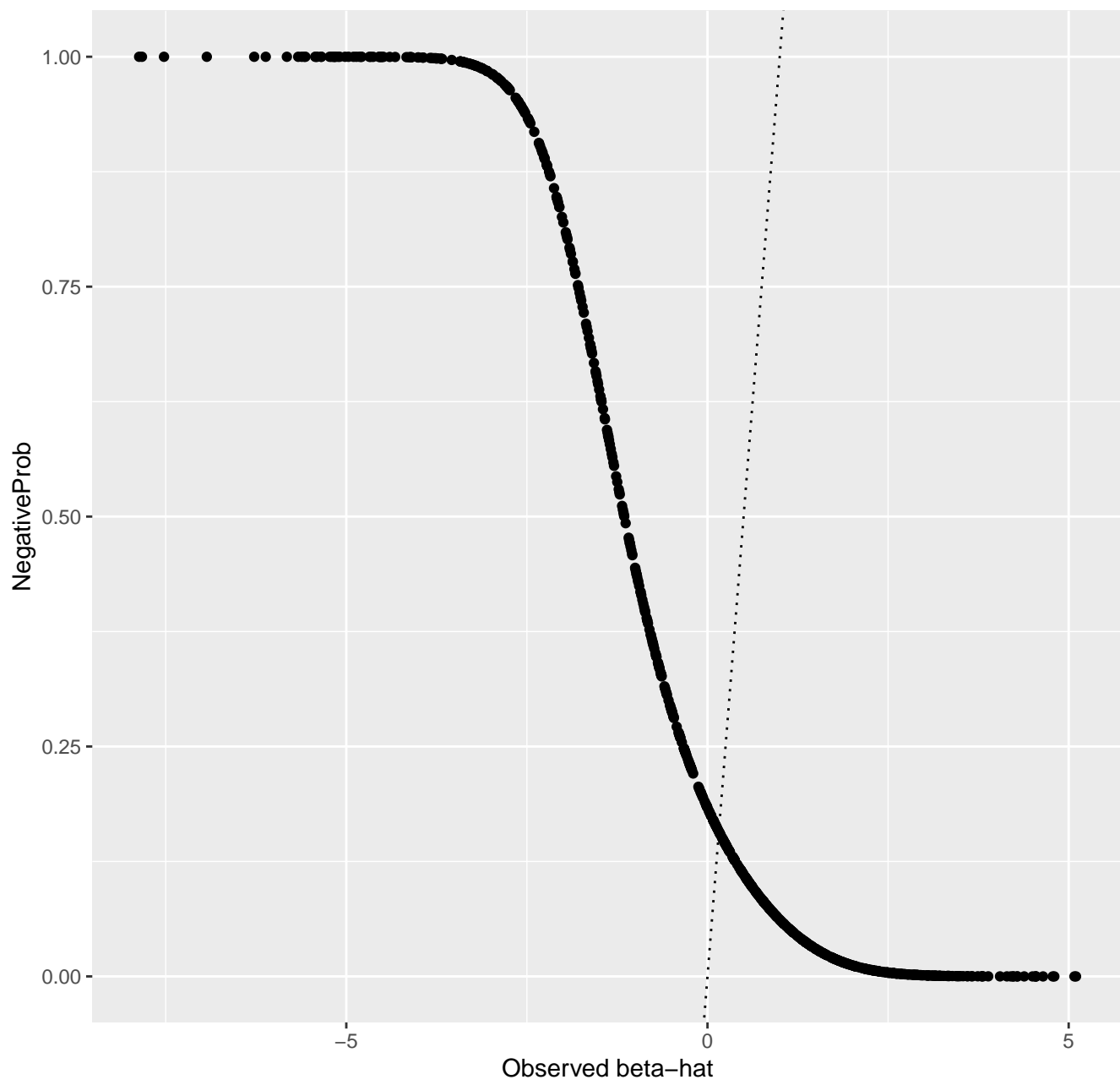
KIF2C_(11004) negative



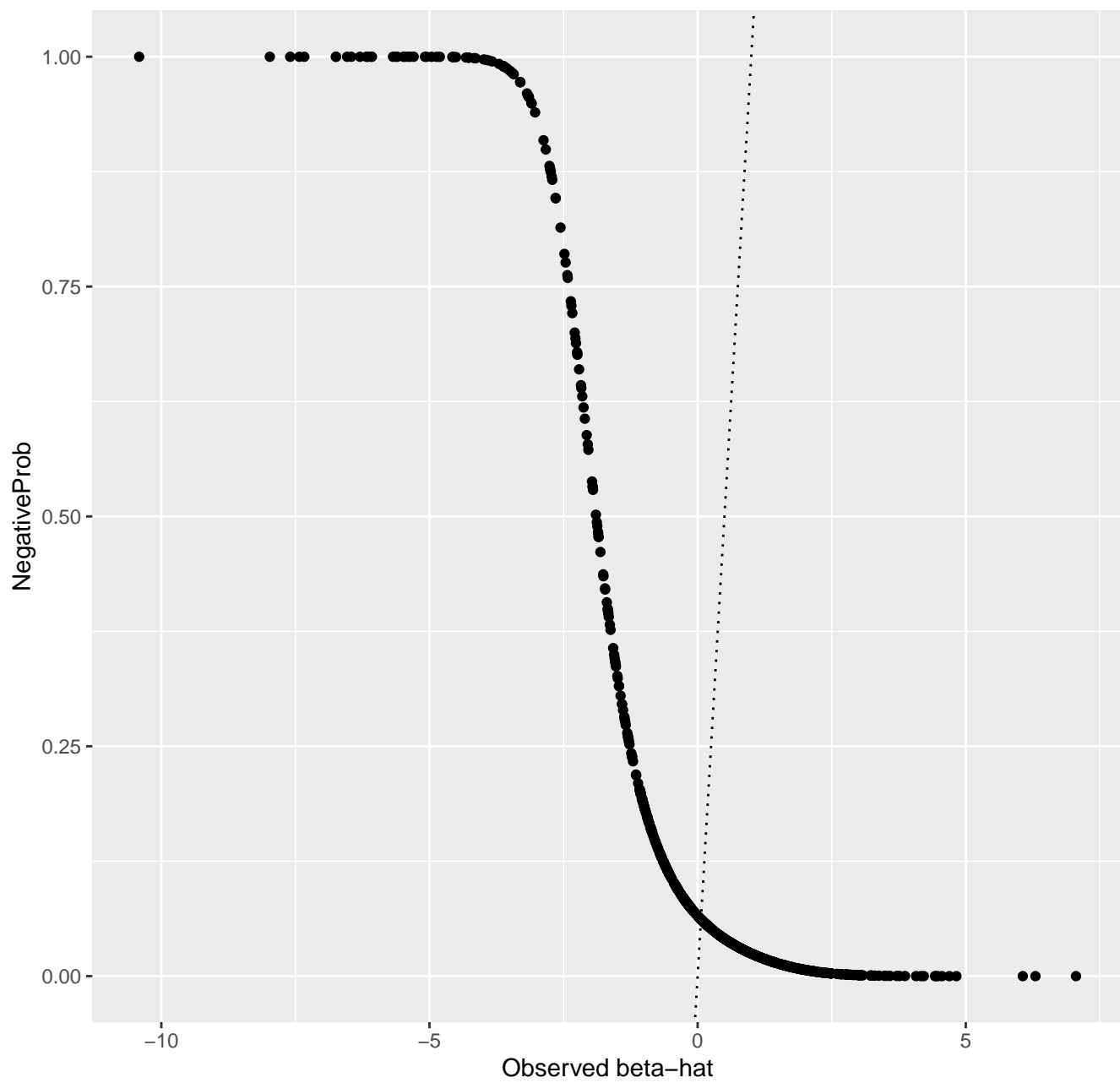
THAP7_(80764) negative



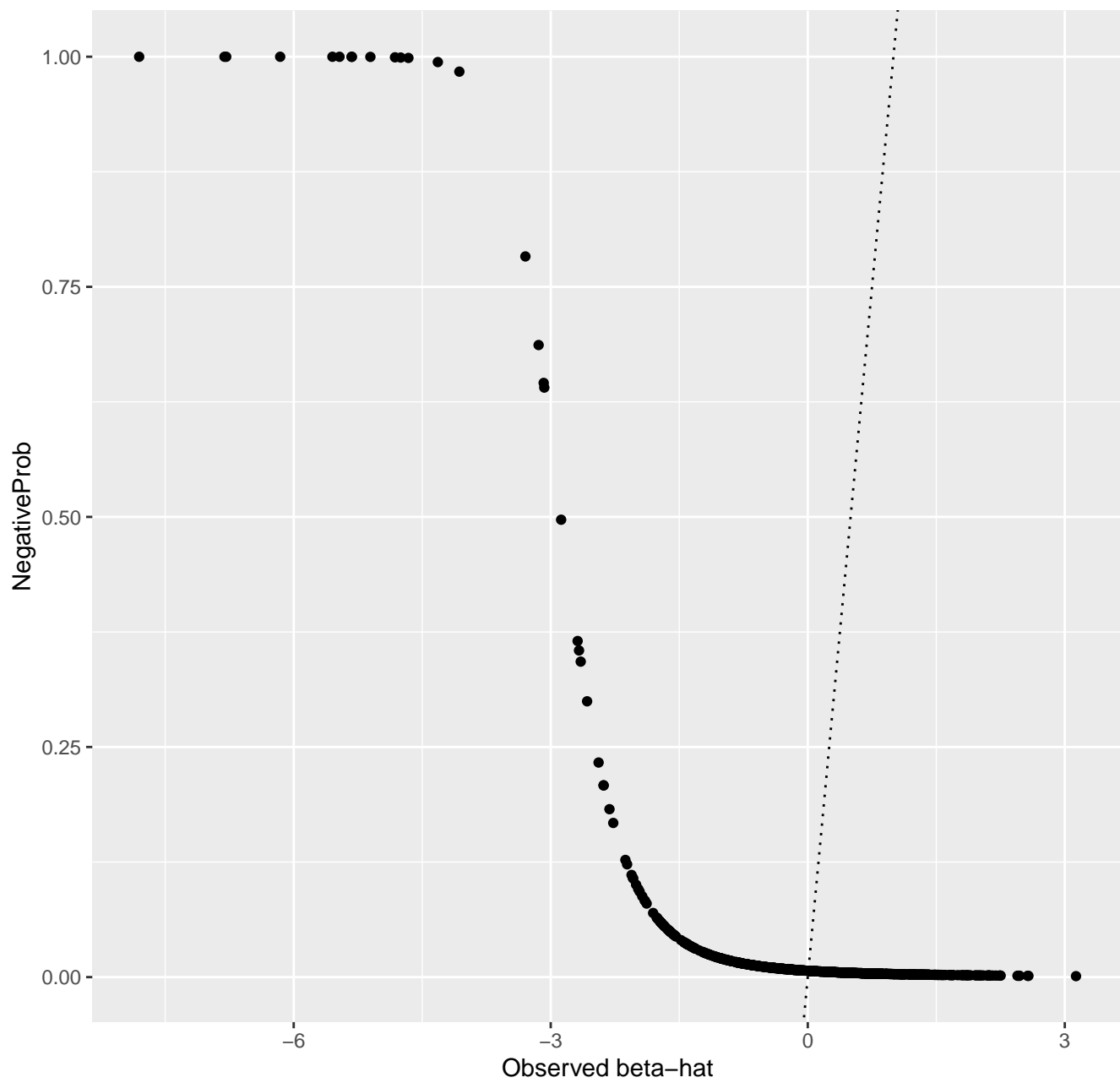
ICE2_(79664) negative



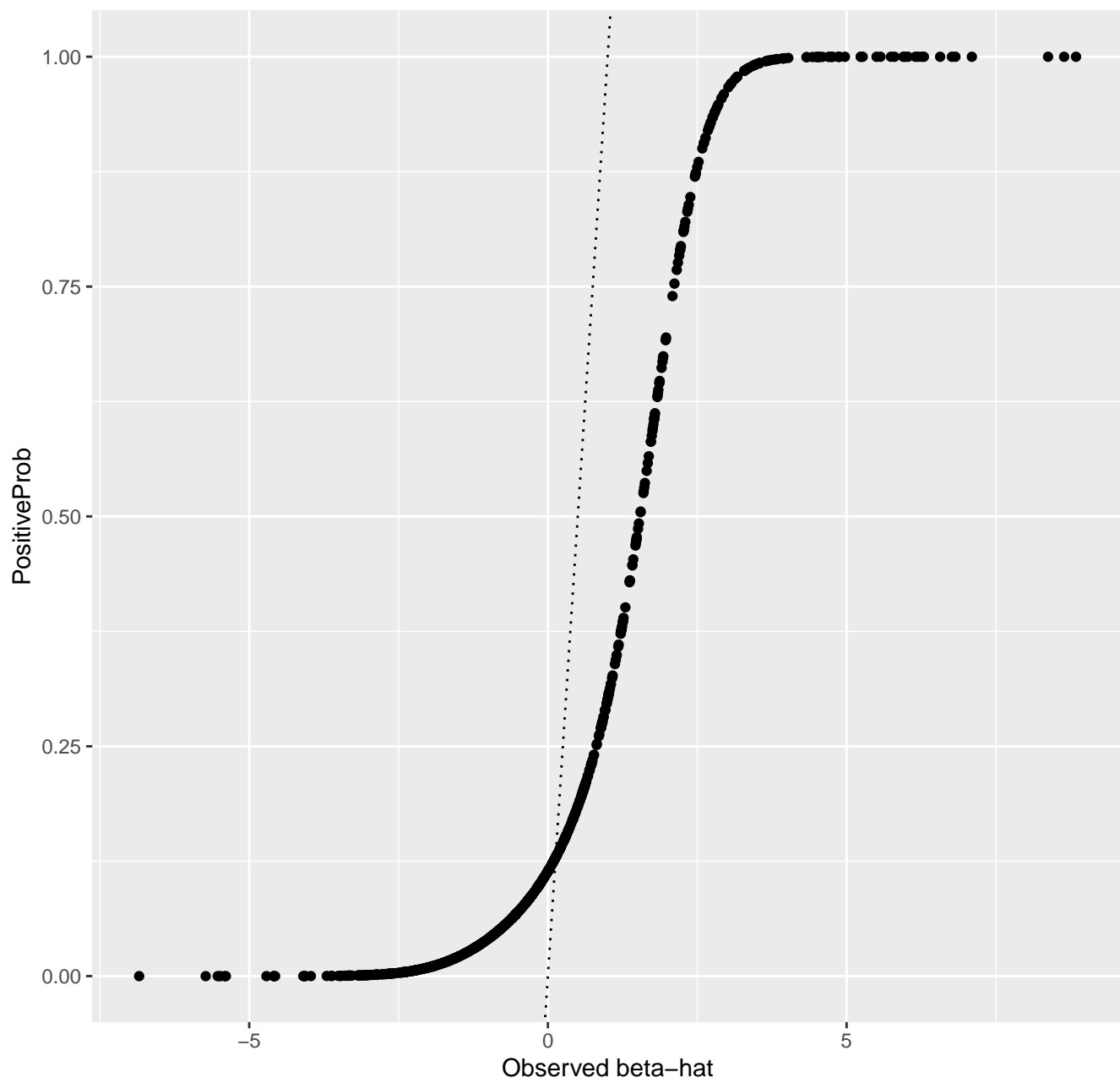
RAB35_(11021) negative



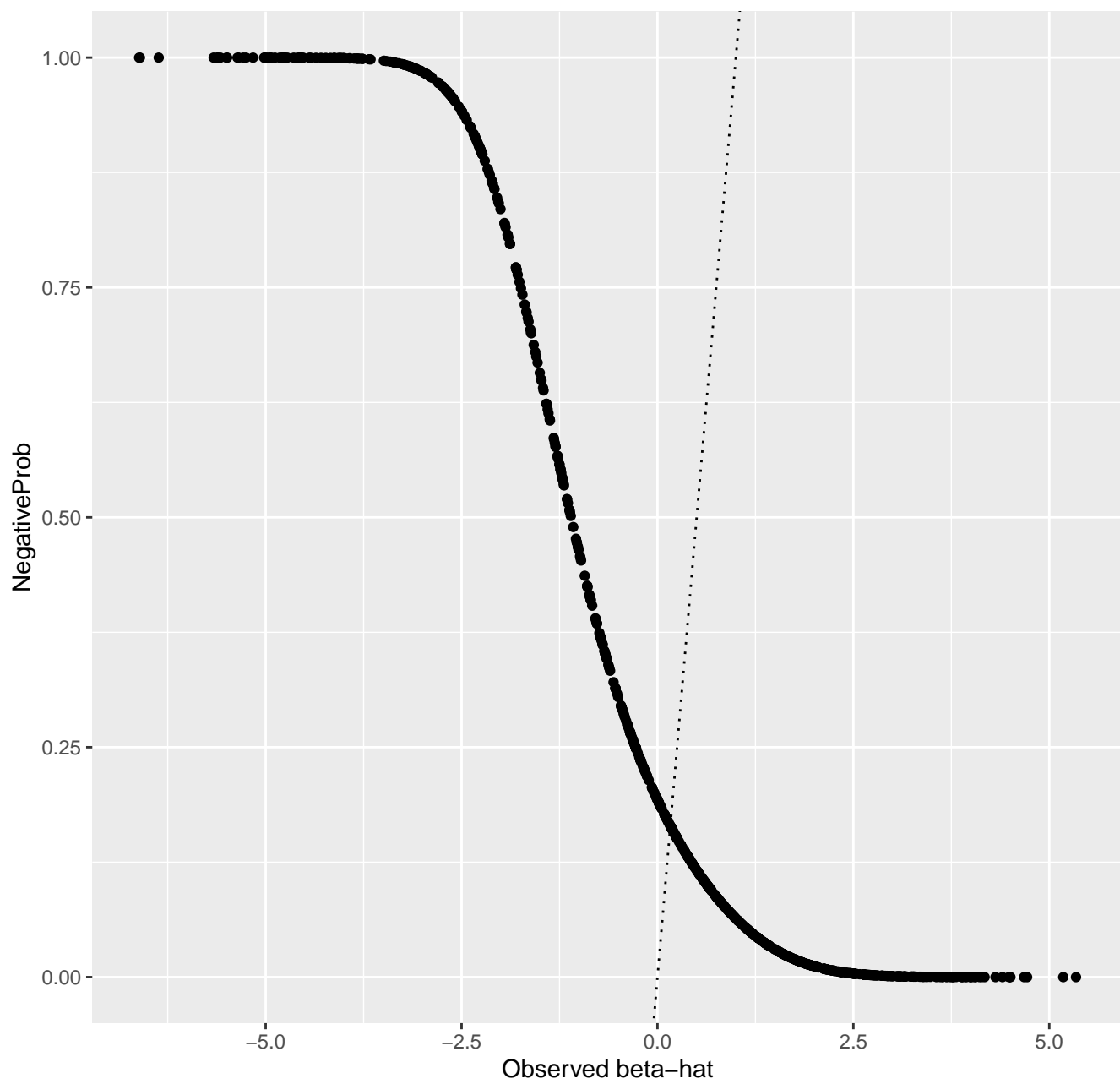
MAP2K1_(5604) negative



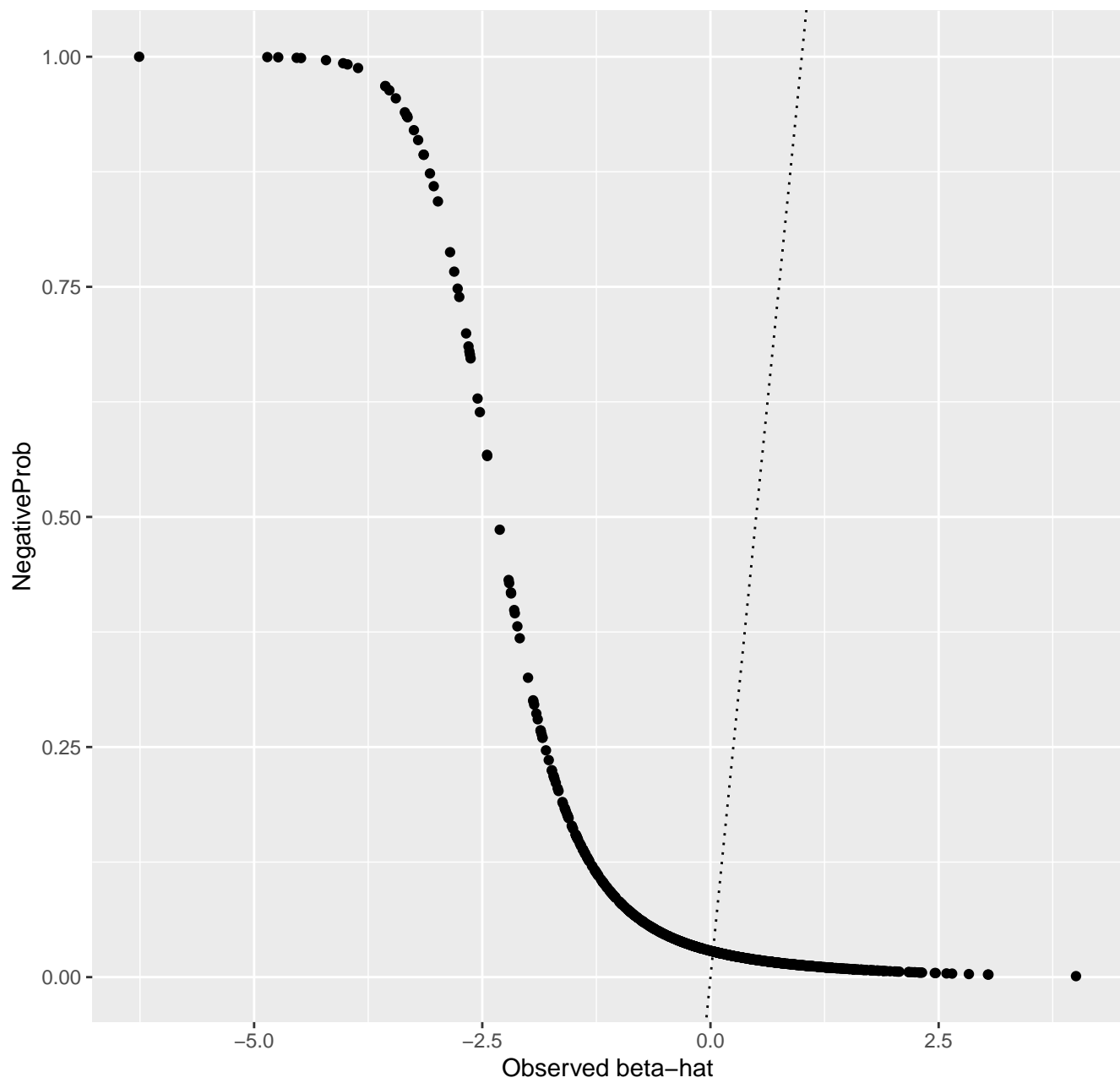
PKMYT1_(9088) positive



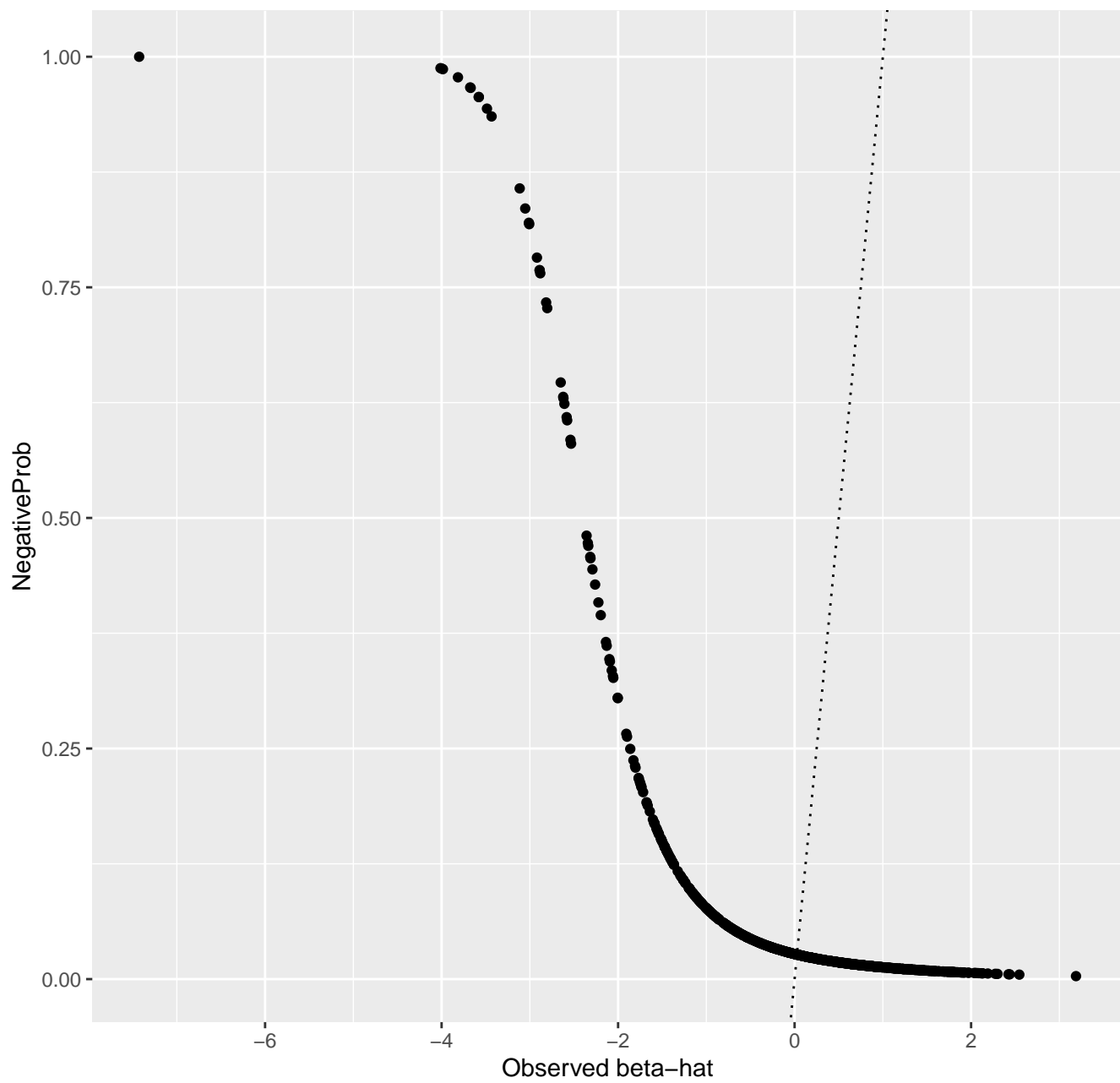
PTK2_(5747) negative



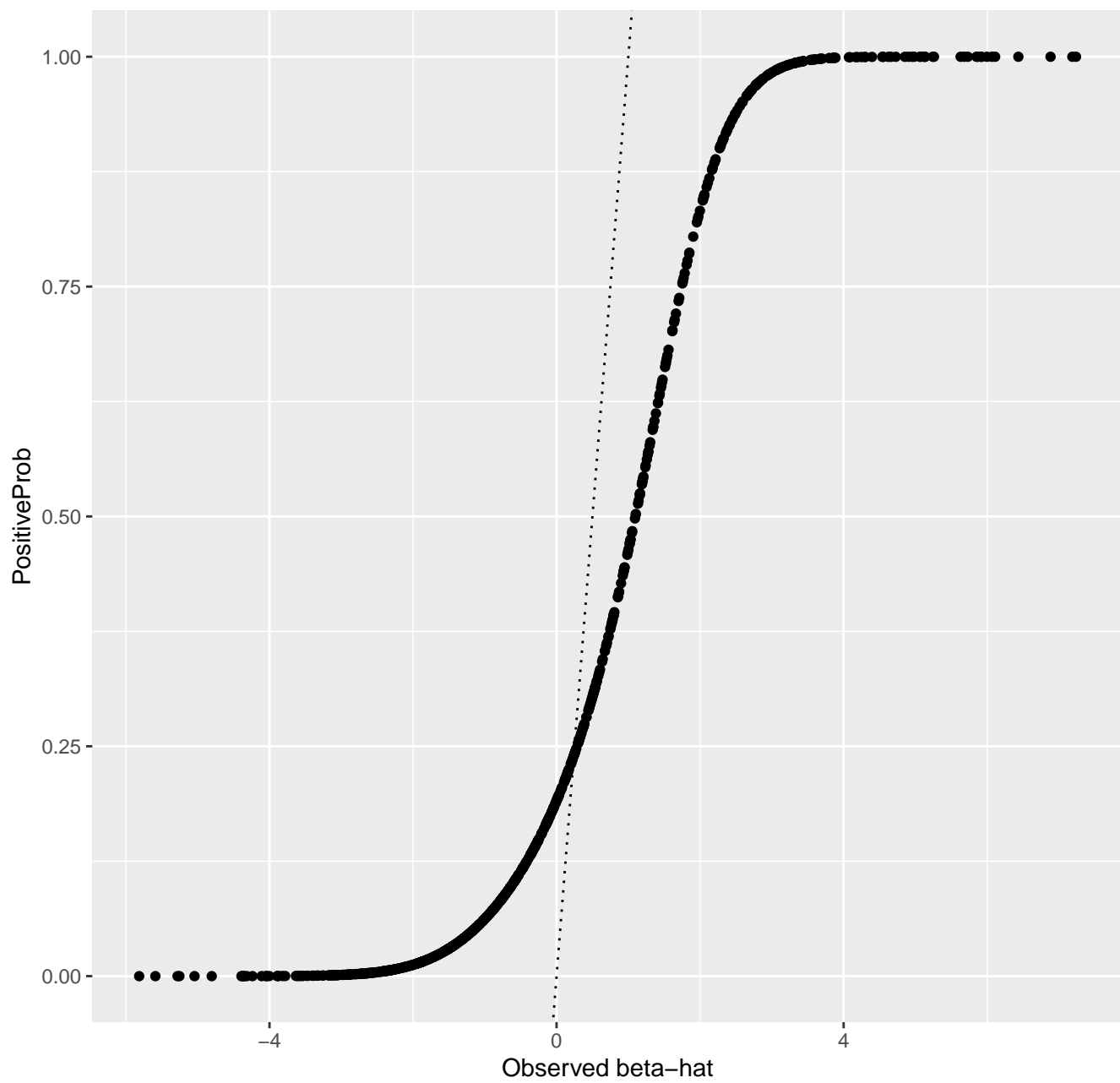
DIAPH3_(81624) negative



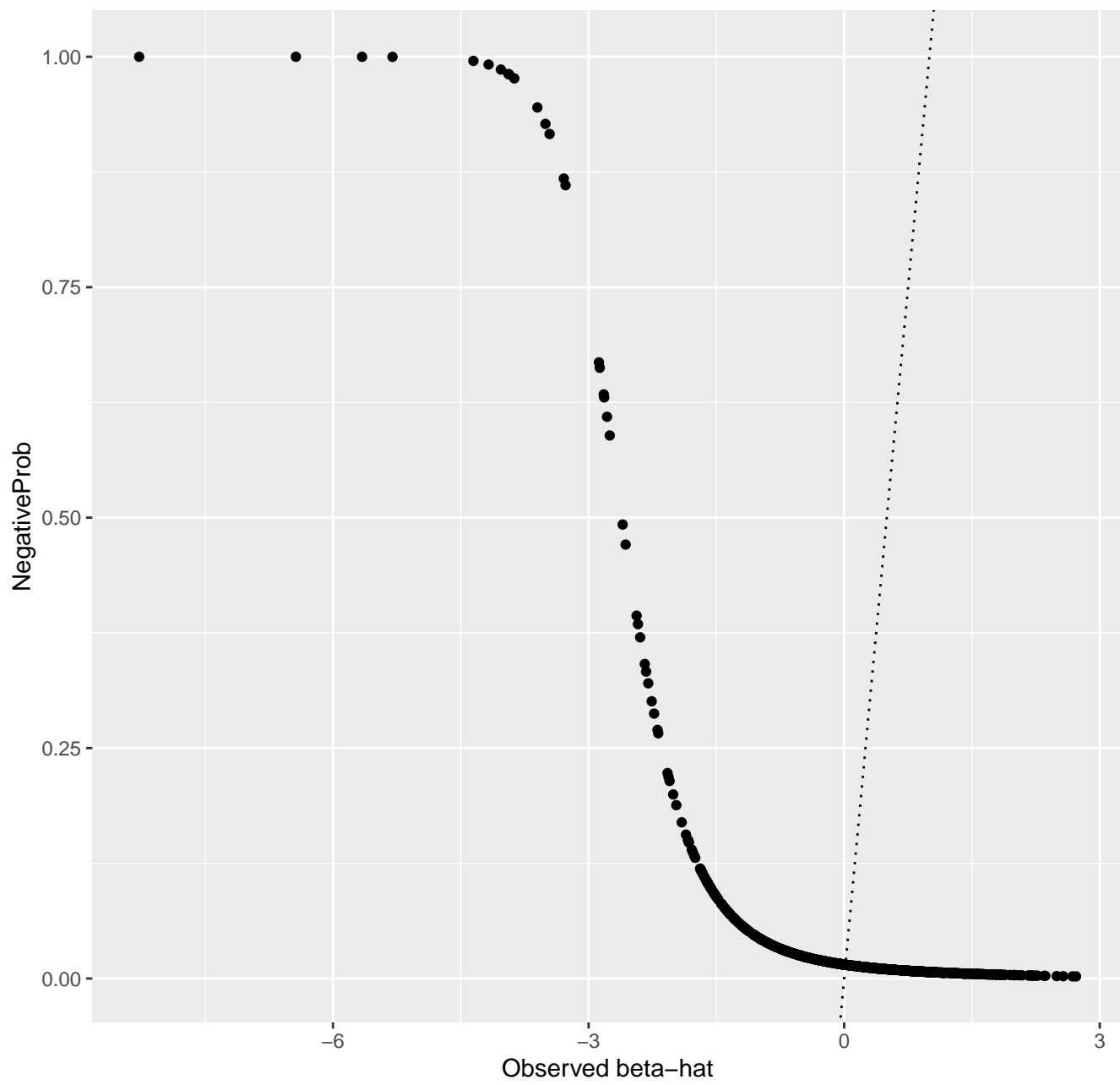
SLC31A1_(1317) negative



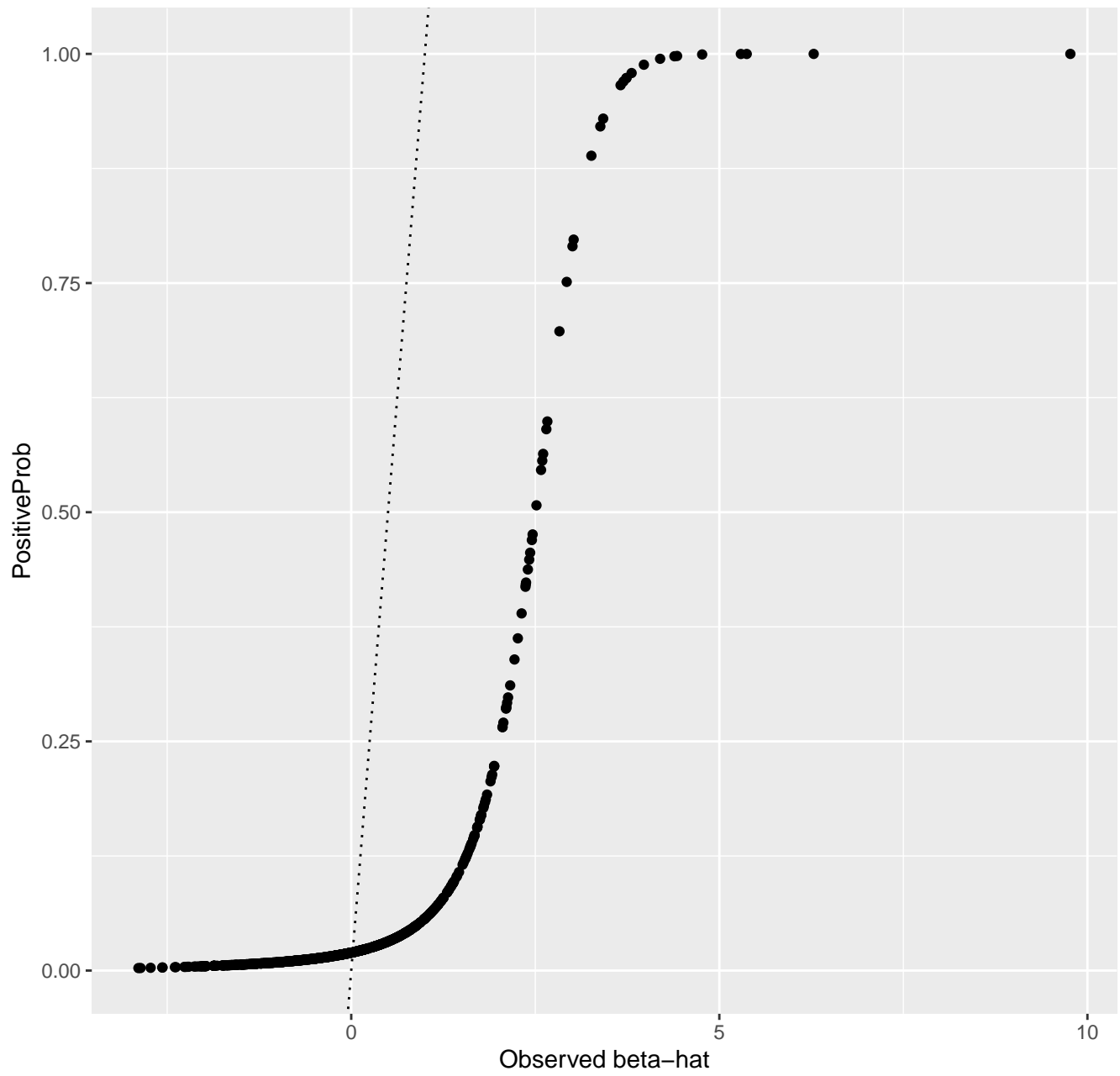
PCYT1A_(5130) positive



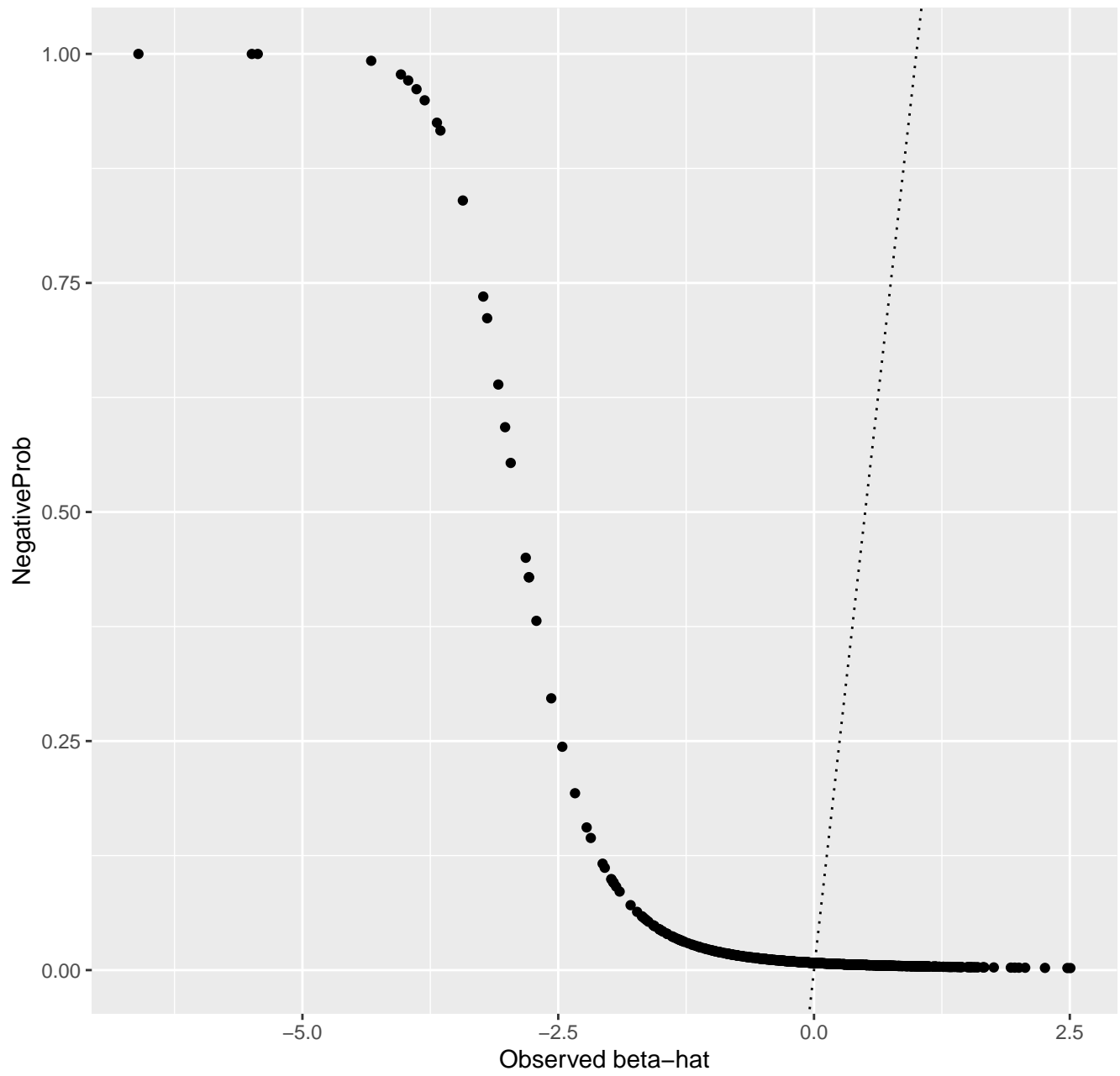
MUS81_(80198) negative



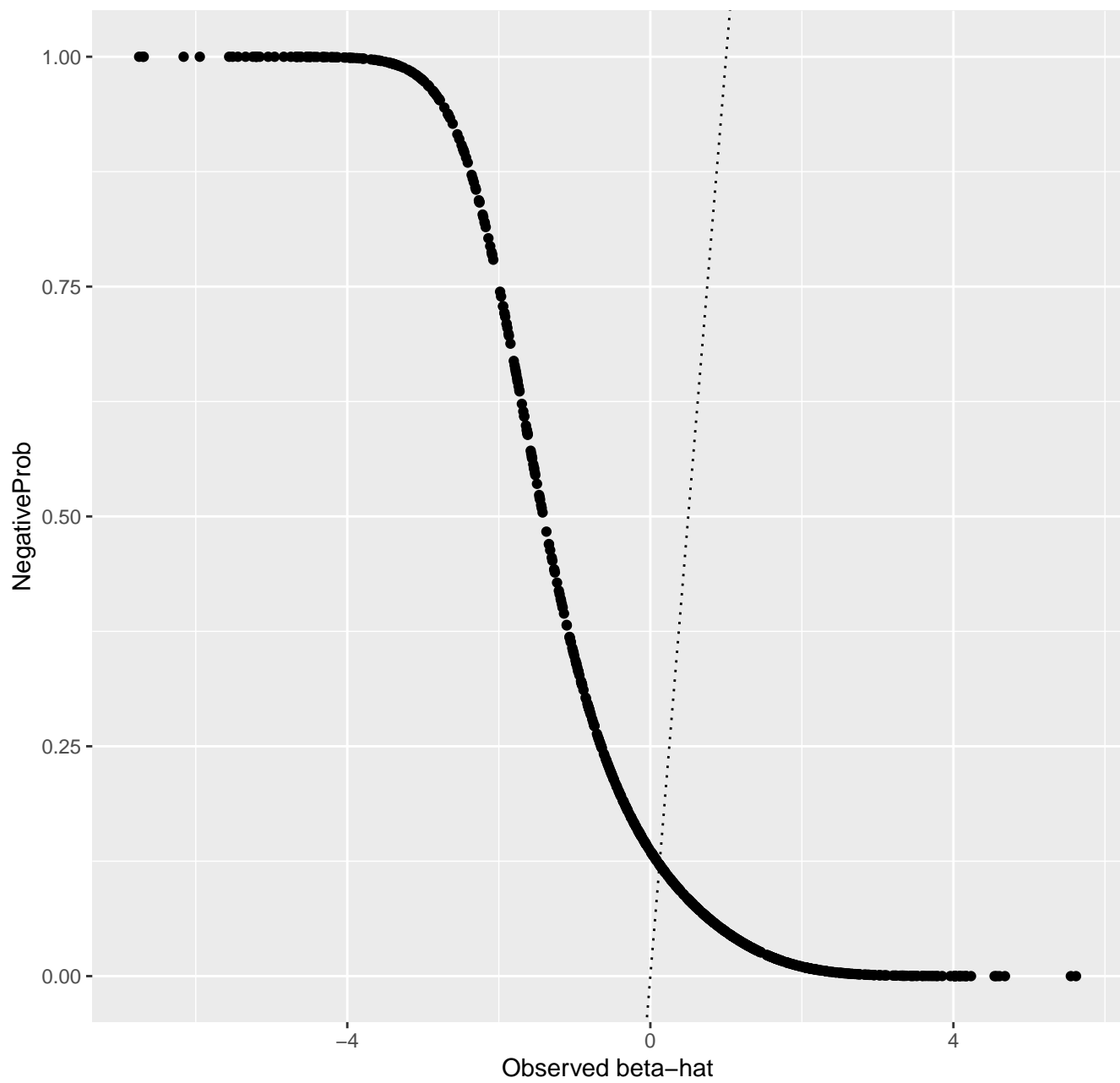
PTPN14_(5784) positive



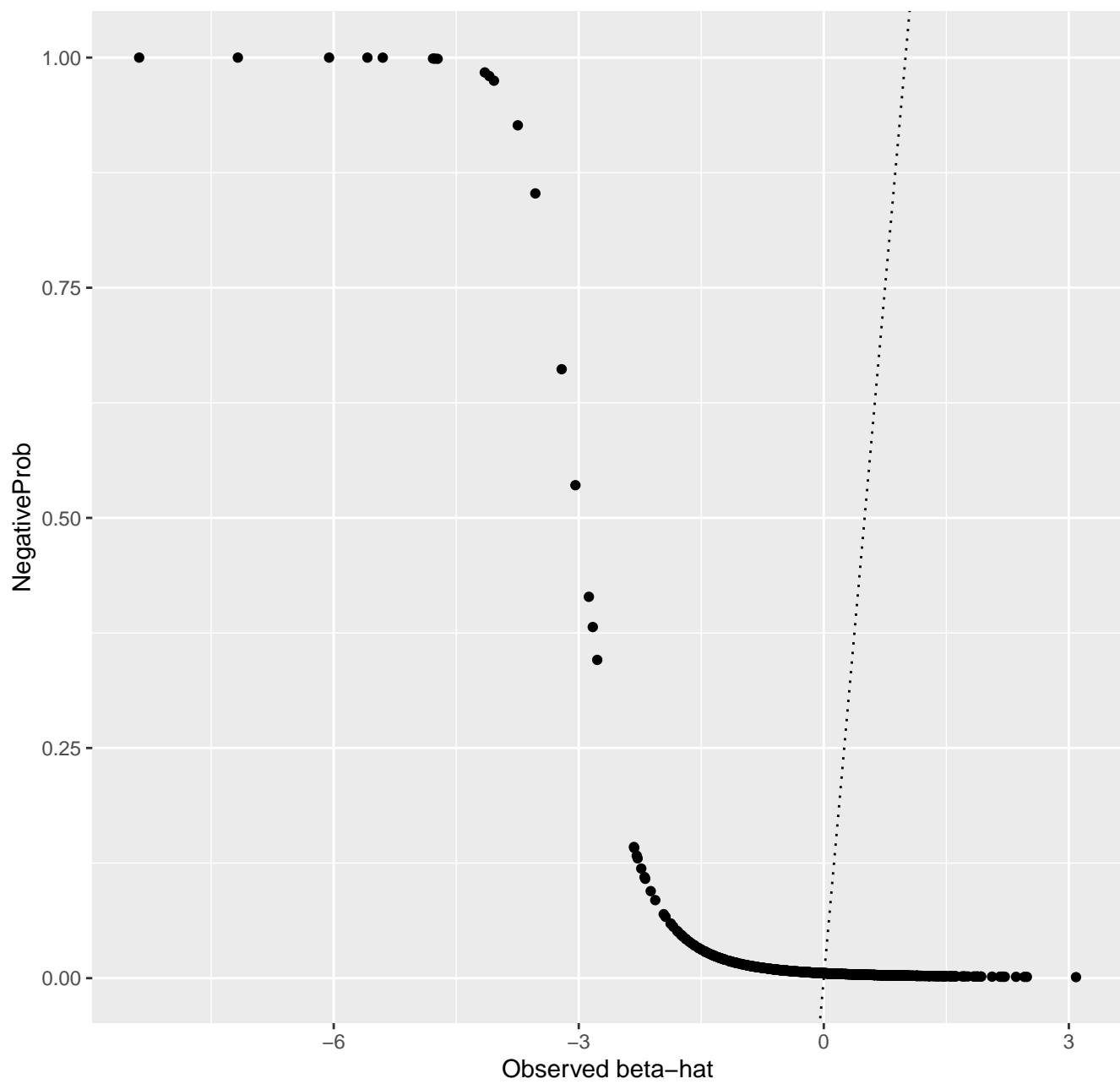
IGF2BP1_(10642) negative



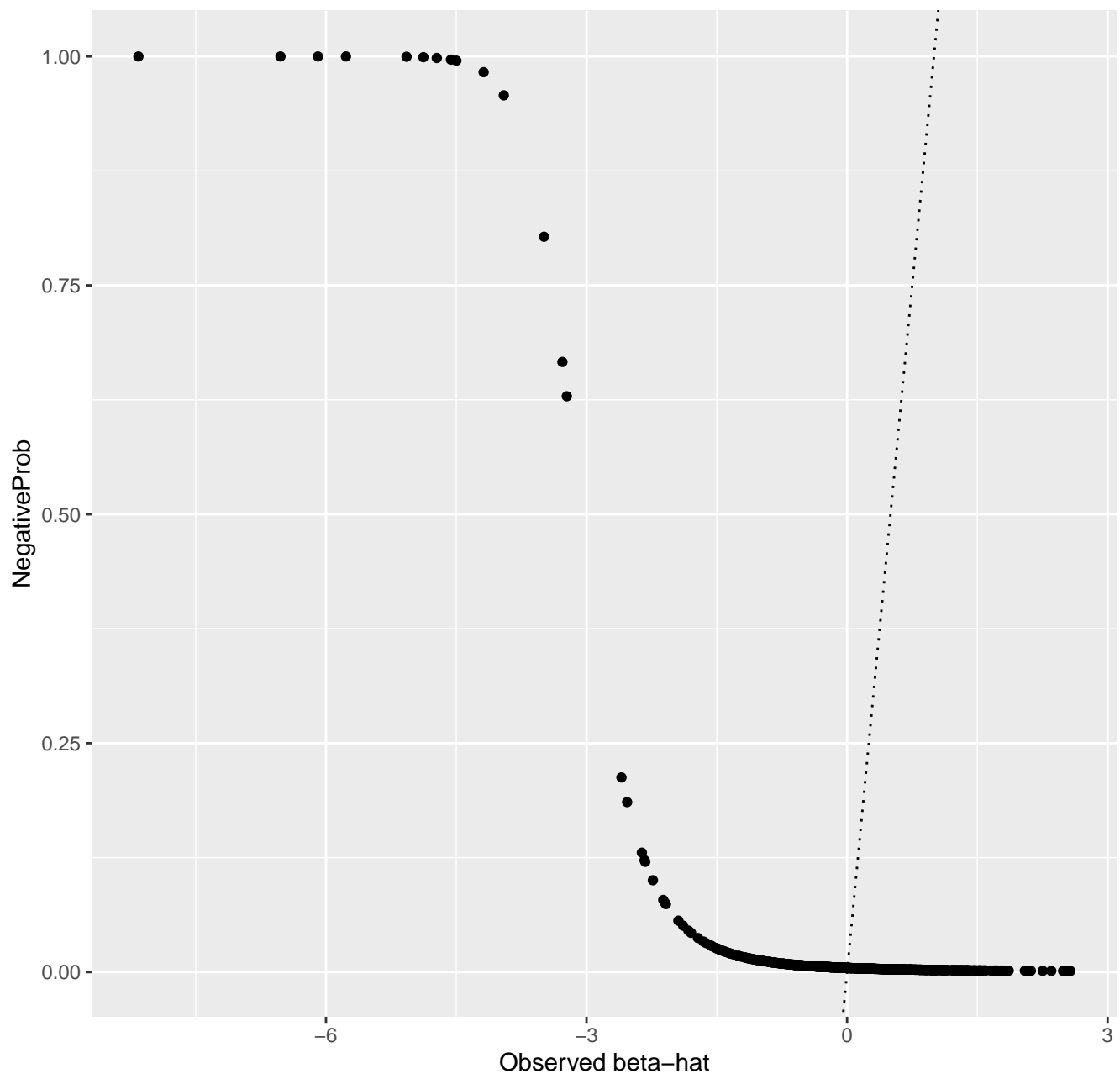
GART_(2618) negative



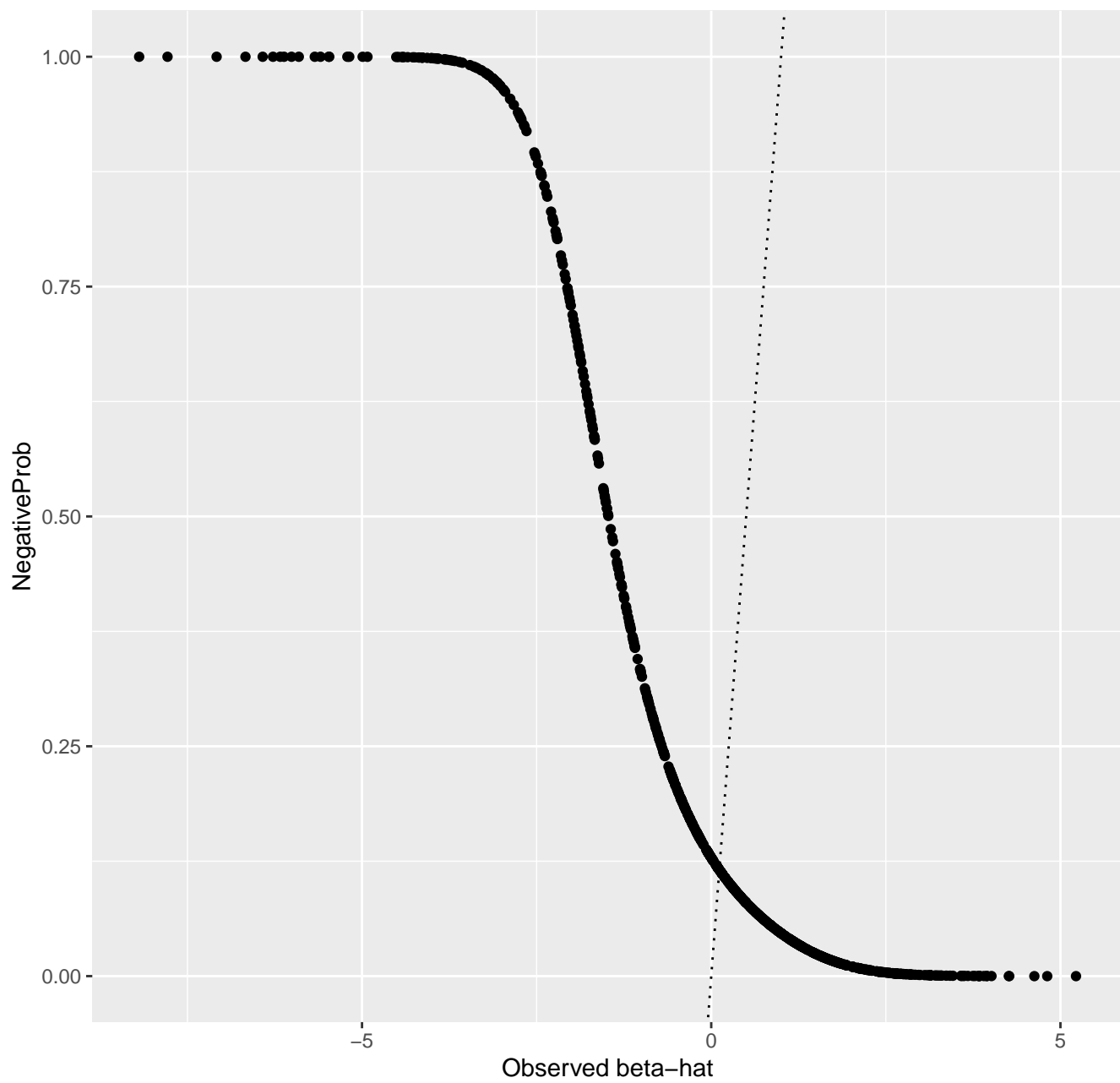
KDELR2_(11014) negative



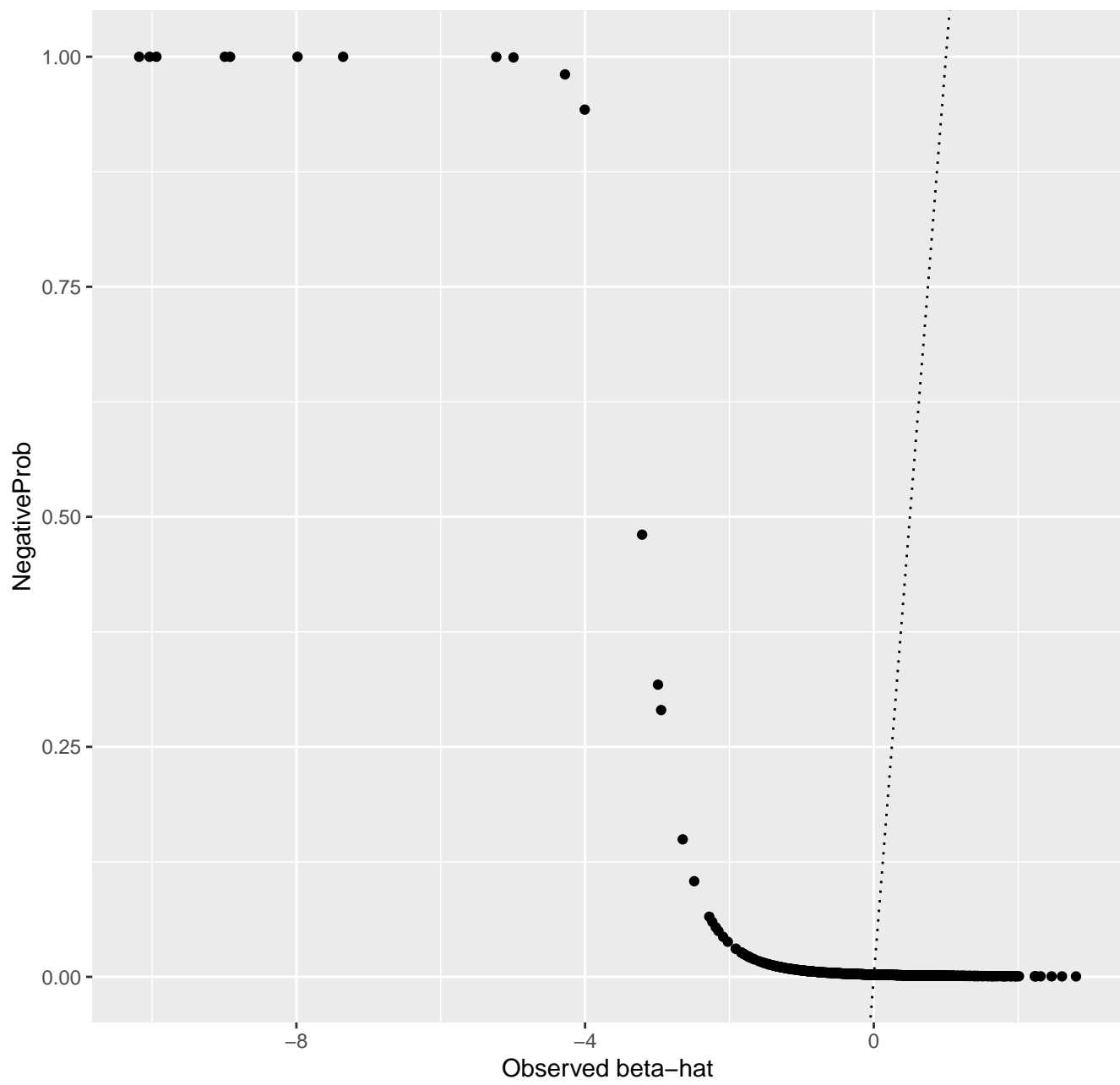
ZFPM1_(161882) negative



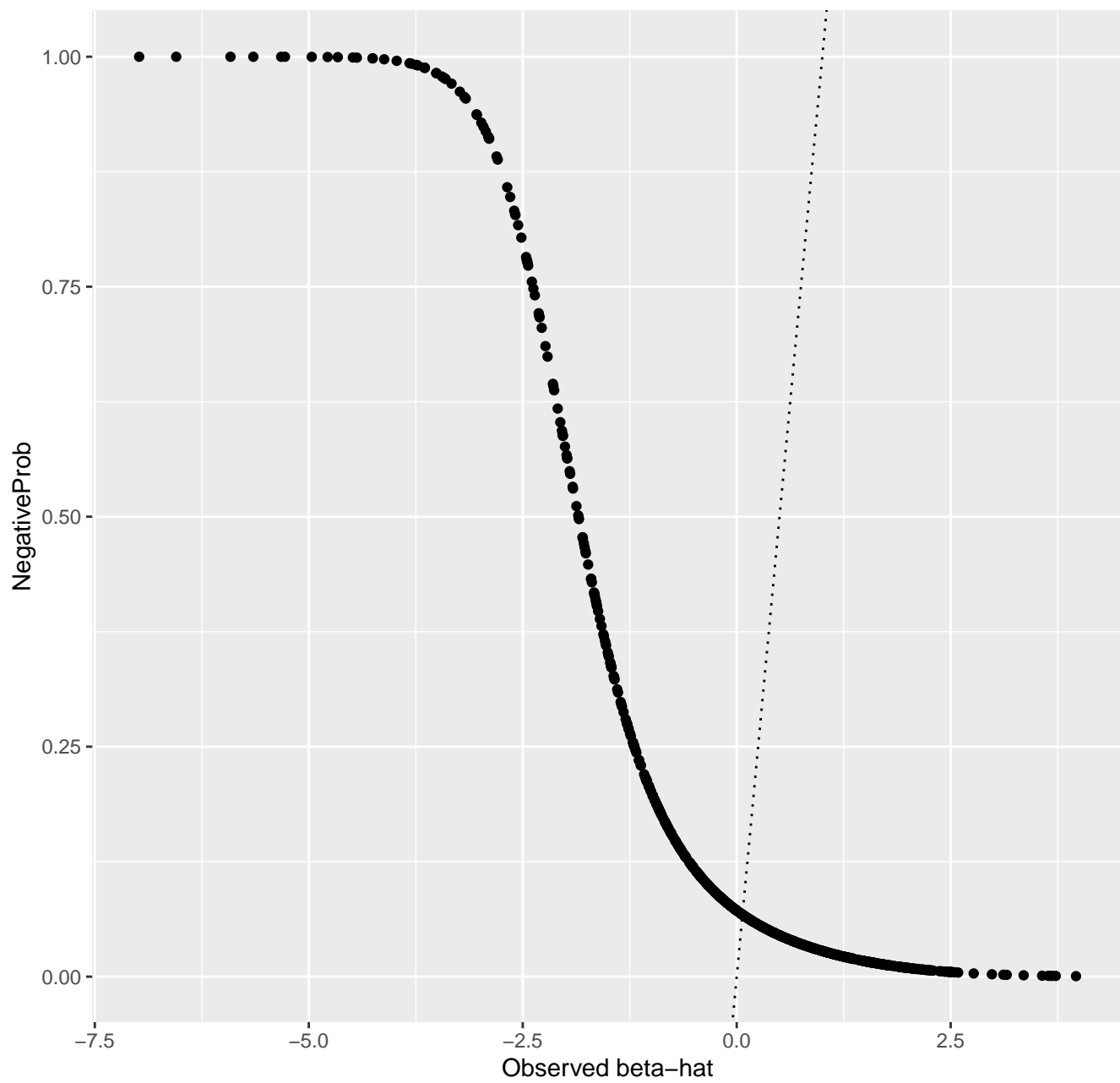
NDUFB10_(4716) negative



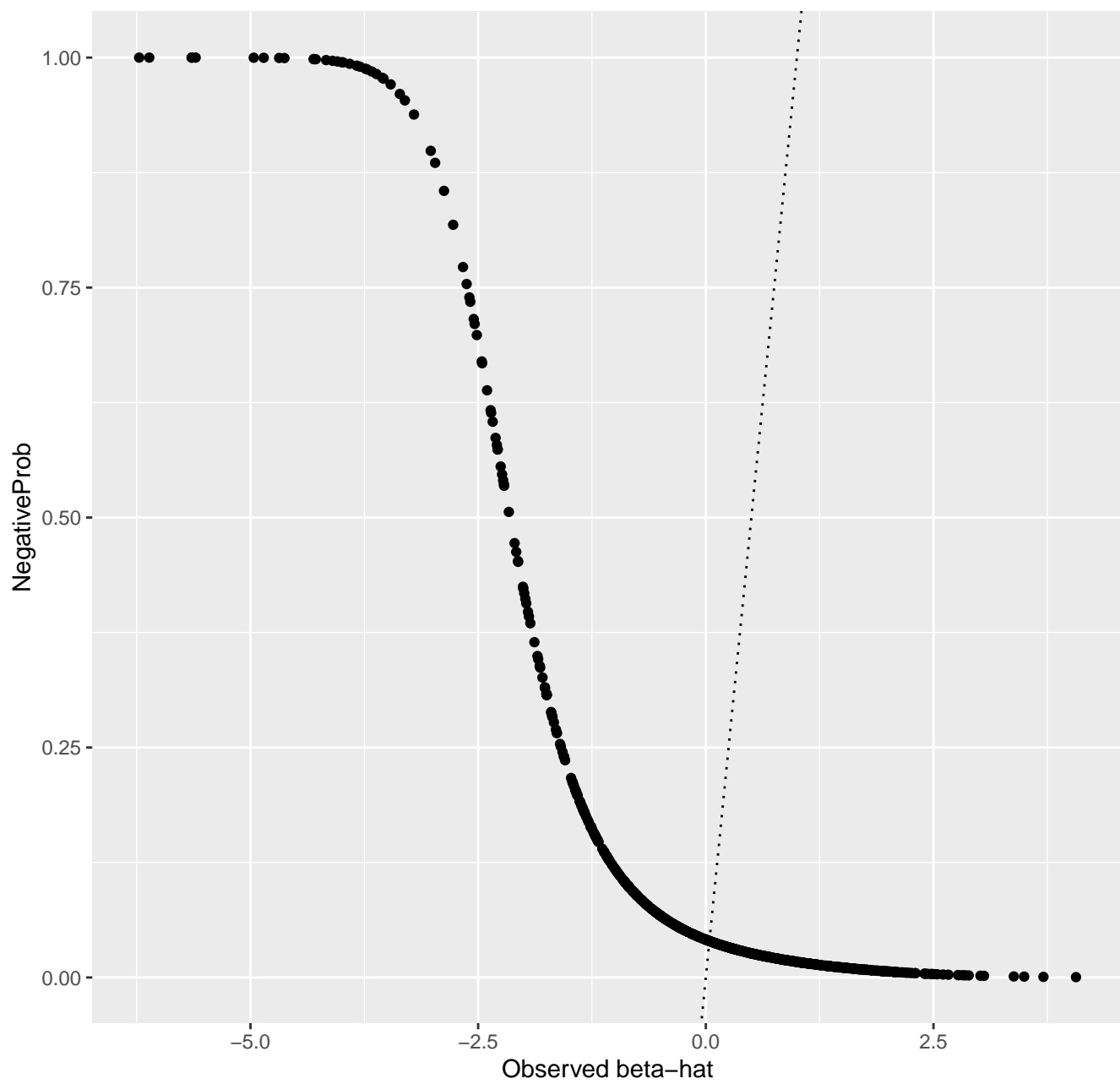
PAX3_(5077) negative



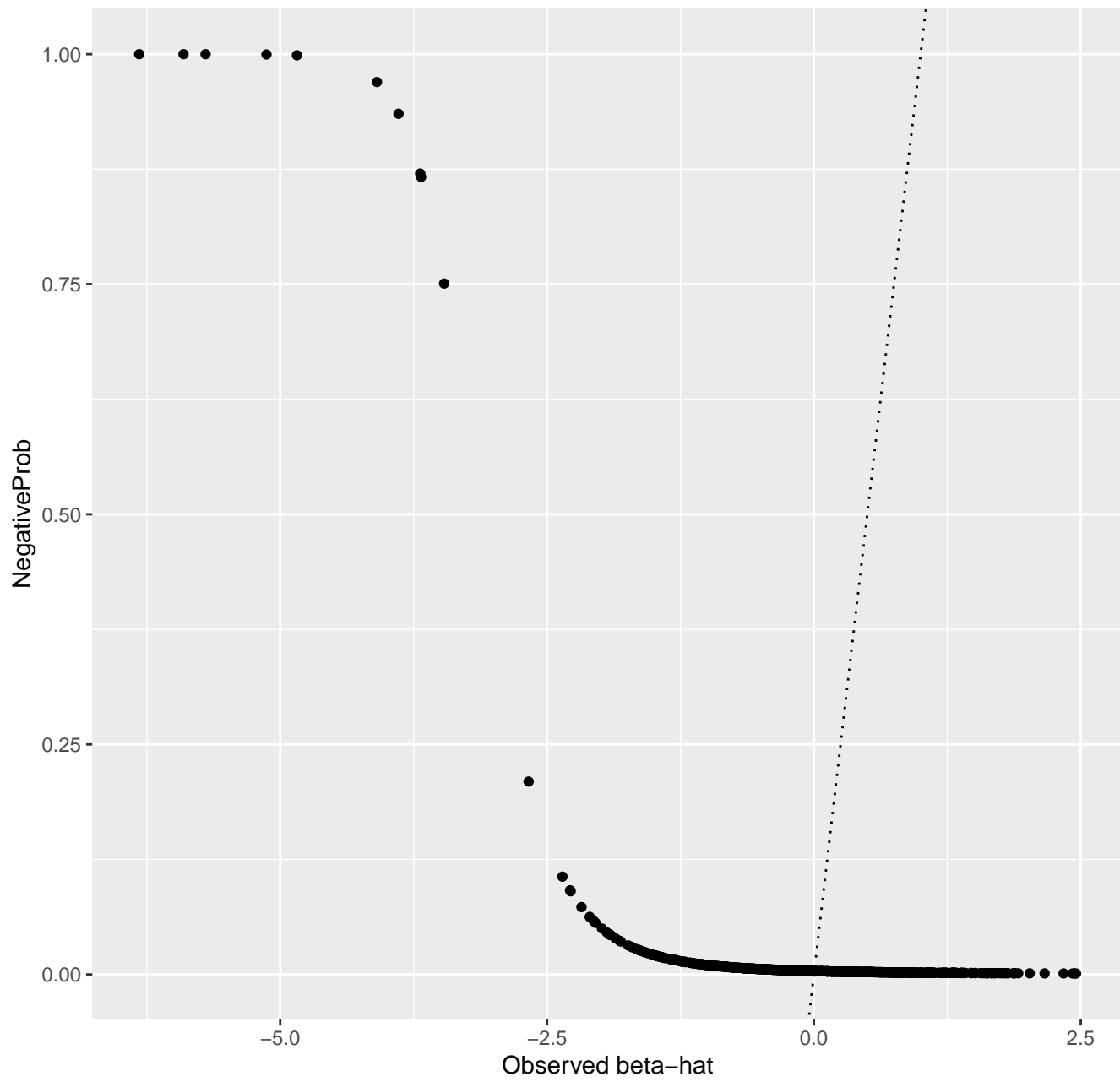
NOA1_(84273) negative



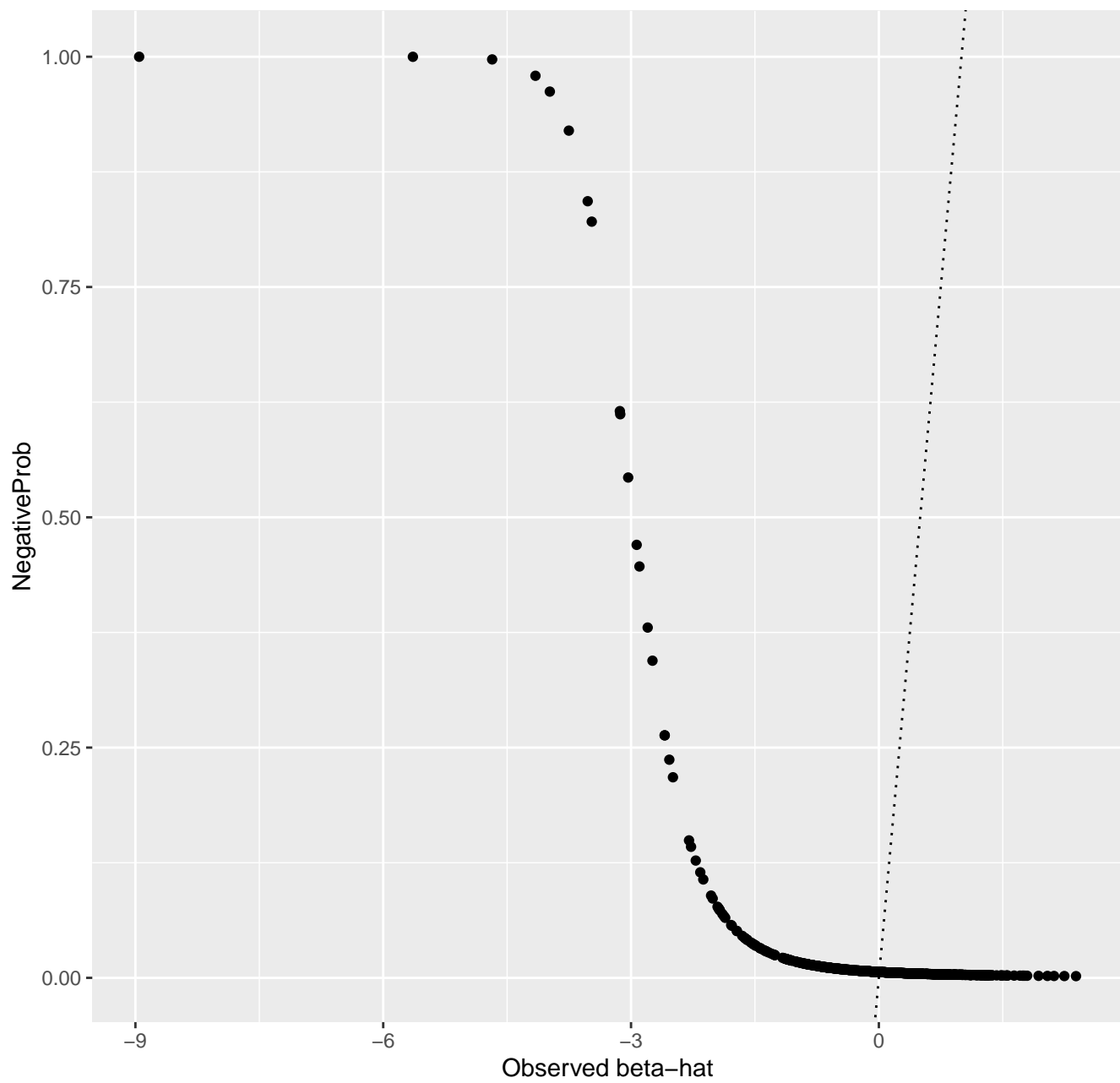
NDUFAF7_(55471) negative



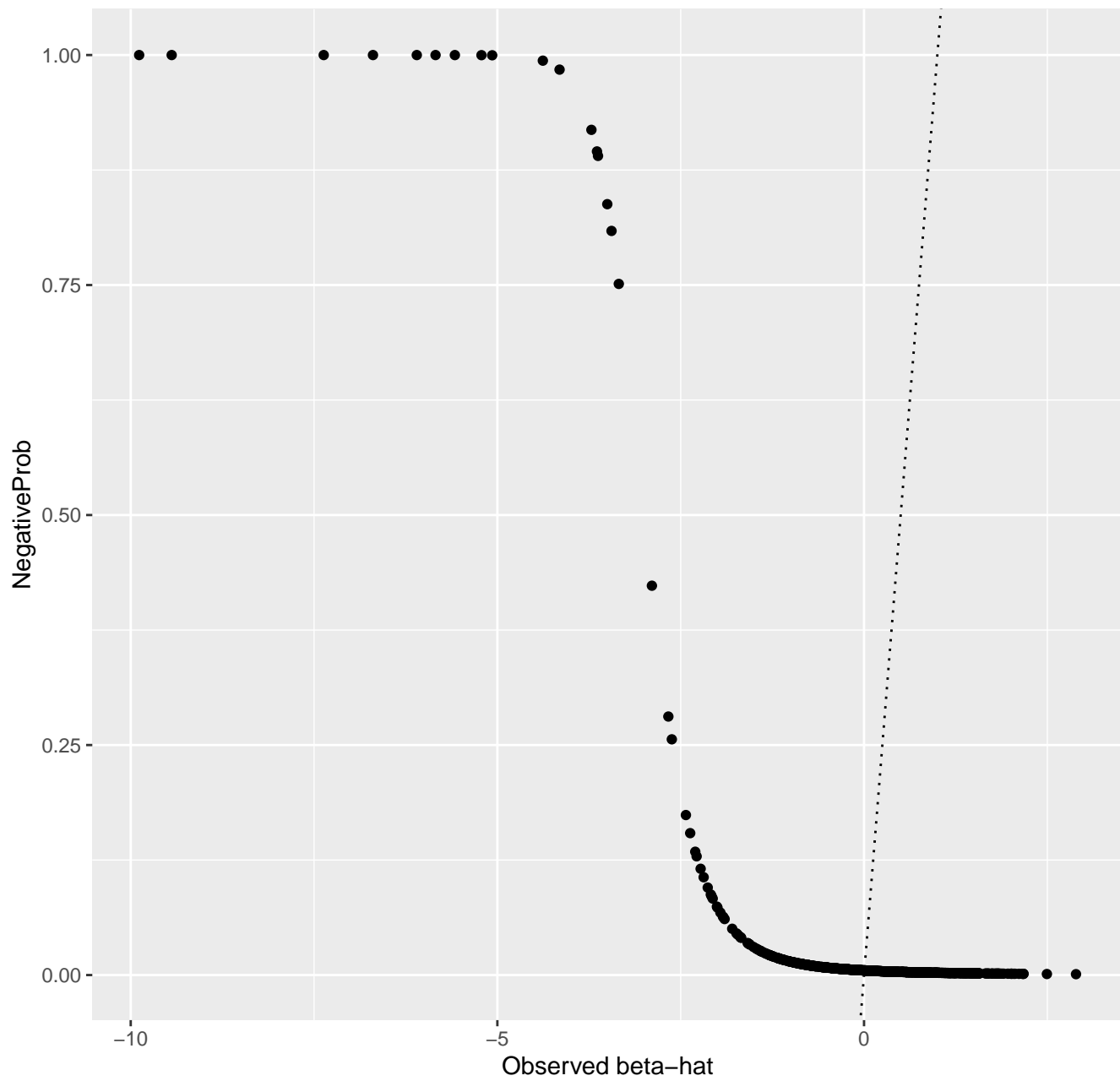
SQSTM1_(8878) negative



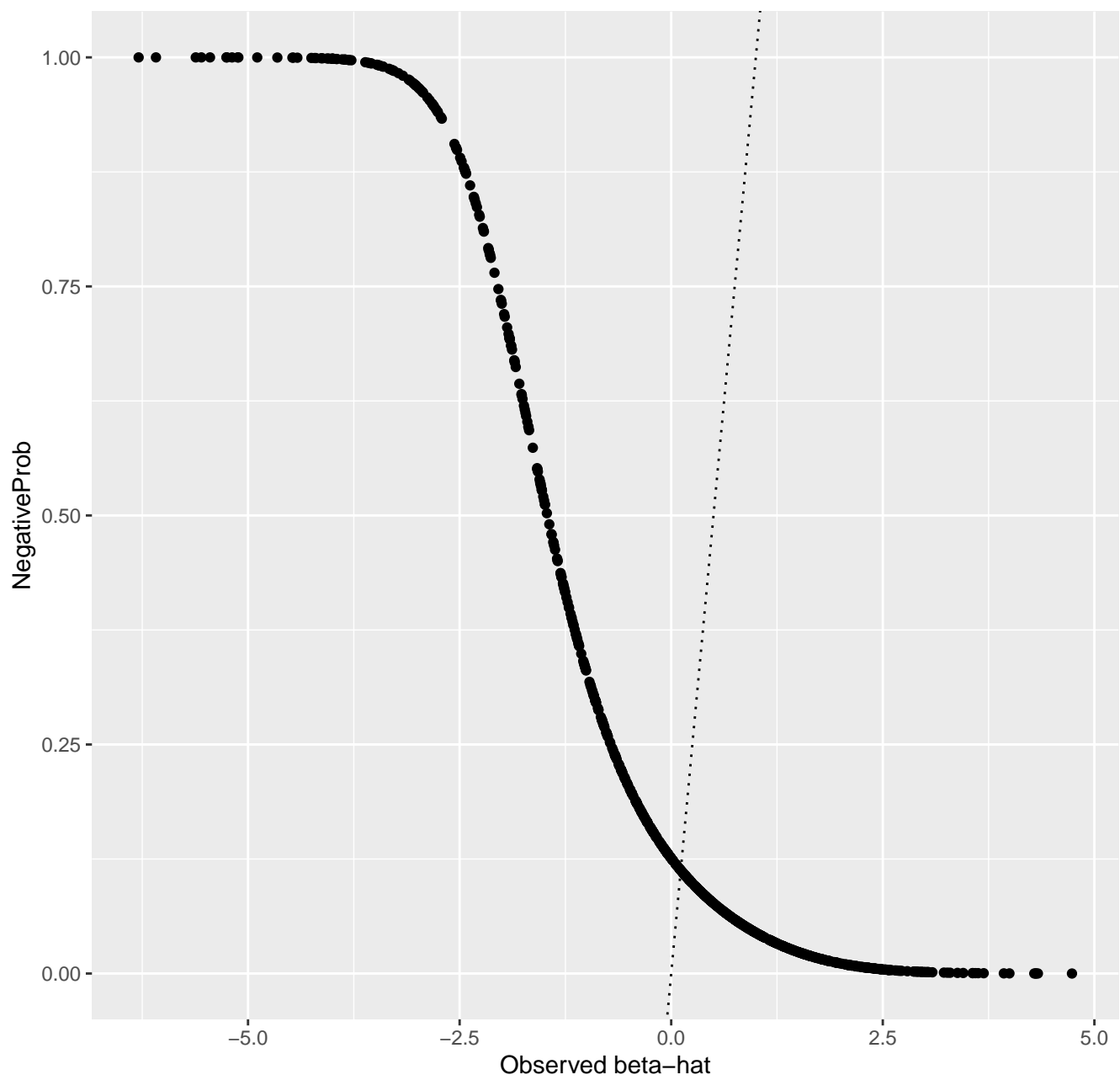
LIN28B_(389421) negative



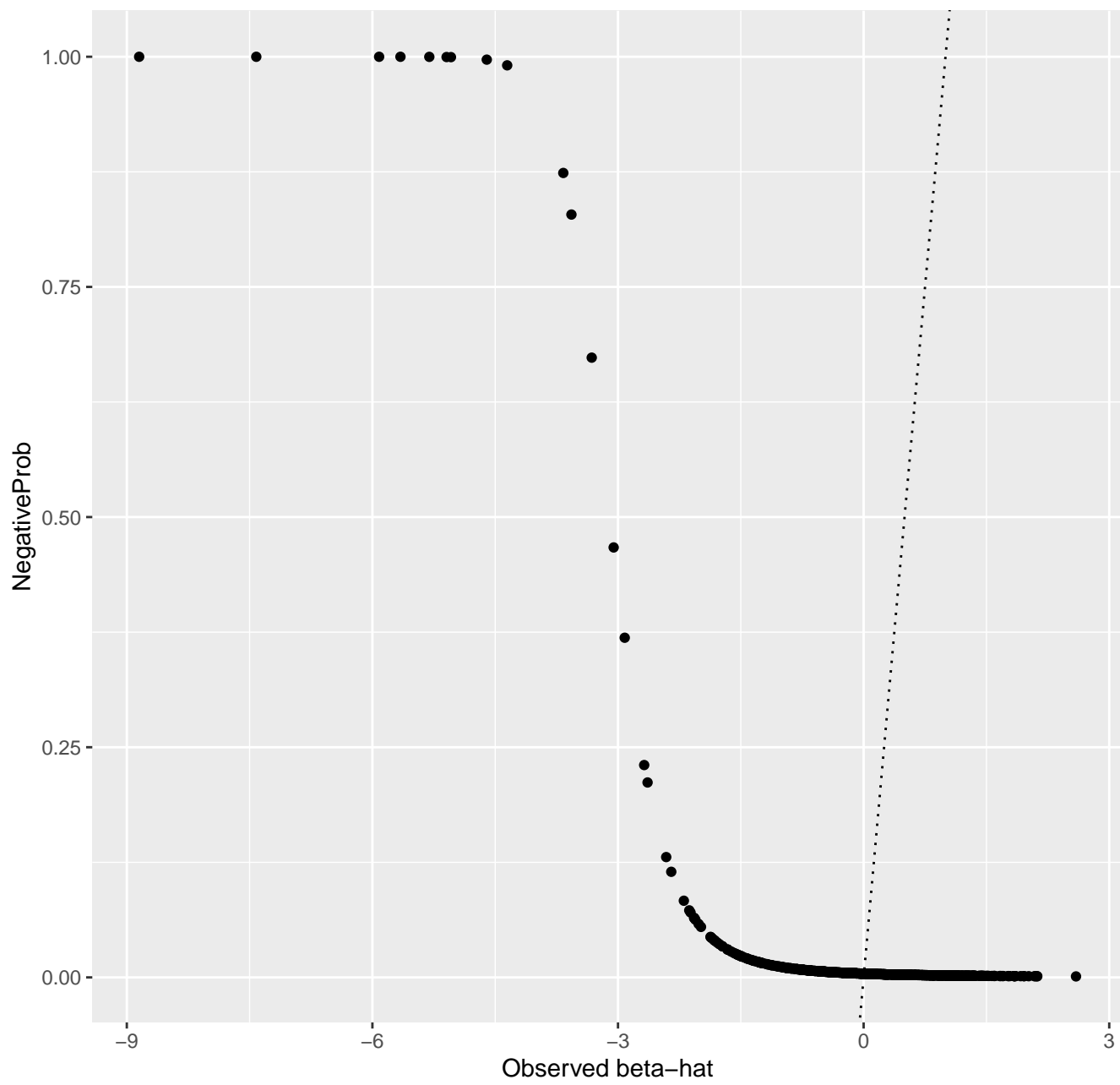
ISG15_(9636) negative



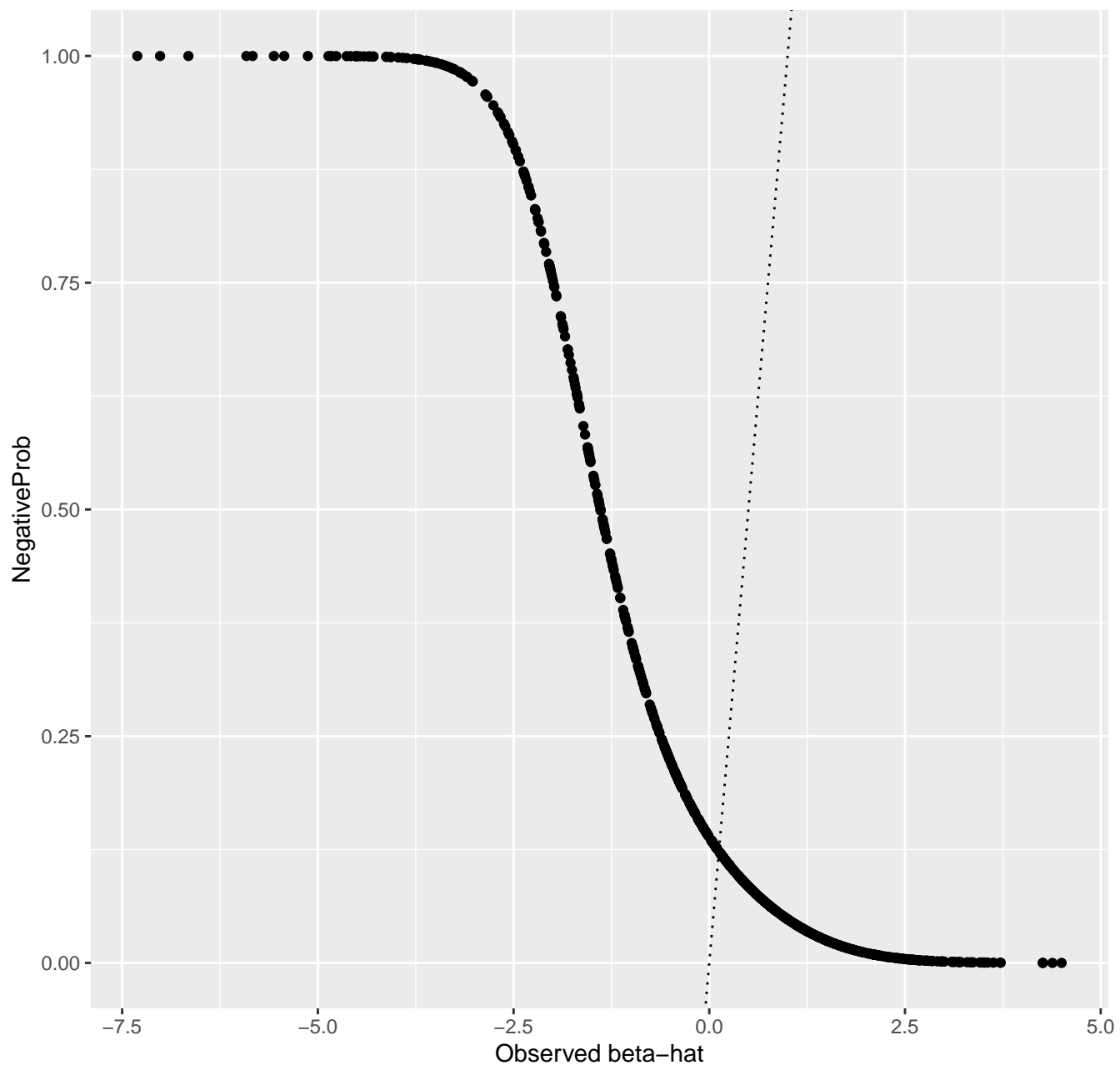
HSD17B12_(51144) negative



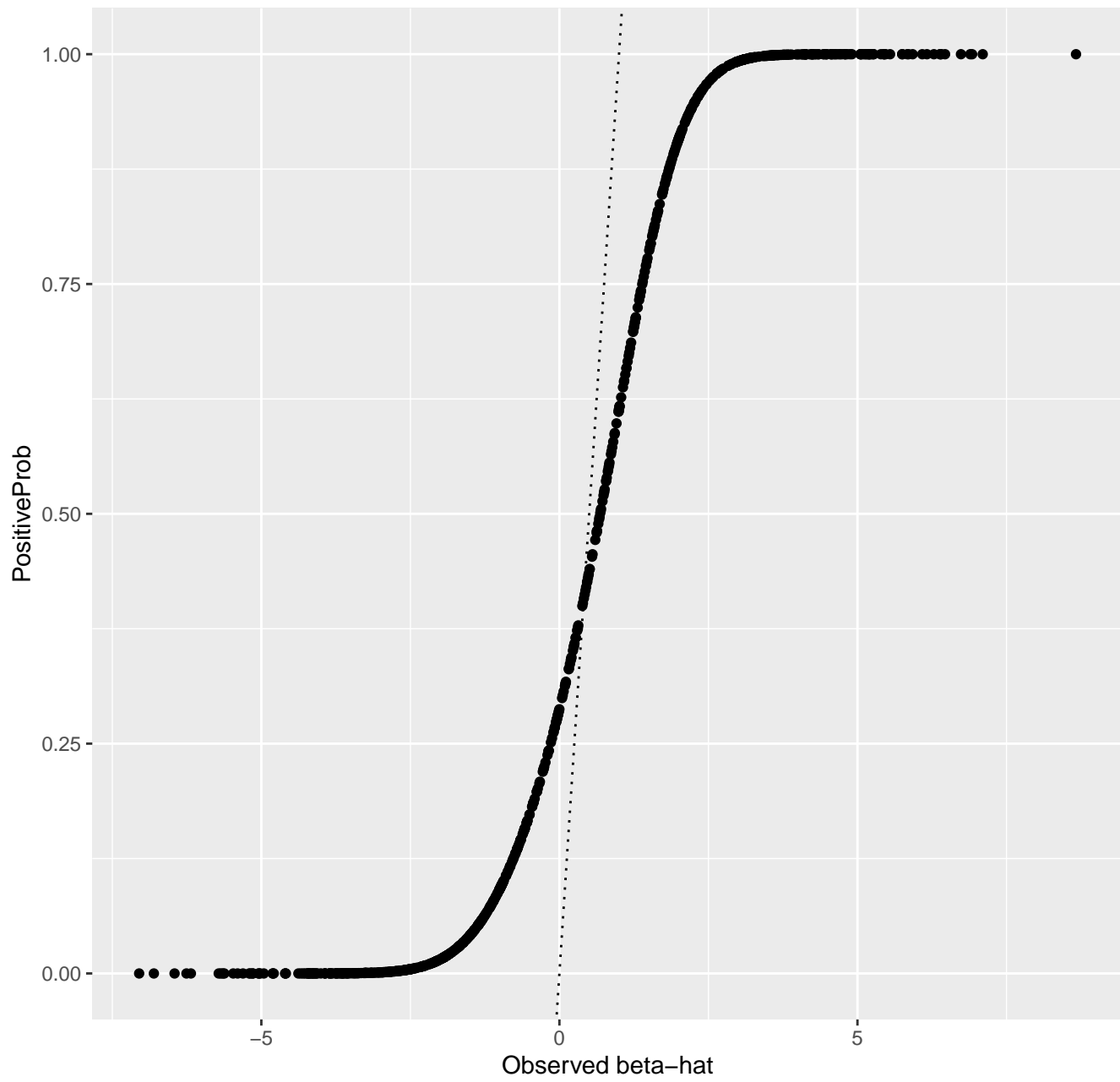
SPDEF_(25803) negative



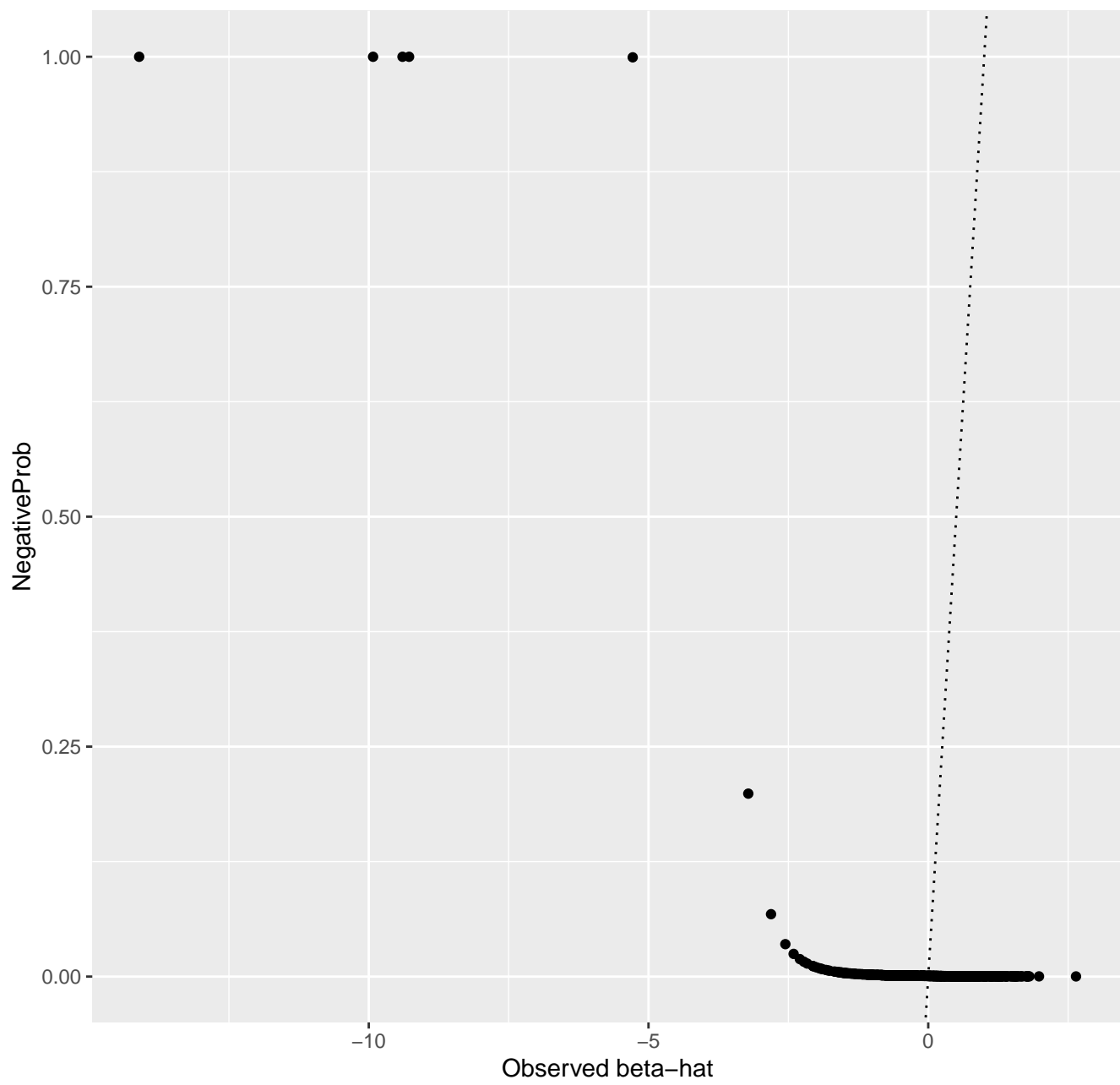
SHOC2_(8036) negative



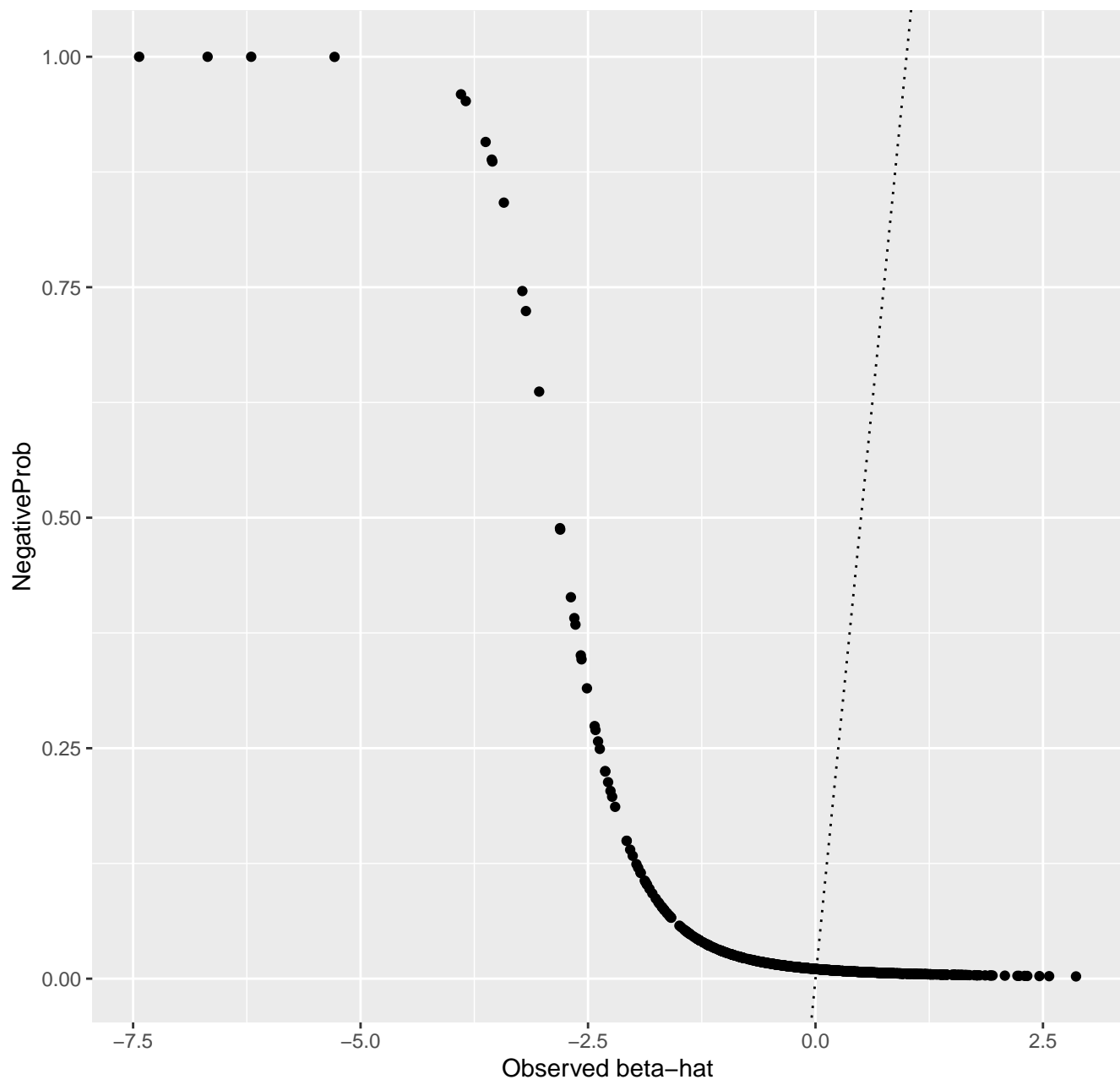
CHMP4B_(128866) positive



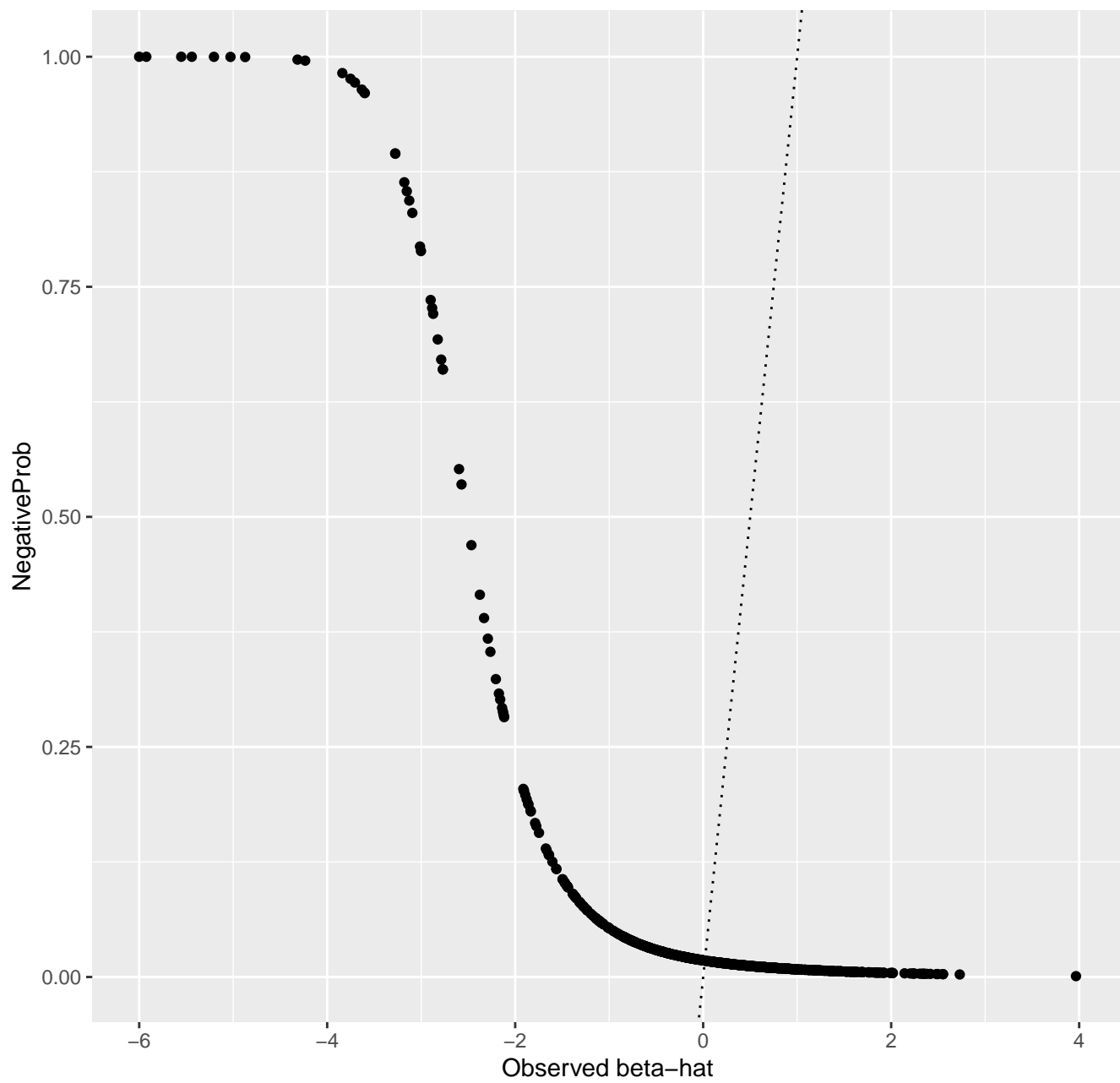
TBXT_(6862) negative



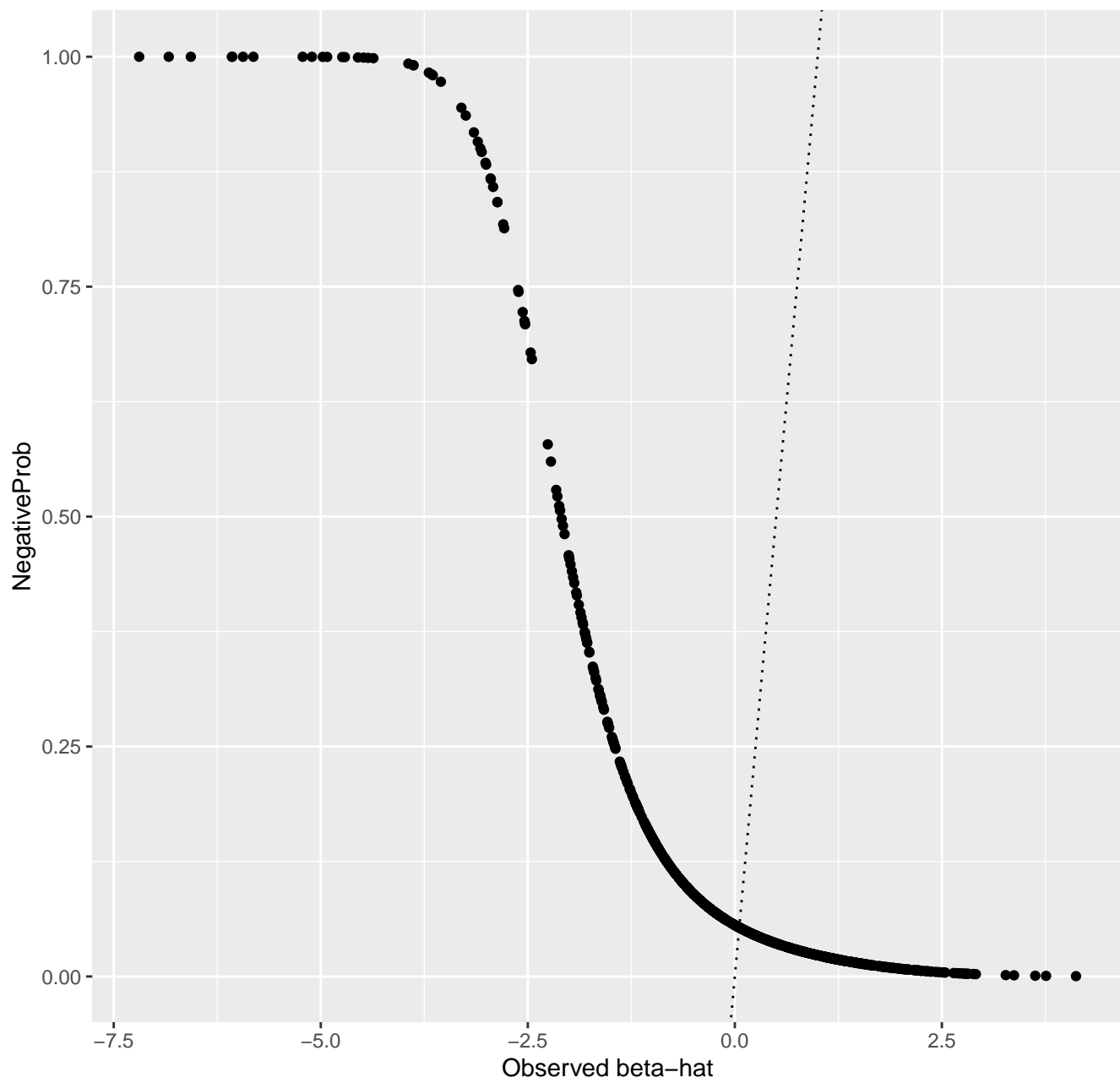
SLC12A7_(10723) negative



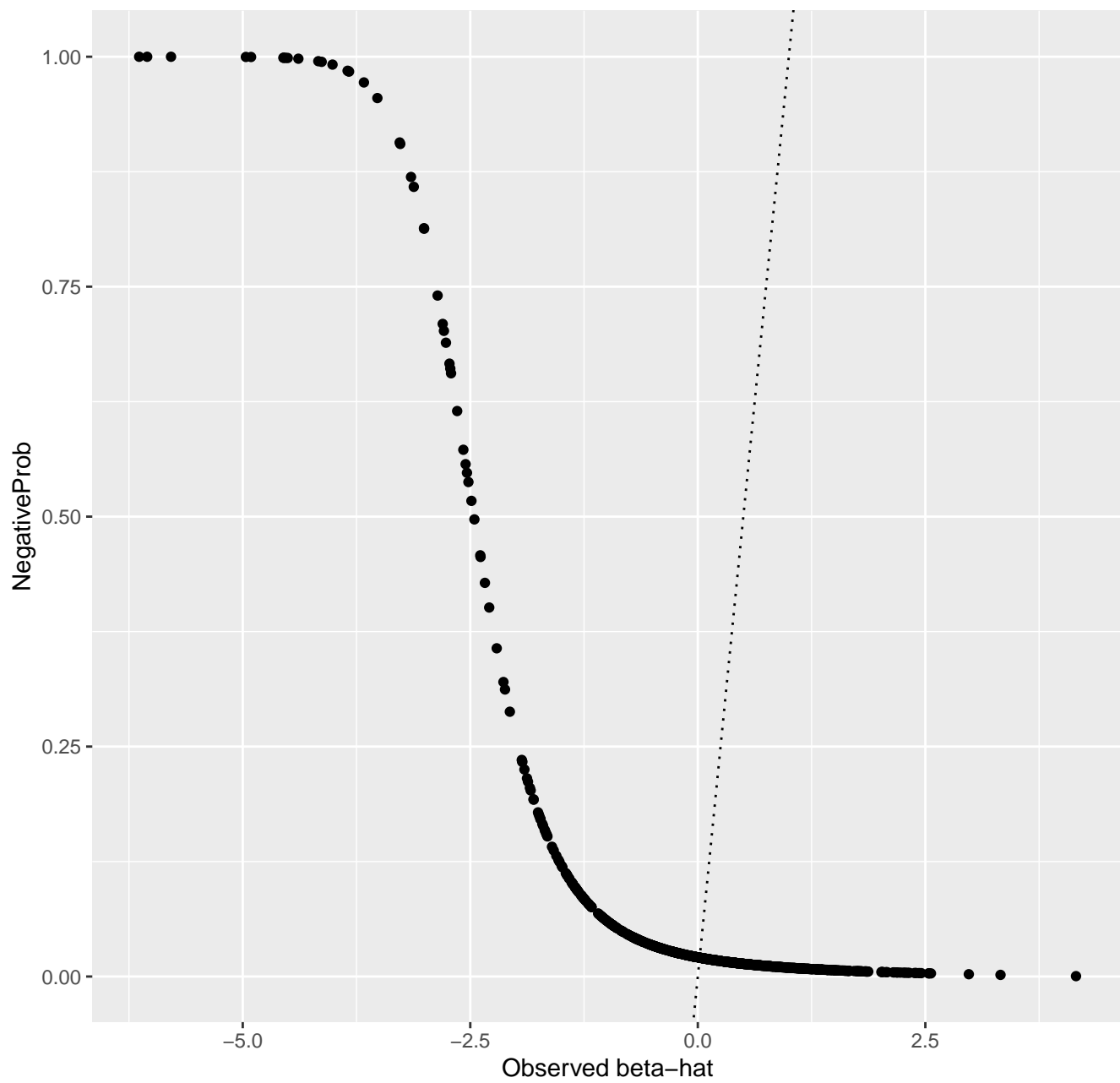
CRTC2_(200186) negative



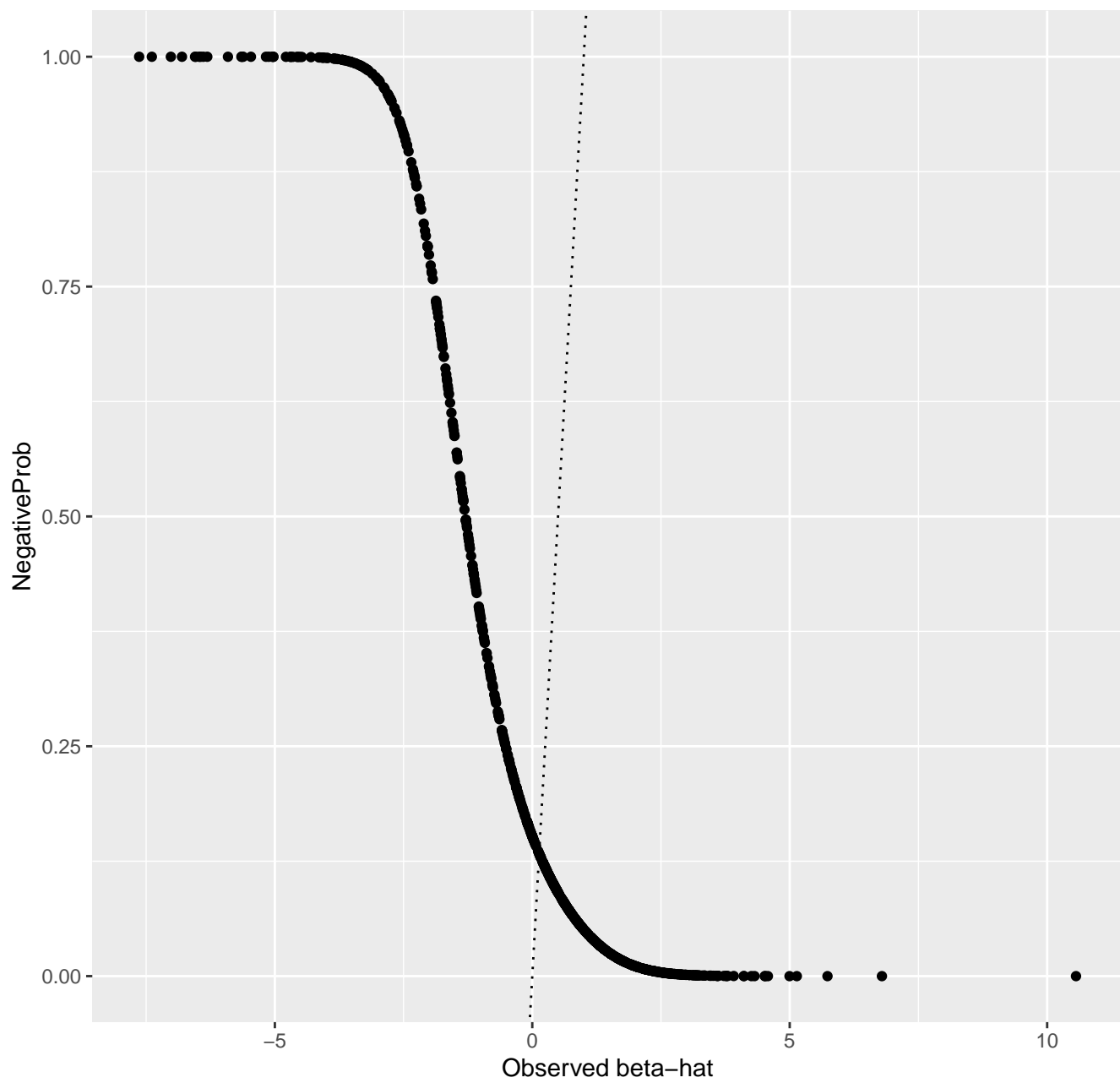
FDX1_(2230) negative



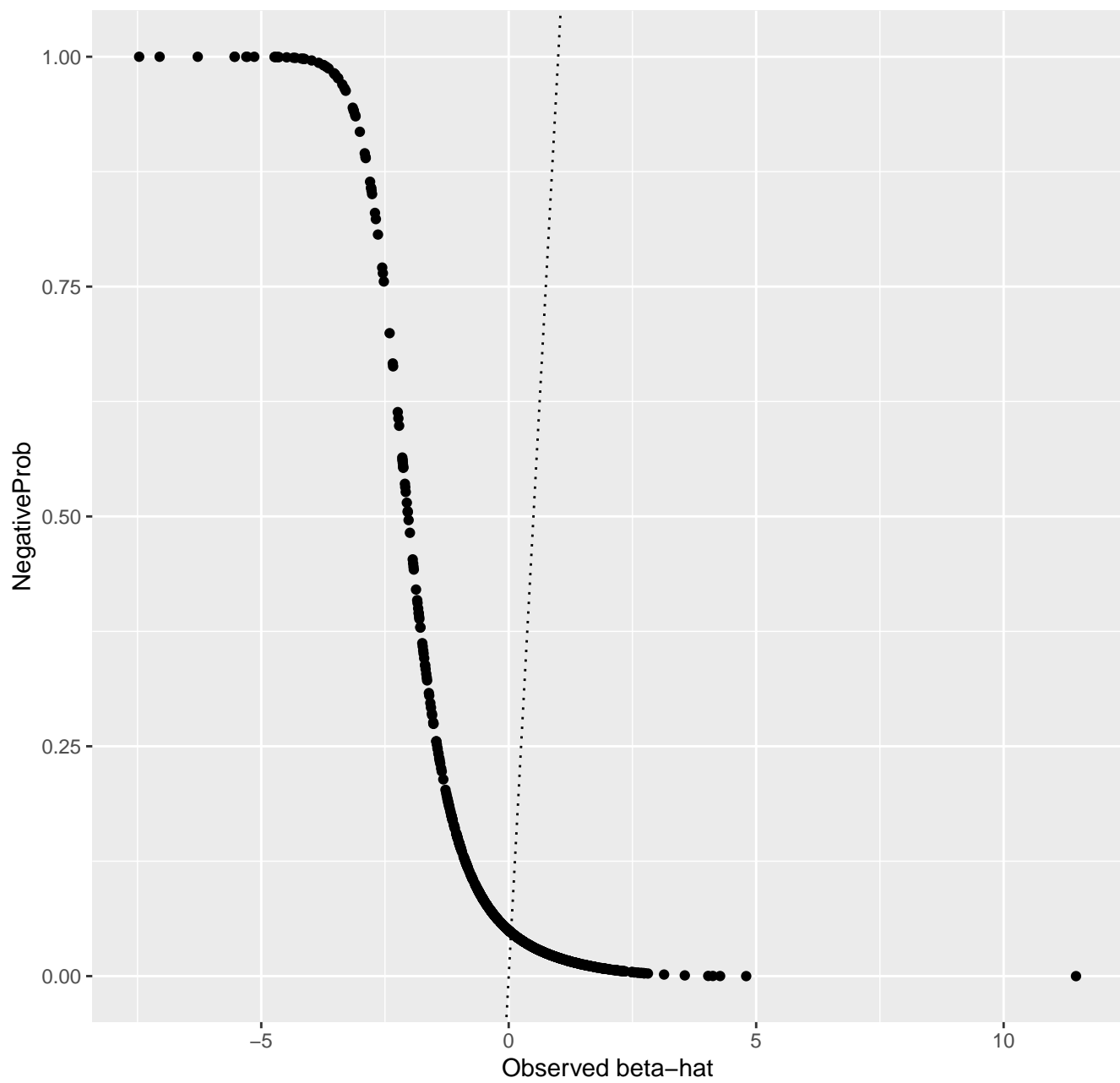
PEX5_(5830) negative



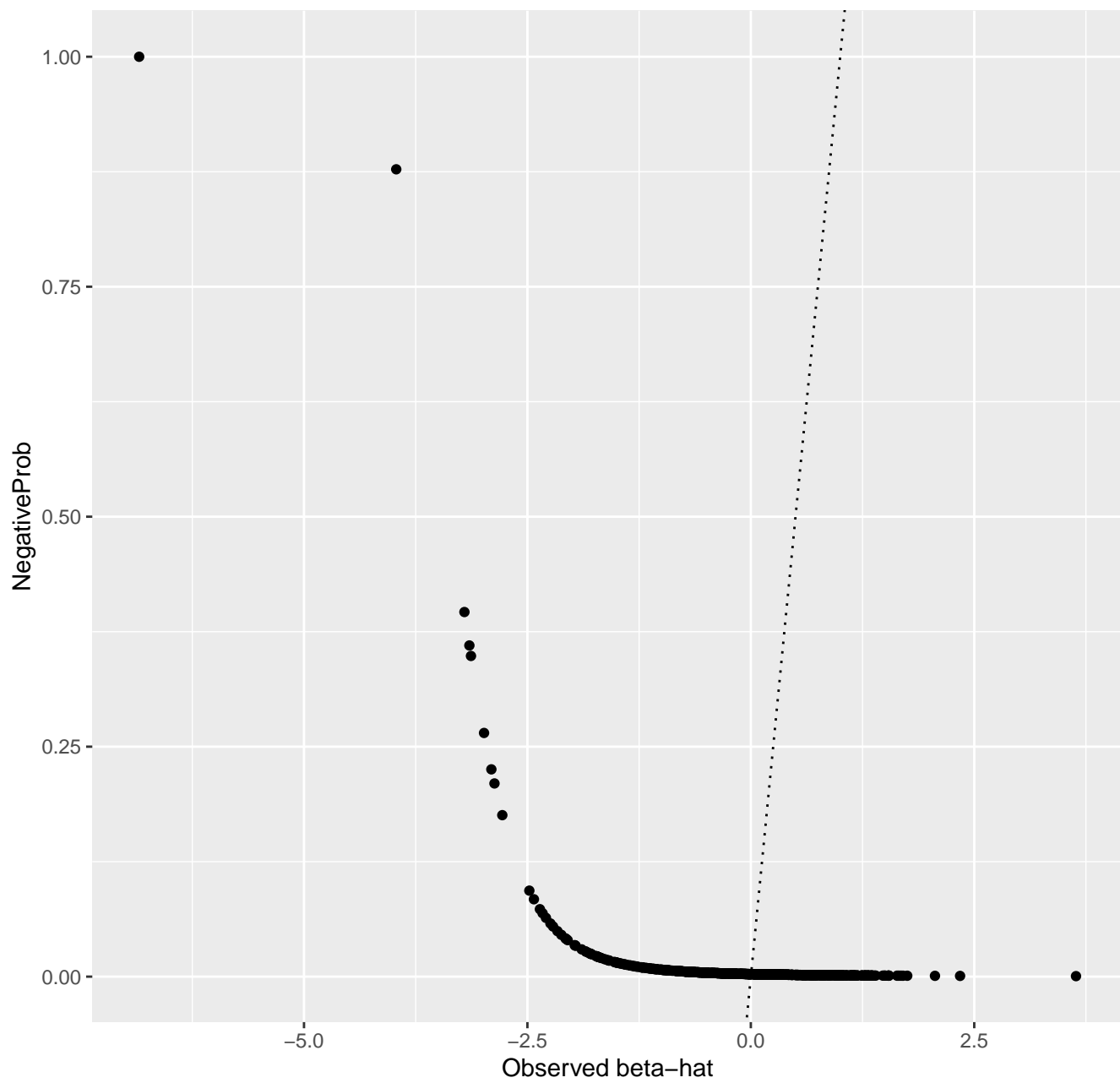
FURIN_(5045) negative



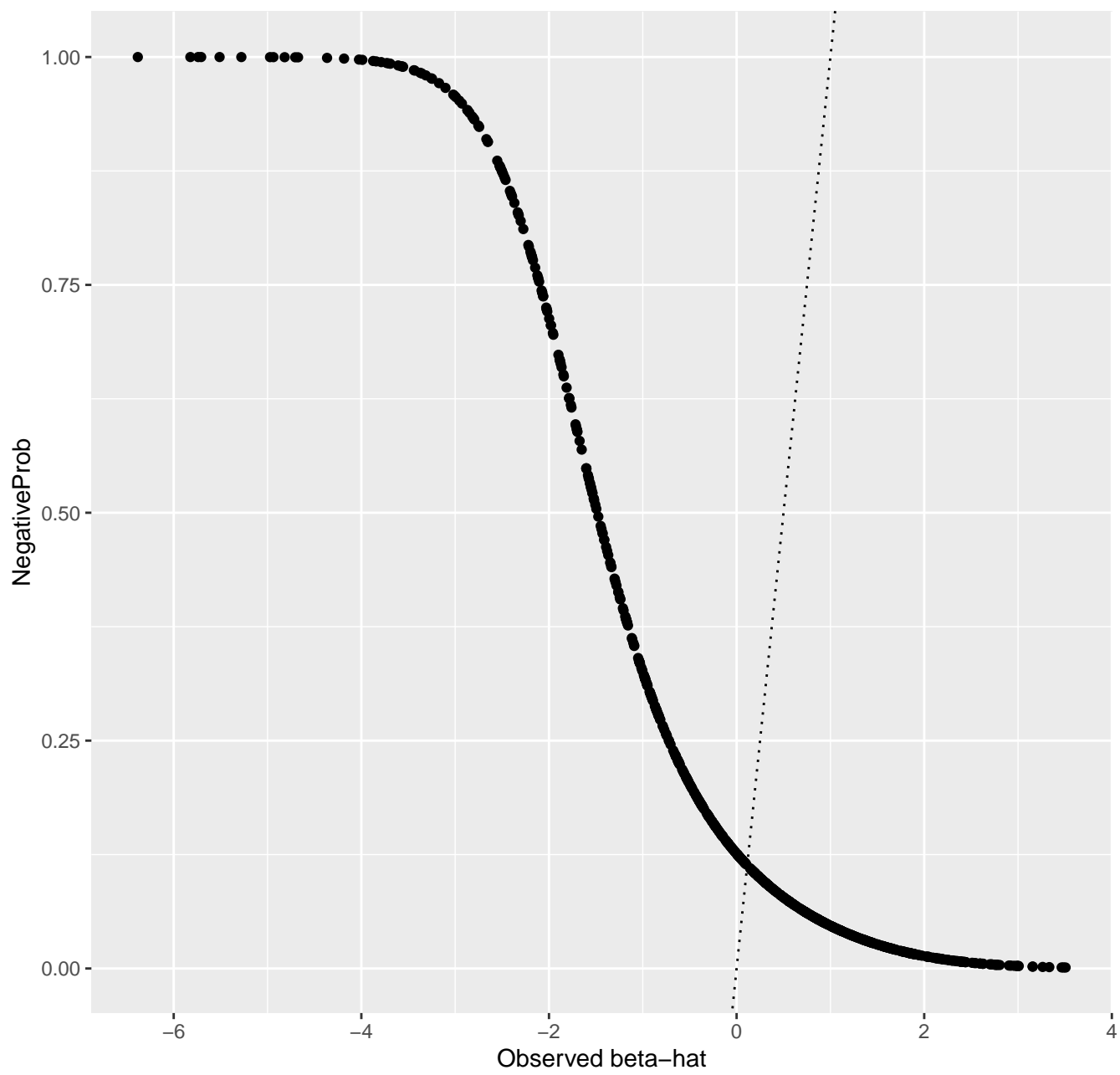
BIRC2_(329) negative



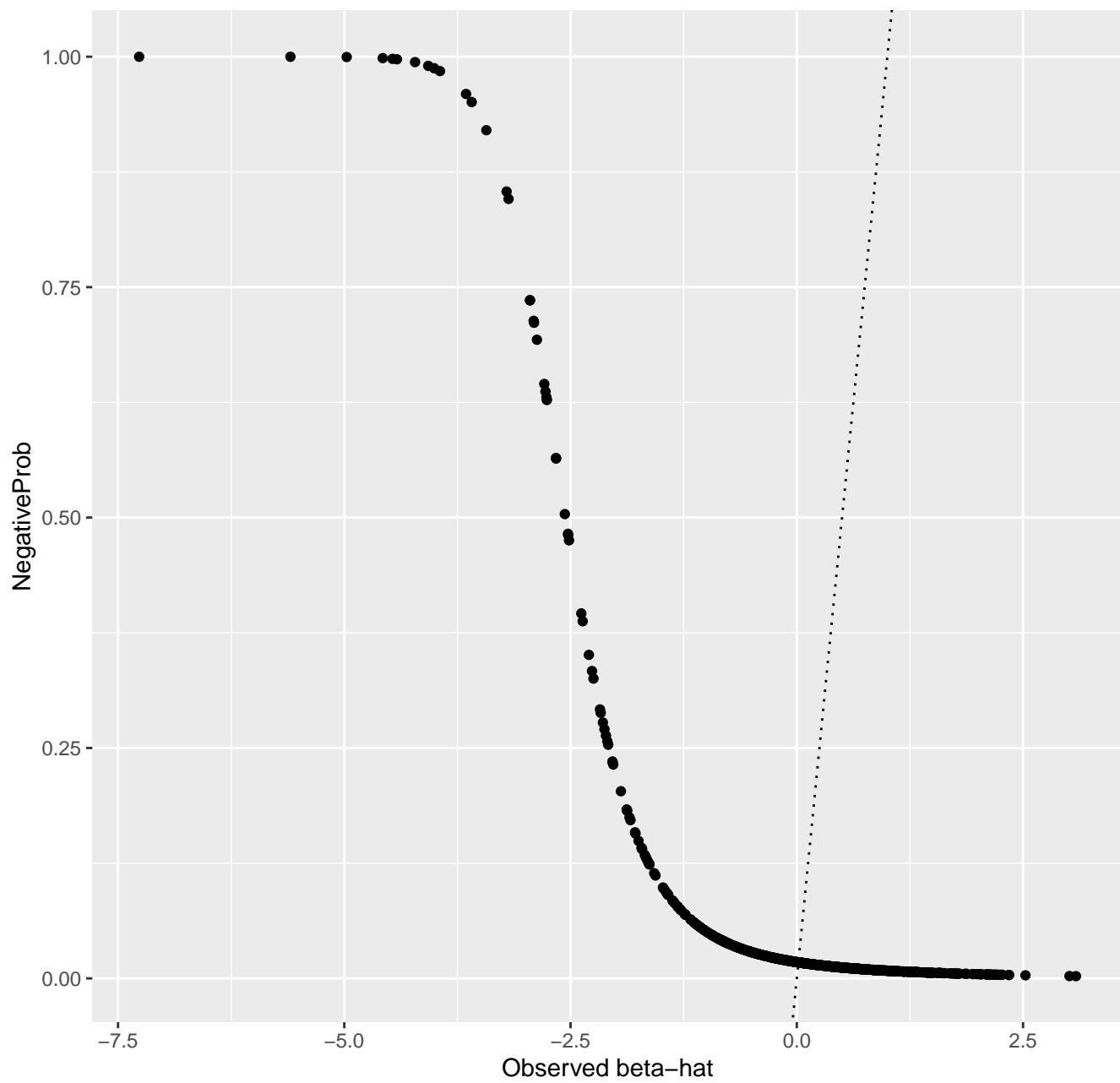
CLMP_(79827) negative



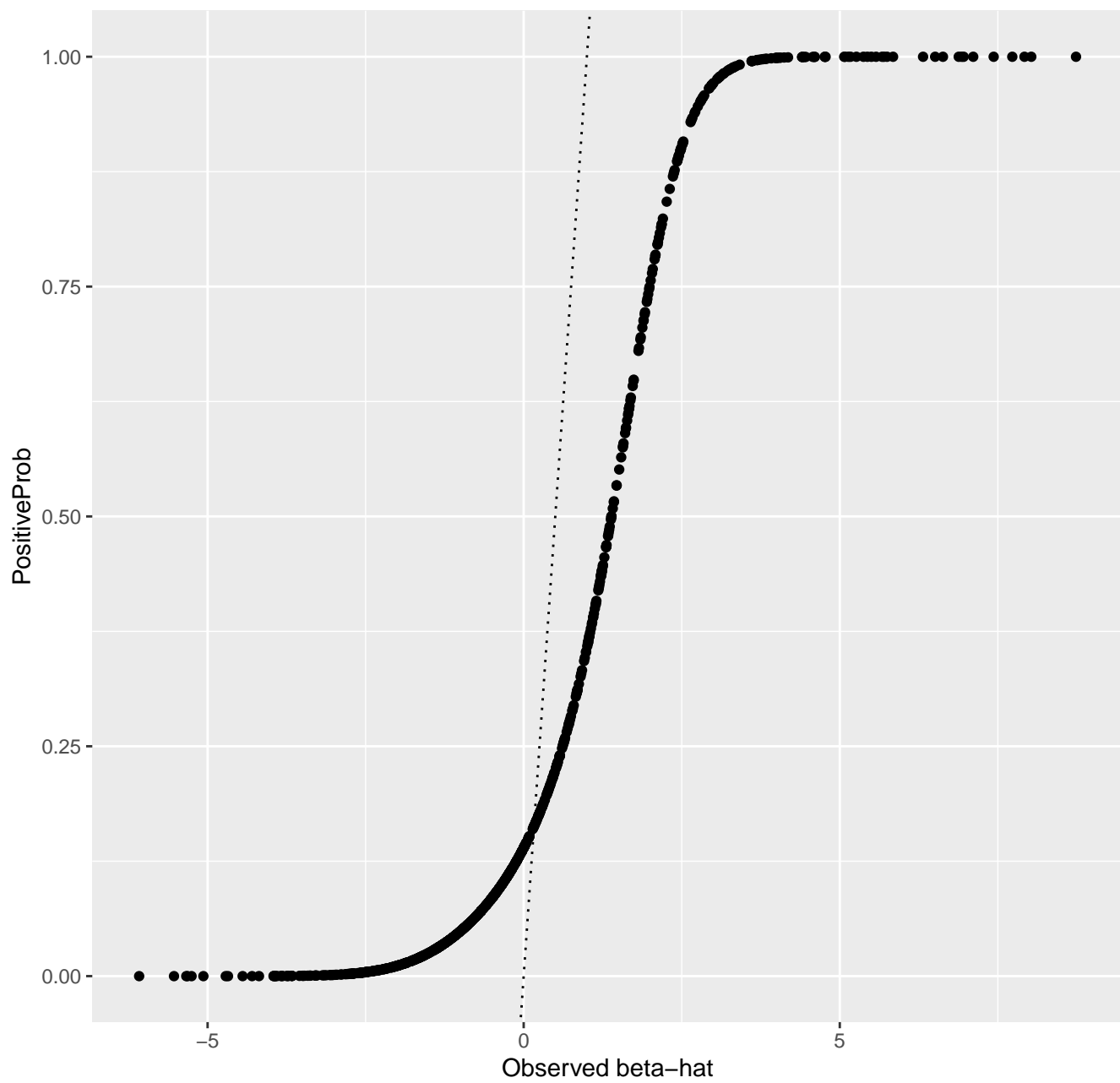
NDUFC1_(4717) negative



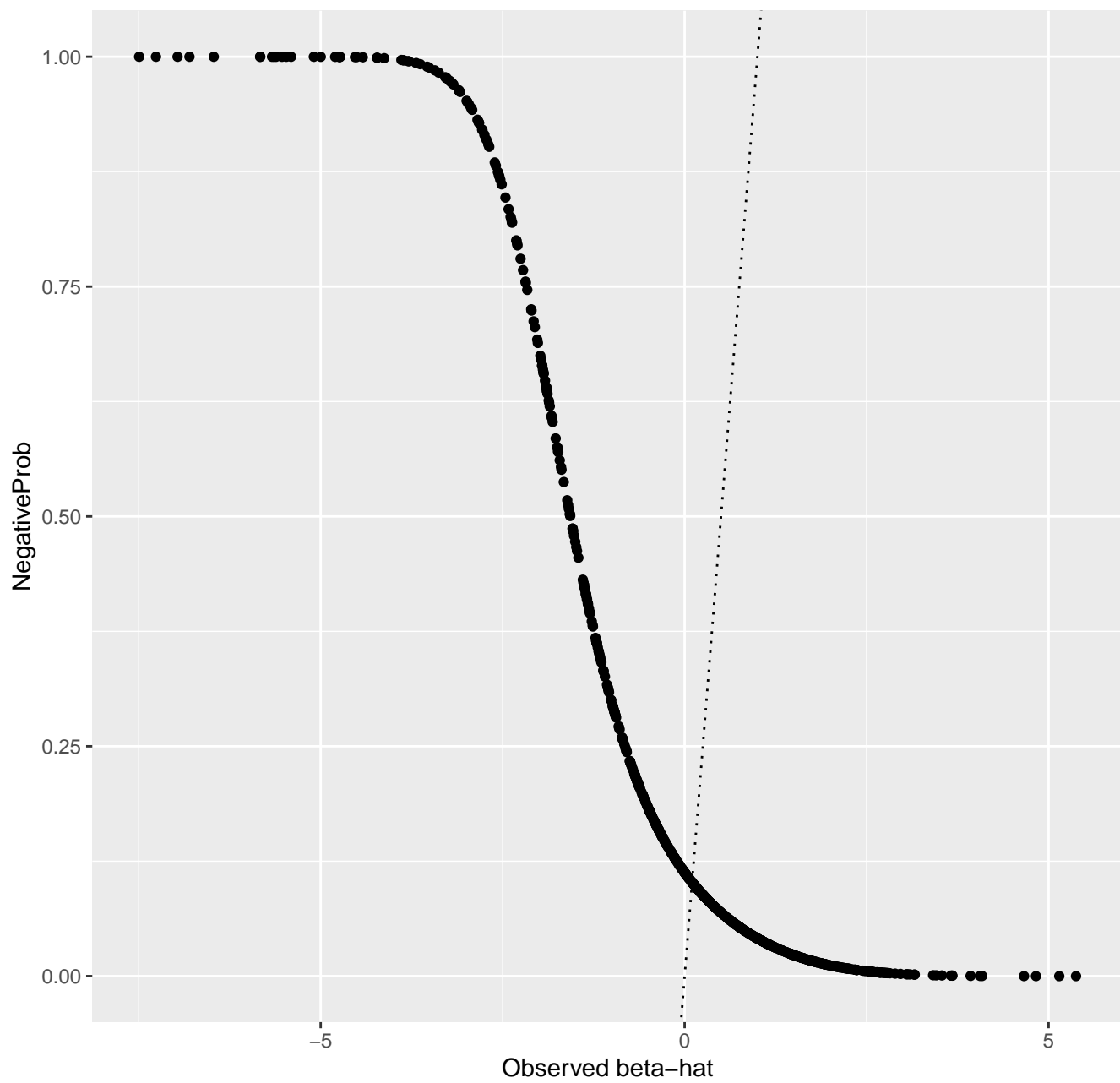
NCDN_(23154) negative



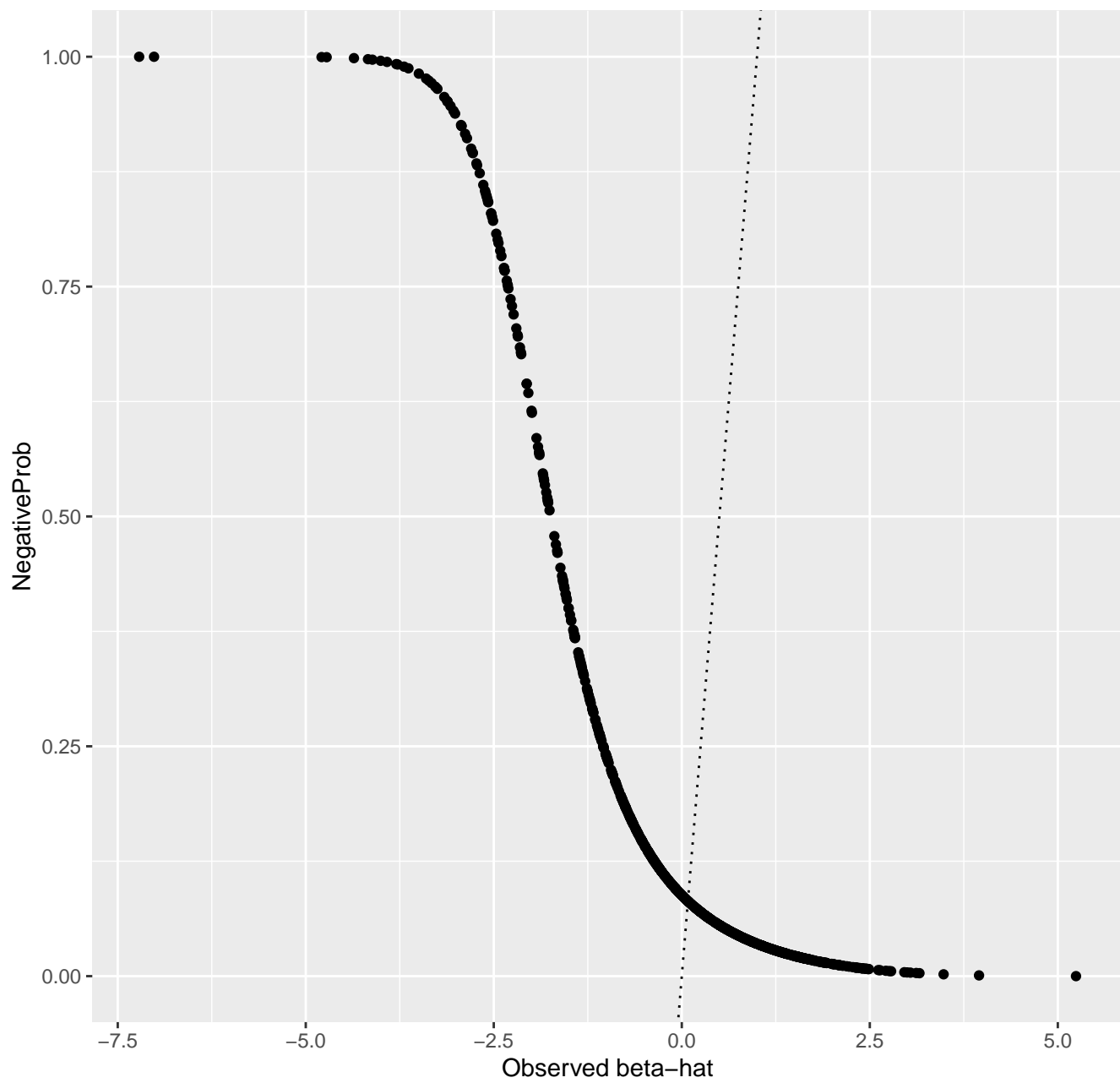
SMARCB1_(6598) positive



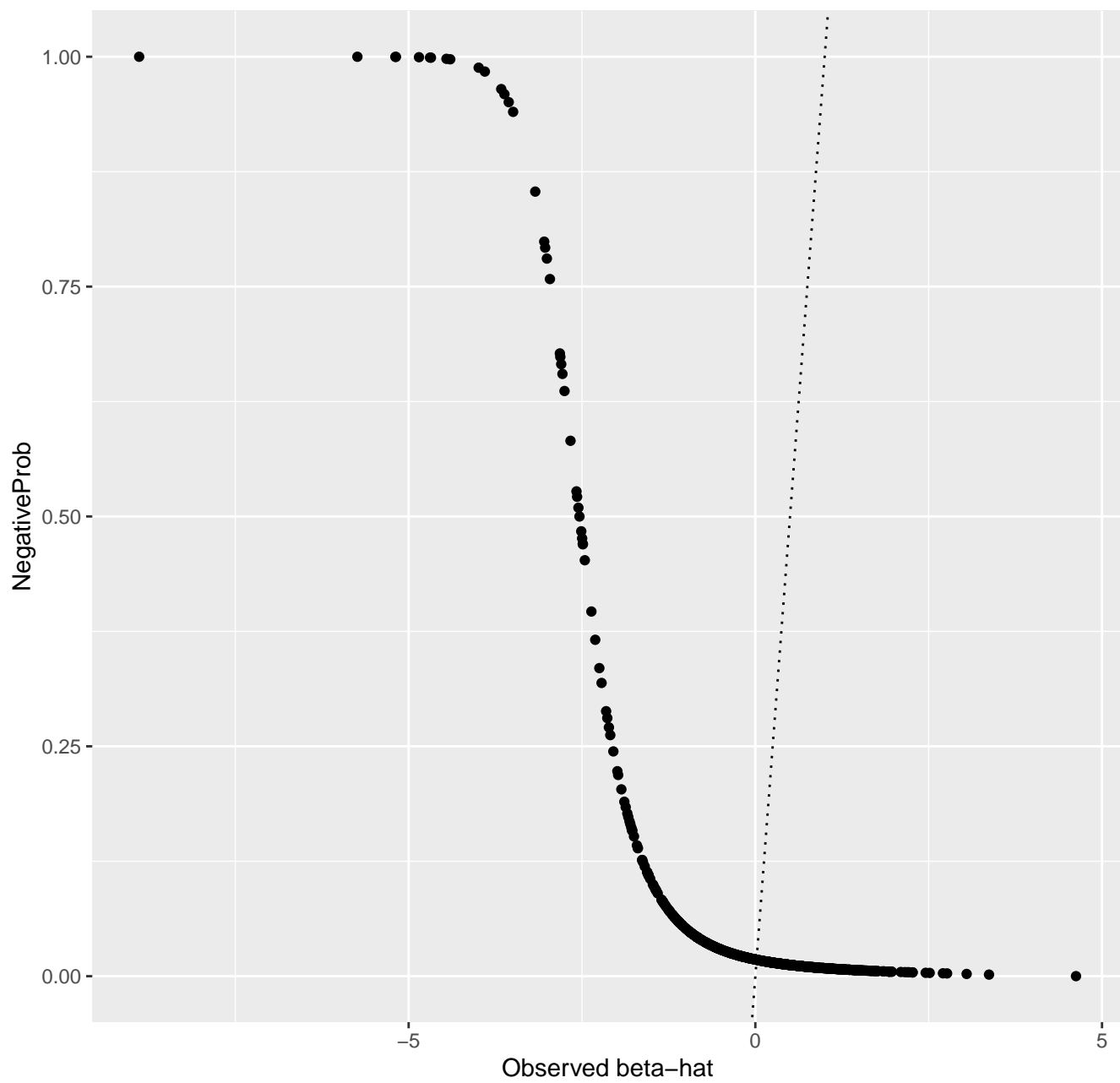
TGIF1_(7050) negative



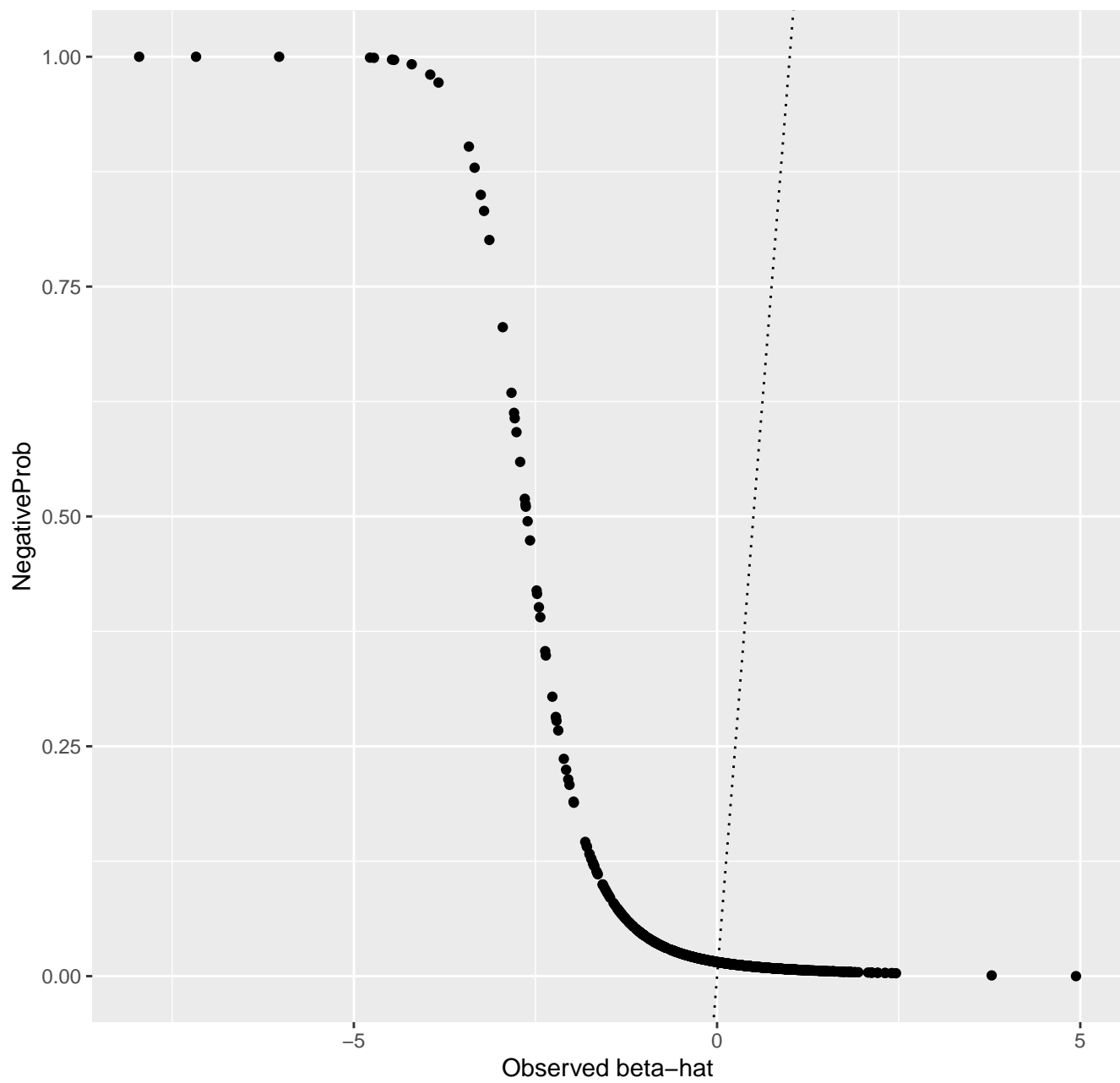
ACTG1_(71) negative



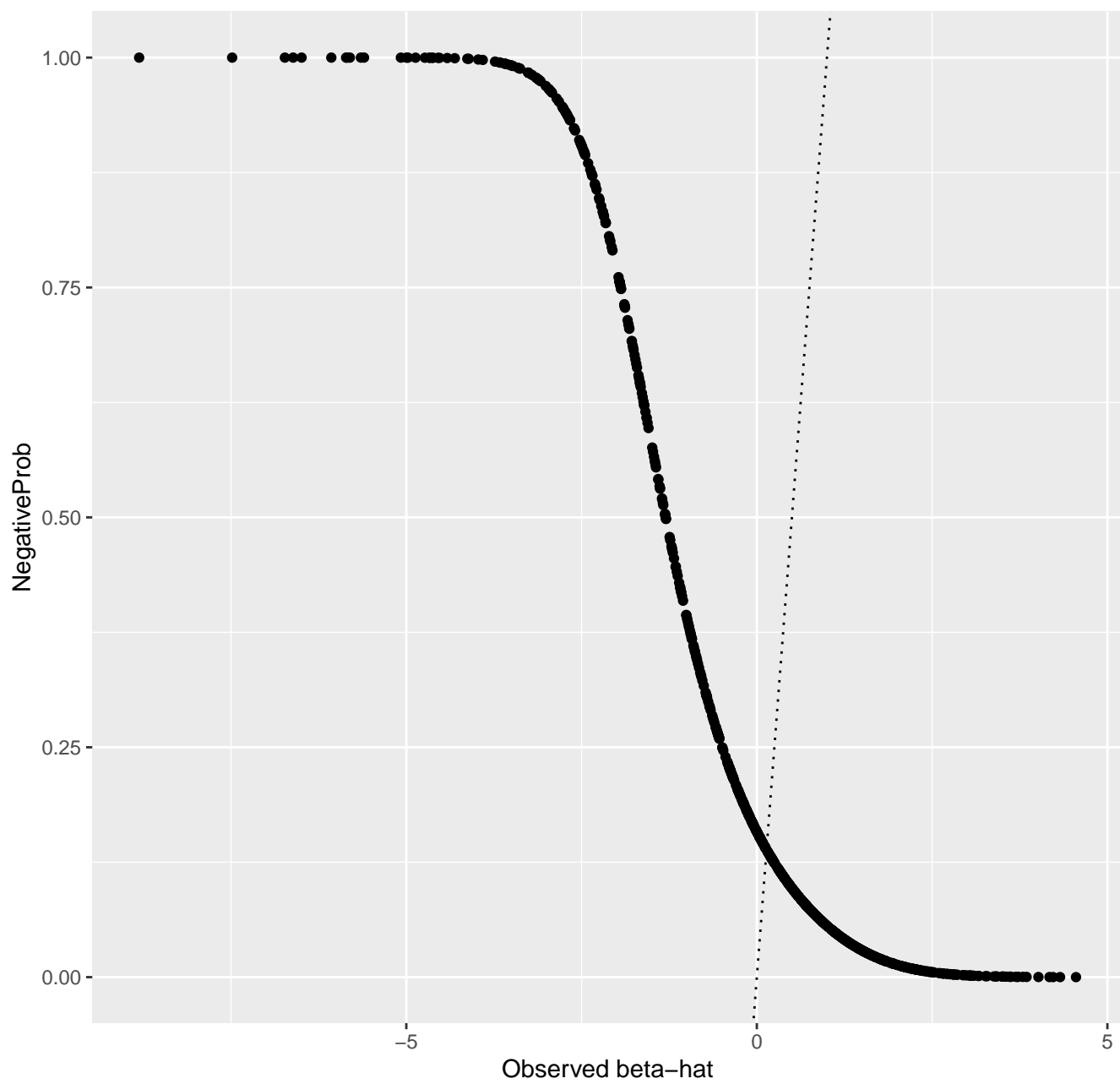
NFIB_(4781) negative



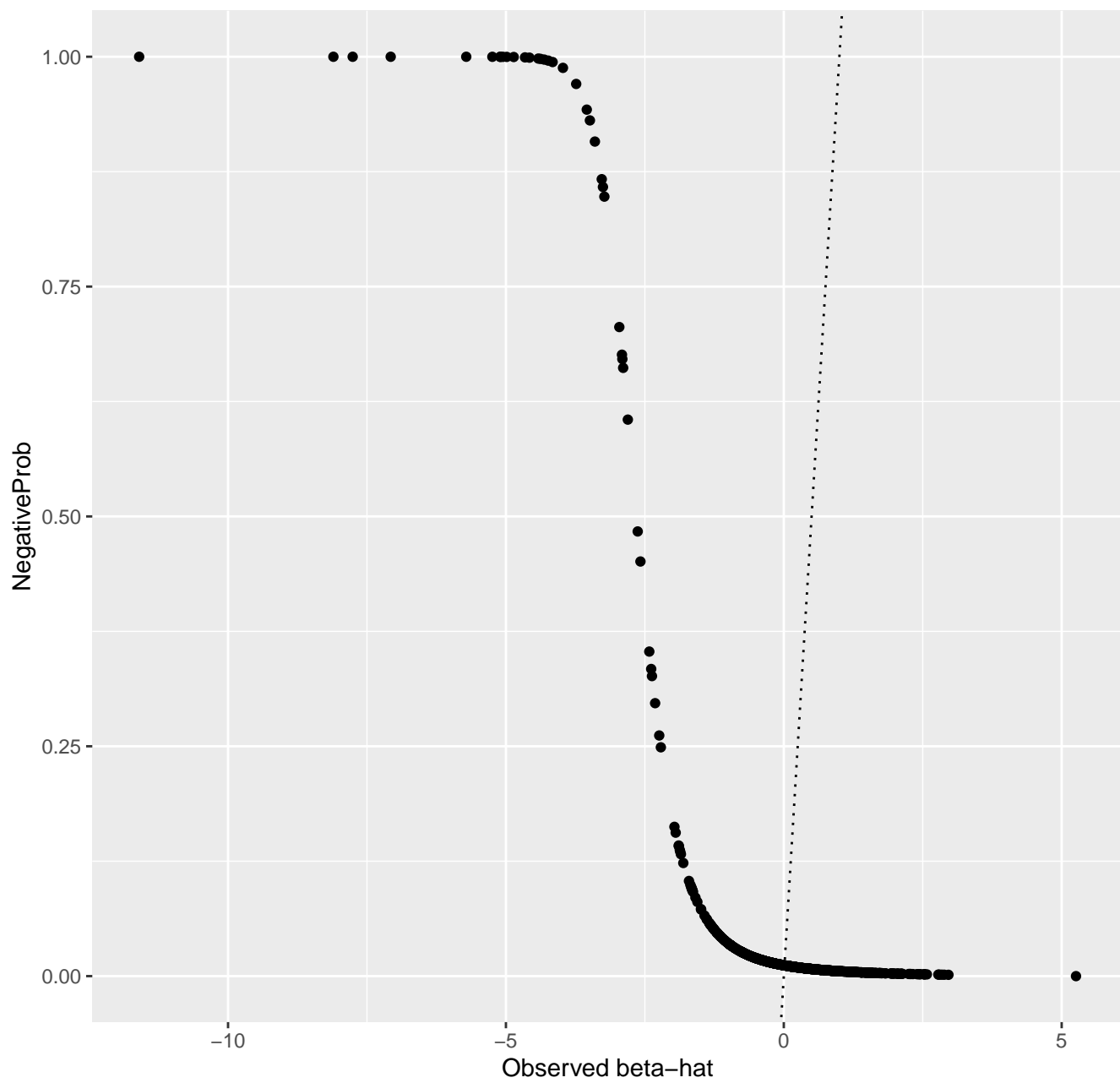
PIK3CD_(5293) negative



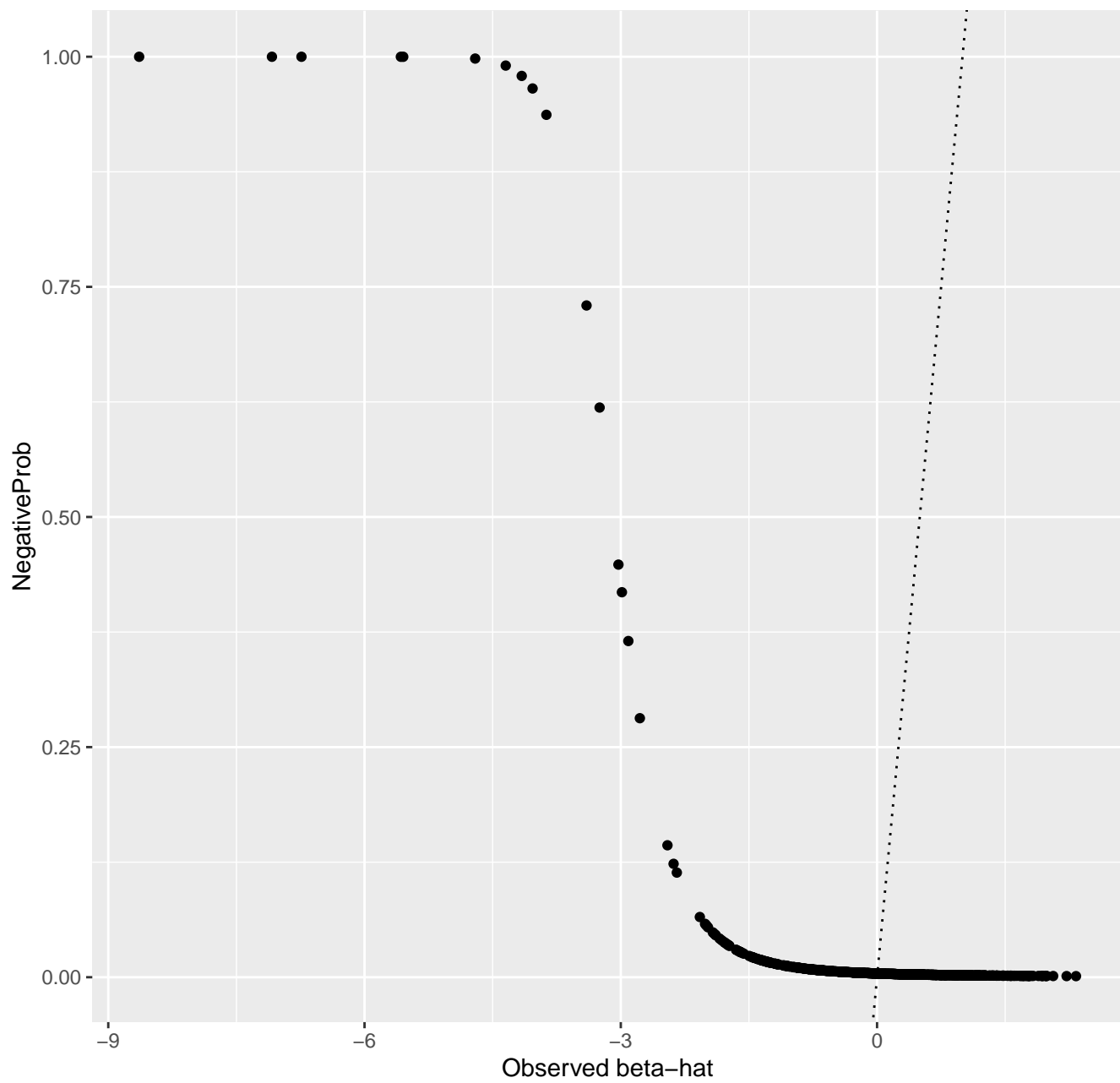
NDUFC2_(4718) negative



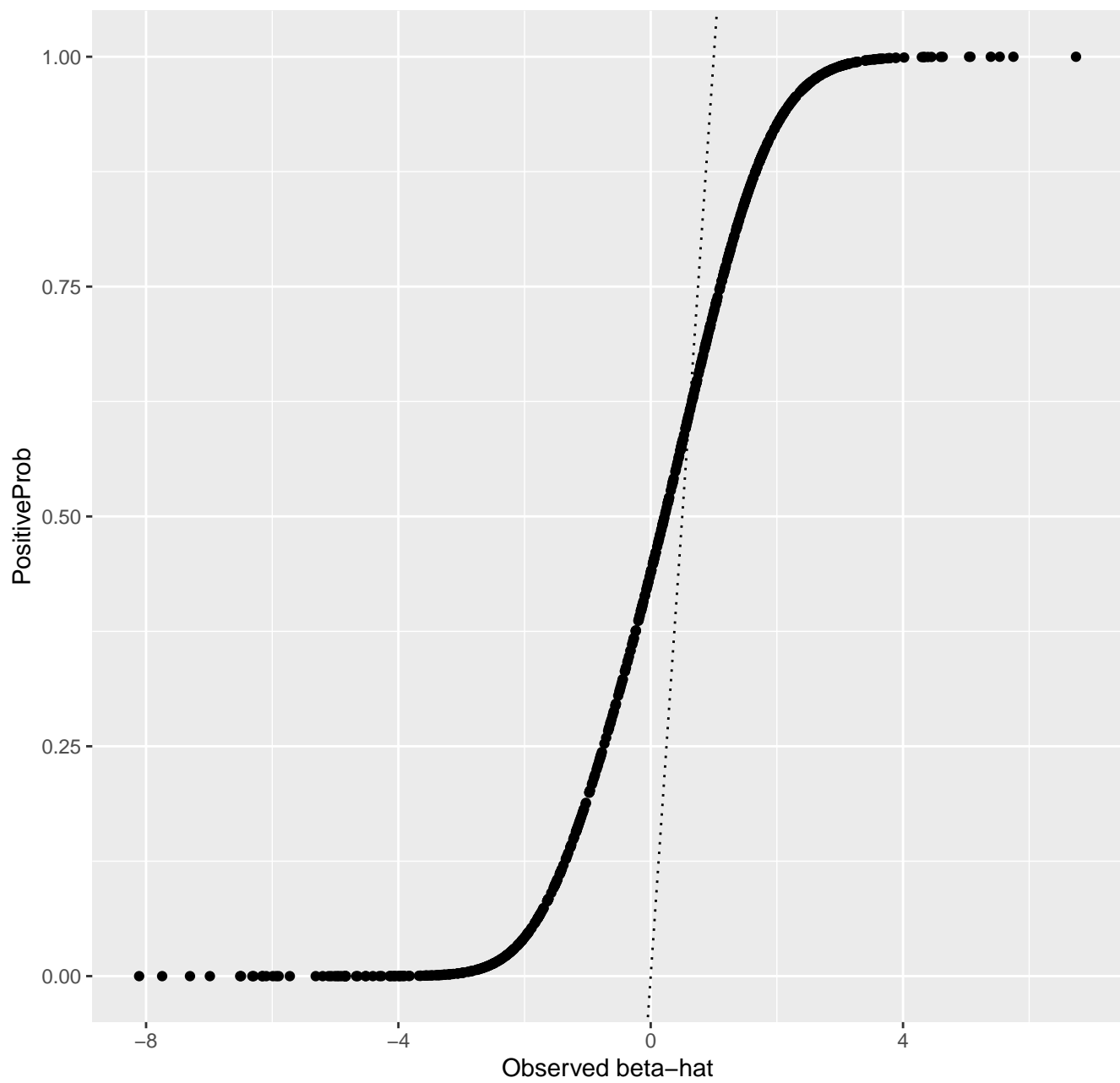
FLI1_(2313) negative



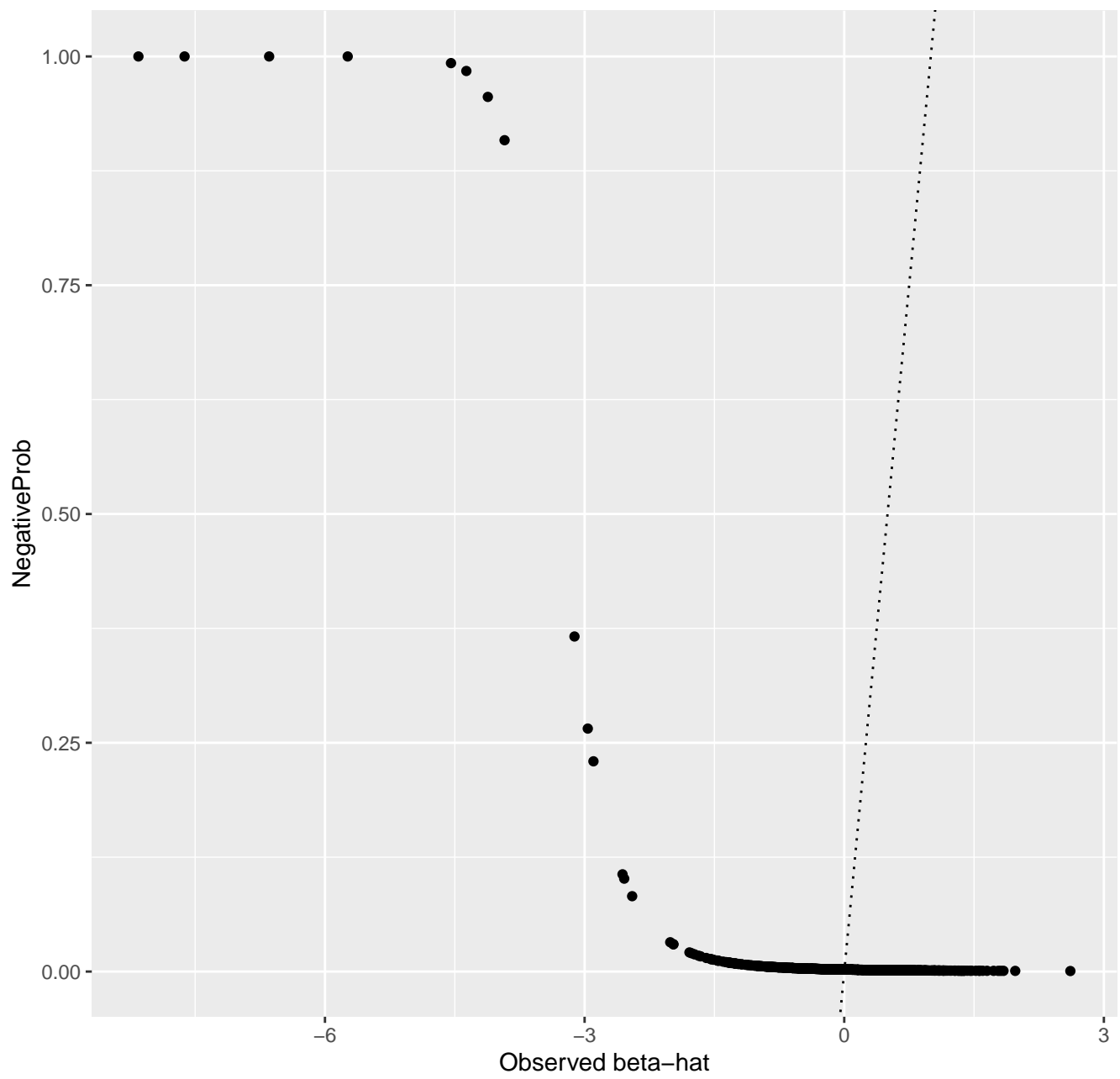
ATP6V0A2_(23545) negative



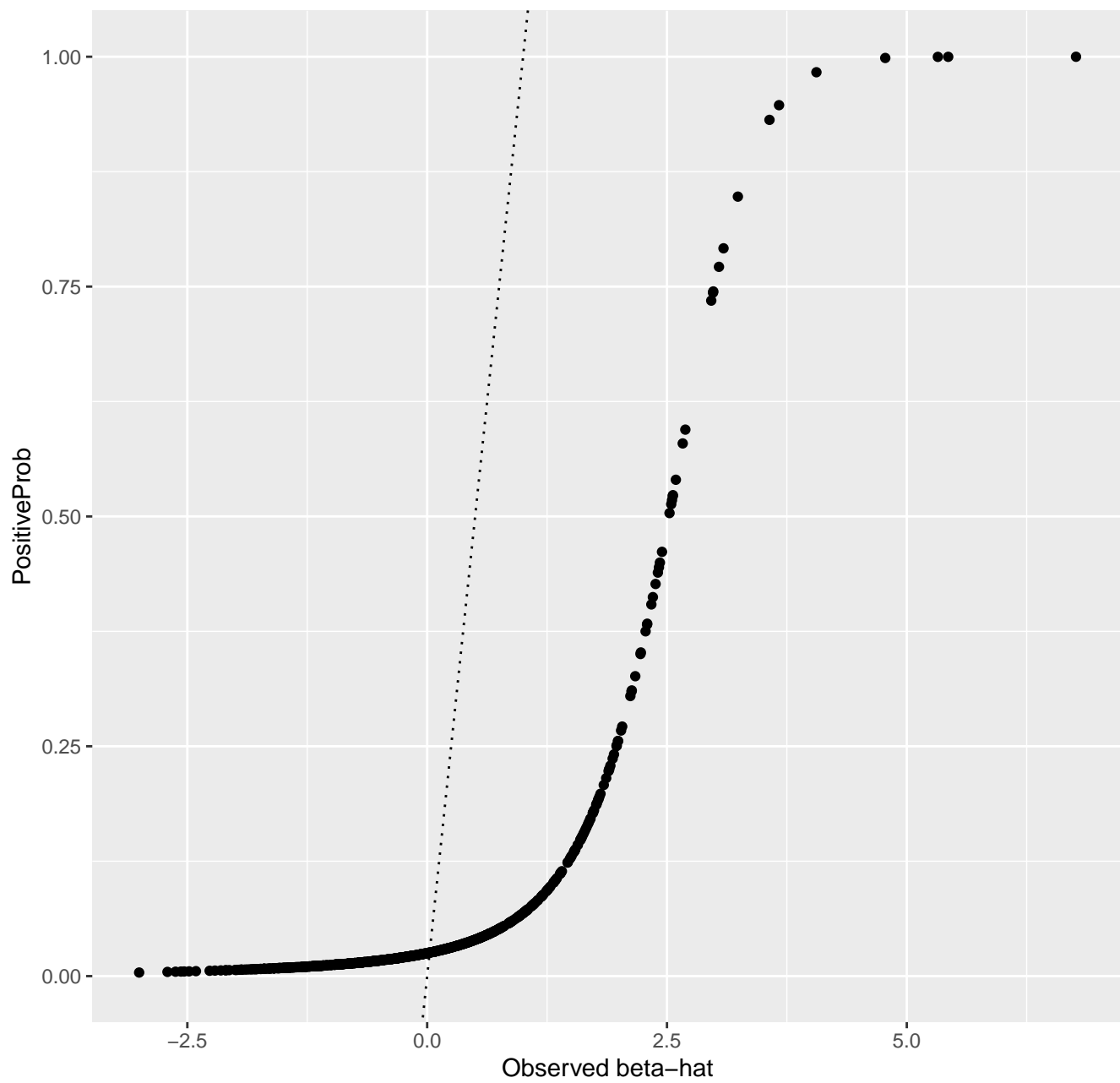
KCTD10_(83892) positive



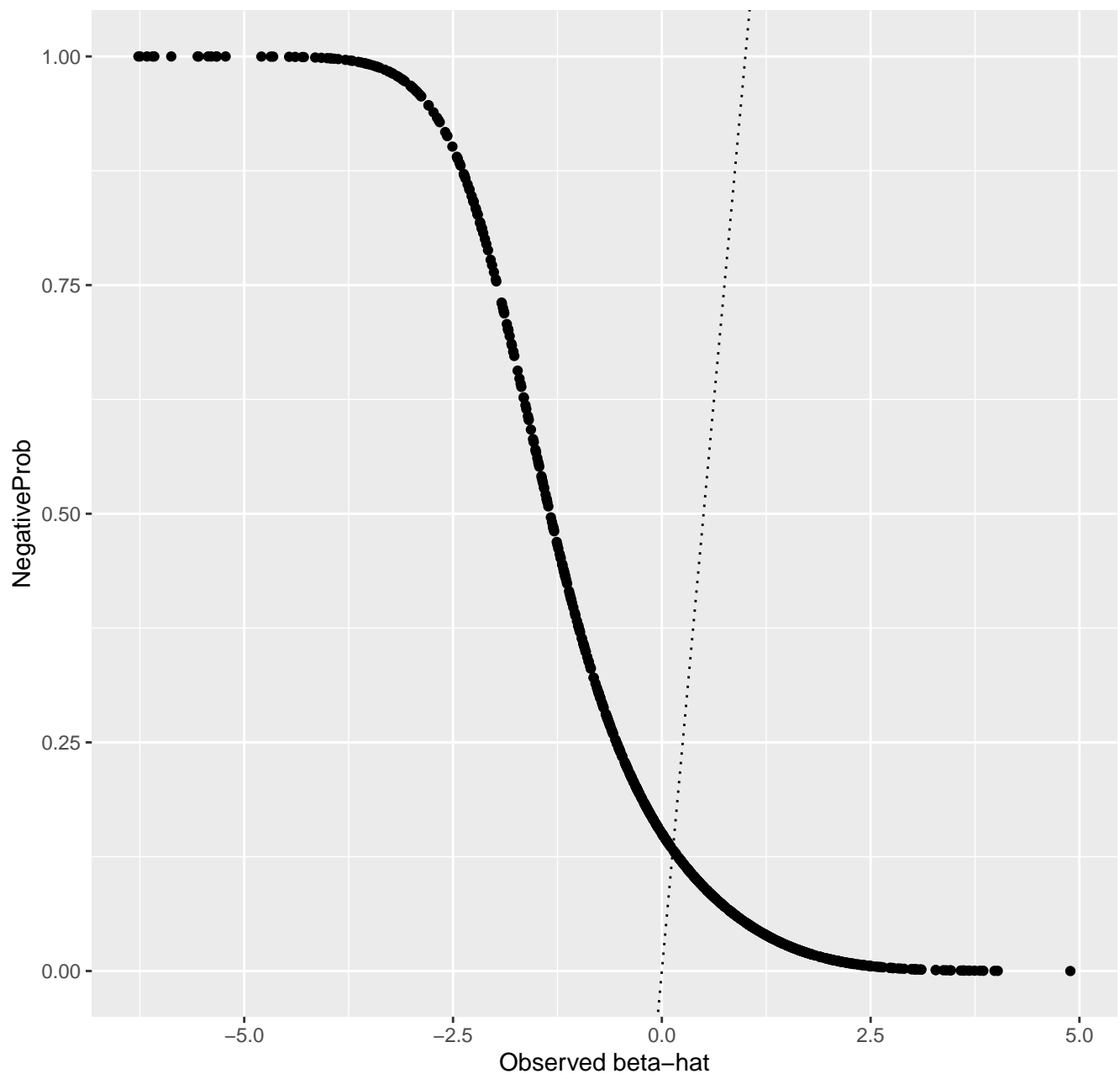
BATF3_(55509) negative



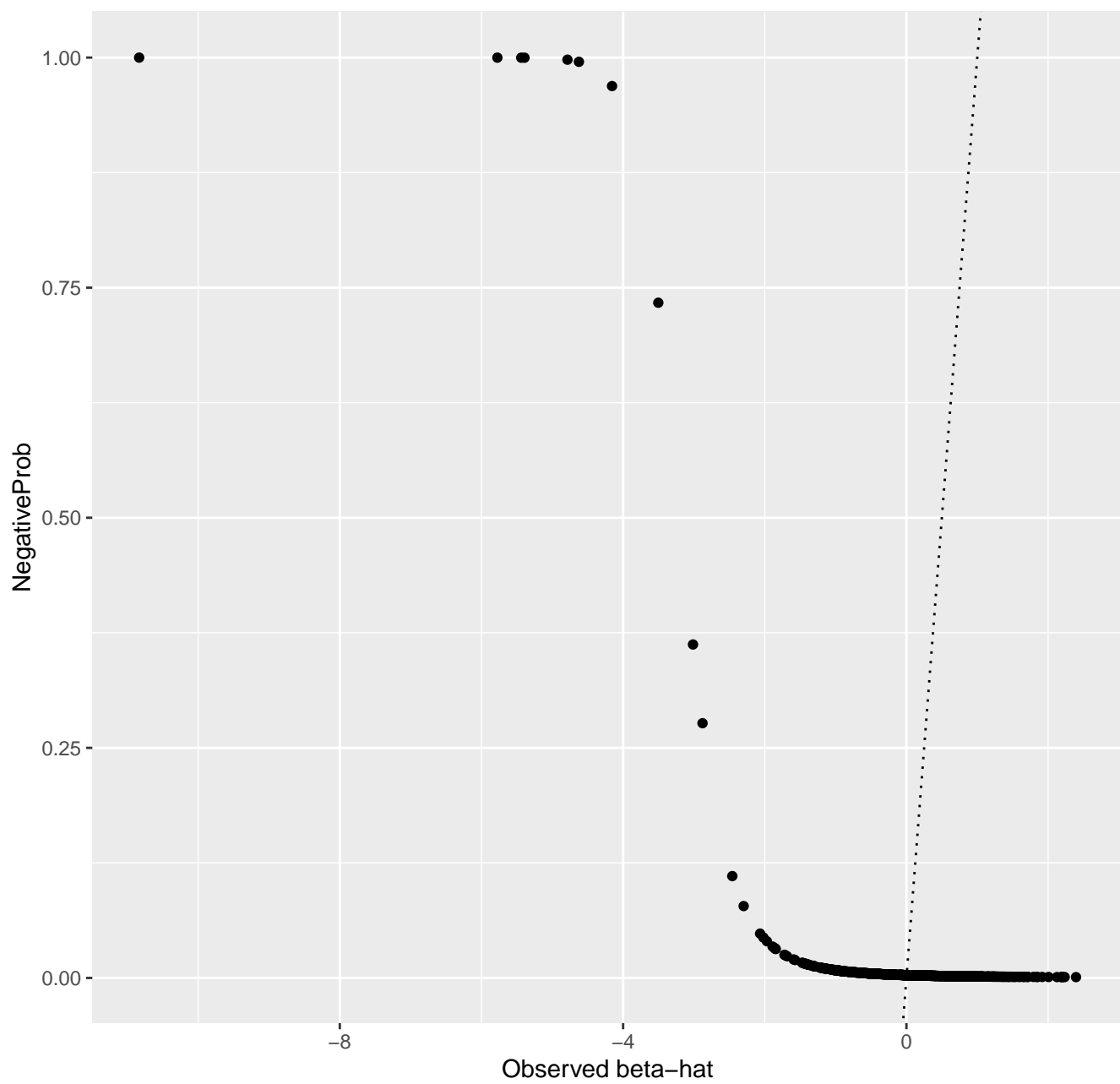
DAPK3_(1613) positive



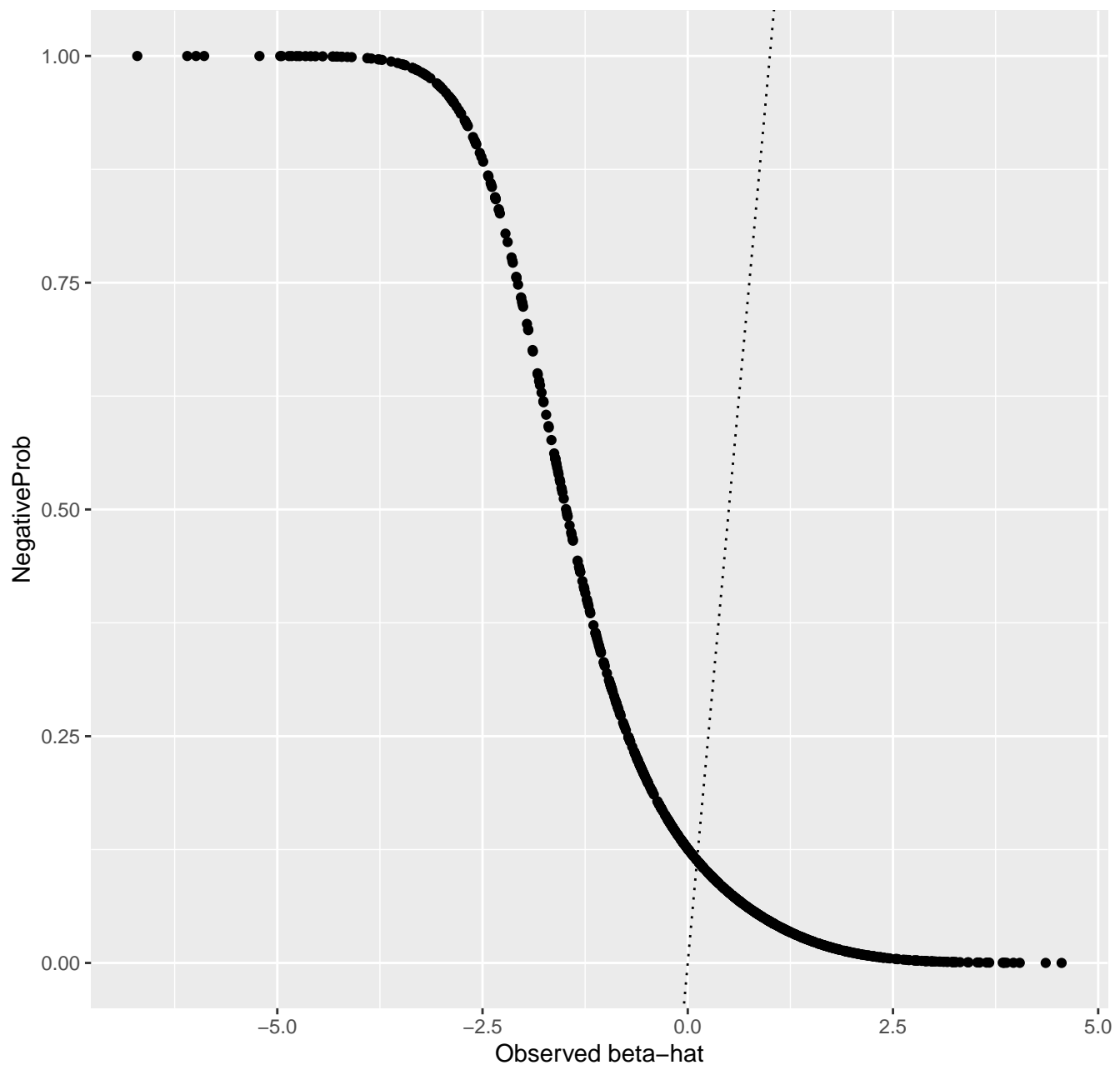
NDUFS8_(4728) negative



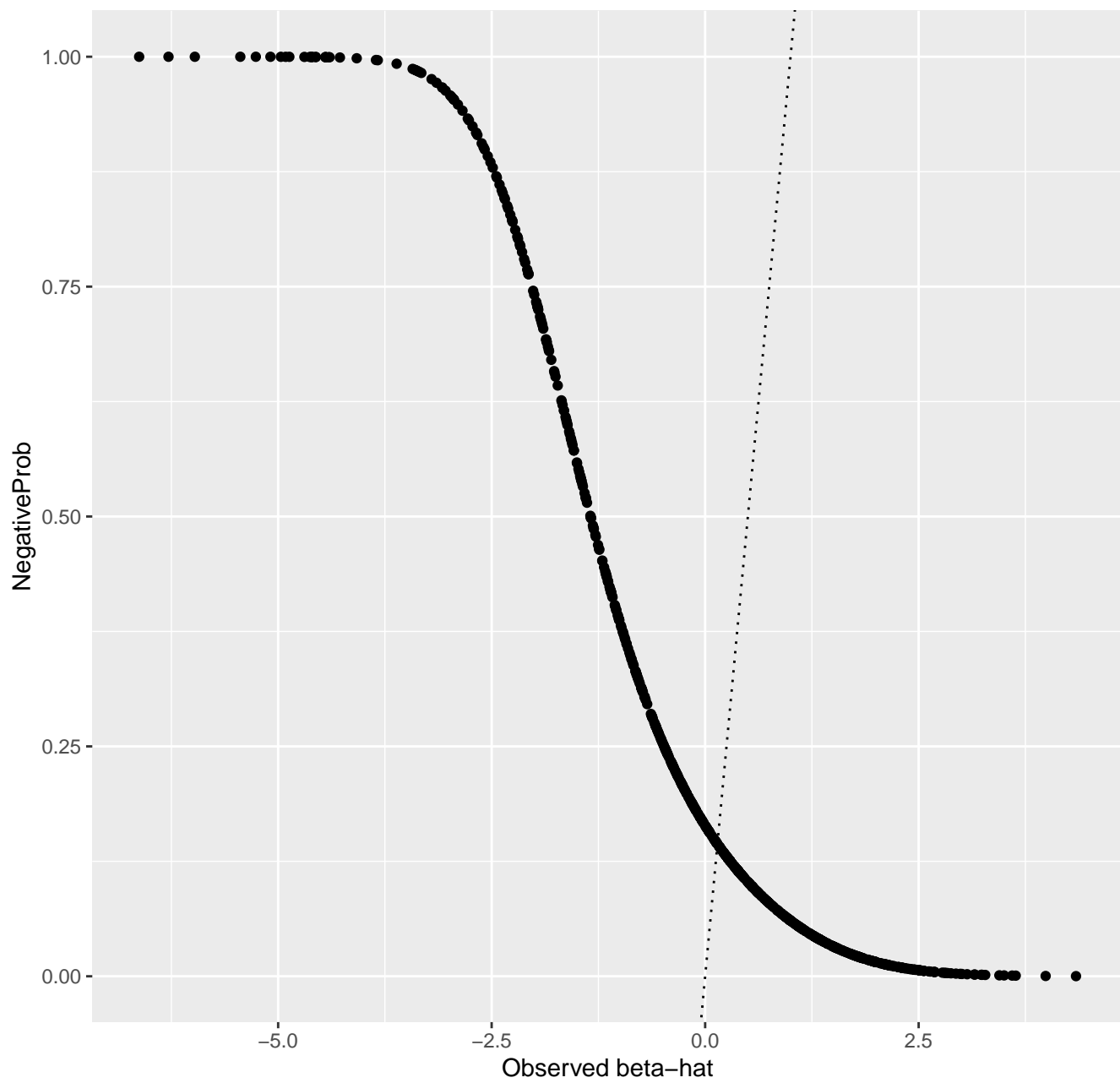
MEF2B_(100271849) negative



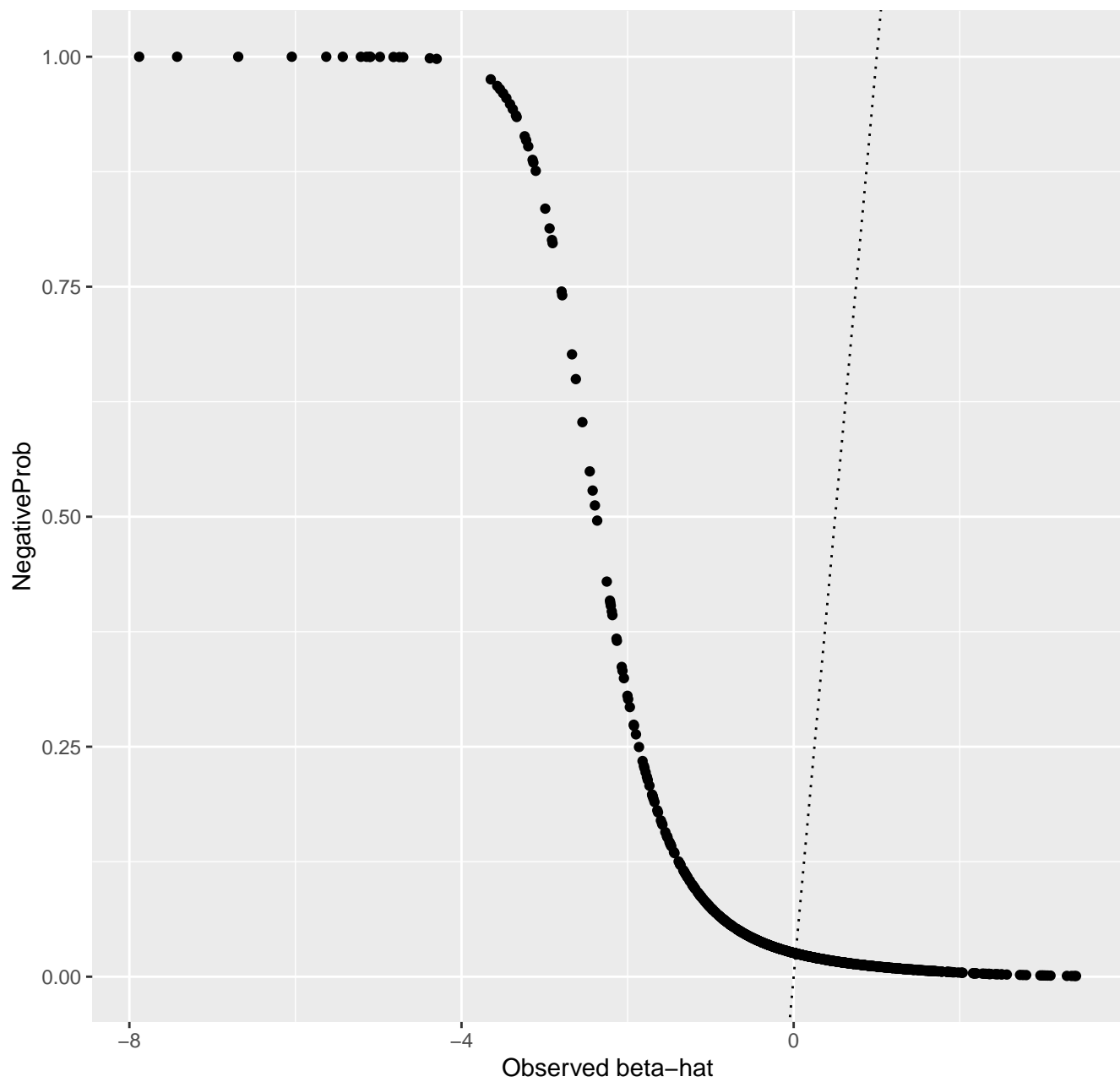
NDUFV2_(4729) negative



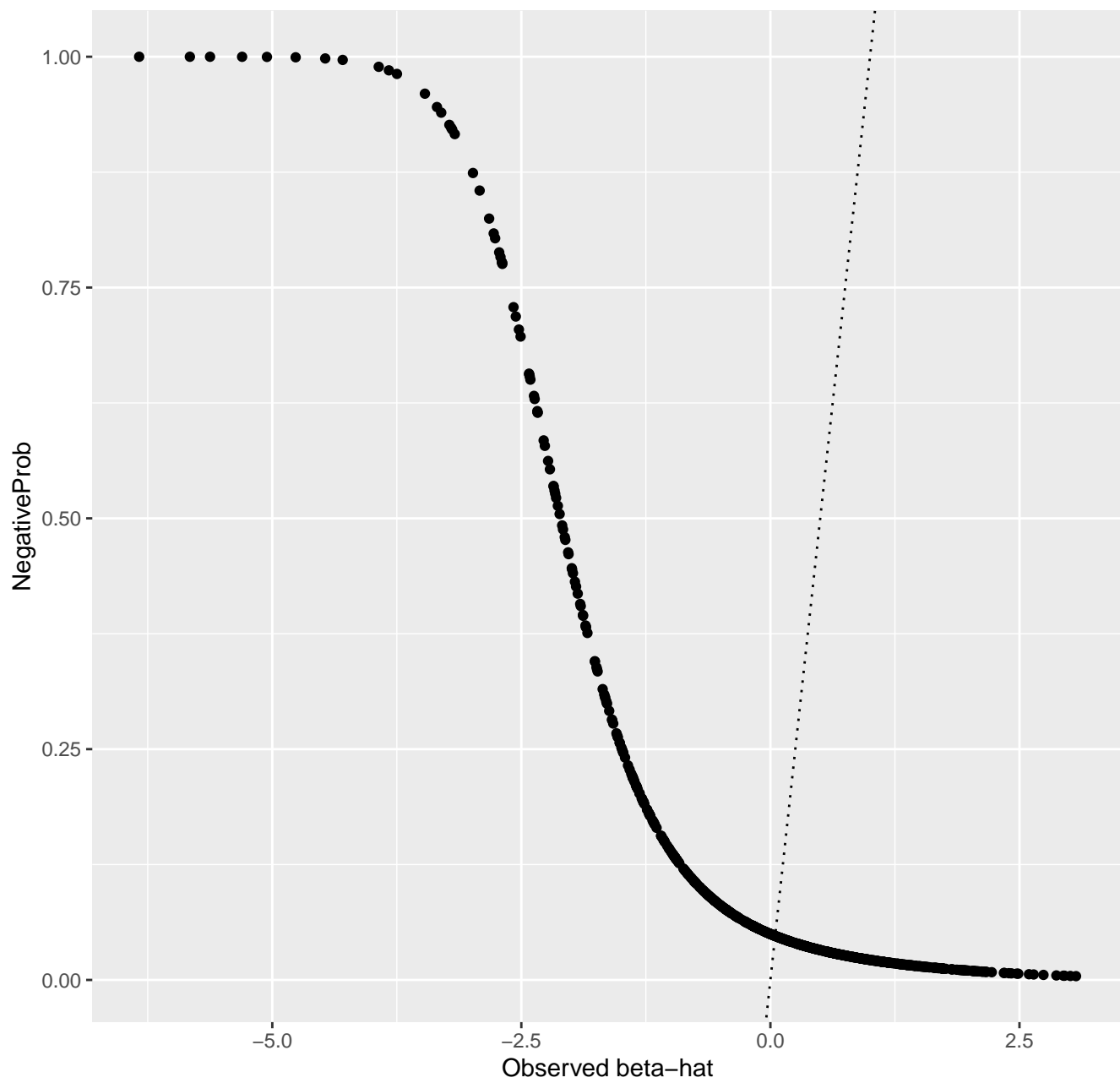
VMA21_(203547) negative



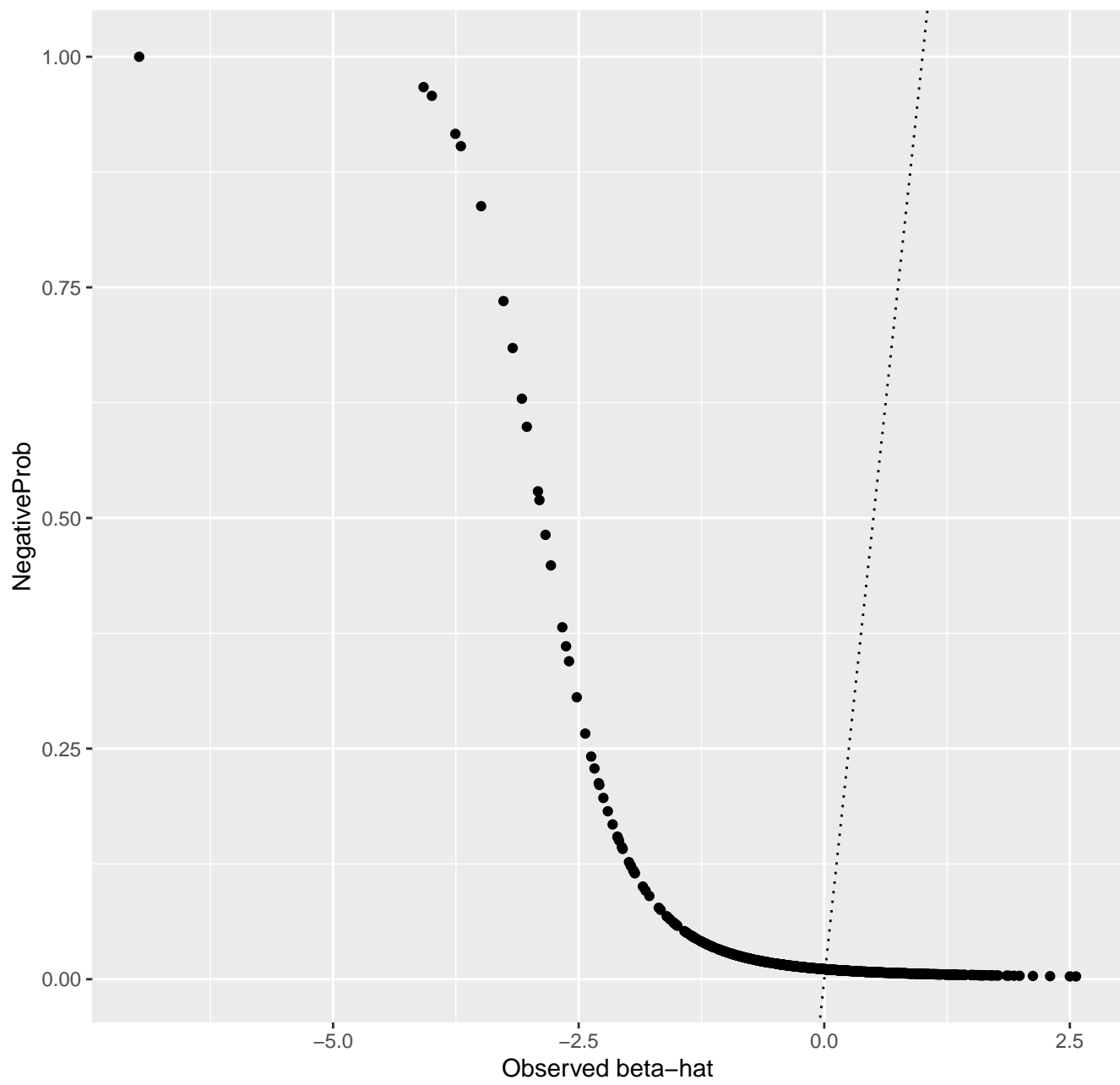
ALAD_(210) negative



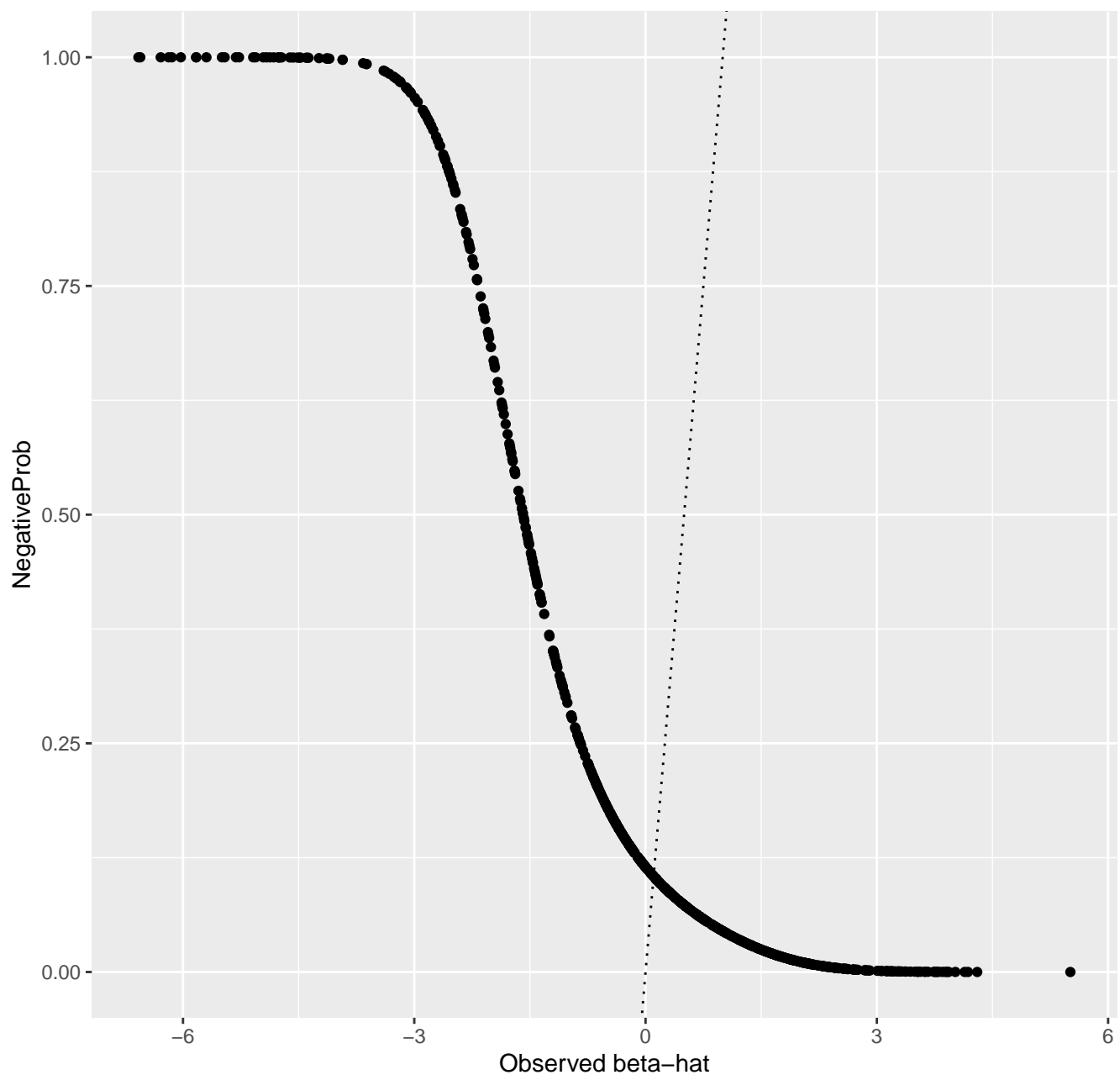
ATPAF2_(91647) negative



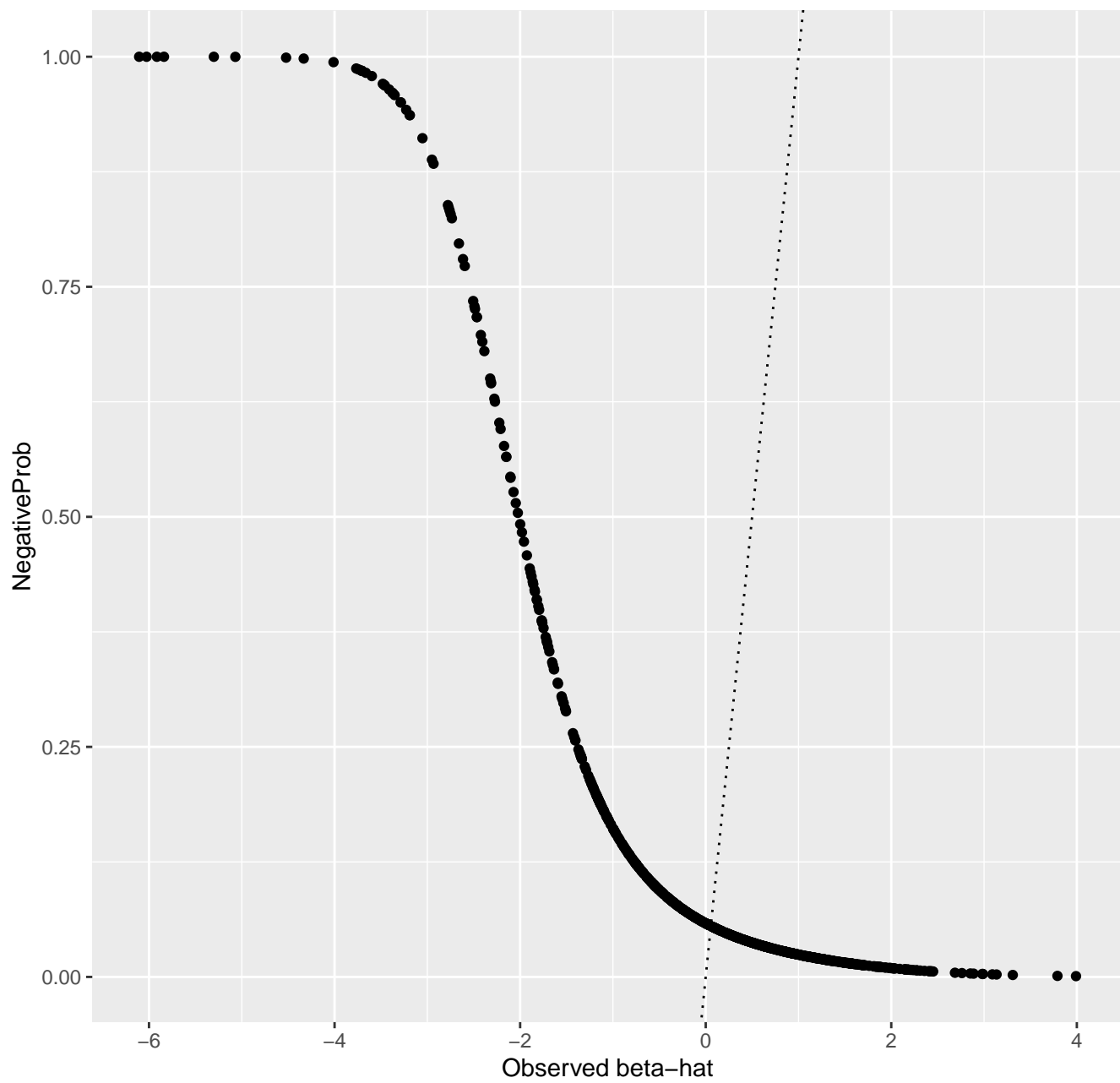
PDXK_(8566) negative



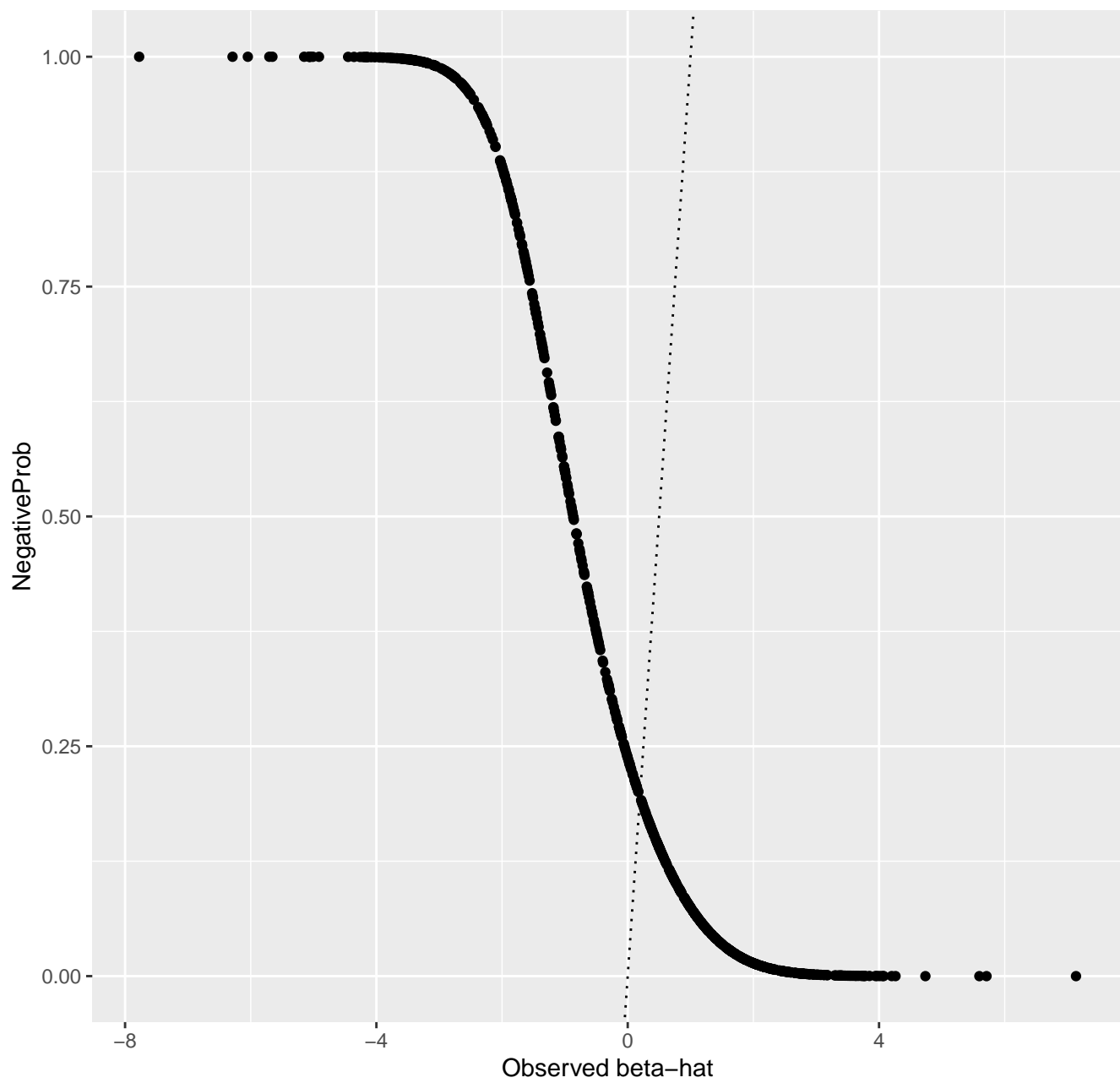
NDUFS1_(4719) negative



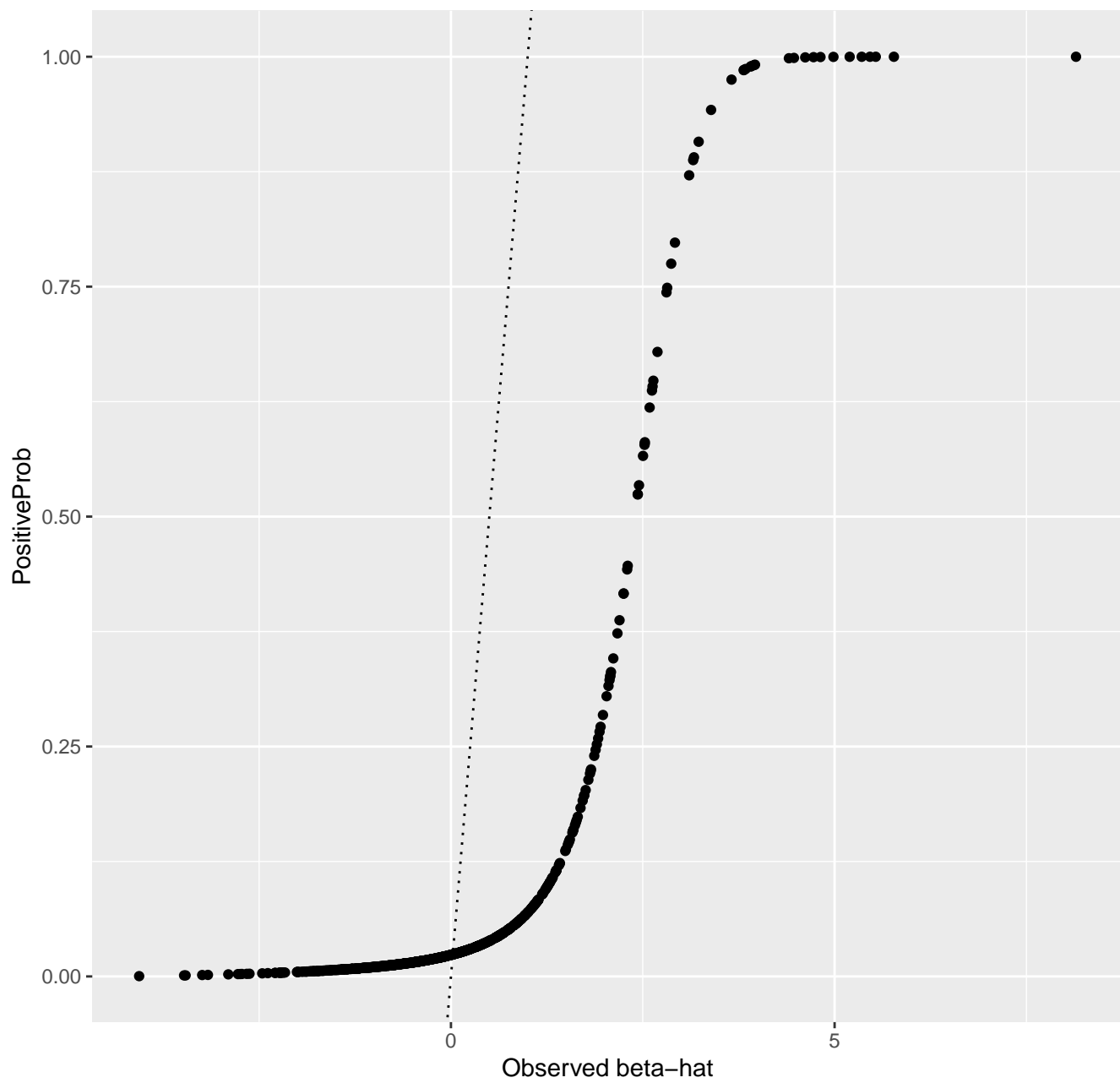
ZNF638_(27332) negative



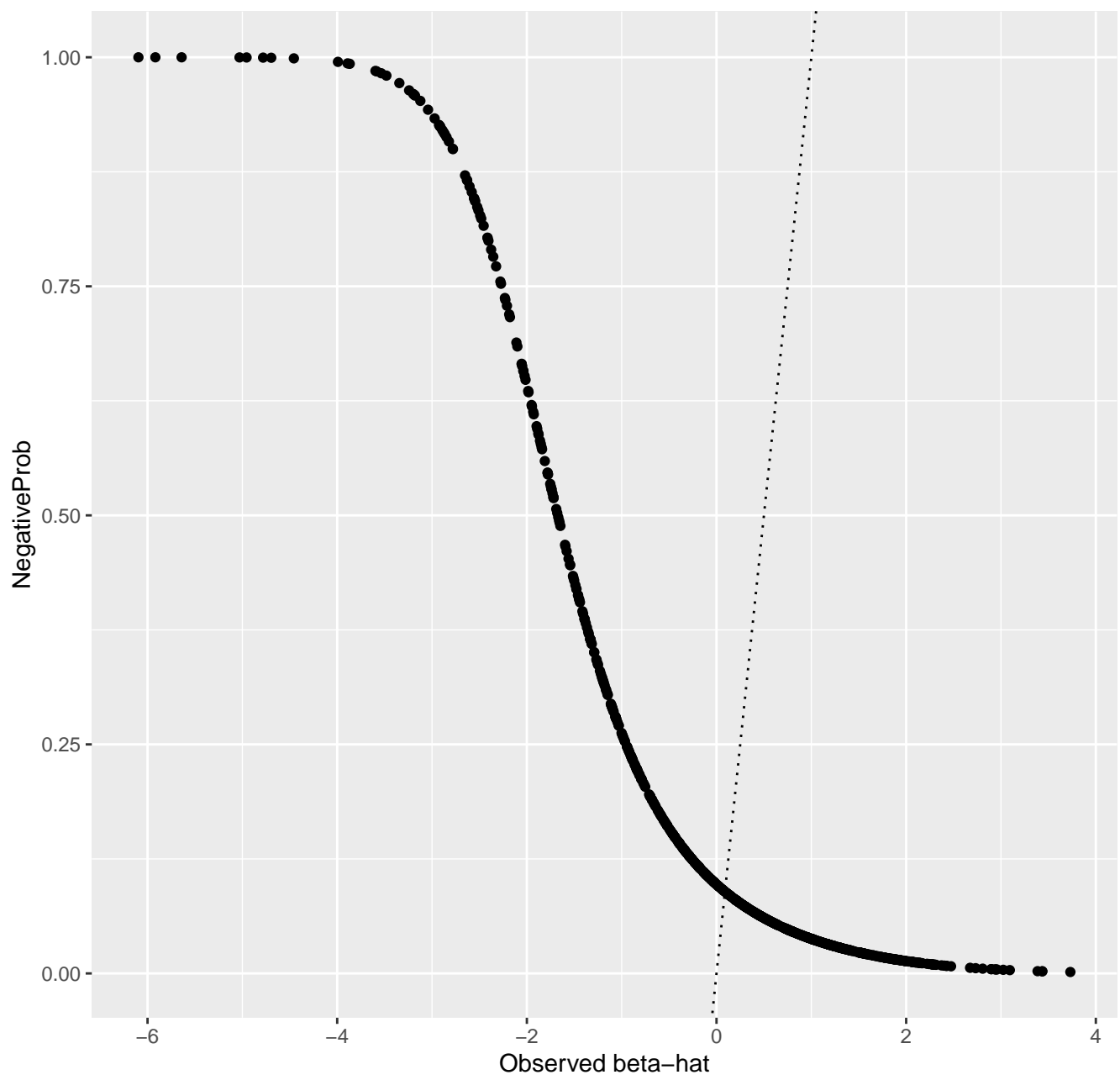
ADSS_(159) negative



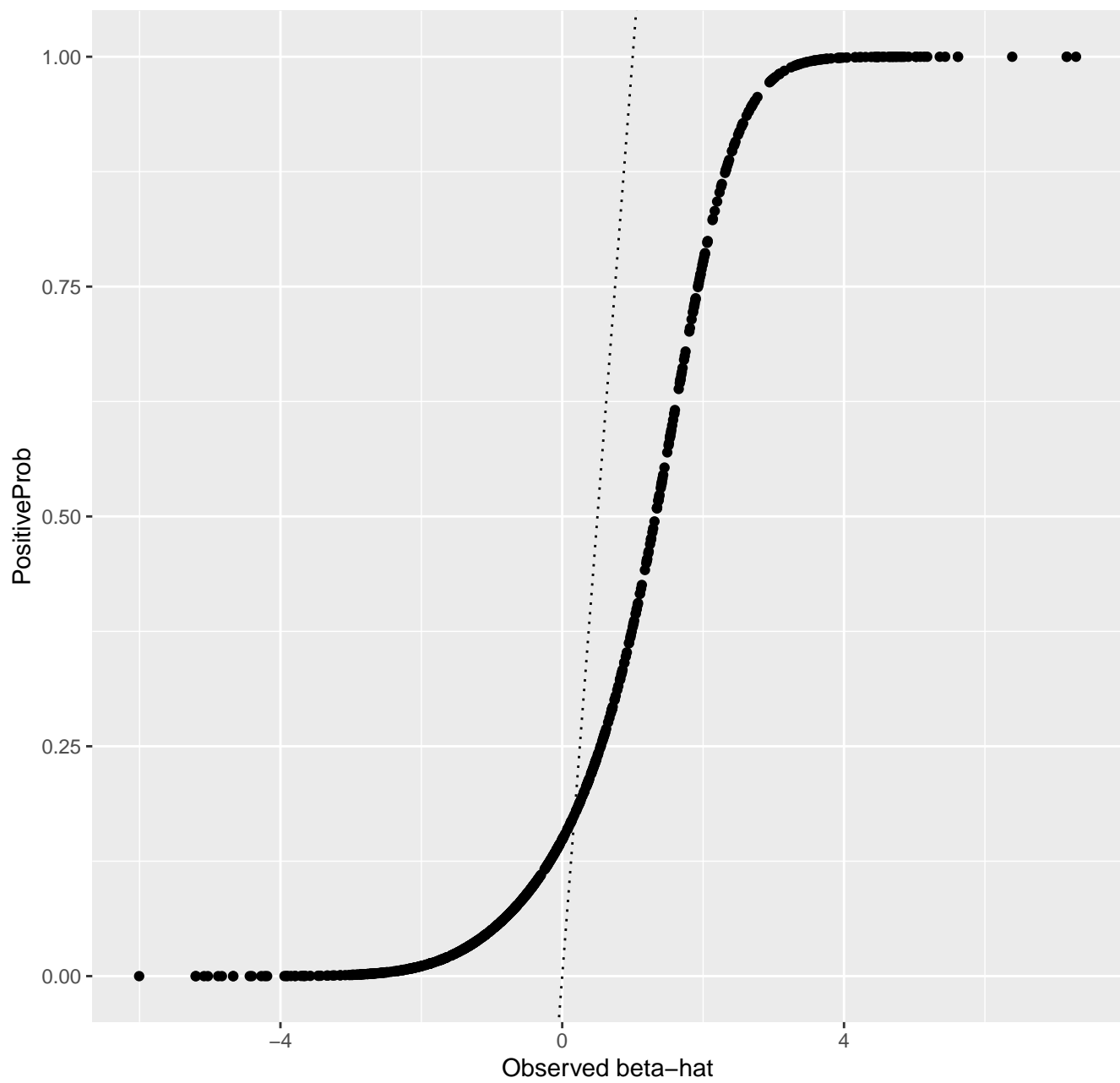
USP28_(57646) positive



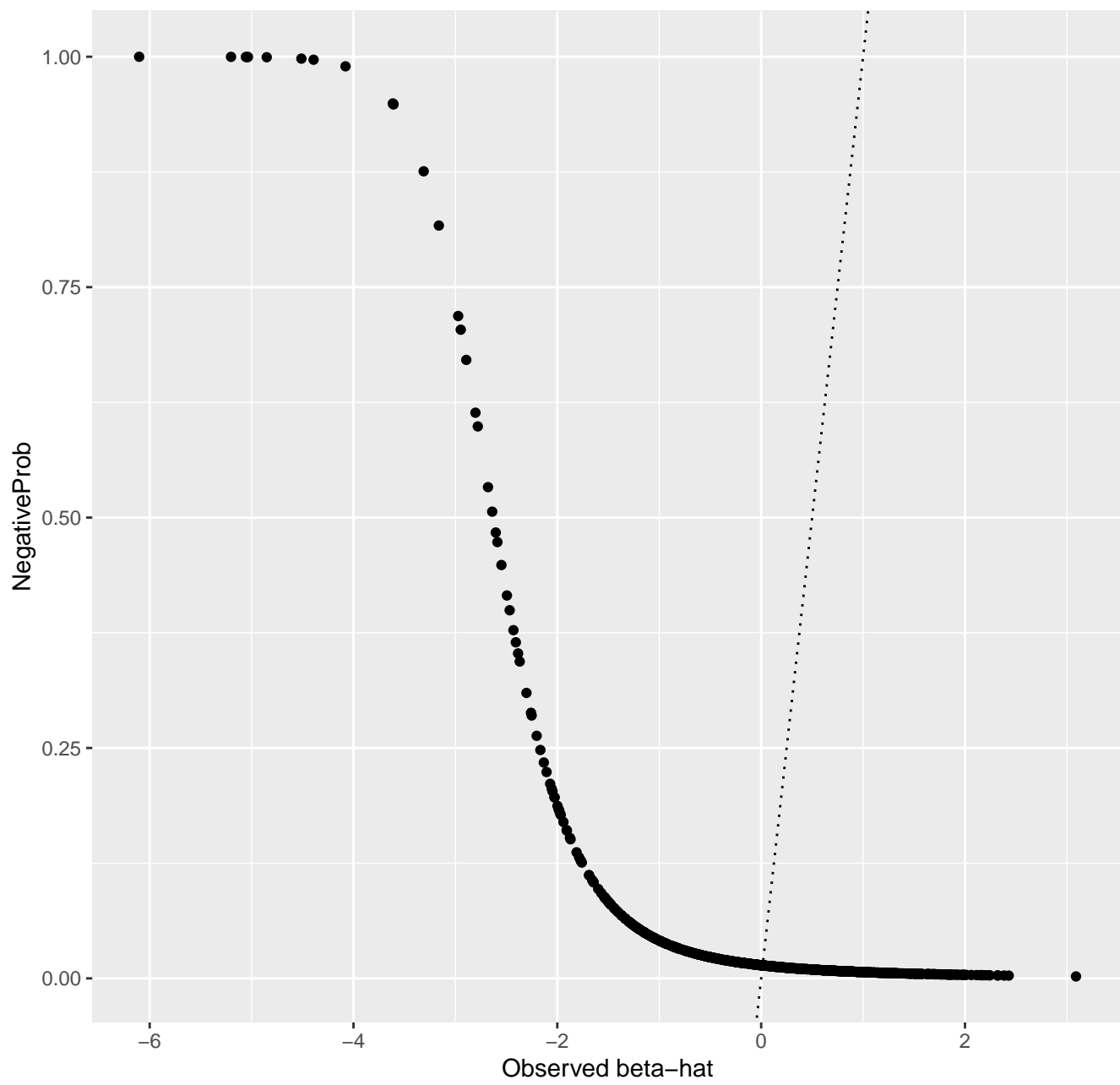
NUPL2_(11097) negative



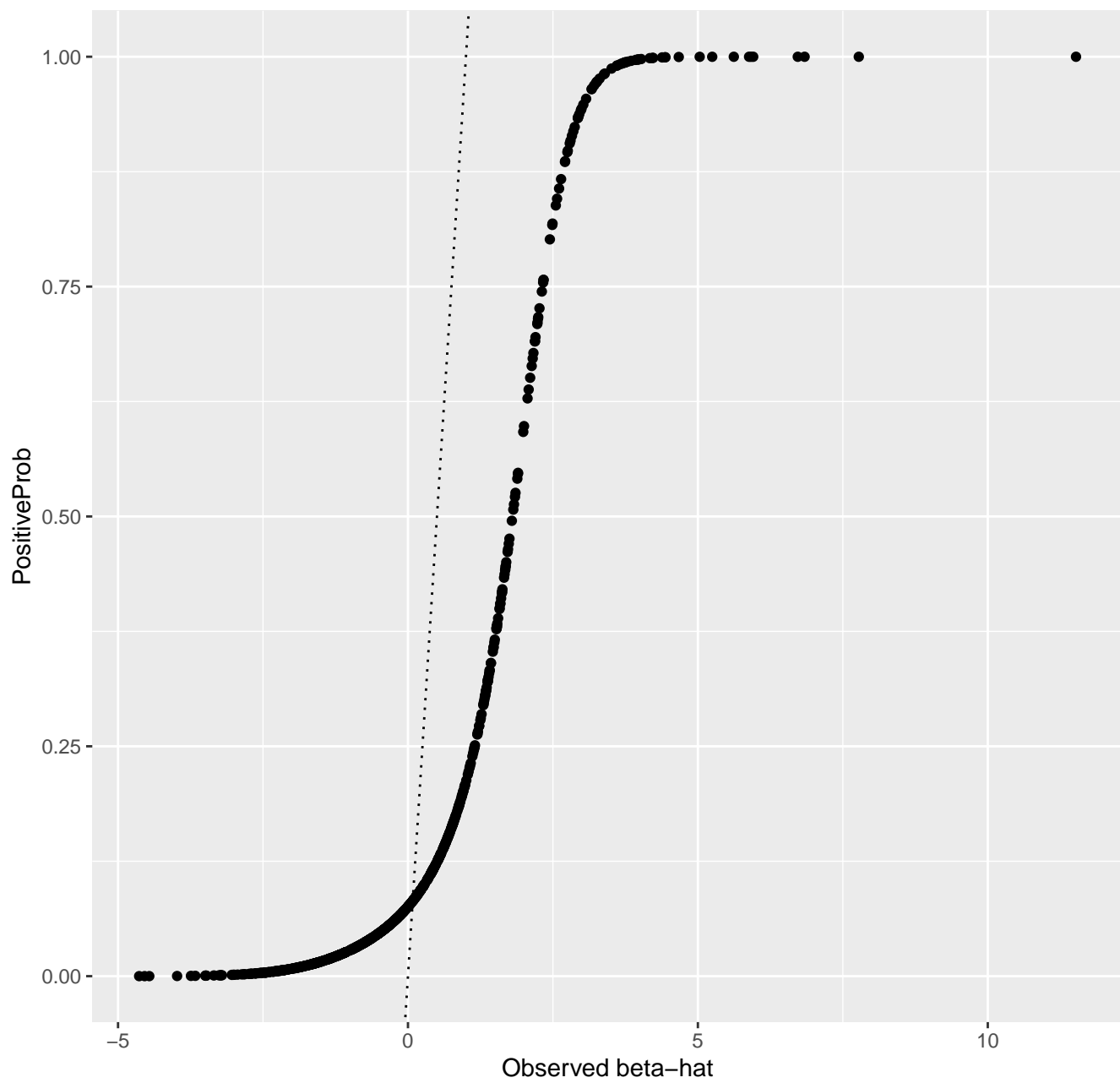
MBTPS1_(8720) positive



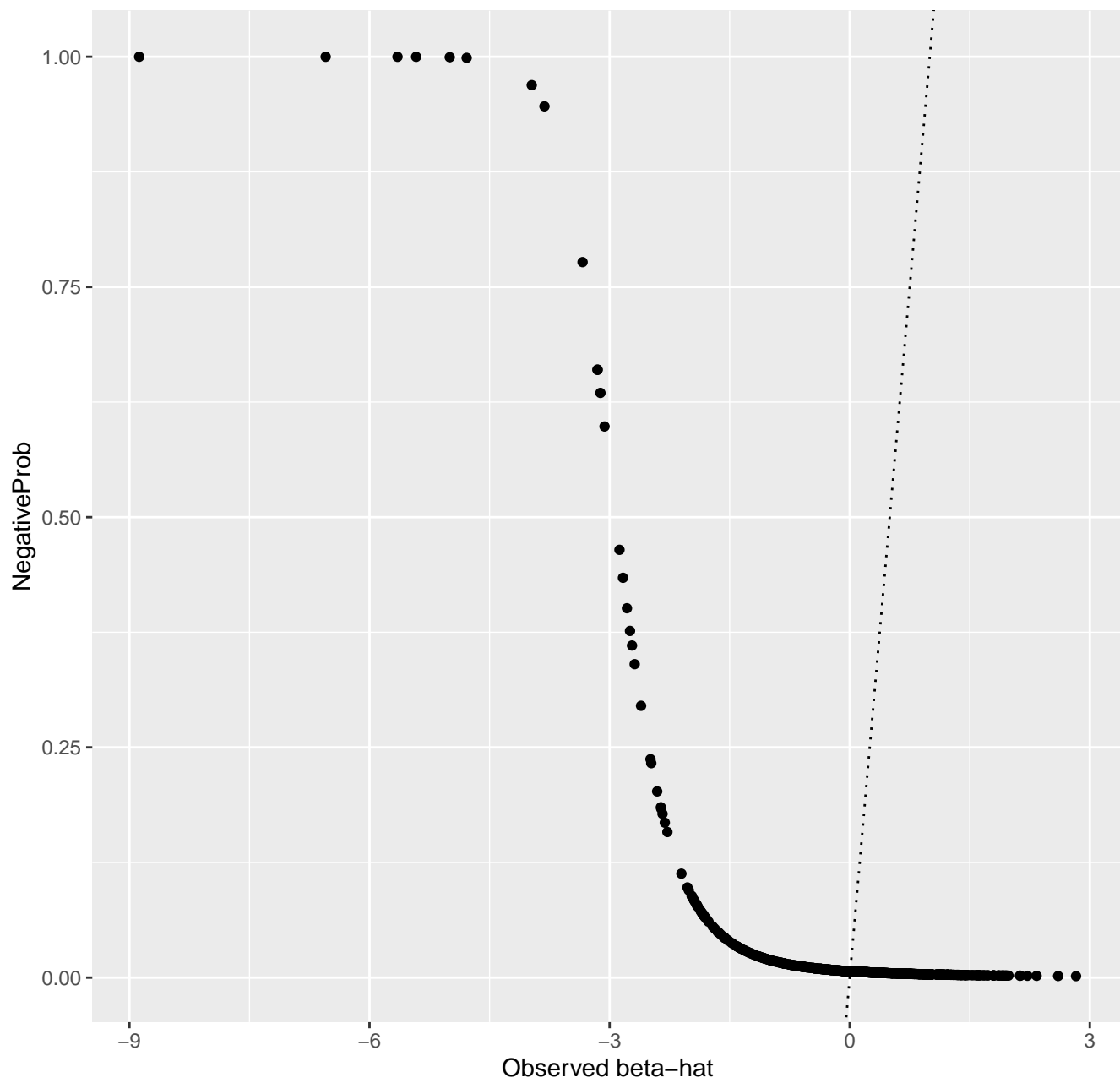
UBE2Z_(65264) negative



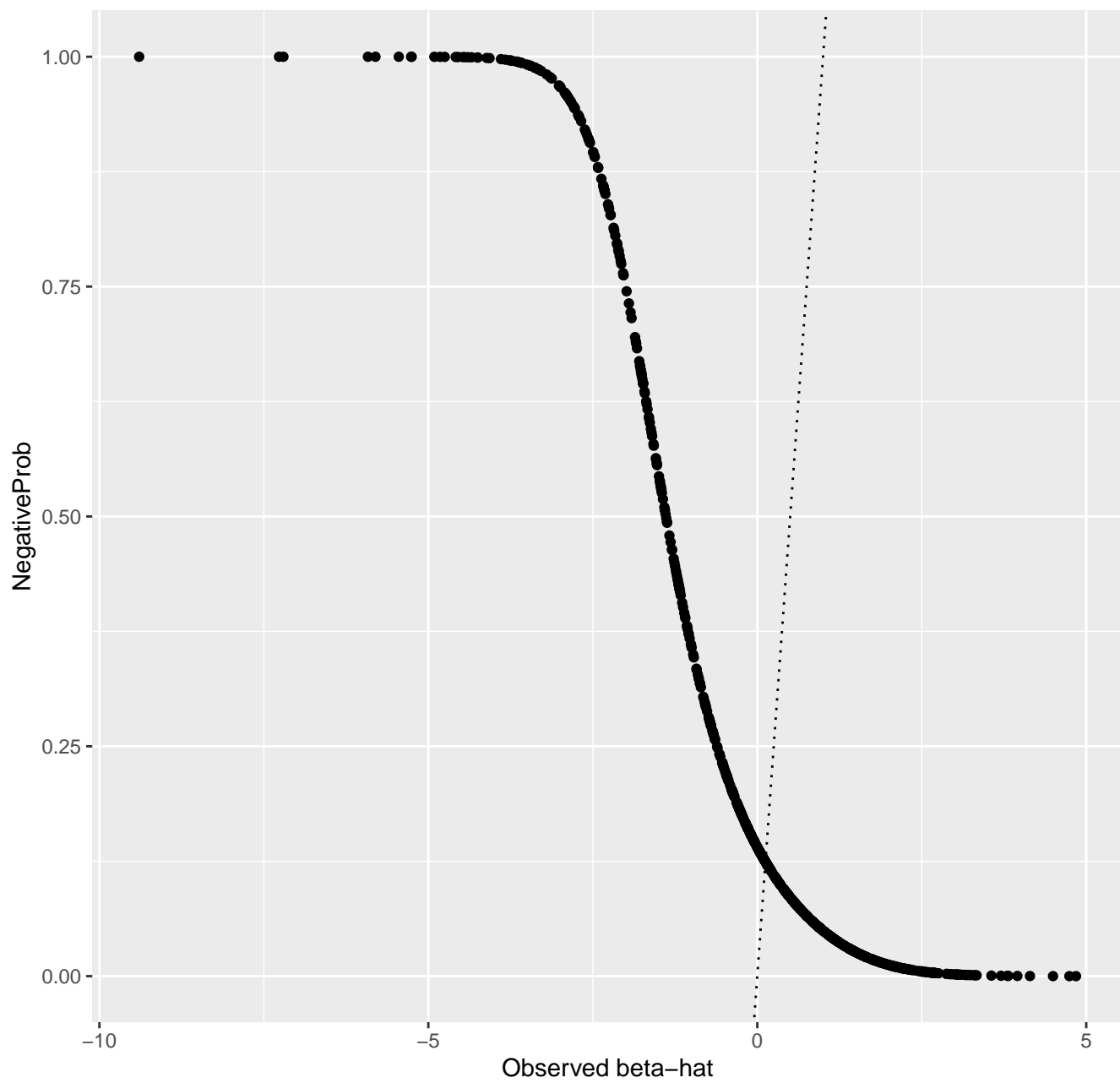
MAP4K4_(9448) positive



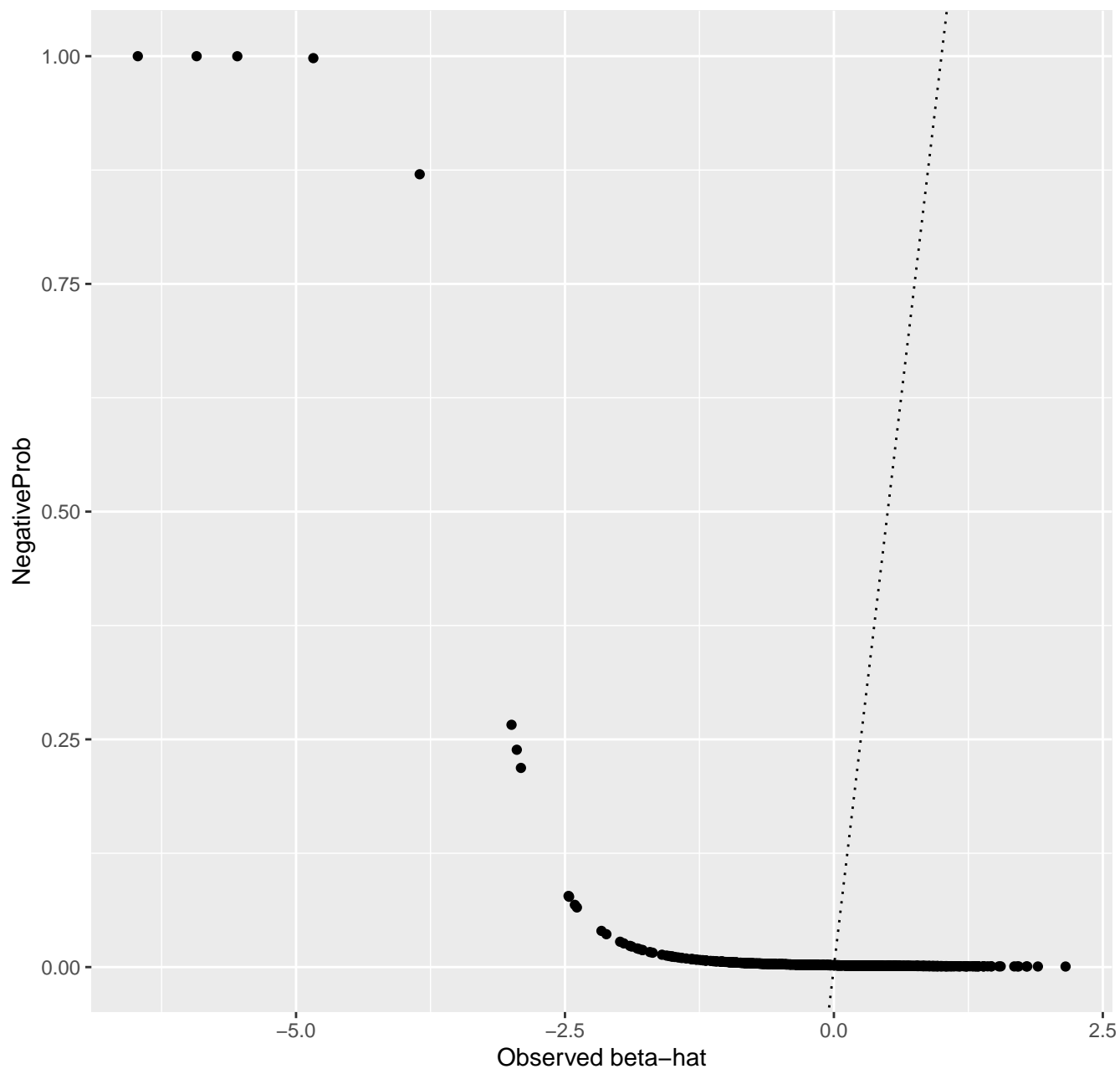
ZNFX1_(57169) negative



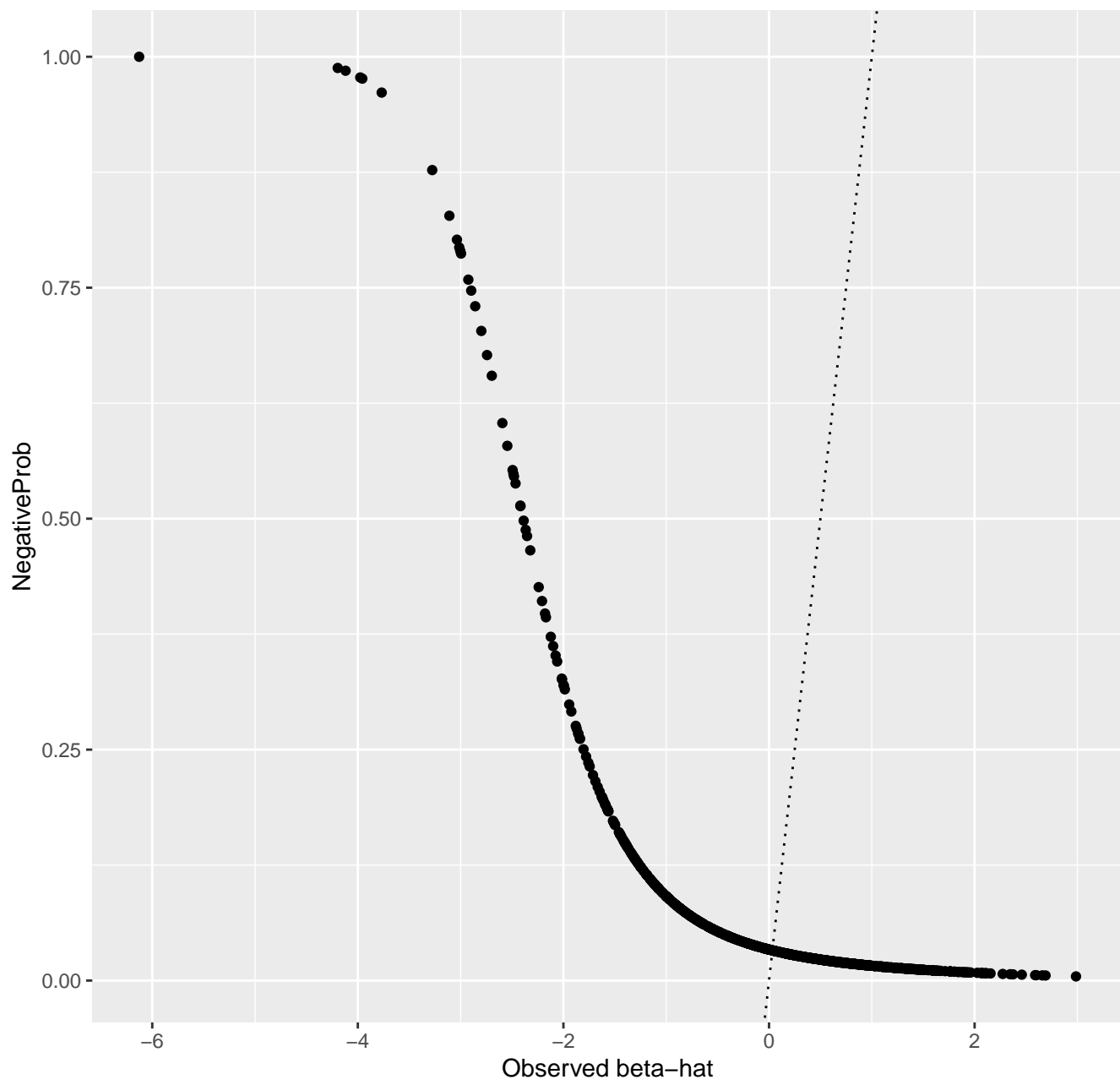
CCNL1_(57018) negative



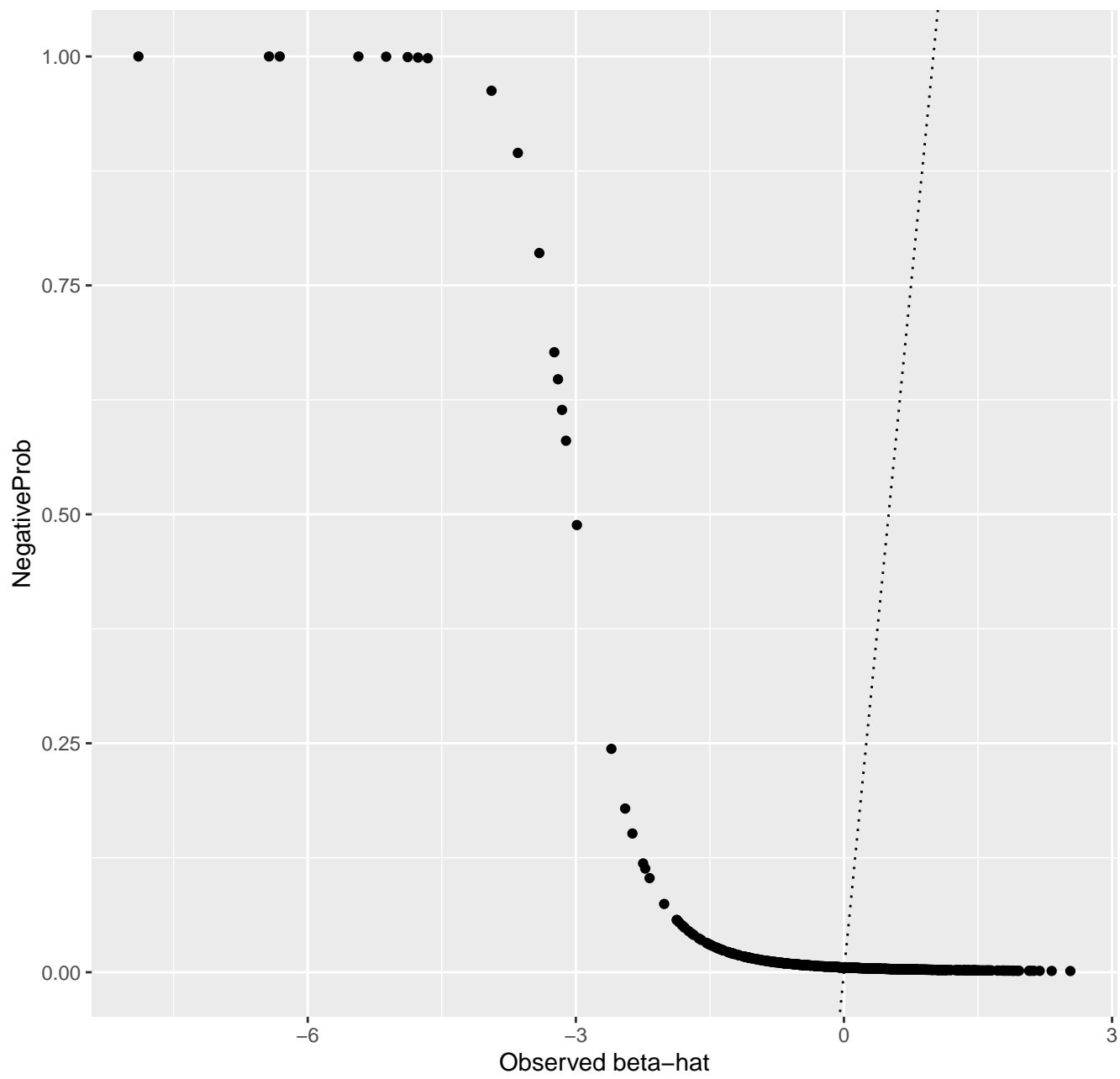
AGR2_(10551) negative



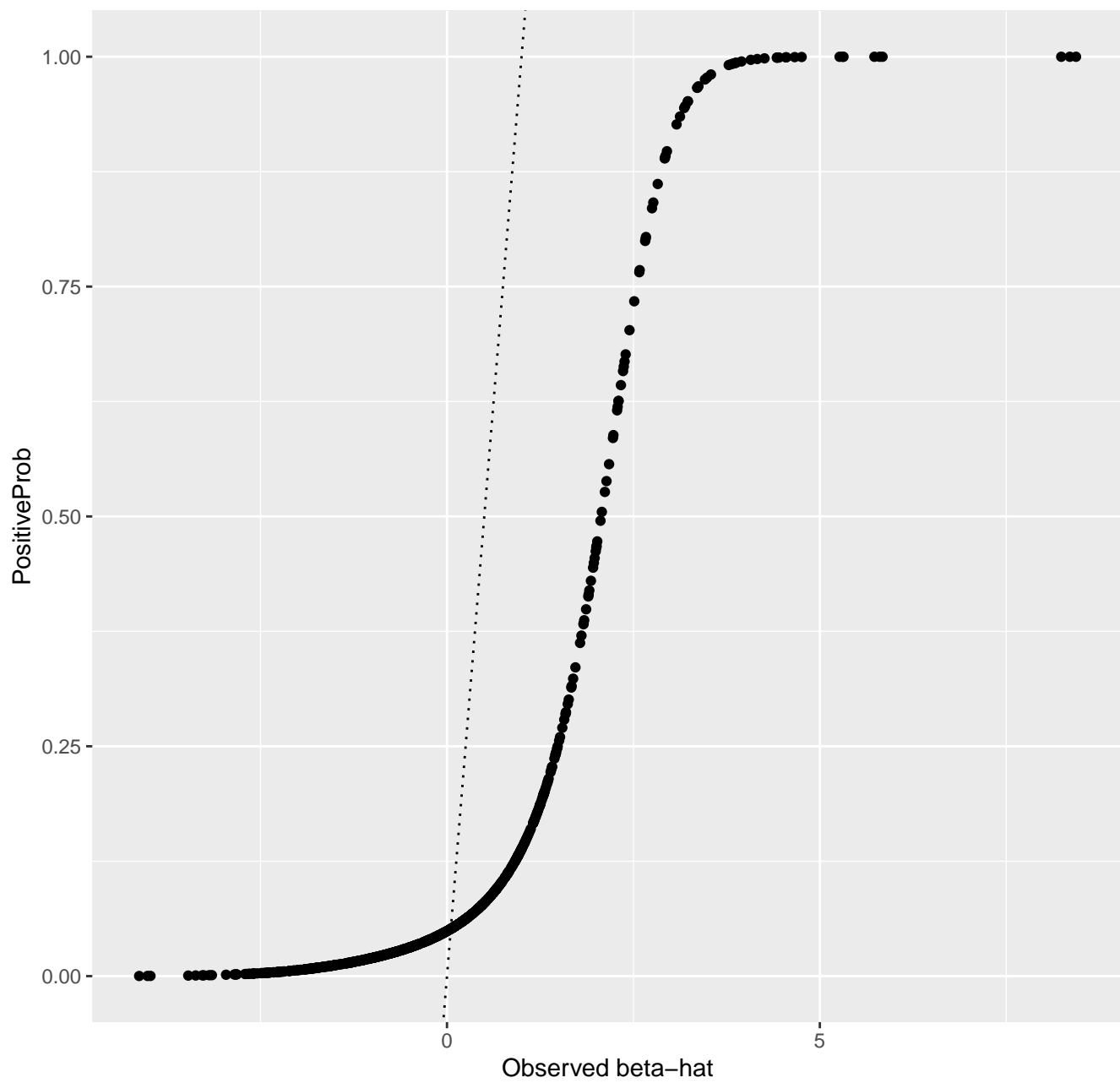
CALM2_(805) negative



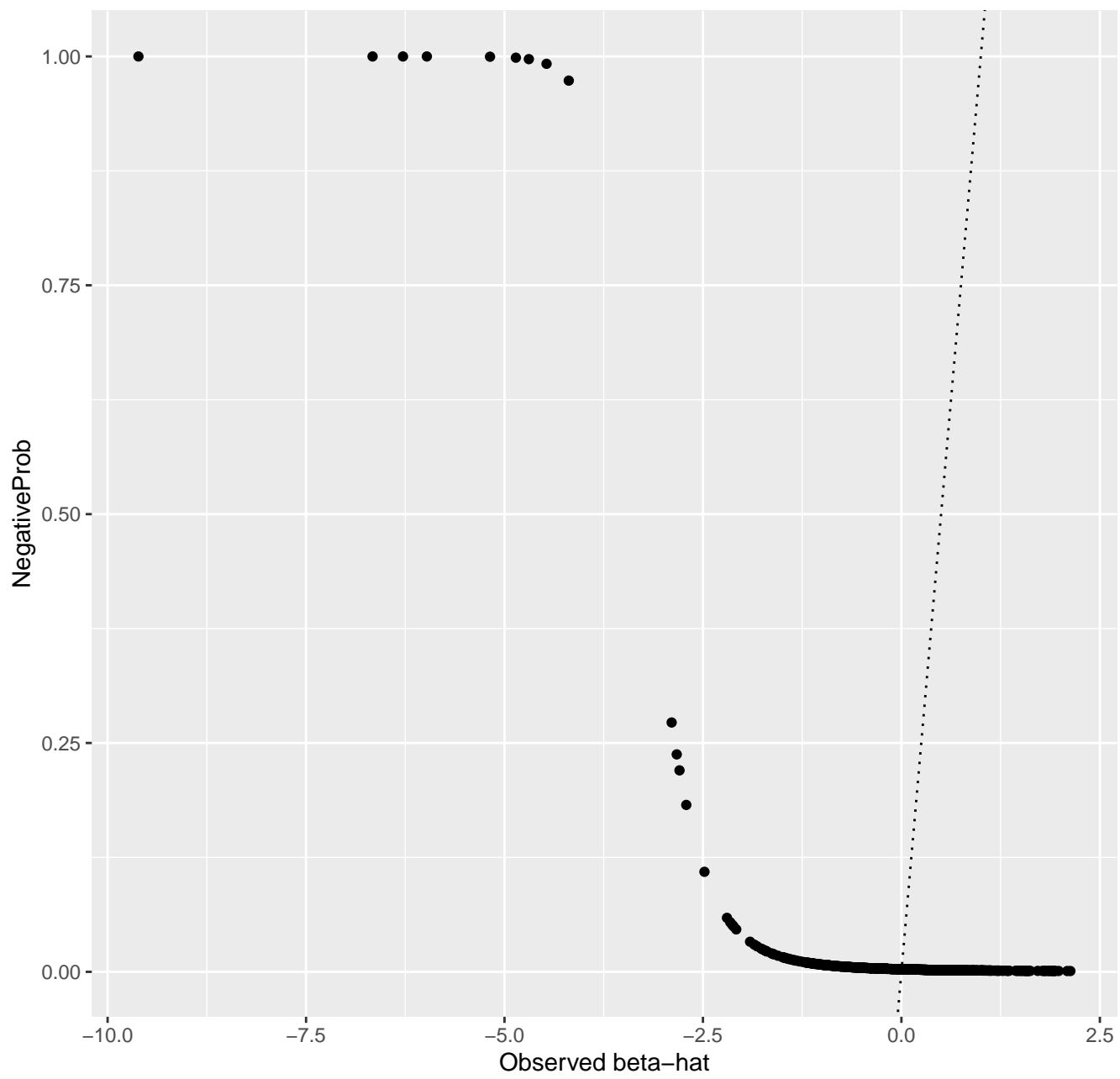
PIM2_(11040) negative



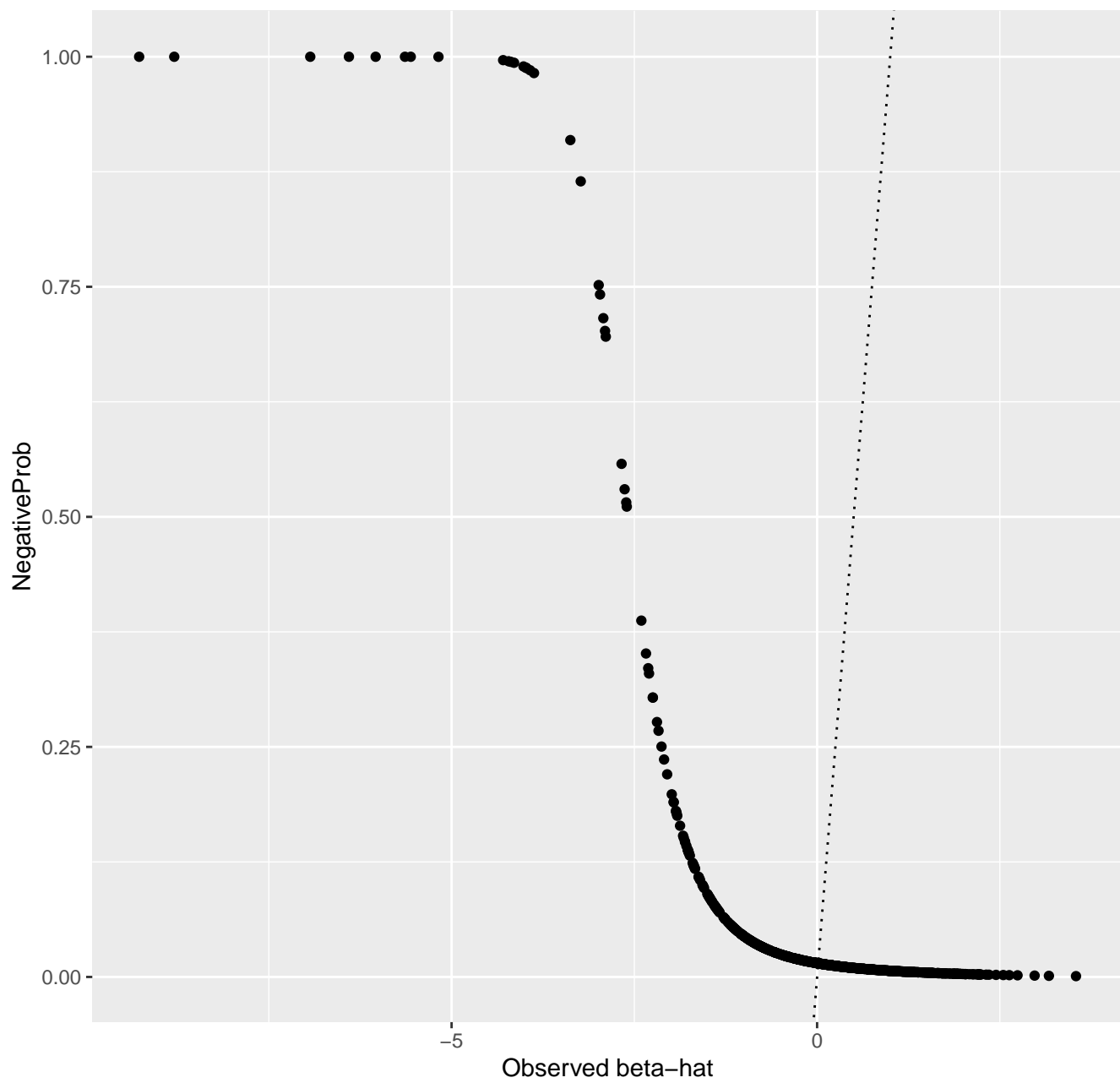
RB1_(5925) positive



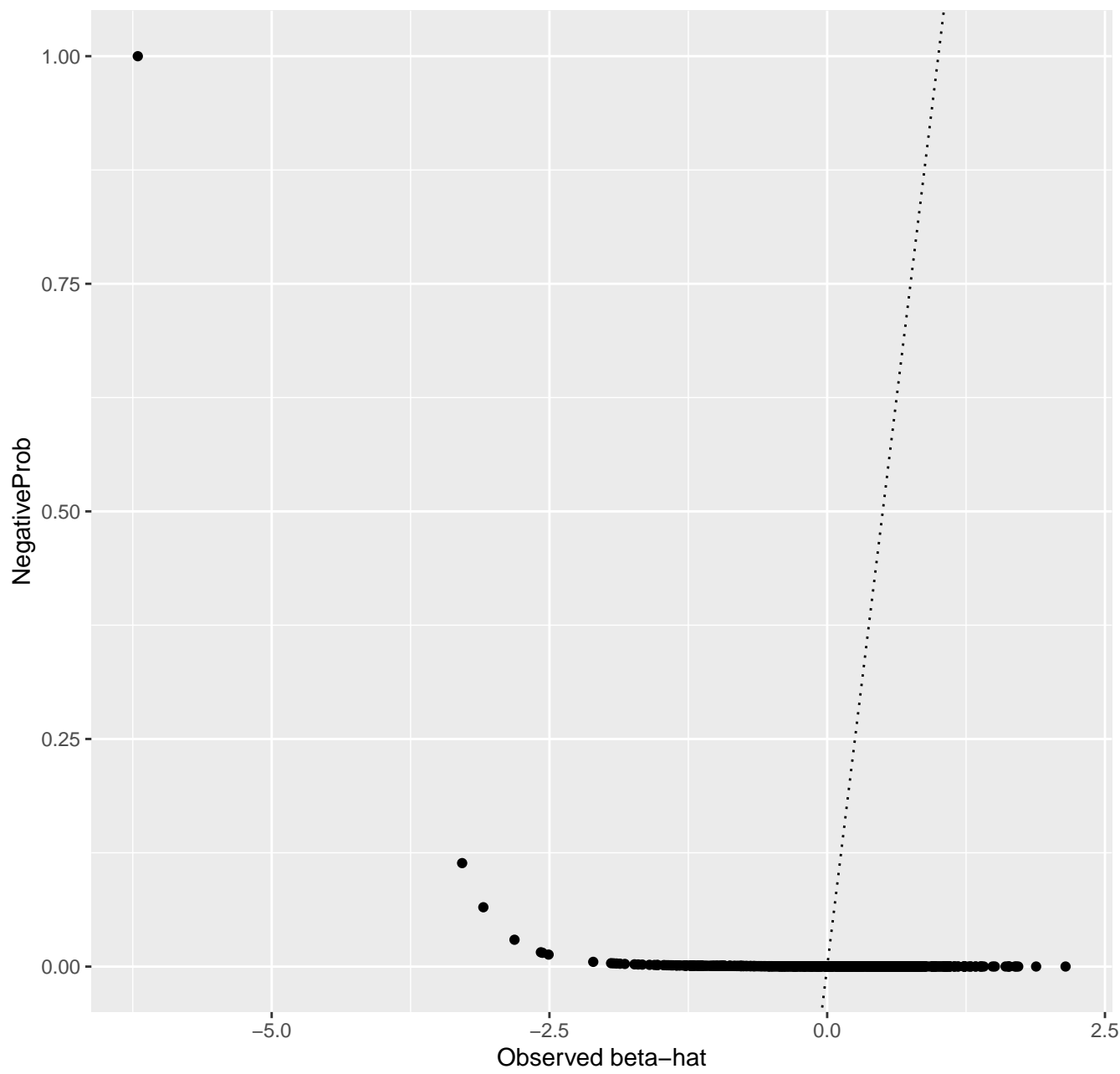
JUP_(3728) negative



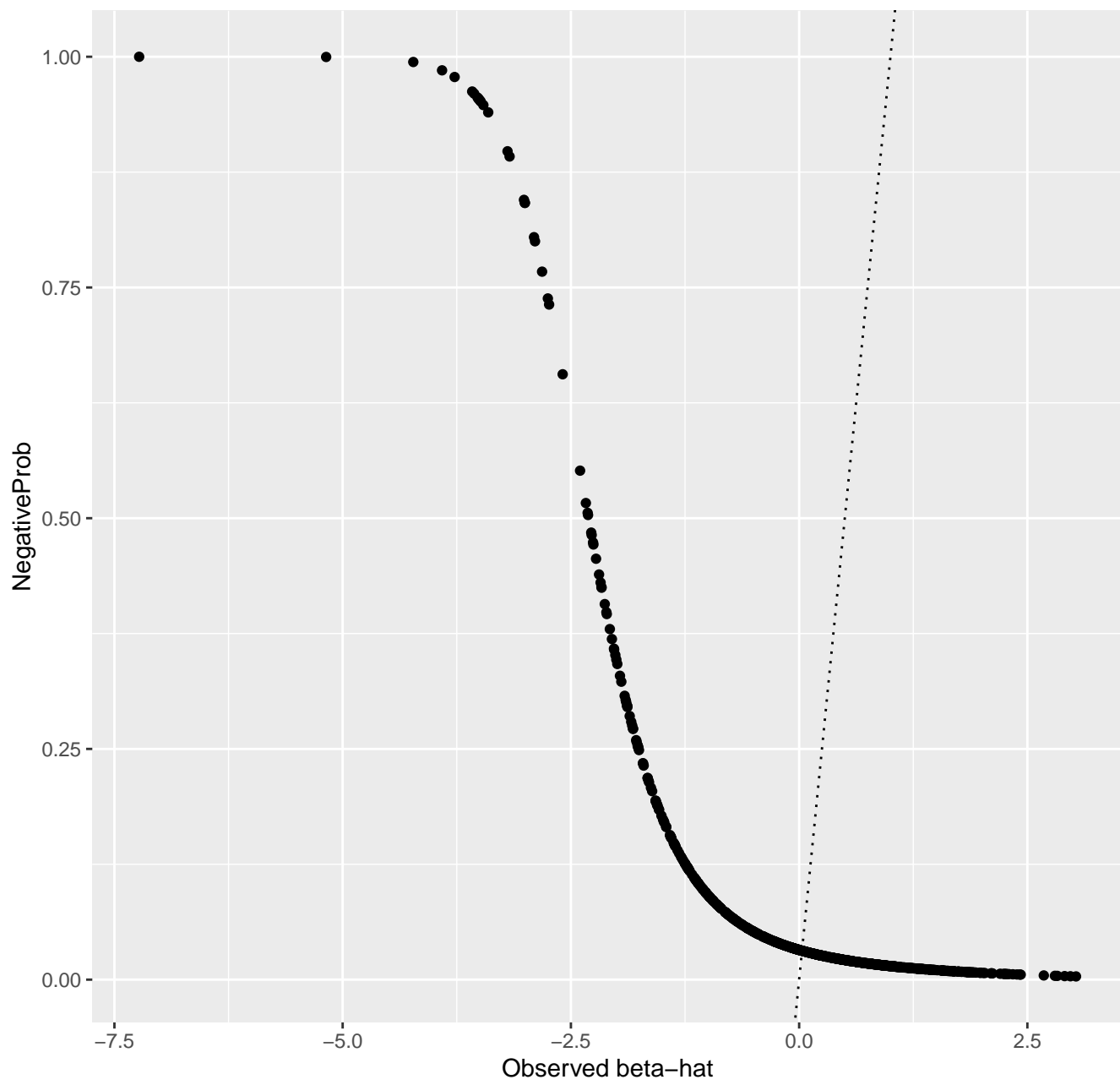
PSMD10_(5716) negative



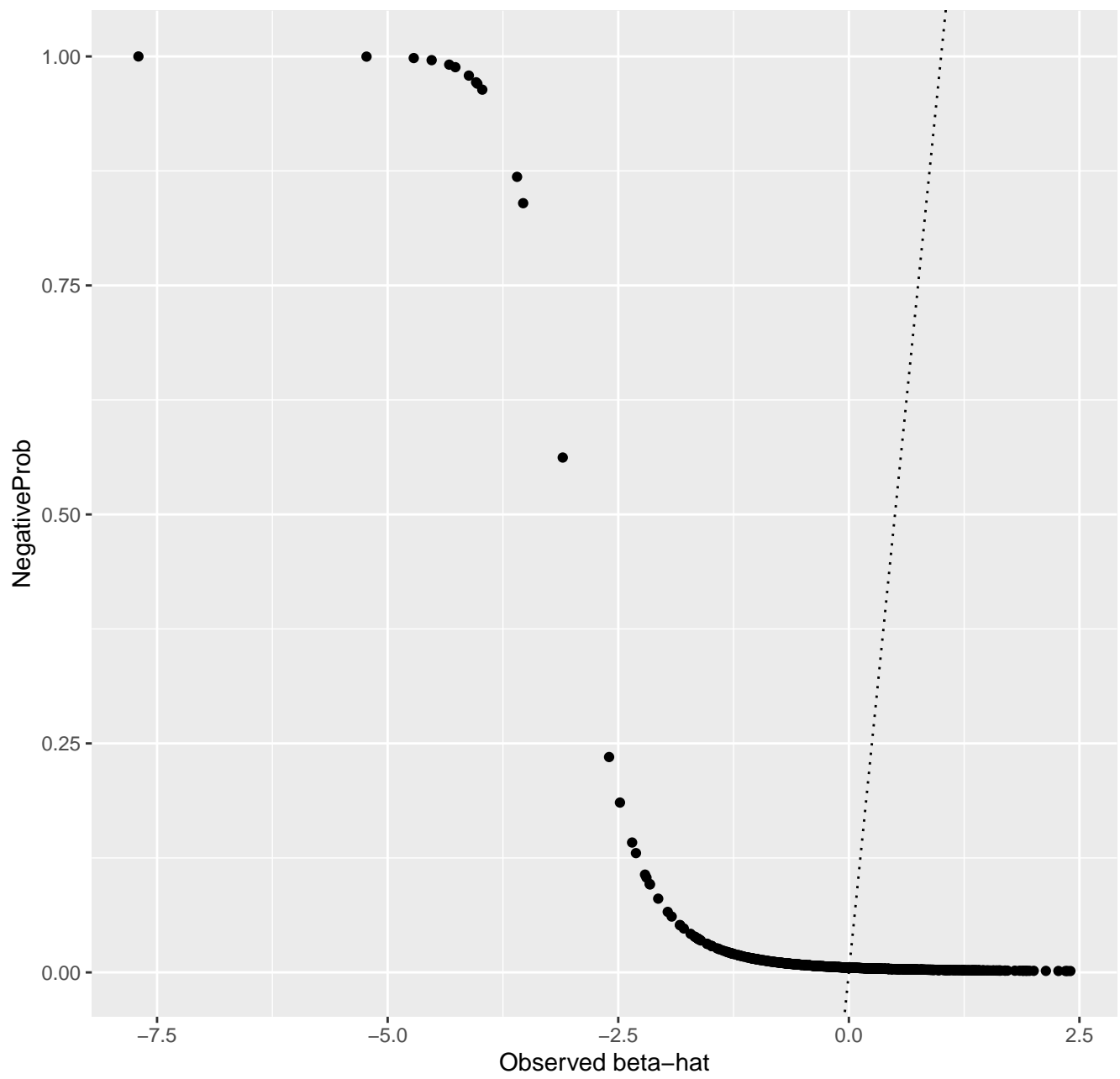
INSL6_(11172) negative



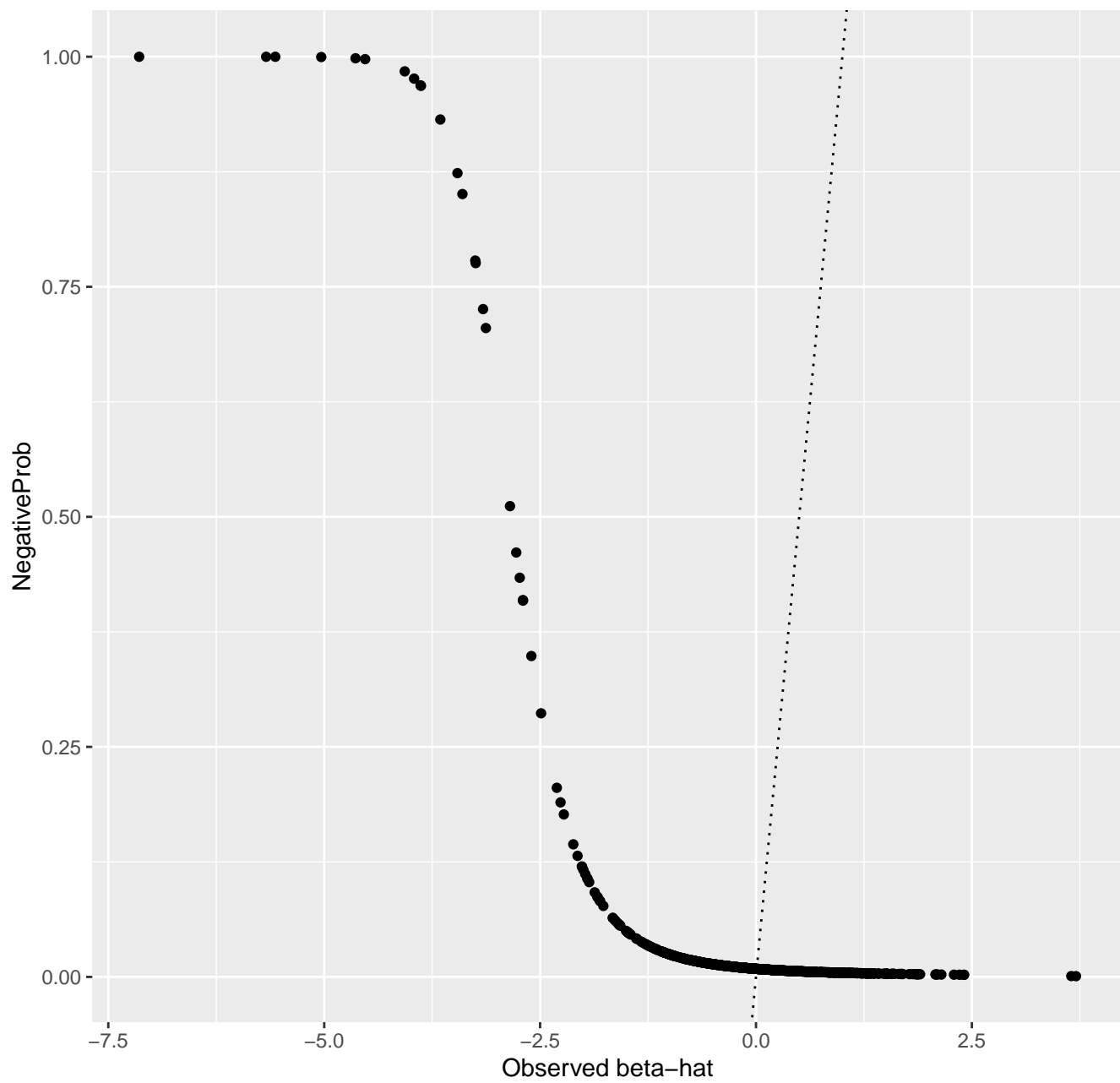
KIF15_(56992) negative



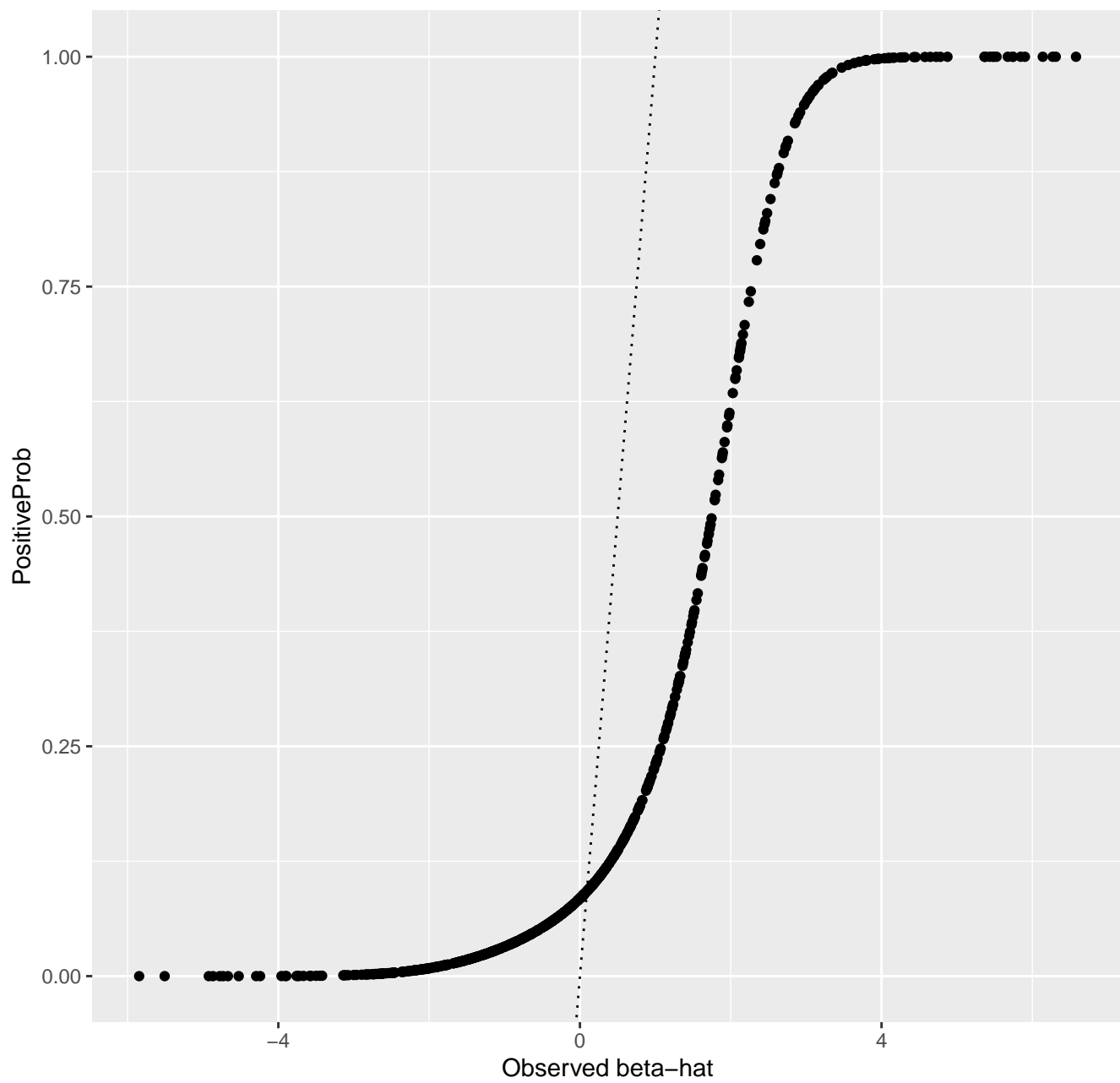
OXSR1_(9943) negative



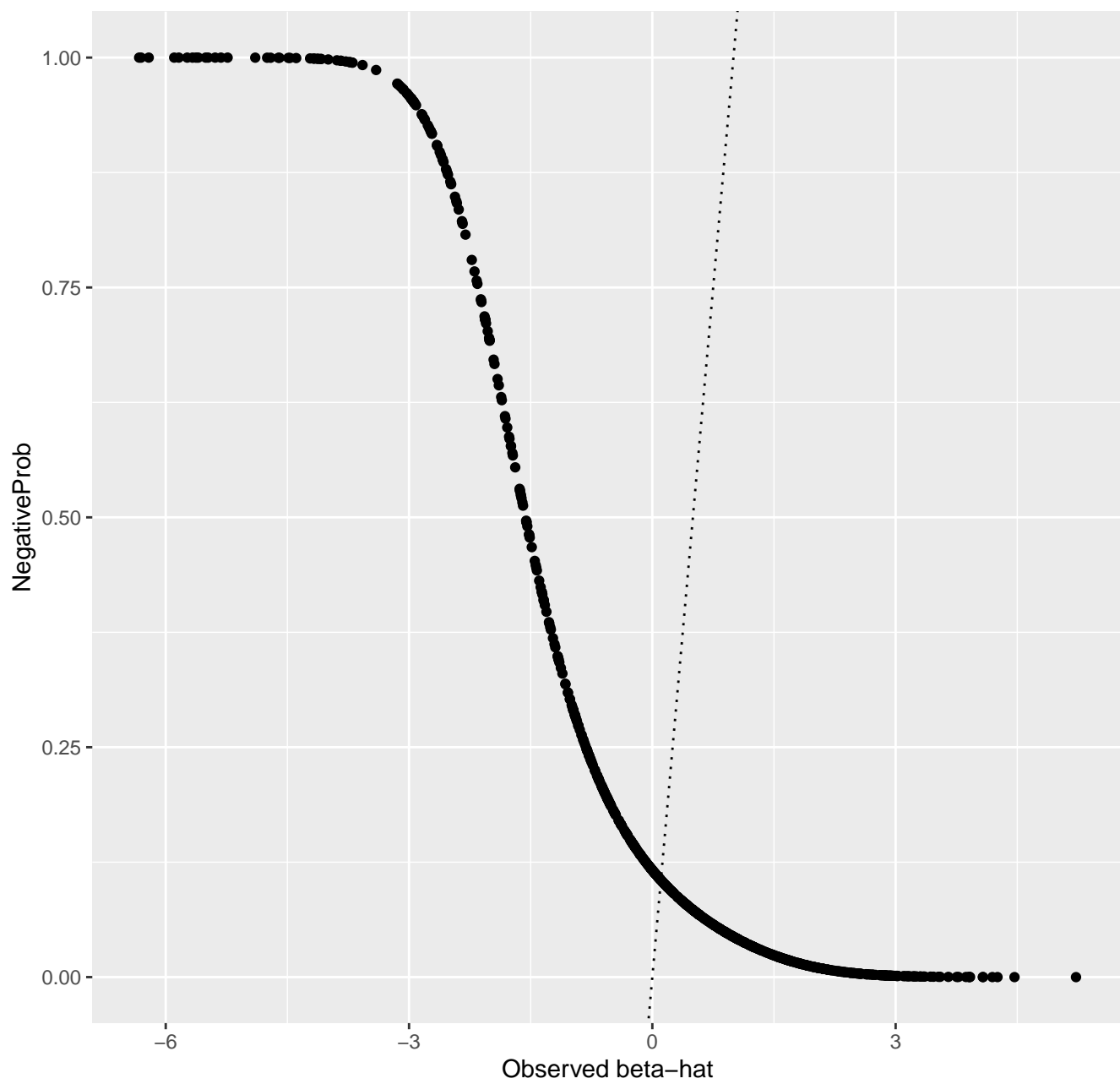
AGPAT5_(55326) negative



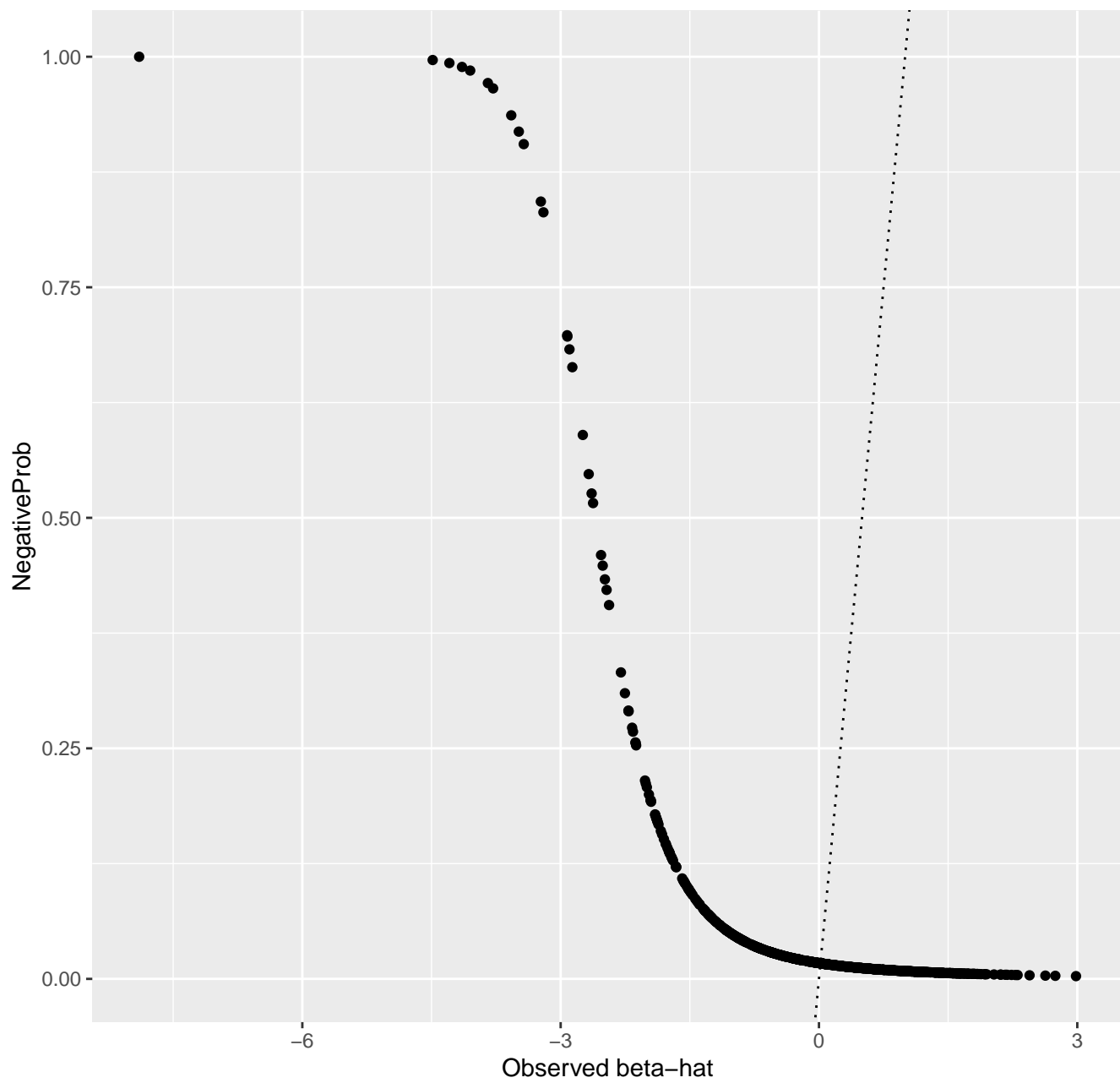
PGK1_(5230) positive



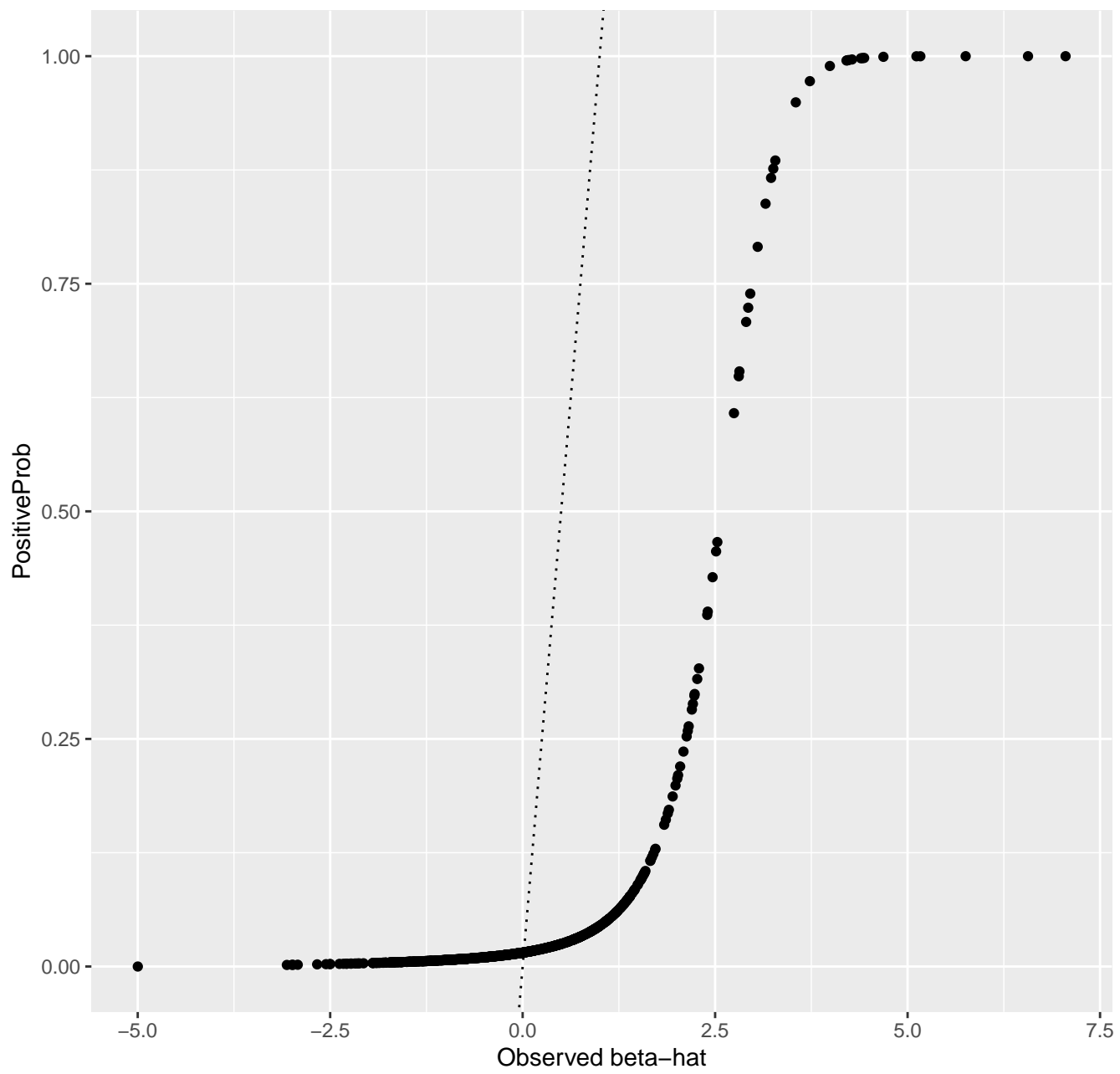
PET117_(100303755) negative



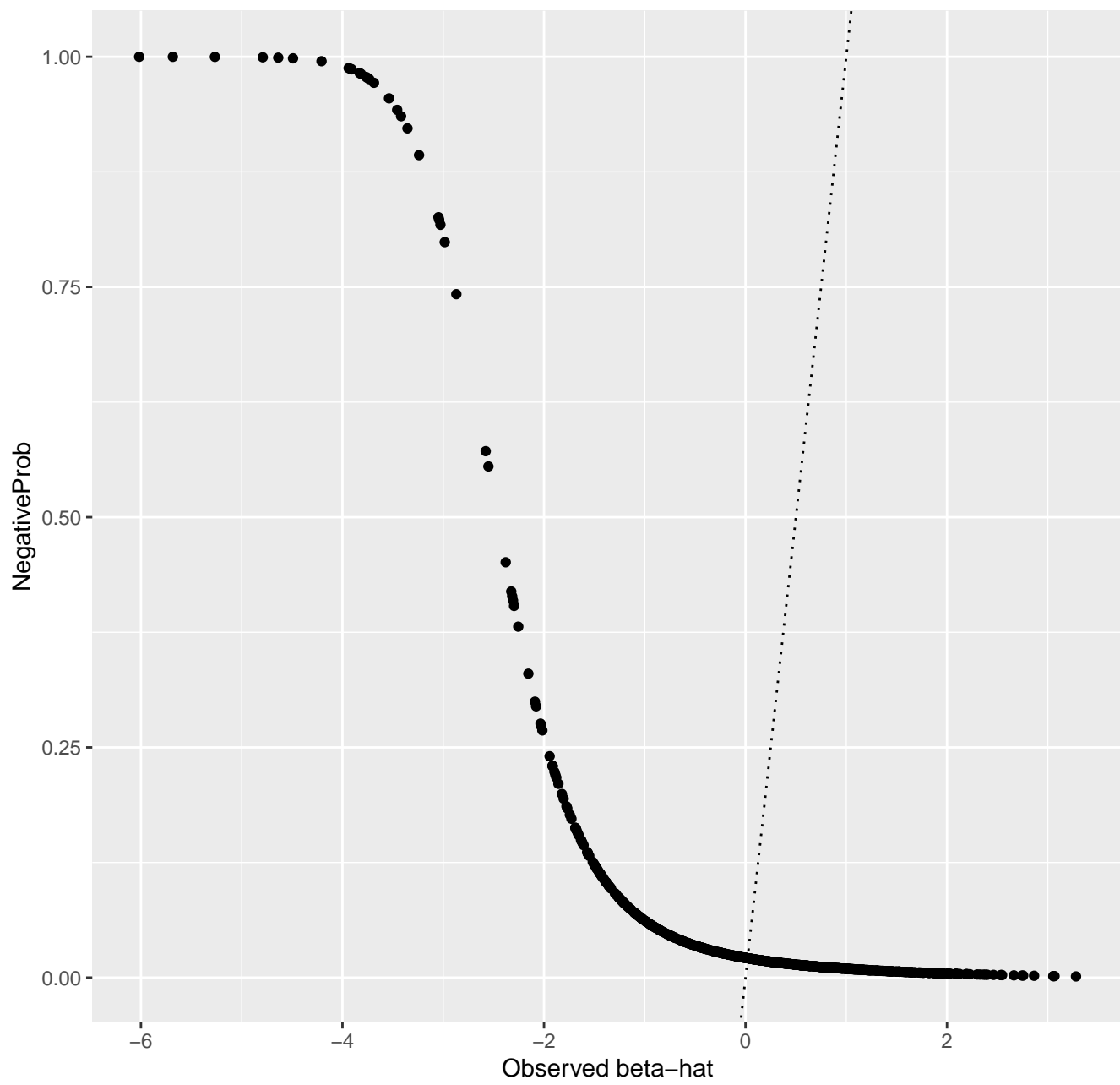
PRUNE1_(58497) negative



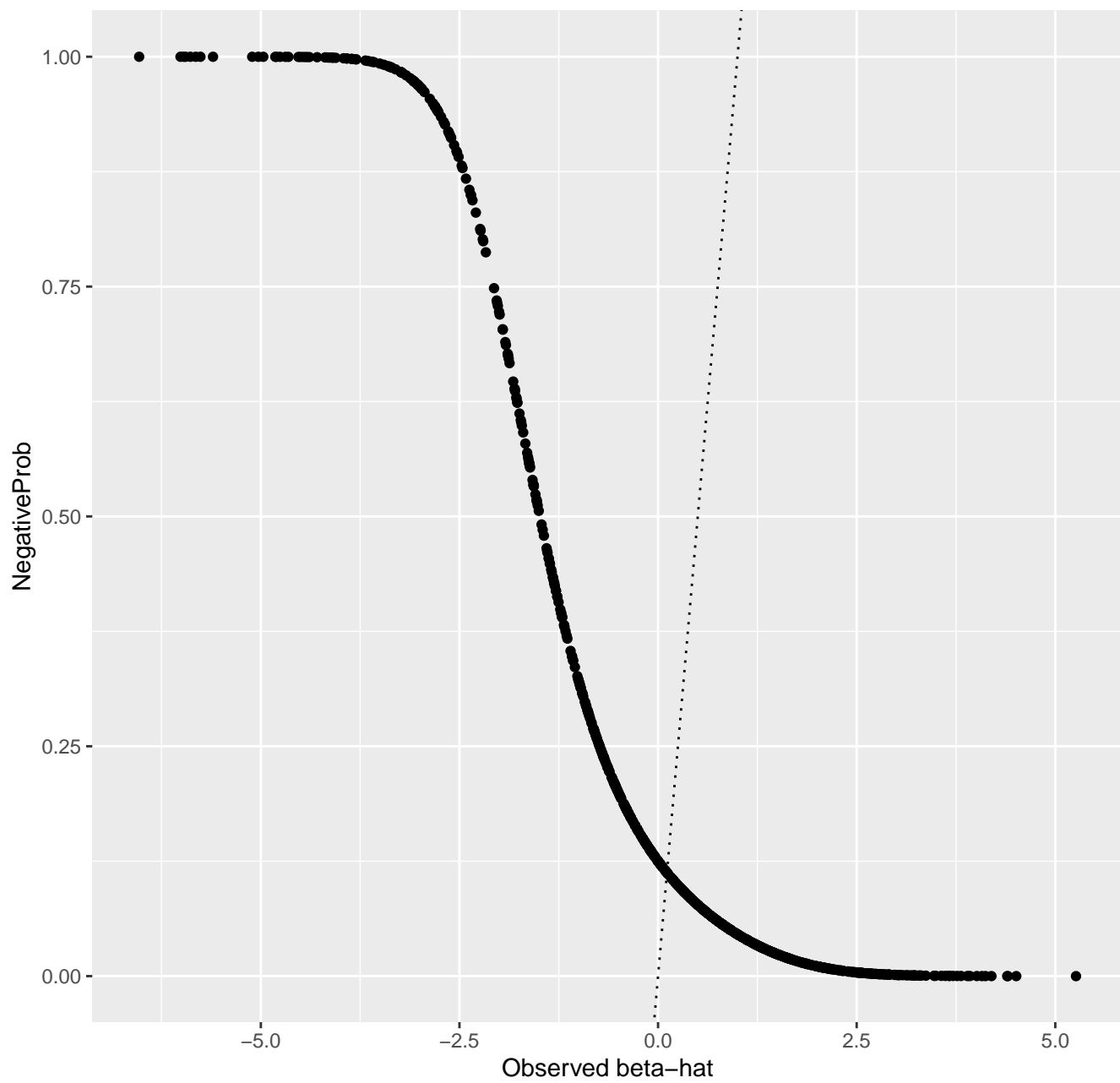
CDKN2A_(1029) positive



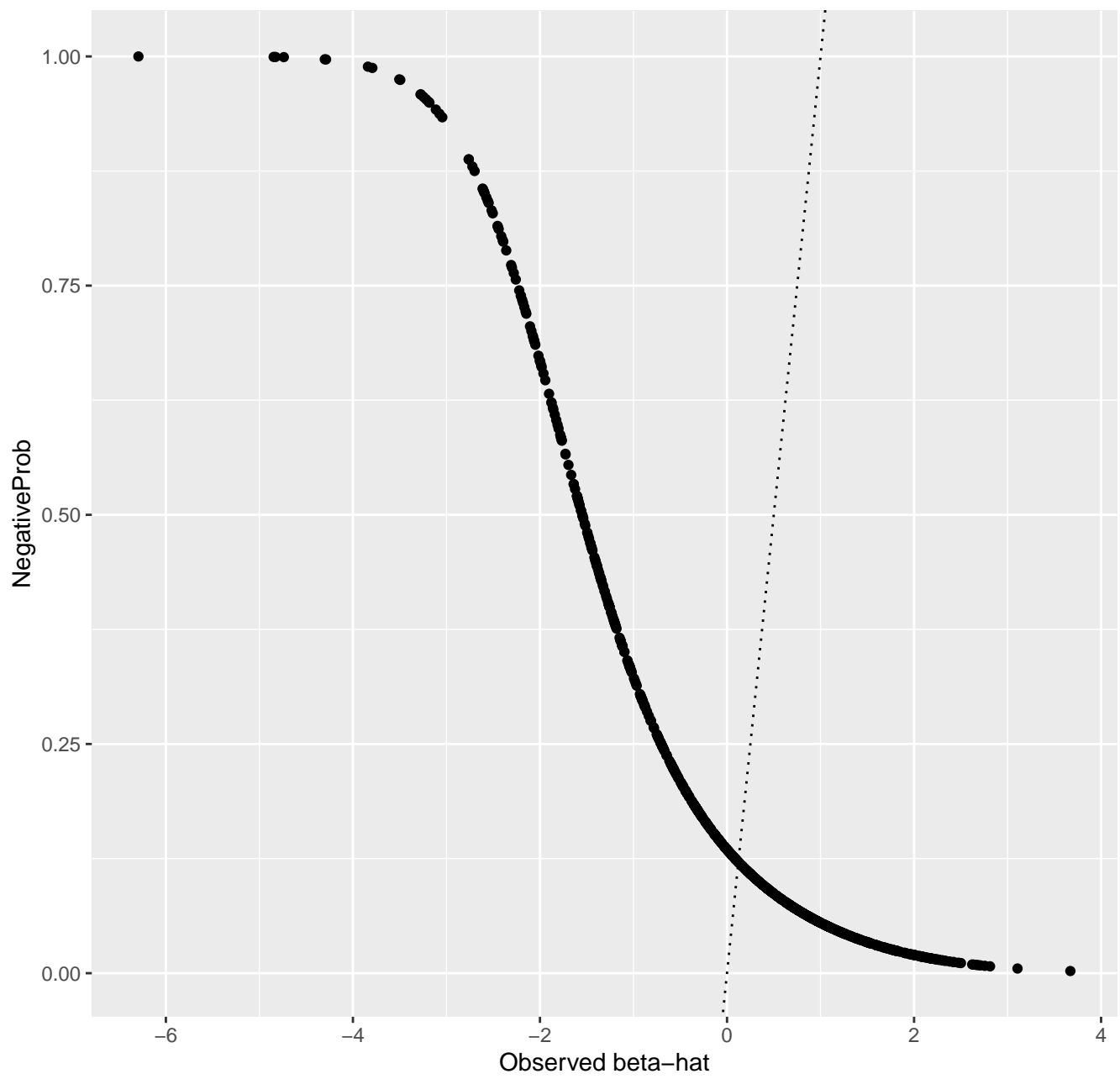
CRTC3_(64784) negative



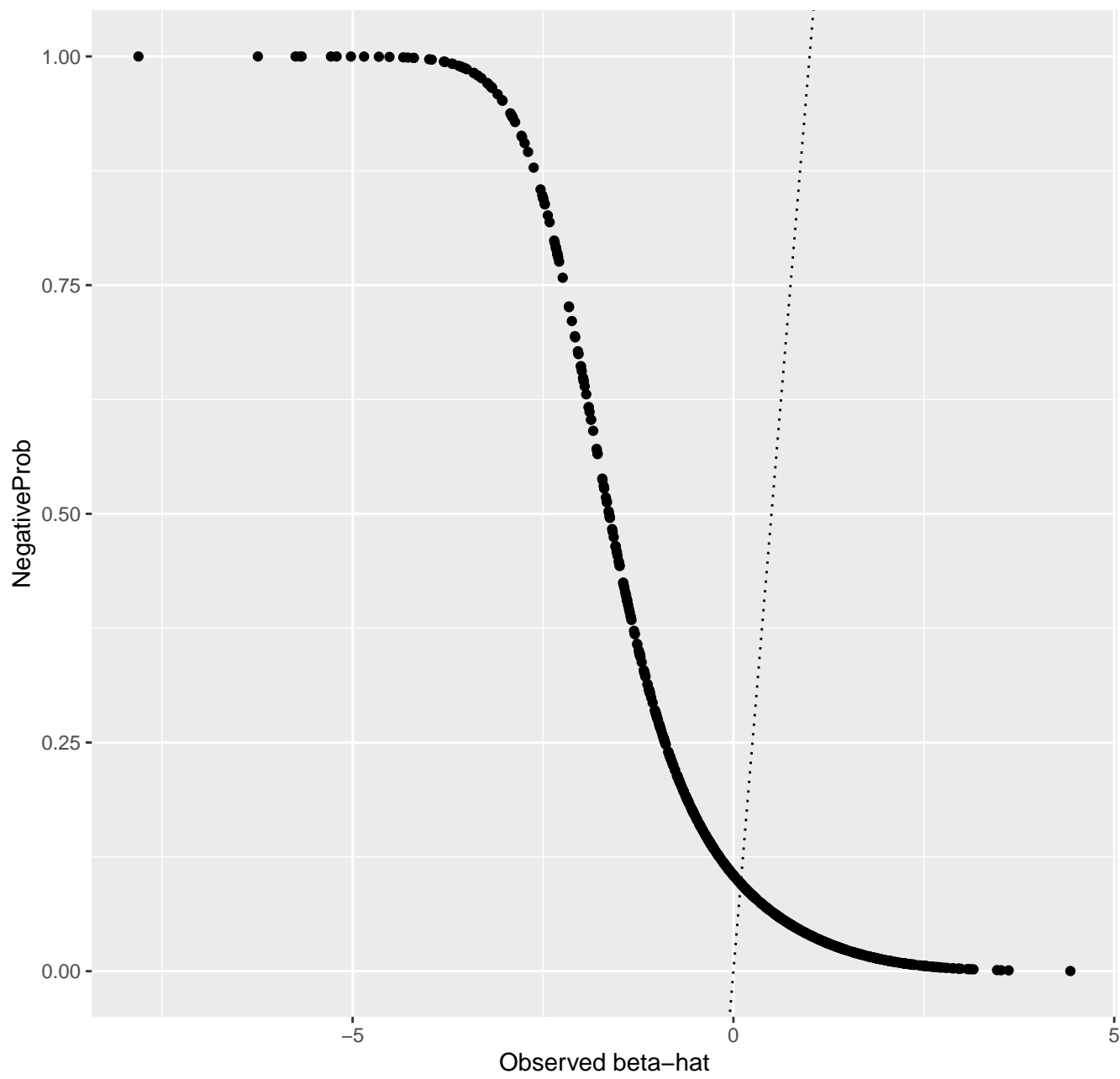
MTHFD1_(4522) negative



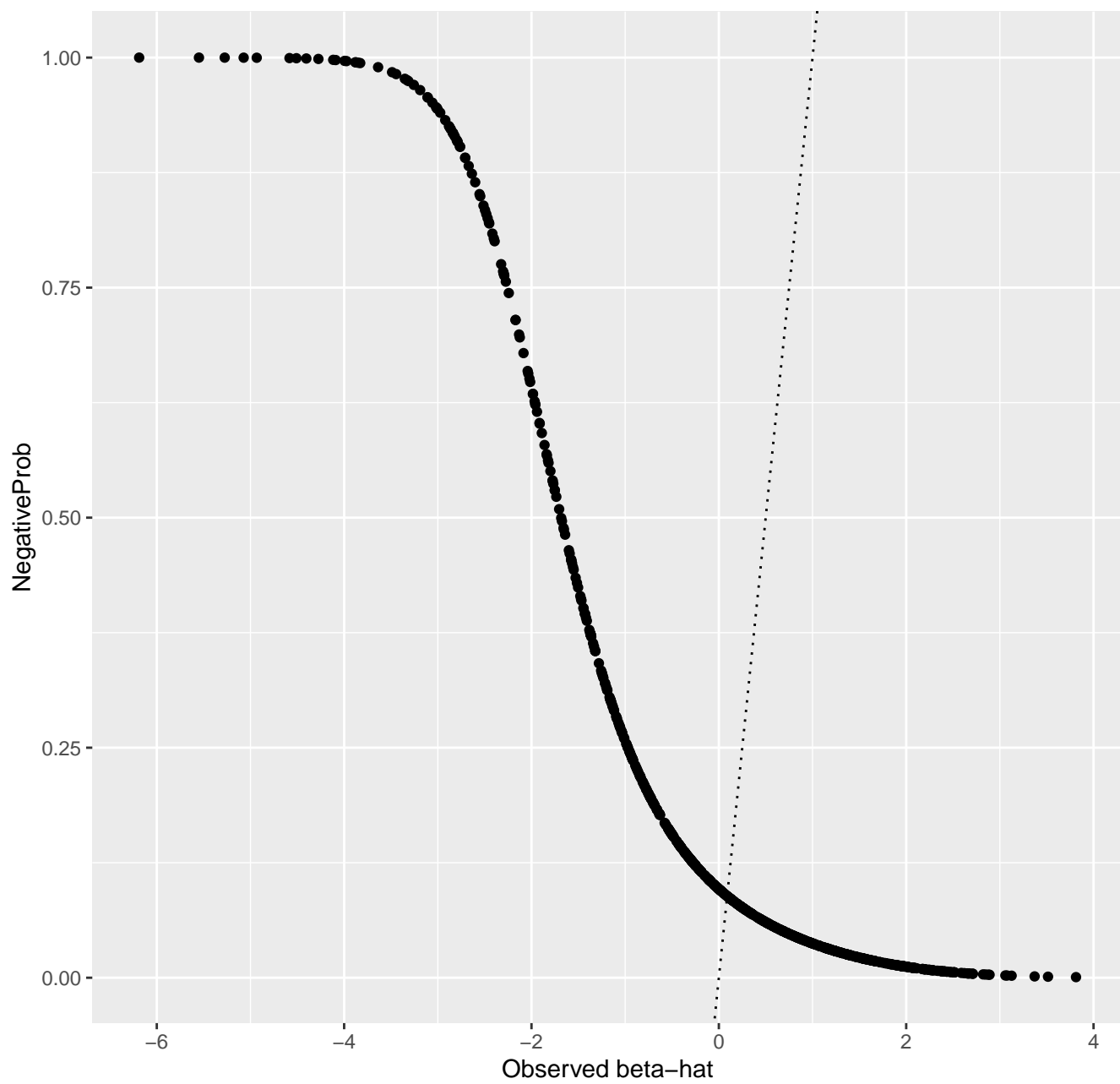
VCL_(7414) negative



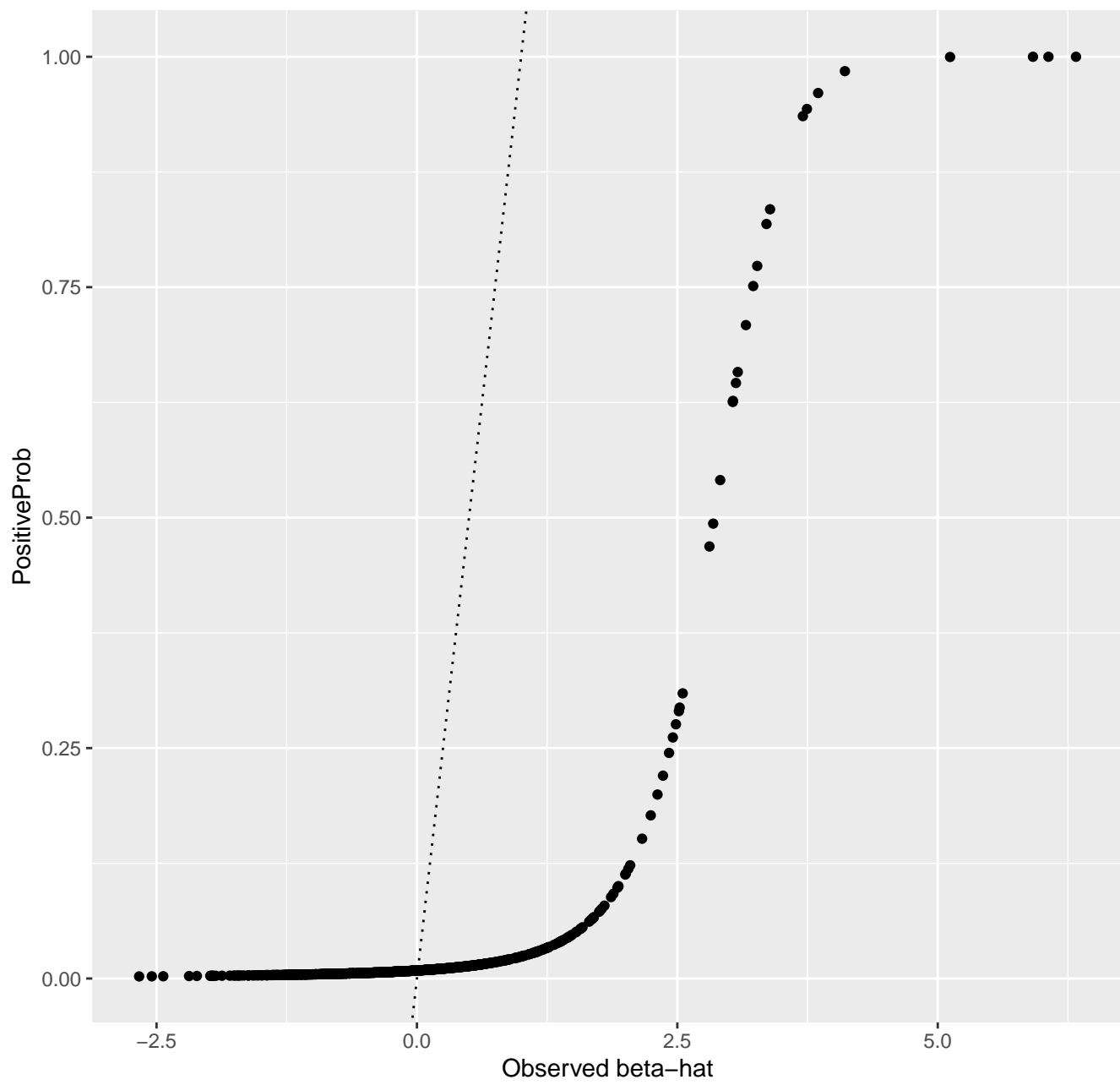
NDUFAF4_(29078) negative



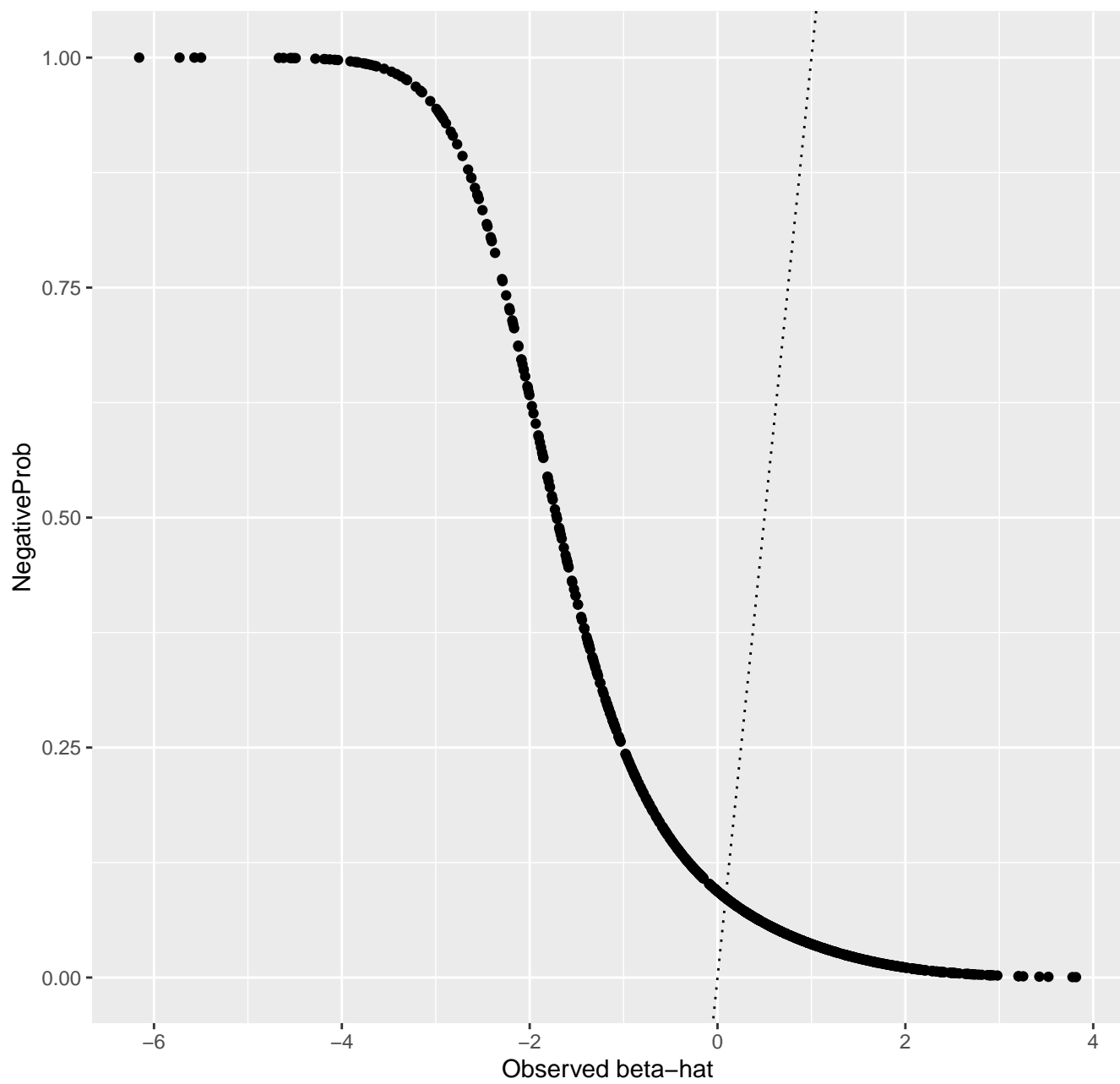
MAD2L1BP_(9587) negative



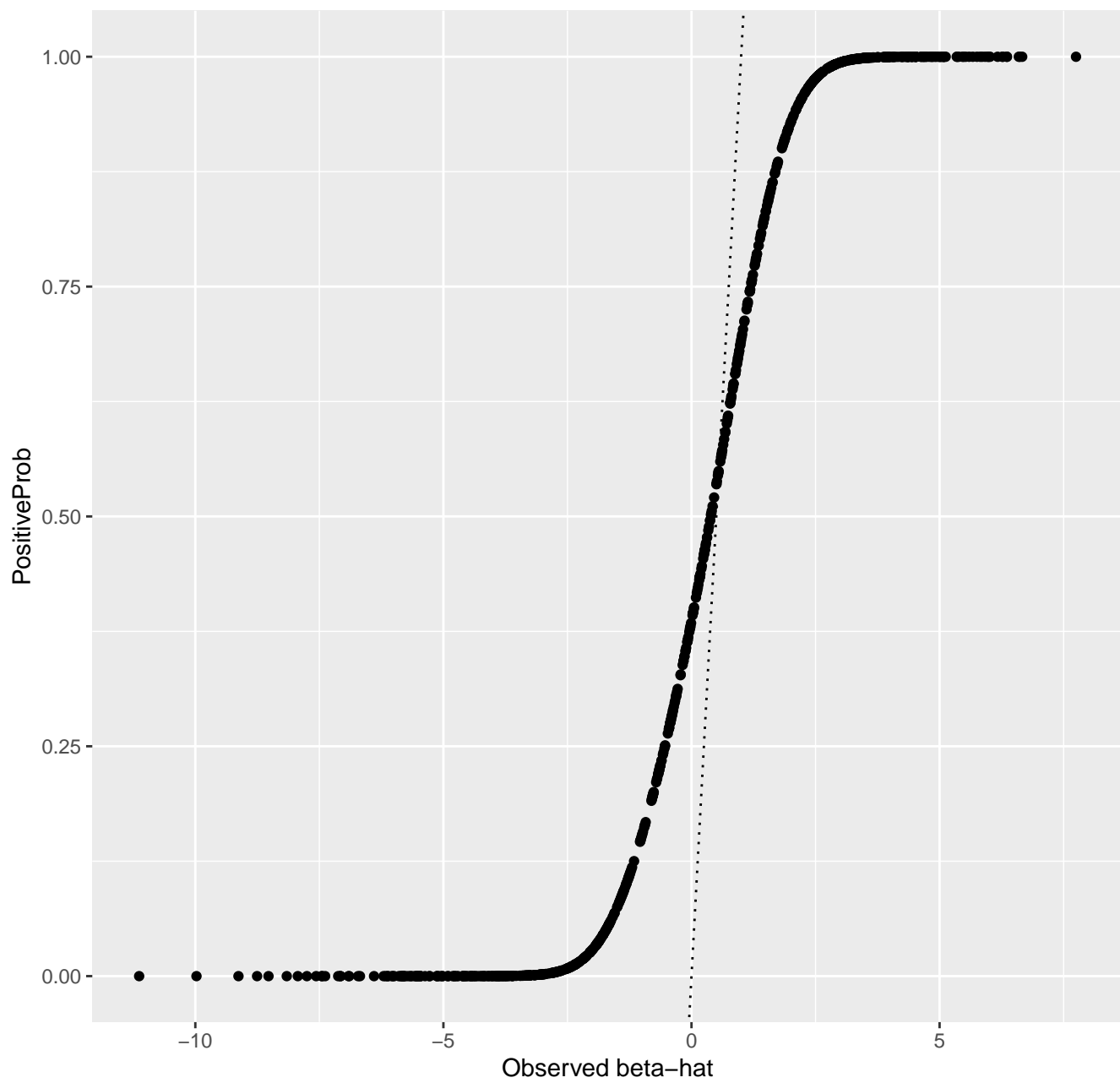
CDKN2C_(1031) positive



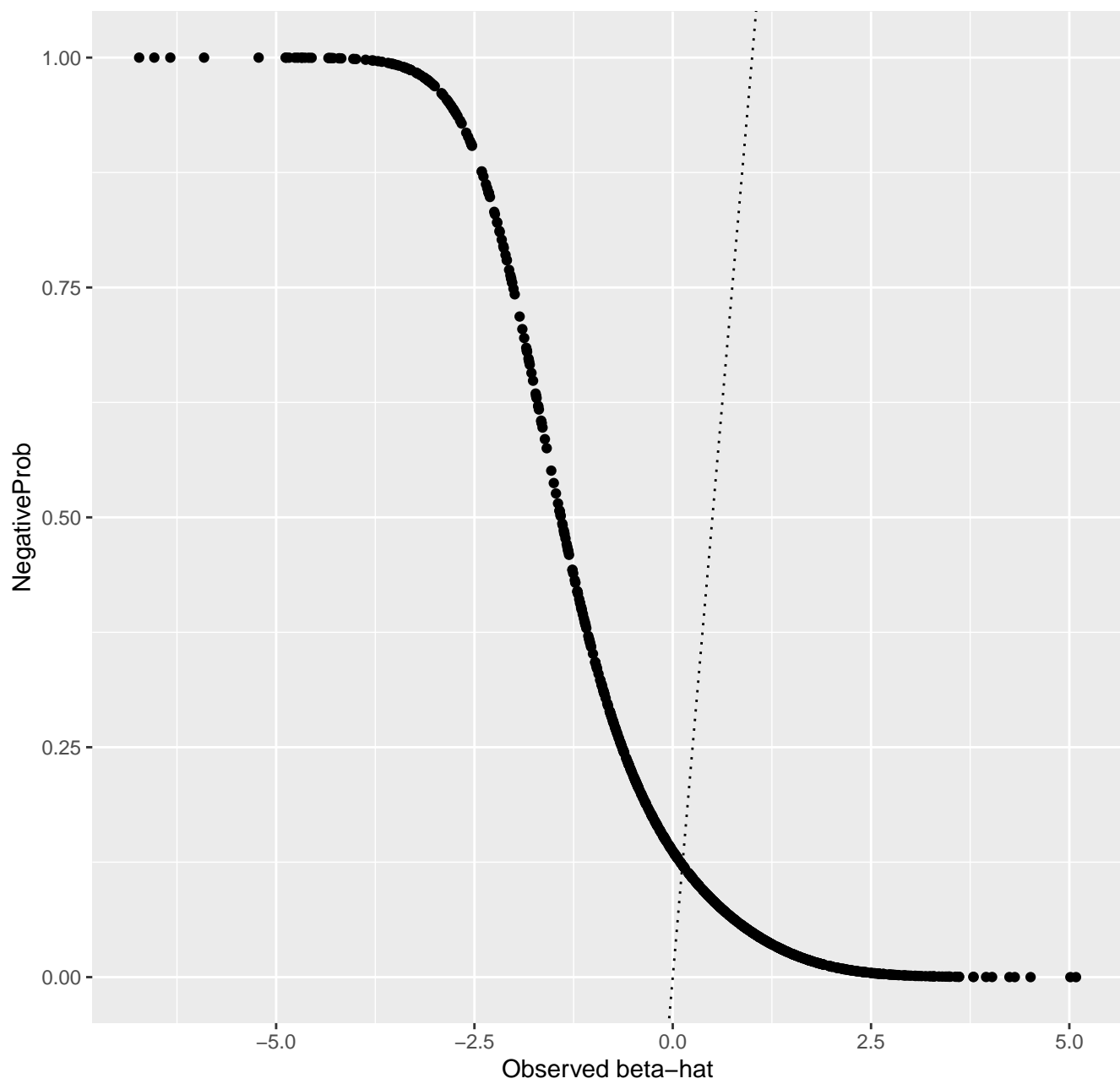
C15orf41_(84529) negative



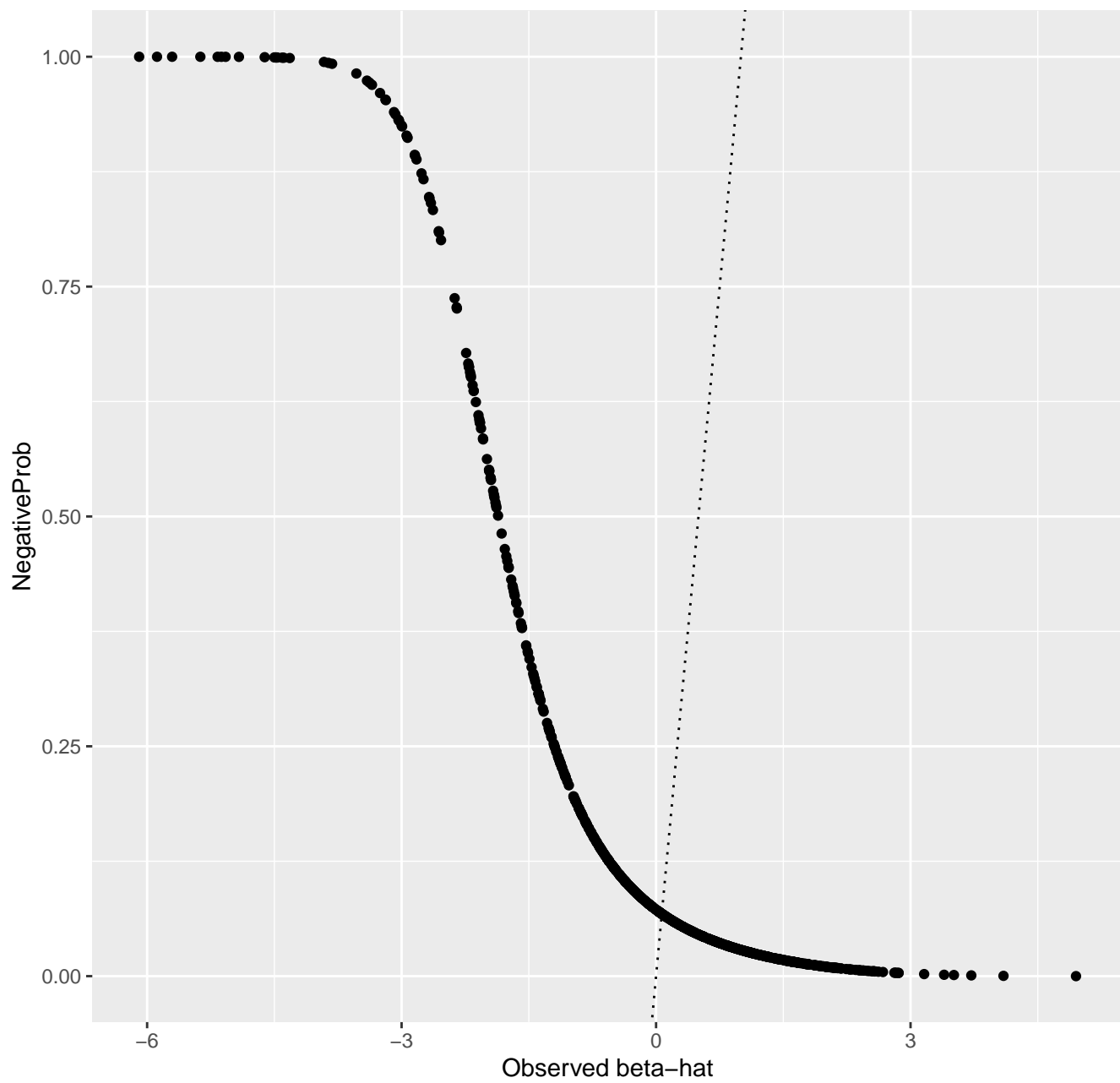
DNAJC9_(23234) positive



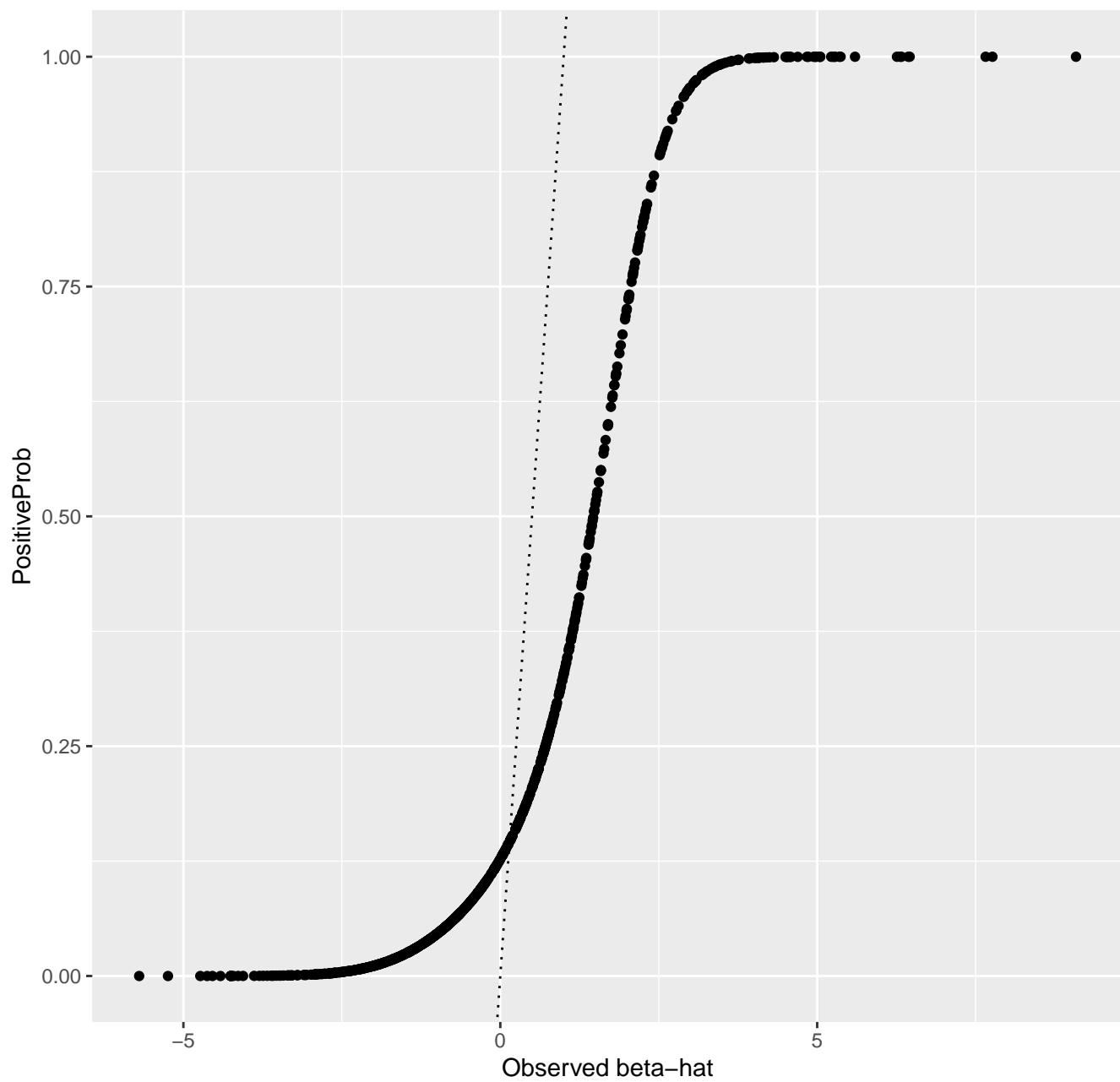
LIAS_(11019) negative



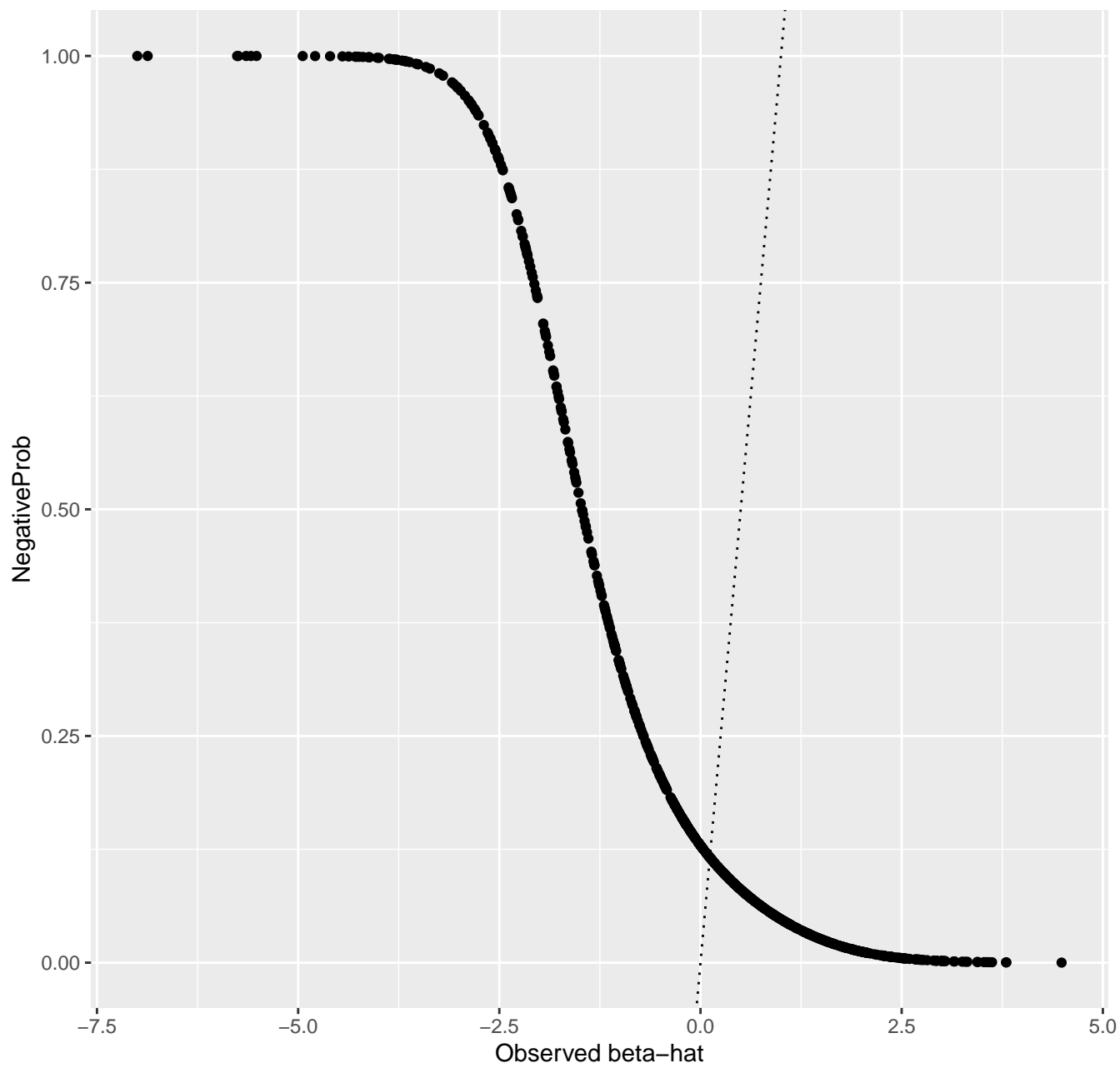
USP1_(7398) negative



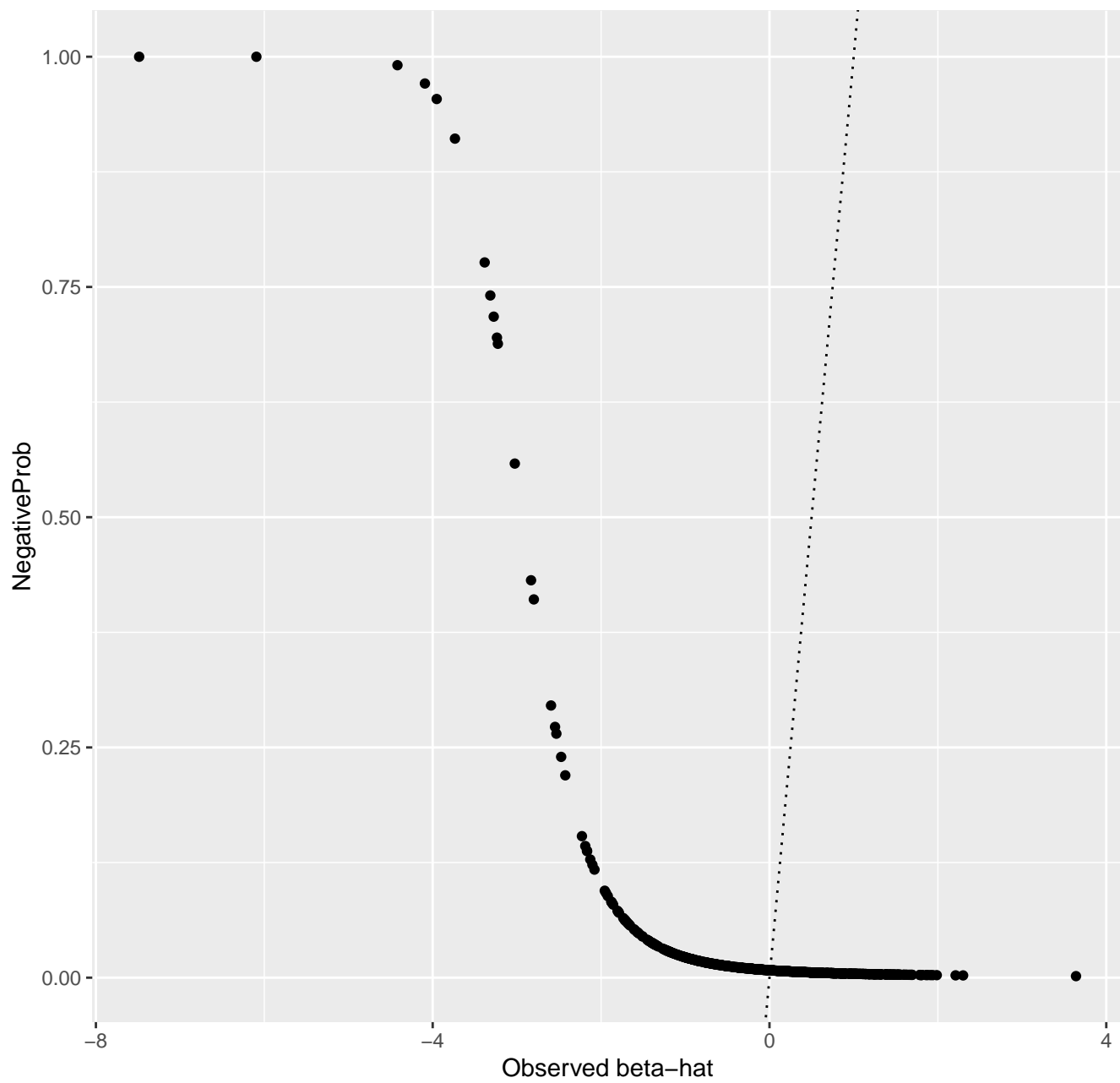
NUP43_(348995) positive



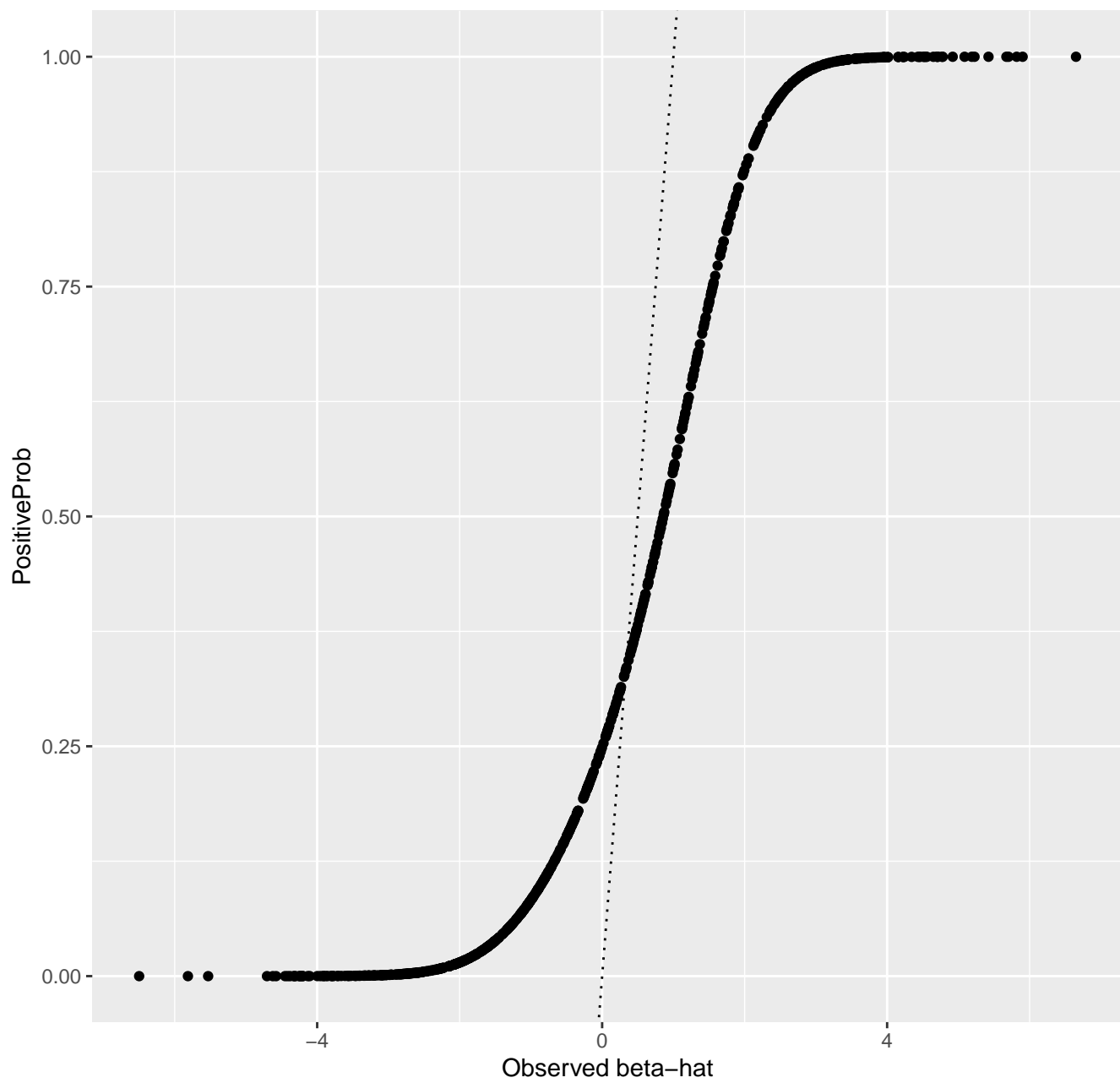
CXorf56_(63932) negative



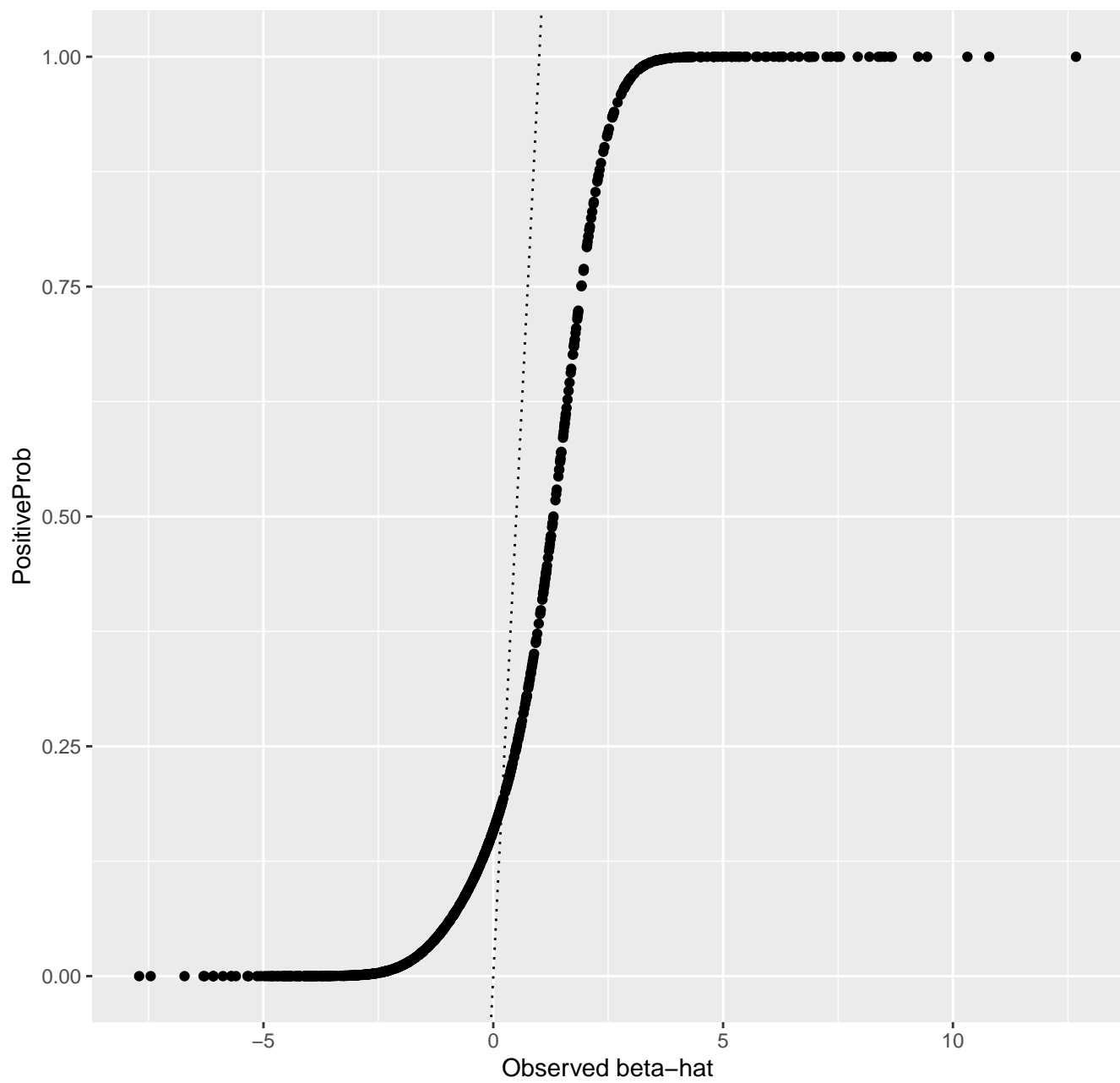
KCNK5_(8645) negative



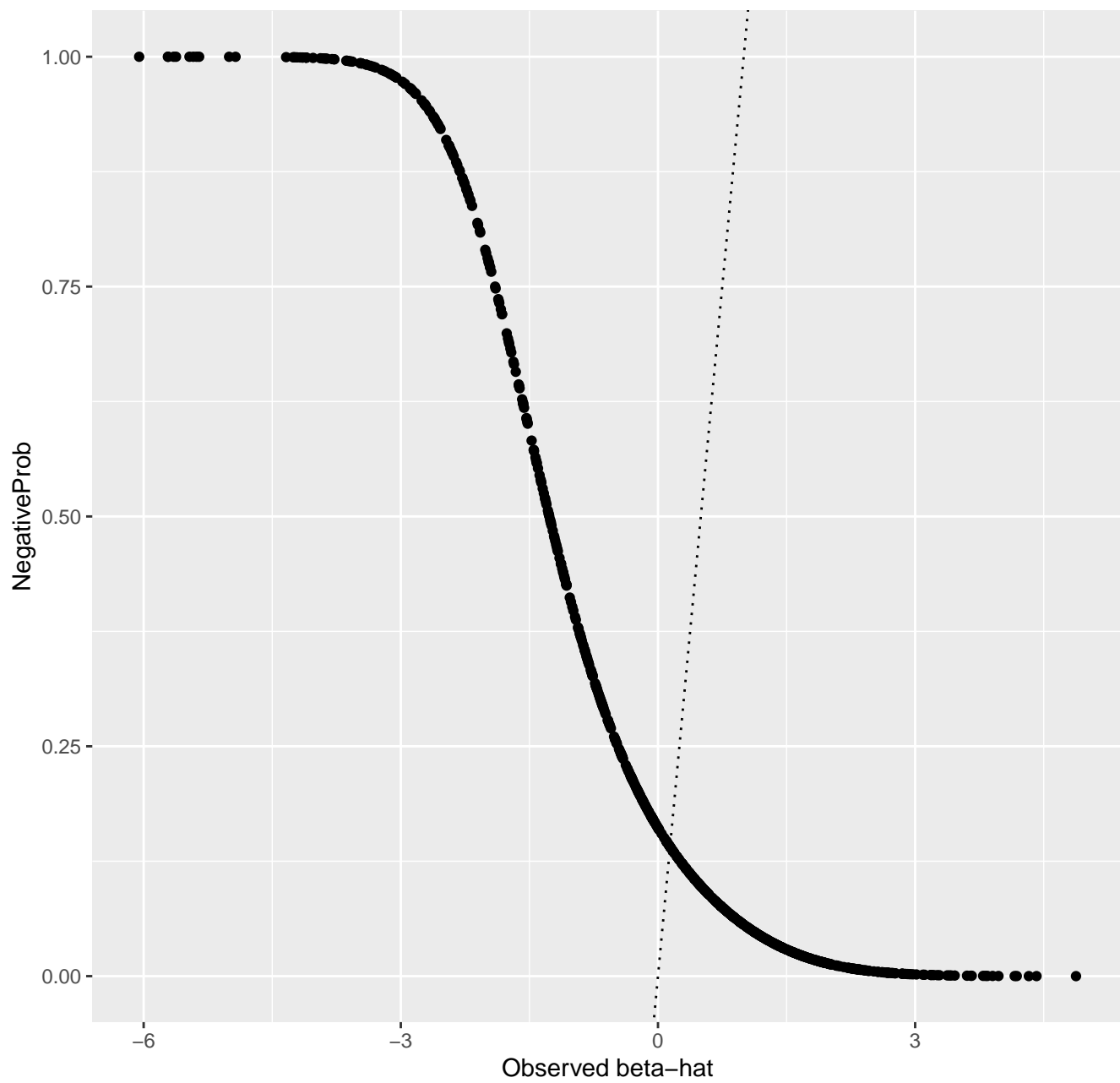
PTPN11_(5781) positive



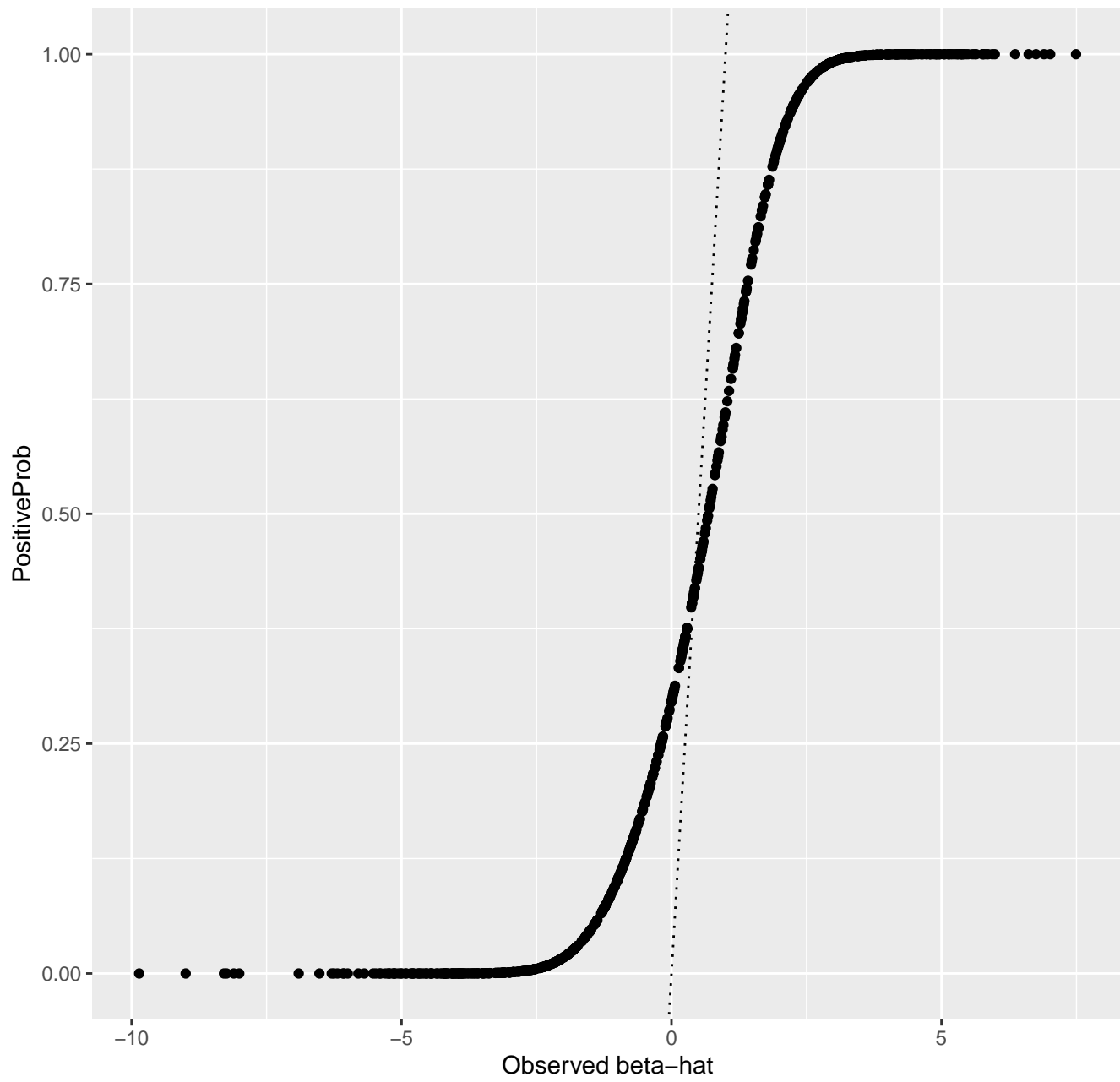
AMBRA1_(55626) positive



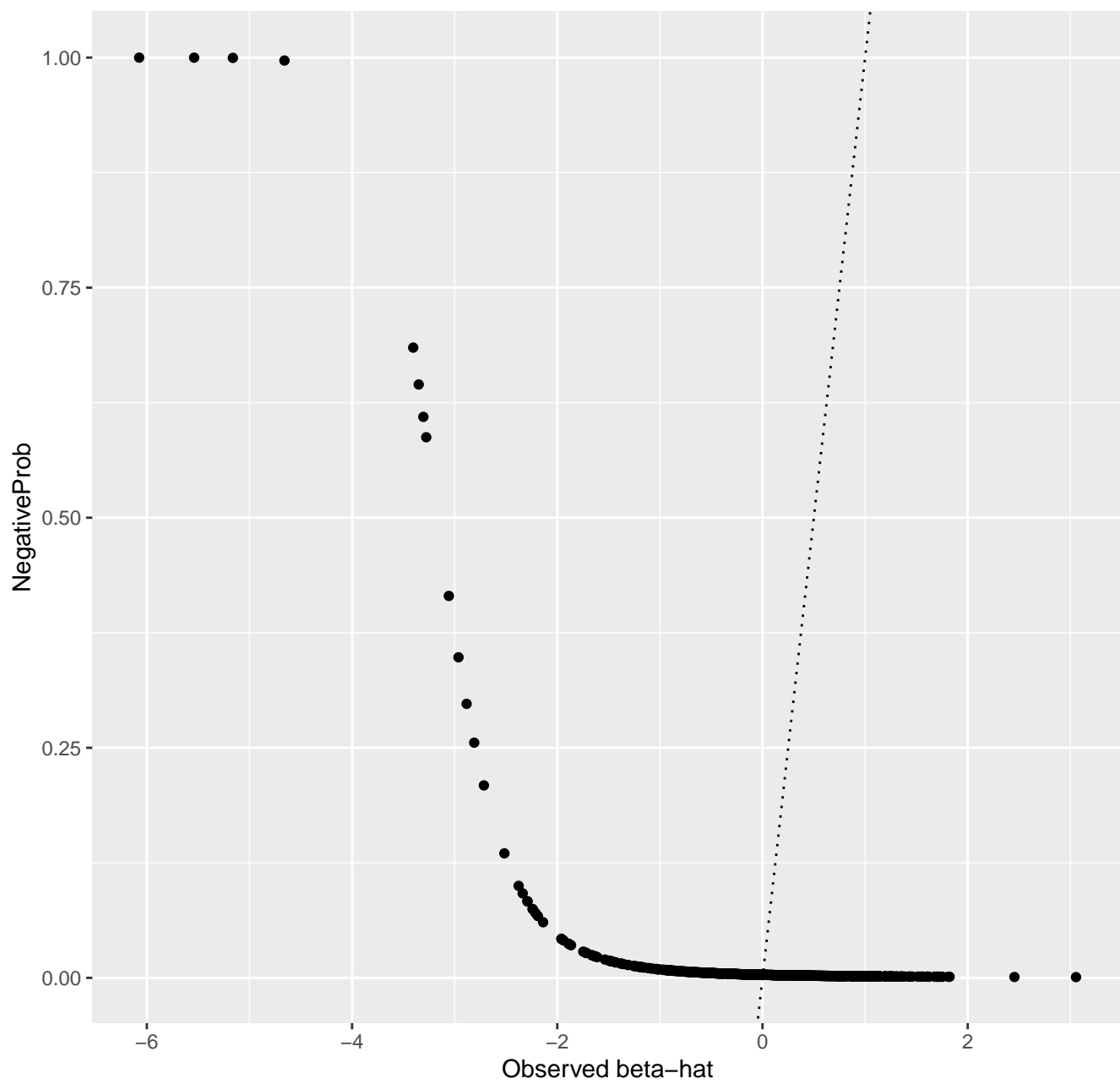
NDUFB6_(4712) negative



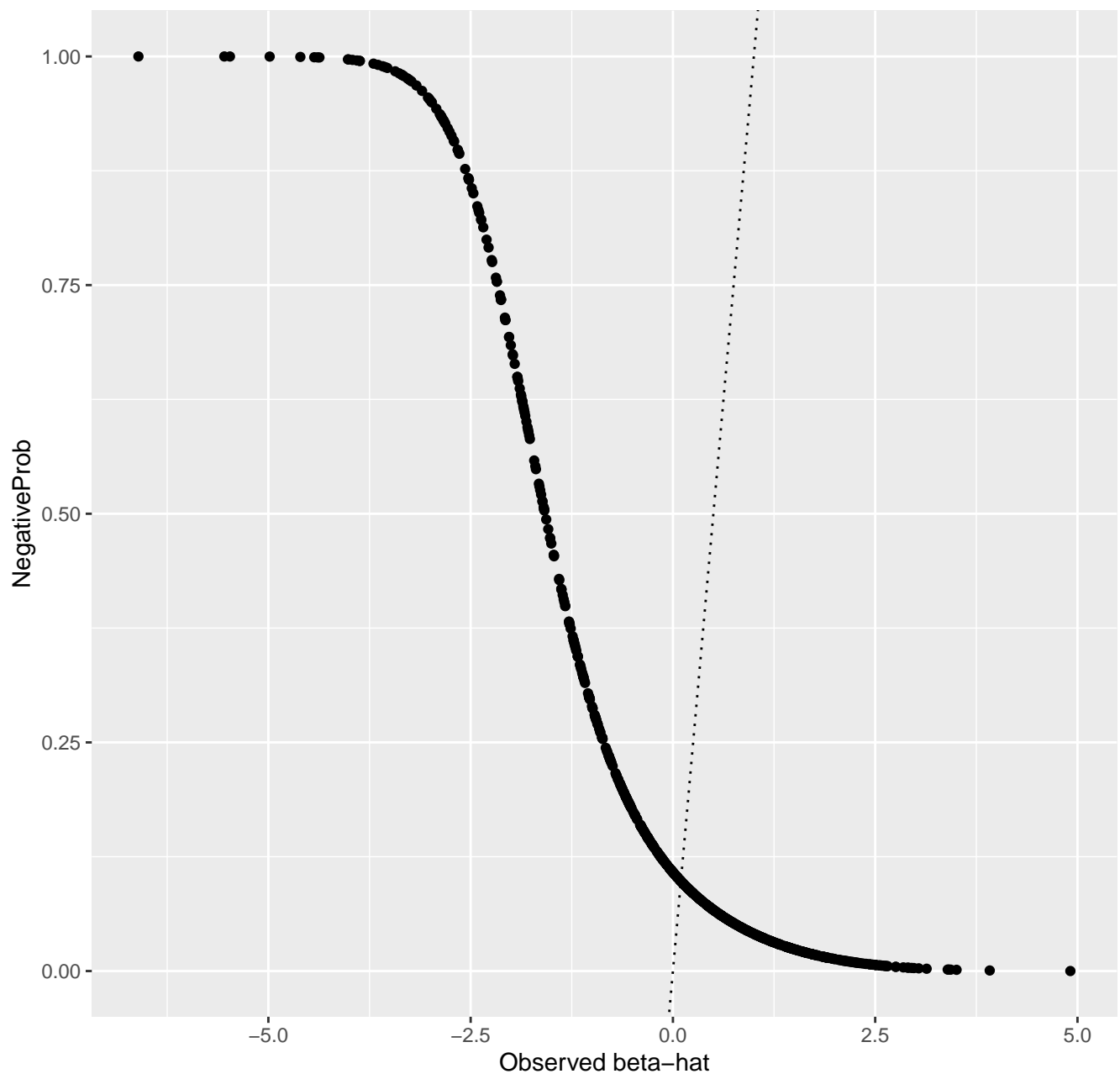
WDR1_(9948) positive



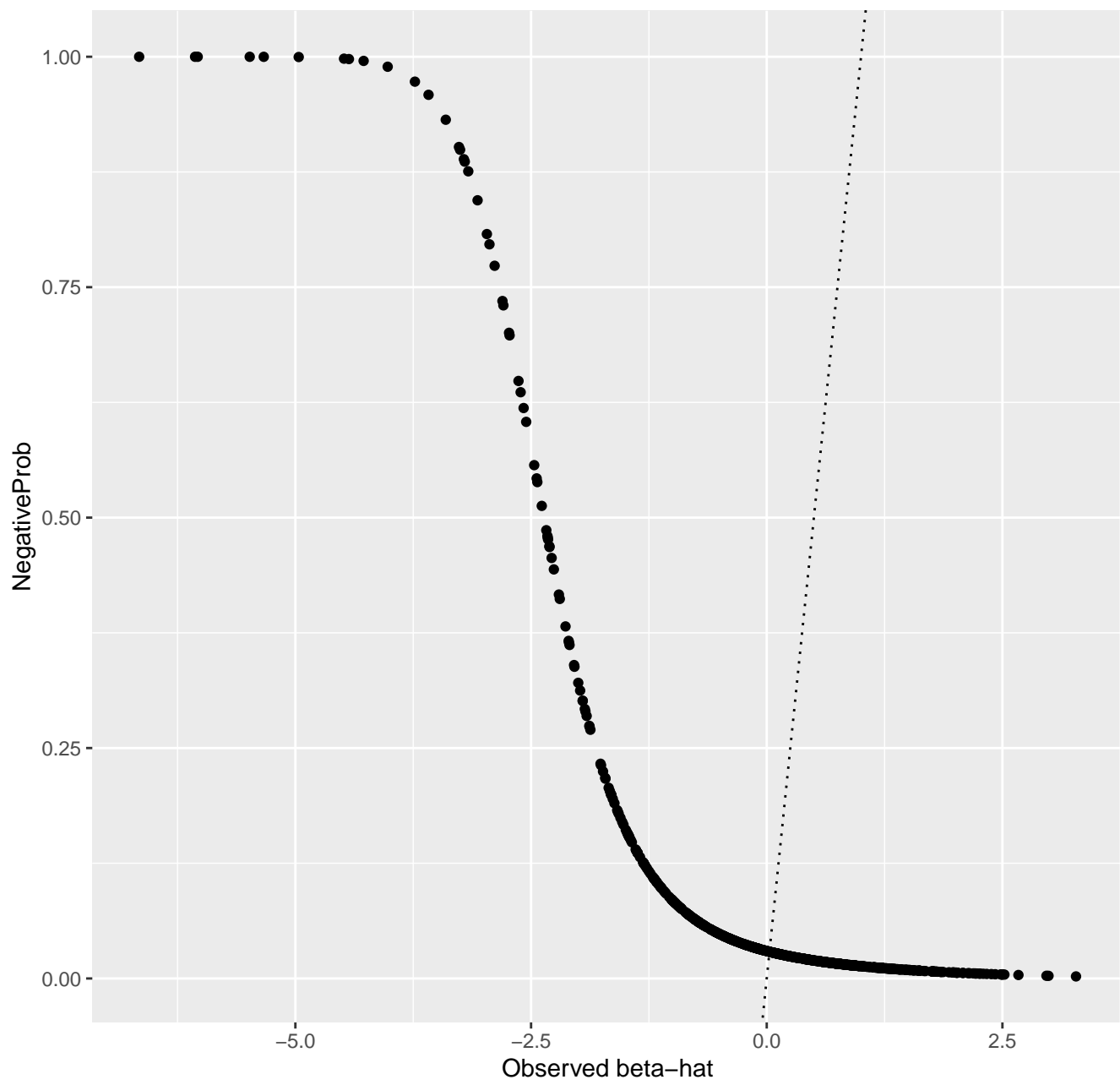
JAGN1_(84522) negative



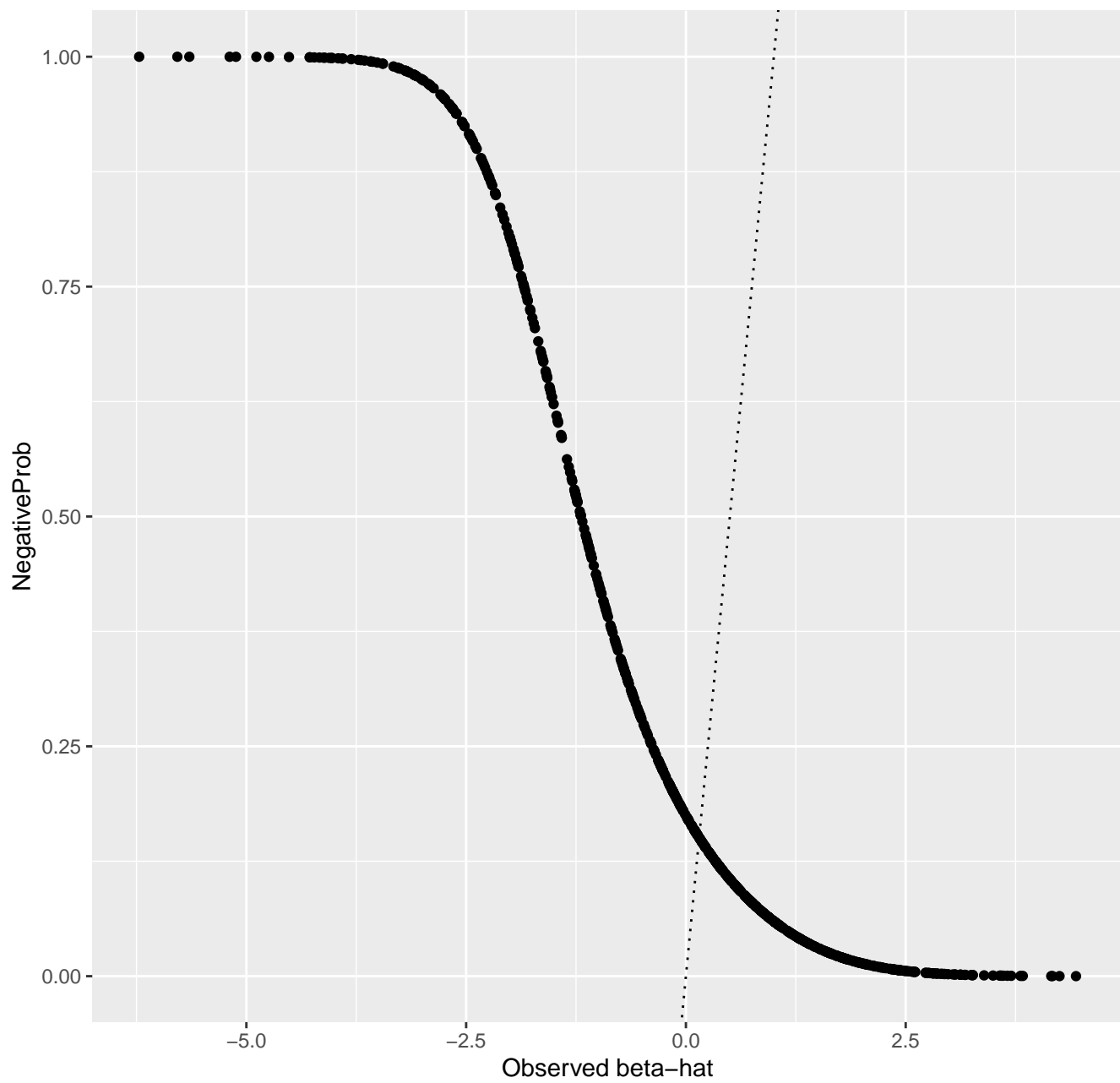
TLK2_(11011) negative



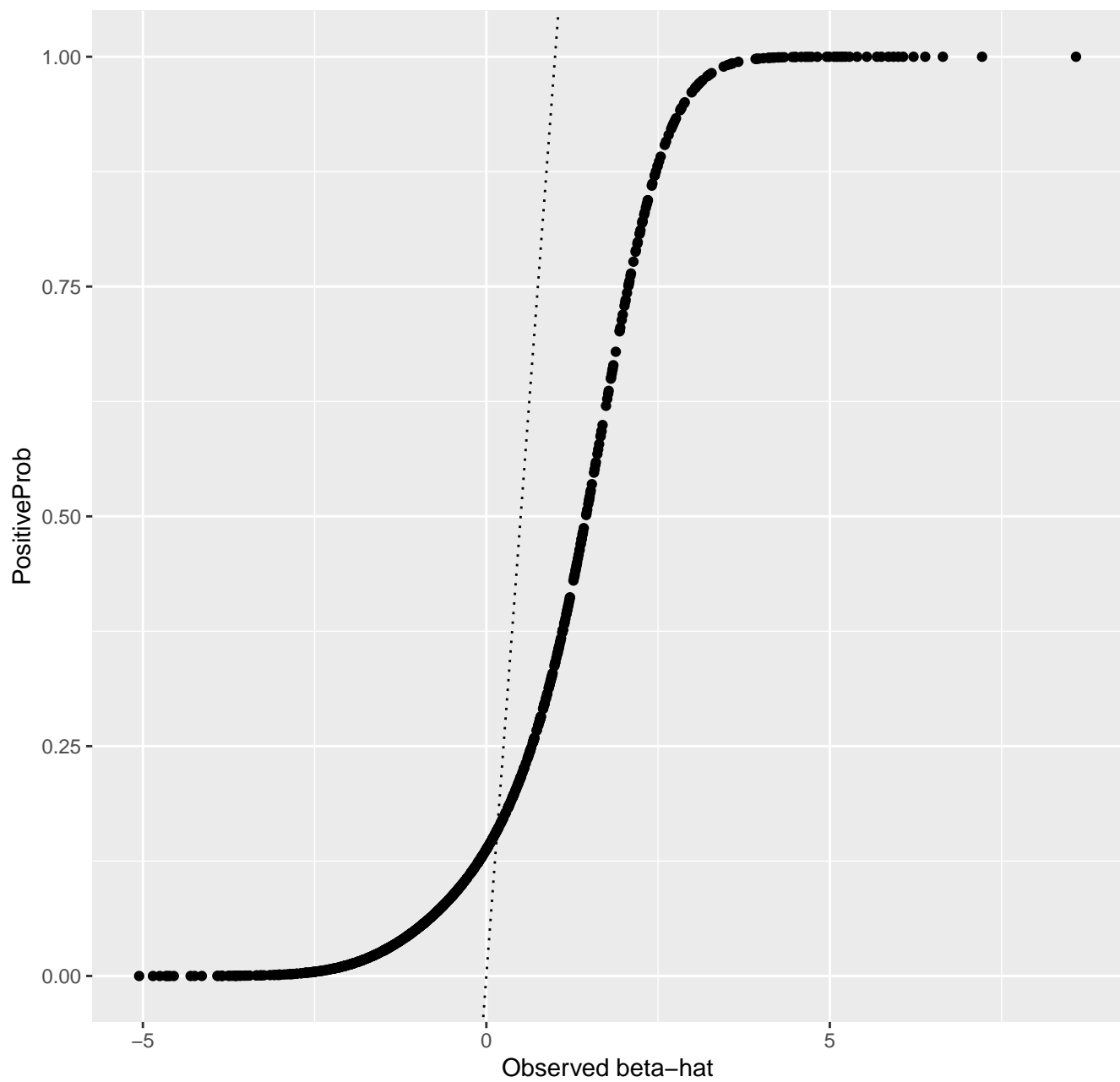
SEC23IP_(11196) negative



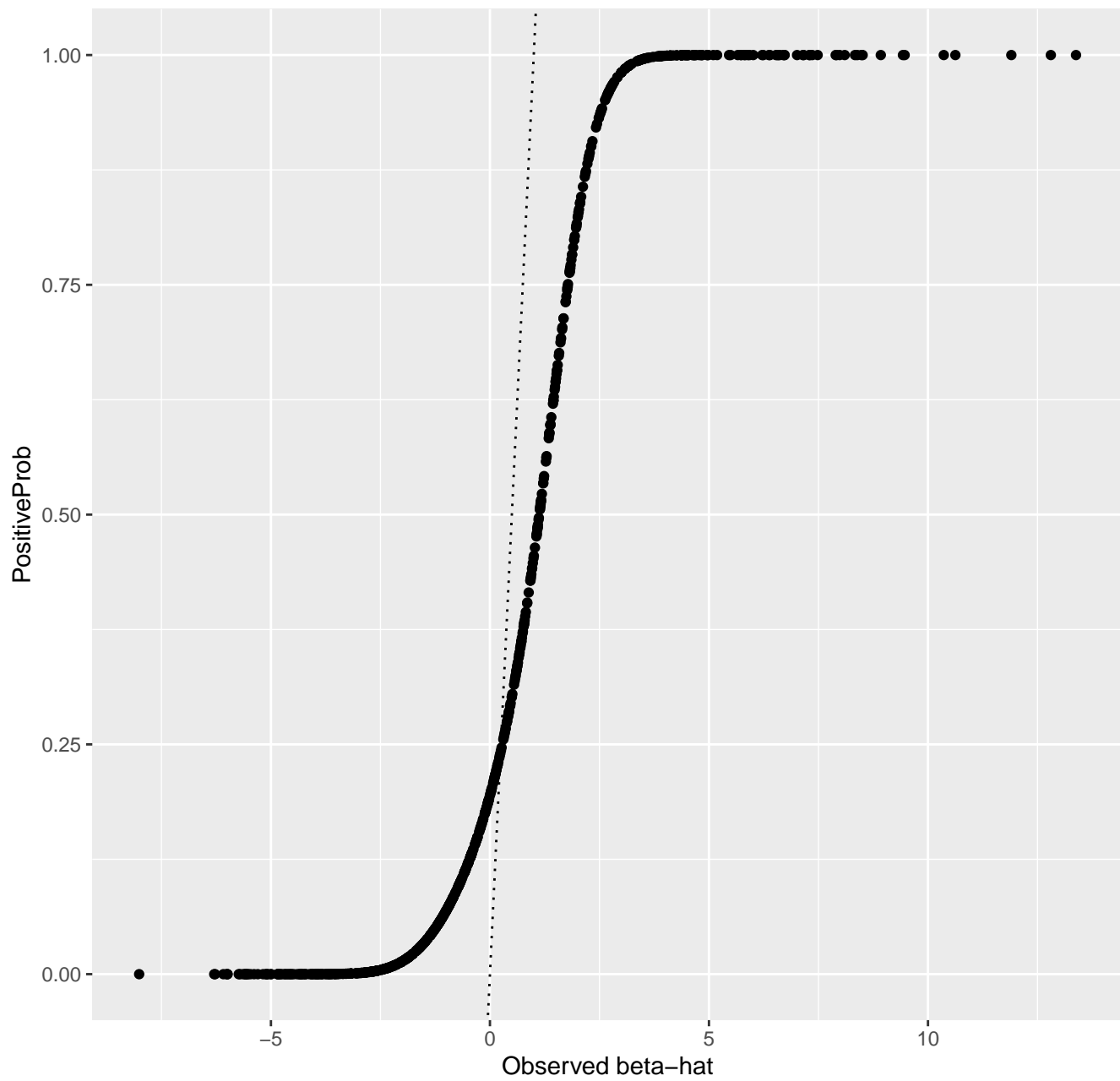
FPGS_(2356) negative



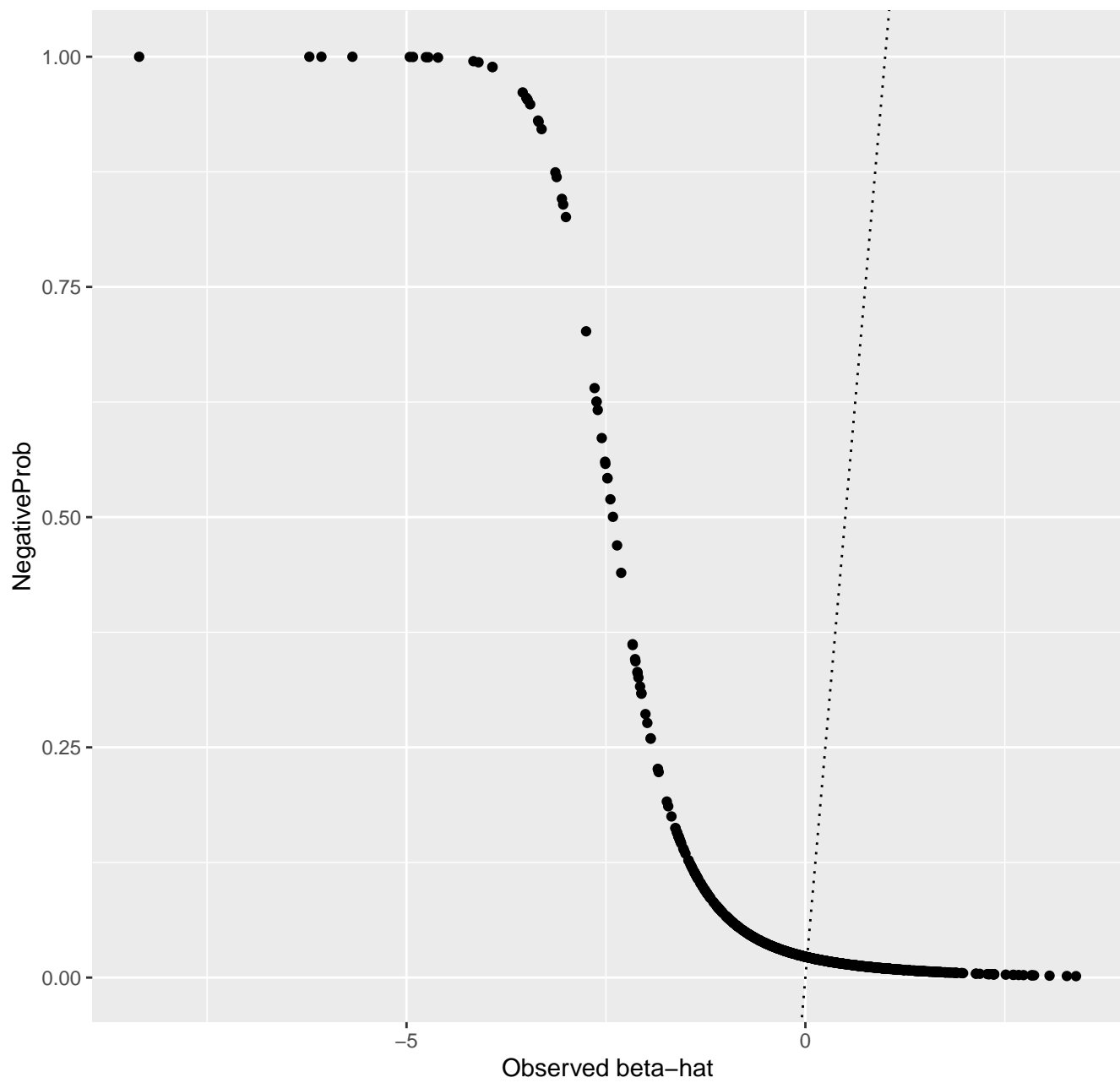
NCAPH2_(29781) positive



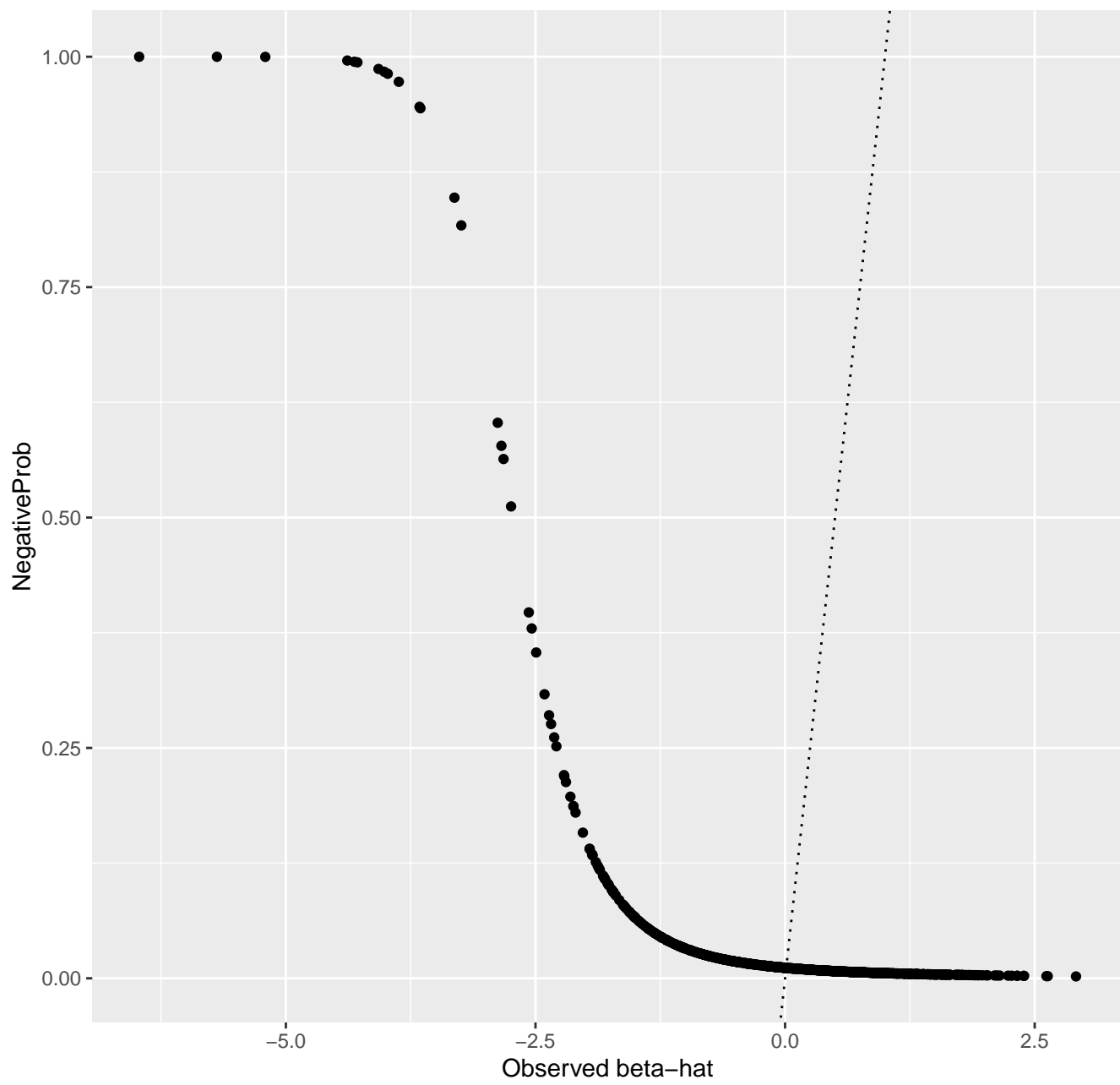
TSC1_(7248) positive



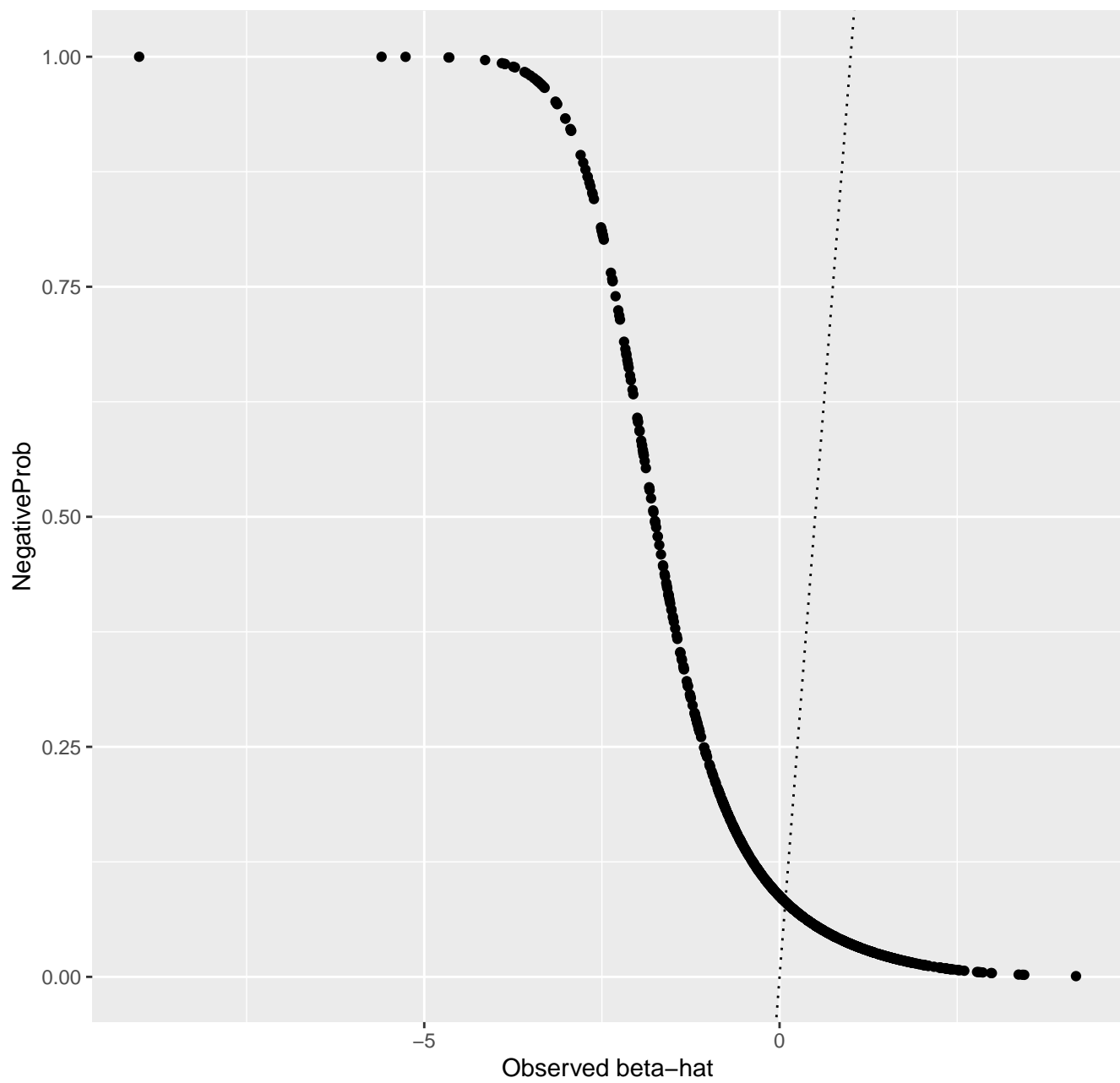
TMEM189_(387521) negative



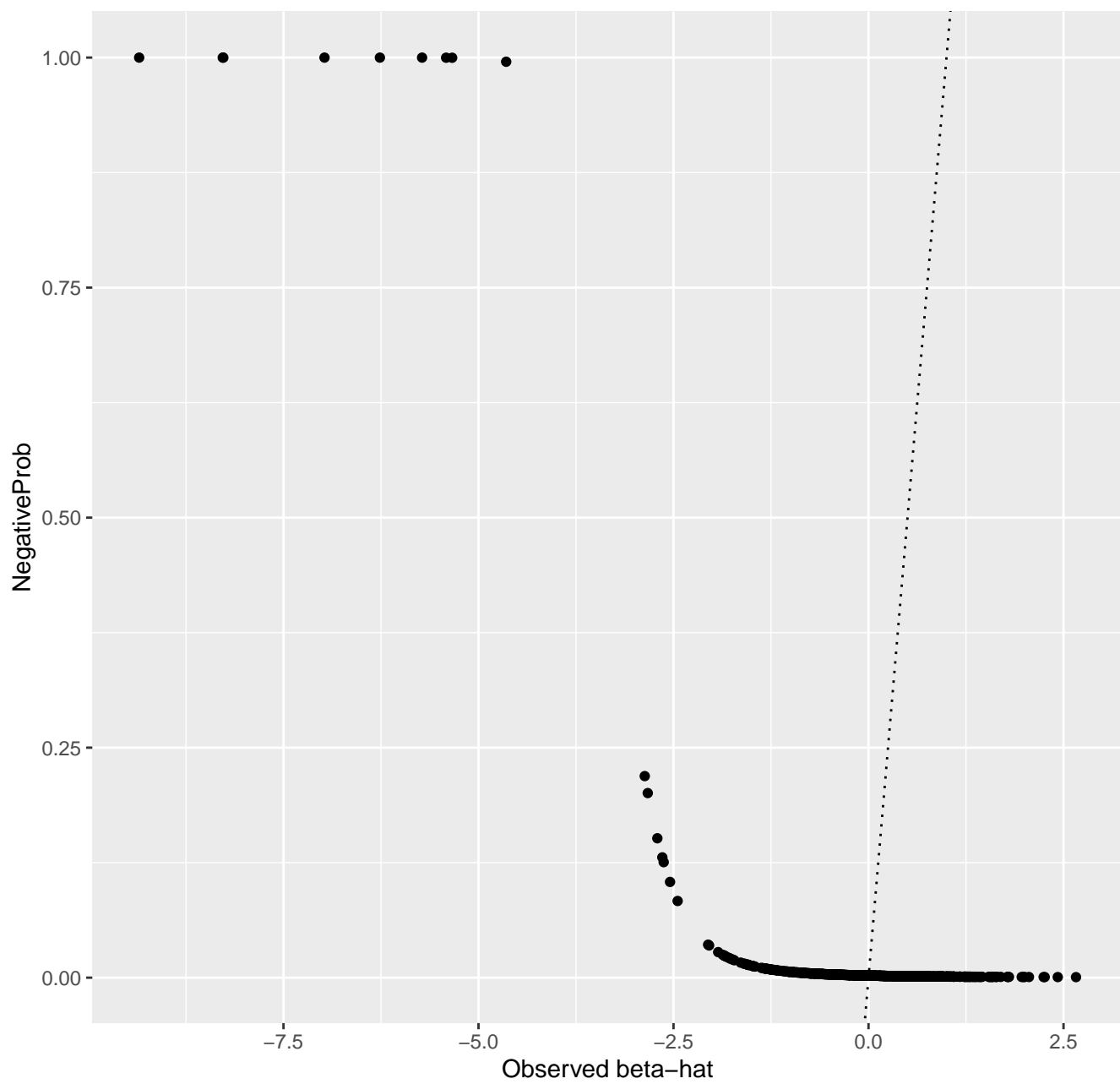
CR1L_(1379) negative



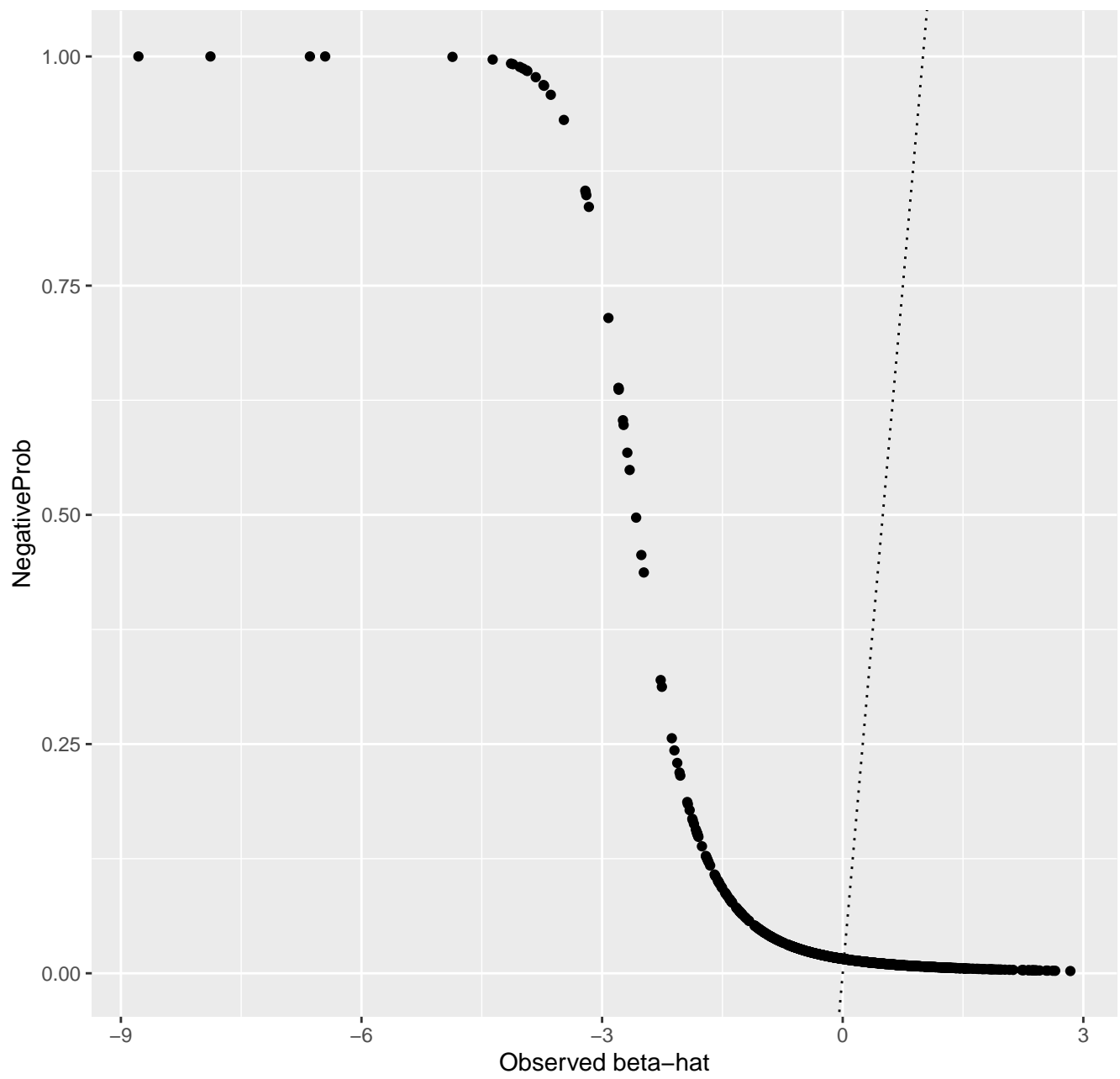
NRDC_(4898) negative



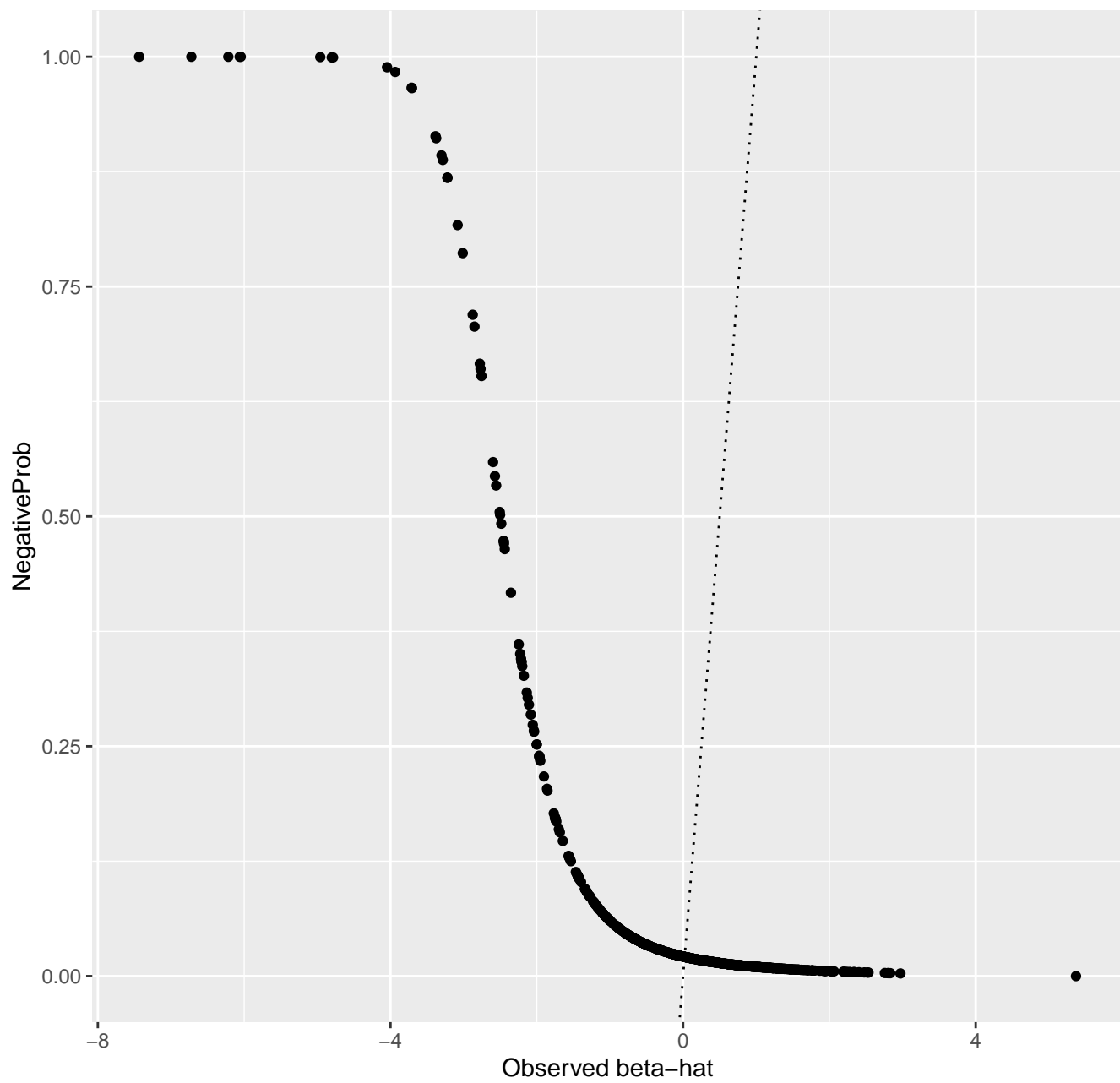
PRPS1_(5631) negative



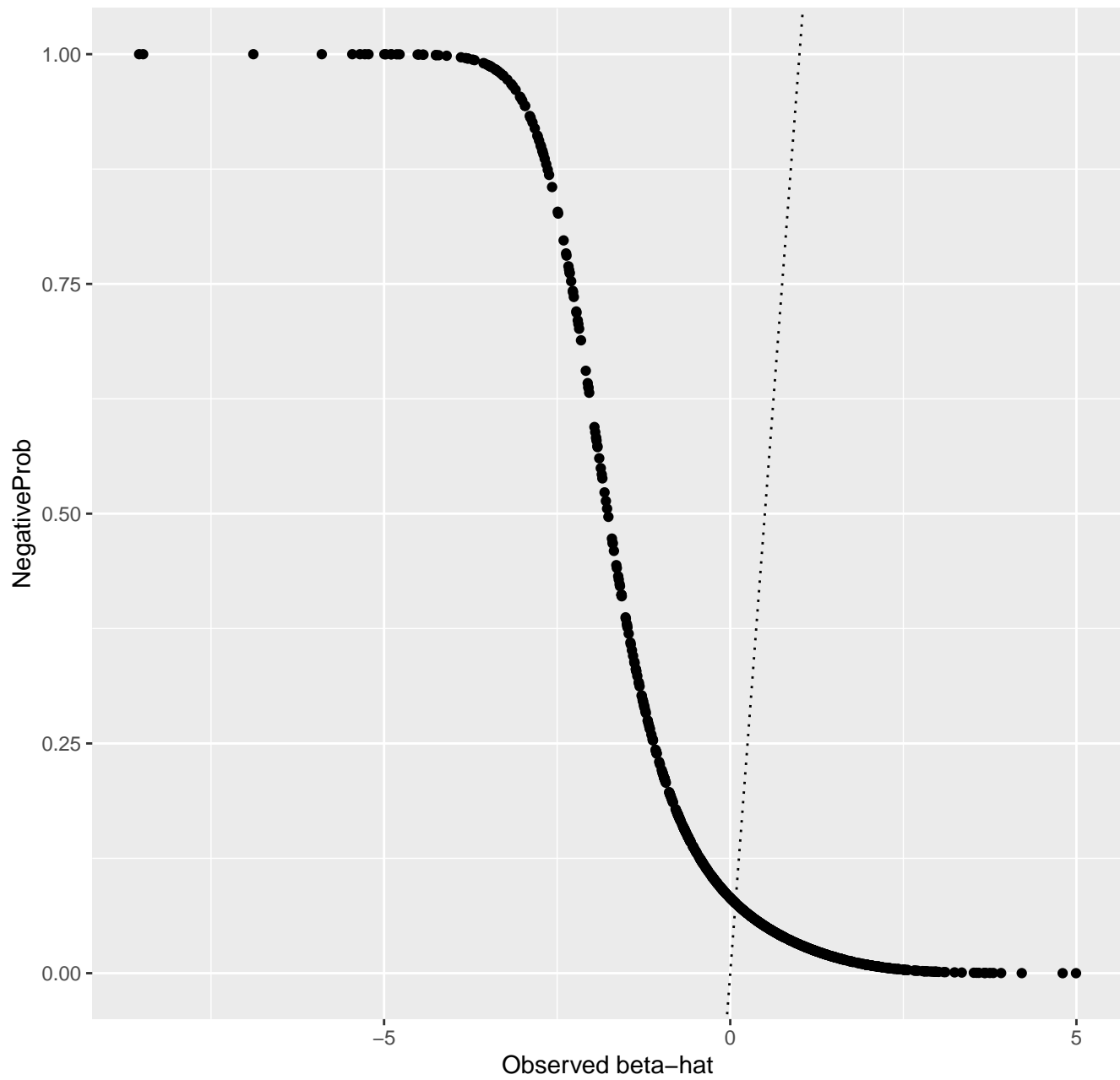
TFAP2C_(7022) negative



MAN1A2_(10905) negative



UBE2S_(27338) negative



FGF4_(2249) negative

